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Modulation of the GABA_A receptor by

CaMK-II dependent phosphorylation.

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Thesis submitted for the degree of Doctor of Philosophy at

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ABSTRACT:

Phosphorylation is an important mechanism for the modulation of ligandgated ion channel function. CaMK-II dependent modulation of GABA_A receptors was investigated through analysis of GABA_A receptor mediated currents recorded by whole-cell patch clamp from non-neuronal and neuronal cell lines and from primary neurones in culture.

Application of pre-activated α -CaMK-II had no effect on recombinant receptors expressed in HEK293 cells. Whole-cell currents recorded from cerebellar granule cells were significantly increased in peak amplitude on application of preactivated α -CaMK-II. Analysis of spontaneous (s)IPSCs recorded from cerebellar granule cells in culture in the presence of pre-activated α -CaMK-II revealed an increase in sIPSC amplitude and decay time constants. Expression of recombinant GABA_A receptors in the neuronal cell line NG108-15 allowed the functional effect of CaMK-II to be studied in a recombinant system. CaMK-II was able to up-regulate the function of $\alpha 1\beta 1\gamma 2S$ and $\alpha 1\beta 3\gamma 2S$ GABA_A receptors but not $\alpha 1\beta 2\gamma 2$ receptors. This modification of function occurred through phosphorylation of Ser³⁸³ on the β 3 subunit and through a downstream activation of a tyrosine kinase and subsequent phosphorylation of the $\gamma 2S$ subunit at previously identified sites of tyrosine kinase phosphorylation, Tyr³⁵⁷ and Tyr³⁵⁹. Tyrosine kinase phosphorylation of γ 2S occurred when co-expressed with $\alpha 1\beta 1$ or $\alpha 1\beta 3$ but not with $\alpha 1\beta 2$. Transfection of different GABA_A receptor subunits into cerebellar granule cells was carried out to confirm the importance of these sites in a neuronal environment. Analysis of sIPSCs recorded from cerebellar granule cell cultures prepared from $\beta 2$ subunit knockout mice revealed α -CaMK-II modification of β 2 subunit-containing receptors at the synapse, resulting in an increase in sIPSC amplitude with no change in decay times. This was in contrast to the lack of any functional modulation of $\alpha 1\beta 2\gamma 2S$ receptors observed in NG108-15 cells.

These findings are potentially important for understanding the mechanisms of CaMK-II dependent modulation of the GABA_A receptor and may also have significant implications for the understanding of inhibitory synaptic plasticity.

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Selected list of abbreviations:

AC, adenylate cyclase; AID, autoinhibitory domain; AKAP, A-kinase anchoring protein; **BAPTA**, 1,2-Bis(2-aminophenoxy)ethane-N,N,N',N'-tetraacetic acid; **BDNF**, brain-derived neurotrophic factor; BDZ, benzodiazepine; CBD, calmodulin binding domain; CaM, calmodulin; CaMK-II, Calcium/calmodulin-dependent protein kinase II; CaN, calcineurin; DARPP-32, dopamine- and cAMP- regulated phosphoprotein: EGTA, Ethylene glycol-bis(2-aminoethylether)-N,N,N',N'-tetraacetic acid; EPSC, excitatory postsynaptic current; GABARAP, GABA_A receptor associated protein; GFP, green fluorescent protein; HEK, human embryonic kidney; LTD, long term depression; LTP, long term potentiation; MAGUK, membrane associated guanylate kinase; MAPK, mitogen-activated protein kinase; m/nAchR, muscarinic/nicotinic acetylcholine receptor; m/sIPSC/P, miniature/spontaneous inhibitory synaptic current/potential; NSF, N-ethylmaleimide sensitive factor; OA, okadaic acid; PCR, polymerase chain reaction; PKA, cAMP-dependent protein kinase; PKB, protein kinase B; PKC, protein kinase C; PKG, cGMP-dependent protein kinase; PLC, phospholipase C; **PP(1/2)**, protein phosphatase 1/2; **PRIP-1**, PLC-related inactive protein type 1; PSD, postsynaptic density; RACK, receptor for activated C kinase;

<u>Chapter One: General Introduction:</u> <u>Modulation of the GABA_A receptor by phosphorylation.</u>

1.1 INTRODUCTION

The γ -aminobutyric acid type A (GABA_A) receptor is a pentameric ligandgated ion channel responsible for the majority of fast synaptic inhibition in the brain. It is related in structure to the cys-loop family of ligand-gated ion channels including the serotonin (5-HT₃) receptors, nicotinic acetylcholine receptors (nAChR) and glycine receptors (Nayeem *et al.*, 1994; Rothlin *et al.*, 1999; Cromer *et al.*, 2002; Korpi *et al.*, 2002). It is formed from five subunits that form a Cl⁻ permeable ion channel (Jenson *et al.*, 2002) that is opened after binding of the amino acid neurotransmitter GABA (Smith and Olsen, 1995; Moss and Smart, 1996; Mehta and Ticku, 1999; Moss and Smart, 2001; Connolly and Wafford, 2004).

1.1.1 Receptor structure.

Numerous different subunits of the GABA_A receptor have been identified which can be divided into subunit families based on sequence homology. These consist of α 1-6, β 1-3, γ 1-3, δ , π , ϵ , θ and ρ (Barnard *et al.*, 1998; Whiting *et al.*, 1999; Korpi *et al.*, 2002). There is typically 30-40 % homology between subunit families and 70-90 % between subunit isoforms (Smith and Olsen, 1995; Korpi *et al.*, 2002). In addition, some of these subunits have multiple splice variants the most commonly expressed being the γ 2S and γ 2L forms of the γ 2 subunit in which the γ 2L variant has an additional eight amino acids at position 338-345 (Moss and Smart, 1996). This heterogeneity of GABA_A receptors allows for a large amount of spatial and temporal diversity with different cells expressing different subunit combinations at different times (McKernan and Whiting, 1996; Whiting et al., 1999; Sieghart et al., 1999).

Each GABA_A receptor subunit contains a large extracellular domain, four transmembrane domains (TM) and a large intracellular loop between TM 3 and 4 (Smith and Olsen, 1995; Moss and Smart, 1996). The TM2 domain lines the channel pore (Xu and Akabas, 1996). The TM3-4 intracellular loop can account for up to 20 % of the total mass of a subunit and is known to contain multiple sites of phosphorylation (Smith and Olsen, 1995; Moss and Smart, 1995; Moss and Smart, 1996; See Fig 1.1).

The subunit stoichiometry of GABA_A receptors in the brain is thought to consist of $2\alpha 2\beta 1\gamma$ (Nayeem et al., 1994; Chang et al., 1996; Tretter et al., 1997; Farrar et al., 1999). In recombinant systems, it has been determined that the β 1 and β 3 subunits can form functional homomeric channels, although this doesn't appear to be the case for $\beta 2$ subunits (Krishek et al., 1996; Connolly et al., 1996a; Connolly et al., 1996b; Wooltorton et al., 1997; Taylor et al., 1999). For the formation of a robust GABA gated Cl⁻ channel it appears that co-expression of an α and a β subunit is required (Connolly et al., 1996a; Taylor et al., 1999). However $\alpha\beta$ channels have different single channel conductance and gating properties than $\alpha\beta\gamma$ channels and they are not modulated by benzodiazepines (Angelotti and Macdonald, 1993; Angelotti et al., 1993a). The inclusion of the γ subunit increases the amplitude of whole-cell currents recorded in a recombinant system and results in an increase in the EC₅₀ for GABA (Angelotti et al., 1993a). The inclusion of $\gamma 2$ in the receptor complex has also been reported to alter the kinetic properties of GABA_A receptors (Boileau et al., 2003). For modulation by benzodiazepines, an α and γ subunit are required (Sigel et al., 1990; Angelotti et al., 1993a).

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Fig 1.1:



Fig 1.1: Illustration of the proposed structure of the GABAA receptor.

(A) The GABA_A receptor is thought to be pentameric consisting of 2α , 2β and 1γ subunit. The channel opens in response to GABA binding and is permeable to Cl⁻ and HCO₃⁻ ions.

(B) Each subunit is proposed to consist of an N-terminal extracellular domain, four trans-membrane domains (TM), a large intracellular loop between TM3-4 and an extracellular N-terminus in which the cys-loop is situated. The large intracellular loop is thought to contain multiple potential sites of phosphorylation. A small external C-terminus is also evident.

(C) The TM2 region of each subunit is thought to line the channel pore.

Adapted from Moss and Smart (1996).

1.1.2 GABA_A receptor trafficking.

GABA_A receptors have been identified in various intracellular compartments within neurones, as well as at the plasma membrane, and it appears that receptors can dynamically shuttle between these locations (Brandon et al., 2002a; Kittler and Moss, 2003). The removal of receptors from the cell surface is mediated by clathrindependent endocytosis (See Fig 1.2). In this context, interaction between β and γ subunits and the adaptin complex, AP2, has been identified. AP2 is critical for the recruitment of receptors into clathrin-coated pits (Kittler et al., 2000). These processes can be modulated by phosphorylation, for example, PKC dependent downregulation of GABA_A receptor function can be partly mediated by a loss of receptors at the cell surface through a block of recycling of receptors back to the cell surface (Connolly et al., 1999). Plic-1 is a GABA_A receptor-associated protein which enhances receptor insertion into the membrane. A block of its interaction with the GABA_A receptor results in a rapid decrease in GABA_A receptor-mediated currents implying that Plic-1 is required for the replacement of receptors lost by internalisation (Bedford et al., 2001). Alteration in trafficking through perturbation of these pathways results in significant effects on GABA_A receptor-mediated currents and inhibitory synaptic transmission (Kittler and Moss, 2003).

The correct targeting of assembled receptors to synaptic sites is crucial for the proper functioning of inhibitory synaptic transmission. It is thought that an interaction between the $\gamma 2$ subunit and the scaffolding protein gephyrin may be involved in receptor targeting and clustering at the synapse (See Fig 1.2). The loss of the $\gamma 2$ subunit results in a large decrease in the number of receptors at synapses despite the fact that $\alpha\beta$ receptors are capable of trafficking to the cell surface. The loss of $\gamma 2$ also results in a subsequent loss of gephyrin, suggesting an important link

between the two proteins (Essrich *et al.*, 1998). Disruption of gephyrin results in the disruption of $\alpha 2/3$ subunit-containing GABA_A receptor clustering and synaptic targeting of $\gamma 2$ subunit-containing receptors. However, in gephyrin knockout mice, some GABA_A receptor subtypes are unaffected. Gephyrin is thought to act as a scaffolding molecule important for receptor clustering of certain subtypes of GABA_A receptor, but evidence suggests there must be additional proteins involved in clustering and trafficking (Kneussel *et al.*, 1999; Kneussel *et al.*, 2001).

The GABA_A receptor associated protein GABARAP has been identified to associate with the $\gamma 2$ subunit and is thought to play a role in intracellular transport and cell surface targeting of the receptor (Chen *et al.*, 2000; Kittler *et al.*, 2001; Kneussel, 2002; Kanematsu *et al.*, 2002; Leil *et al.*, 2004). The correct targeting of receptors to synapses is significant given the large amount of heterogeneity of GABA_A receptors and the apparent precise control of GABA_A receptor subtype expression and localisation (Koulen *et al.*, 1996; Brünig *et al.*, 2002).



Fig 1.2: Trafficking of the GABA_A receptor and interactions at the plasma membrane.

(A) $GABA_A$ receptor clustering in some cases is thought to be mediated by the protein gephyrin. Associations between gephyrin, the $GABA_A$ receptor and the cytoskeleton have been proposed to anchor and cluster $GABA_A$ receptors at the synapse.

Anchoring proteins can target protein kinases to the GABA_A receptor and can modulate, or be critical for kinase function (e.g. PKC and RACK-1).

(B) GABARAP has been implicated in the trafficking of GABA_A receptors to the plasma membrane through an interaction with the γ subunit of the GABA_A receptor, microtubules and N-ethylmaleimide sensitive factor (NSF). NSF is key protein involved in membrane fusion and is critical for intracellular transport.

(C) GABA_A receptors at synaptic sites undergo constitutive endocytosis via clathrinmediated endocytosis. This is mediated by an interaction between β and γ subunits and adaptor proteins (e.g. AP2) essential for the recruitment of protein cargo into clathrin-coated pits.

Adapted from Moss and Smart (2001).

1.1.3 GABA_A receptor localisation.

Immunocytochemistry and in situ hybridisation have allowed the temporal and spatial heterogeneity of GABA_A receptors within the brain to be revealed. There are distinct patterns of $GABA_A$ receptor subunit expression in different regions of the brain at different points in development (Persohn et al., 1992; Wisden et al., 1992; Laurie et al., 1992a; Laurie et al., 1992b). Different cell types often express a particular subset of the many subunits. Even within the same cell it appears that different subunits are targeted to precise locations (Koulen et al., 1996; Connolly et al., 1996b; Brünig et al., 2002). There is also evidence that certain subunits are preferentially expressed together (Benke et al., 1994). Certainly the number of receptor subtypes in the brain is believed to be significantly less than the theoretical number of possible combinations (McKernan and Whiting, 1996). The β 2 subunit is widely expressed and accounts for 60 % of all β subunits with a relatively smaller contribution of $\beta 1/3$ (Benke et al., 1994). However, recent evidence suggests that $\alpha 1\beta 2$ subunits prefer to co-assemble and that $\alpha 2/3\beta 3$ subunits also preferentially coassemble (Benke et al., 1994; Ramadan et al., 2003; Ortinski et al., 2004; Vicini and Ortinski, 2004). The $\alpha 1\beta 2\gamma 2$ receptor and the $\alpha 2\beta 3\gamma 2$ receptor together account for 75-85 % of the diazepam-sensitive receptors in the brain (Benke et al., 1994). Although these associations are not mutually exclusive, they suggest a broad division between these two receptor subtypes with each form being targeted to specific cellular locations (Whiting et al., 1999; Vicini and Ortinski, 2004).

1.1.4 Physiological role and pathology.

Activation of $GABA_A$ receptors in the adult brain causes the rapid influx of Cl^{-} ions which results in a hyperpolarisation of the cell. As a consequence they are

critical in the control of neuronal excitability (Moss and Smart, 1996). In immature neurones, GABA is excitatory resulting in cell depolarisation and the generation of action potential firing. Independent of the net flow of Cl⁻ ions, the opening of GABA_A receptors is often enough to shunt excitatory synaptic currents (Ruiz *et al.*, 2003). The shift from excitation to inhibition occurs when the KCC2, K^+ / Cl^- co-transporter is expressed changing the electrochemical gradient for Cl⁻ (Ben-Ari, 2002).

The GABA_A receptor is also the site of action of a number of important, clinically relevant drug families. The most widely used and characterised of these is the benzodiazepine family, which acts as an allosteric modulator of GABAA receptors enhancing their function in response to GABA. Disruptions to the normal function of GABA_A receptors have been linked to many pathological conditions (Mehta and Ticku, 1999). The clinical use of benzodiazepines is particularly widespread in anxiety and epilepsy (Lüddens et al., 1995; Basile et al., 2004). One of the most widely studied pathological conditions associated with GABAA receptors is epilepsy in which disruption or alteration of normal GABAergic synaptic transmission has been observed in a number of different experimental models (Kardos, 1999; Jones-Davis and Macdonald, 2003). Analysis of familial epilepsy has revealed mutations in GABA_A receptor subunit genes which result in altered functional behaviour of these receptors (Bianchi et al., 2002; Buhr et al., 2002; Fisher, 2004; Macdonald et al., 2004). GABA_A receptors are also thought to play an important role in alcohol dependence (Young et al., 2004) and alterations in the normal physiology of inhibitory synaptic transmission has been linked to schizophrenia (Lewis et al., 2005), Alzheimer's disease (Armstrong et al., 2003) and a number of anxiety disorders (Nemeroff, 2003; Basile et al., 2004).

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GABA_A receptors have been demonstrated to be critical for the formation of network oscillations crucial to behaviour (Nusser *et al.*, 2001; Gaiarsa *et al.*, 2002). The function of GABA_A receptors is critical to the development and maintenance of normal neuronal network activity. As a result it is of great importance to understand how GABA_A receptor function is regulated within the cell. The cellular signalling mechanism of phosphorylation has been shown to modulate the function of many ligand-gated ion channels including the GABA_A receptor. There is a substantial amount of evidence that phosphorylation can modulate GABA_A receptor function in physiologically relevant ways, however, there is not yet a full understanding of the mechanisms and the significance of this form of cellular signalling and modification.

1.2 Phosphorylation of the GABA_A receptor: recombinant systems.

The process of phosphorylation involves the covalent modification of receptor structure through the transfer of a highly charged phosphate group from adenosine triphosphate (ATP) to a serine, threonine or tyrosine residue within a substrate. This covalent modification is catalyzed by a series of enzymes known as protein kinases. The subsequent cleavage of the phosphate group from a substrate (dephosphorylation) is catalysed by protein phosphatases (Moss and Smart, 1996). The phosphorylation state of a given substrate is often bi-directionally regulated and the functional consequences are determined by a balance between kinase and phosphatase activities. The most common class of protein kinases are the serine/threonine second messenger dependent kinases such as cAMP-dependent protein kinase A (PKA) and protein kinase C (PKC). The second major class of

kinases are those that specifically phosphorylate tyrosine residues (Moss and Smart, 1996; Smart, 1997).

1.2.1 Phosphorylation in vitro.

Protein kinases catalyse phosphorylation acting at specific sites that can be recognised by the sequence surrounding the site of phosphorylation, which is often specific to a particular kinase. Consensus sites for a number of different kinases have been identified on $GABA_A$ receptor subunits, most commonly on the TM3-4 intracellular loop. However, consensus sites do not act as a guarantee that phosphorylation will actually occur in a living cell. Phosphorylation can occur at sites that do not conform to consensus sequences. As a result consensus sites can only be used as a rough guide for the potential targets and sites of phosphorylation (Moss and Smart, 1996; Sedelnikova and Weiss, 2002).

Consensus sites for a number of Ser/Thr kinases, such as PKA / PKC / cGMP-dependent protein kinase (PKG) and calcium/calmodulin dependent protein kinase II (CaMK-II) have been identified on the large intracellular loop between TM 3 and 4 in a range of GABA_A receptor subunits (Moss and Smart, 1996). These sites, to date, have all been identified on the β and γ subunits. The intracellular domains of GABA_A receptor subunits have been expressed as glutathione-S-transferase (GST) fusion proteins in *E-coli* which have been used to identify the individual sites of phosphorylation *in vitro* (See Table 1.1). The β 1 fusion protein can be phosphorylated by PKC/PKG/PKA/CaMK-II at Ser⁴⁰⁹ (Moss *et al.*, 1992a; McDonald and Moss, 1994). The β 2 fusion protein can be phosphorylated at the corresponding site, Ser⁴¹⁰, by the same kinases (Moss *et al.*, 1992a; McDonald and Moss, 1997) and the β 3 fusion protein can be phosphorylated at Ser⁴⁰⁹ by

PKA/PKC/PKG/CaMK-II (McDonald and Moss, 1997). CaMK-II can also phosphorylate additional sites in the β 1 and β 3 fusion proteins at Ser³⁸⁴ and Ser³⁸³, respectively (McDonald and Moss, 1994; McDonald and Moss, 1997). Both splice variants of γ 2 can be phosphorylated by CaMK-II at Ser³⁴⁸ and Thr³⁵⁰, and PKC can phosphorylate γ 2S/L at Ser³²⁷ (Moss *et al.*, 1992a; McDonald and Moss, 1994). The γ 2L subunit can also be phosphorylated by PKC and CaMK-II at Ser³⁴³ within the 8 amino acid insertion (Machu *et al.*, 1993; McDonald and Moss, 1994).

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GST

Subunit	Residue	Kinase	Reference
β1	Ser ³⁸⁴ Ser ⁴⁰⁹	CaMK-II PKA/PKC/PKG/CaMK-II	McDonald and Moss (1994)
β2	Ser ⁴¹⁰	PKA/PKG/PKC/CaMK-II	Moss <i>et al.</i> , (1992)
β3	Ser ³⁸³ Ser ⁴⁰⁸ Ser ⁴⁰⁹	CaMK-II PKC PKA/PKG/PKC/CaMK-II	McDonald and Moss (1997)
γ <mark>2</mark> S	Ser ³²⁷ Ser ³⁴⁸ Thr ³⁵⁰	PKC CaMK-II CaMK-II	McDonald and Moss (1994)
γ2L	Ser ³²⁷ Ser ³⁴³ Ser ³⁴⁸ Thr ³⁵⁰	PKC PKC/CaMK-II CaMK-II CaMK-II	McDonald and Moss (1994) Moss <i>et al.</i> , (1992) Machu <i>et al.</i> , (1993)

I/C Loop TM3-4

(P)

Table 1.1: A summary of the main sites of phosphorylation identified on the TM3-4 loop of the β and γ subunits of the GABA_A receptor.

Each of these potential sites of phosphorylation was identified by expression of GST-fusion proteins of the TM3-4 intracellular loop for different GABA_A receptor subunits and co-incubation, *in vitro*, with various protein kinases. Specific residues that were phosphorylated were identified by site-specific mutagenesis.

1.2.2 Phosphorylation of recombinant GABA_A receptors: PKC

The study of phosphorylation in neurones has been hampered by the heterogeneity of GABA_A receptors between cell types and within the same cell. As a result phosphorylation of GABA_A receptors has been studied in a number of different recombinant systems to attempt to determine the functional consequence of phosphorylation of different receptor subunits. Transfection of relevant subunit cDNA's into a recombinant cell line with no endogenous GABA_A receptor subunits allows the precise subunit combination of the receptor to be controlled within reasonable limits. However, even within recombinant systems there is a considerable degree of variation in the reported effects of phosphorylation (Song and Messing, 2005).

Phosphorylation of the recombinant subunit combination $\alpha 1\beta 2\gamma 2S$ expressed in *Xenopus oocytes* by PKC has been reported to decrease the amplitude of wholecell currents recorded. This modification was abolished by the site-specific mutation of the residues Ser⁴¹⁰ site on $\beta 2$ and Ser³²⁷ on the $\gamma 2S$ subunit to alanine (Kellenberger *et al.*, 1992). Both sites were thought to be required for a full functional effect. A similar functional result of PKC activation on $\alpha 1\beta 2\gamma 2S$ receptors expressed in *Xenopus oocytes* was also reported that was not mediated by phosphorylation of $\beta 2$ Ser⁴¹⁰. In this case it was thought that PKC dependent phosphorylation resulted in a decrease in the number of GABA_A receptors at the cell surface (Chapell *et al.*, 1998). PKC activity has also been reported to block receptor recycling of $\alpha 1\beta 2\gamma 2$ receptors expressed in HEK293 cells back to the cell surface thereby reducing receptor number at the cell surface (Connolly *et al.*, 1999).

PKC dependent phosphorylation of $\alpha 1\beta 1\gamma 2S$ expressed in HEK293 cells and in *Xenopus oocytes* results in a decrease of the amplitude of whole-cell currents

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recorded that was found to be dependent on phosphorylation of Ser⁴⁰⁹ in the β 1 subunit, Ser³²⁷ in the γ 2S subunit and Ser³⁴³ in γ 2L (Leidenheimer *et al.*, 1992; Krishek *et al.*, 1994). However, phosphorylation of similar residues in the β 1 and γ 2L subunits has also been shown to mediate a PKC dependent enhancement of the amplitude of whole-cell currents recorded from L929 fibroblasts transfected with $\alpha 1\beta 1\gamma 2L$ receptors (Lin *et al.*, 1994; Lin *et al.*, 1996).

The discrepancies between these different effects may be in part due to the mechanism of activation of PKC. It is difficult to rule out the possibility that activation through different mechanisms may activate different signalling pathways. It is always possible that another downstream kinase or signalling pathway is involved in, or masks an apparent effect of, phosphorylation. In addition, the cell type may alter the response to phosphorylation as different cell lines may contain different endogenous phosphatases and kinases that may alter the basal phosphorylation state of the receptor.

Many kinases have multiple substrates within a cell and play a role in many different cellular processes. There is increasing evidence that the localisation of a kinase, possibly through specific kinase anchoring proteins, has a profound effect on the functional effects of phosphorylation (Pawson and Scott, 1997). Some differences in the response to phosphorylation between cell lines may therefore reflect differences in the anchoring proteins present. The receptor for activated <u>C</u> kinase-1 (RACK-1) has been shown to facilitate phosphorylation and functional modulation of GABA_A receptors in HEK293 cells. In this case PKC dependent phosphorylation of β 1 Ser⁴⁰⁹ resulted in a depression of whole-cell currents. Blocking the binding of RACK-1 to the β 1 subunit (with a dominant negative peptide) abolished this PKC dependent effect (Brandon *et al.*, 2002b).

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1.2.3 Phosphorylation of recombinant GABAA receptors: PKA

One further factor that is important concerns the receptor subunit composition. PKA dependent phosphorylation of recombinant receptors has been demonstrated to be dependent on the β subunit present. Within HEK293 cells, PKA increased the amplitude of whole-cell currents recorded from $\alpha 1\beta 3\gamma 2S$ receptors through phosphorylation of Ser^{408,409}. PKA dependent phosphorylation of β 1 at Ser⁴⁰⁹ resulted in a depression of whole-cell currents recorded from HEK293 cells expressing $\alpha 1\beta 1\gamma 2S$ (Moss *et al.*, 1992b). The mutation of Ser⁴⁰⁸ on the $\beta 3$ subunit converted the observed enhancement into a depression similar to that seen for $\beta 1$ as well as reproducing the phosphorylation profile of the β 1 subunit. The β 2 subunitcontaining receptors were not functionally modulated at all by PKA in this study (McDonald et al., 1998). Chronic activation of PKA in C α 12 fibroblast cells (L929 cell line over-expressing the active catalytic subunit of PKA) led to an enhancement of $\alpha 1\beta 1\gamma 2S$, receptors, in contrast to the depression observed after acute activation of PKA (McDonald et al., 1998). This enhancement was also dependent on Ser⁴⁰⁹ but in this case was also dependent on the presence of the $\gamma 2S$ subunit (Angelotti et al., 1993b). This implied that acute transient activation of a protein kinase may have a different functional effect to that of chronic activation. In addition over-expression of this catalytically active form of PKA (cAMP-independent) in this cell line could be expected to produce levels of PKA activity that far exceeded the levels reached using other protocols.

The effects of PKA dependent phosphorylation on desensitisation have also been studied in a recombinant system through rapid application of GABA to HEK293 cells expressing $\alpha 1\beta 1\gamma 2L$ and $\alpha 1\beta 2\gamma 2L$. The residues Ser⁴⁰⁸ in $\beta 1$ and Ser⁴⁰⁹ in $\beta 3$ were again identified as critical for PKA dependent modulation. In this case PKA phosphorylation of both β 1 and β 3 resulted in alterations in desensitisation and deactivation that prolonged the duration of inhibitory postsynaptic current (IPSC)–like currents (Hinkle and Macdonald, 2003) suggesting that phosphorylation may be capable of modulating IPSC duration in neurones. The relatively less well studied protein kinase B (PKB/Akt), has also been demonstrated to phosphorylate the β 2 residue Ser⁴¹⁰ in HEK293 cells. This appeared to increase the number of GABA_A receptors at the cell surface (Wang *et al.*, 2003b).

1.2.4 Phosphorylation of recombinant GABA_A receptors: tyrosine kinases.

In addition to Ser/Thr kinases, tyrosine kinases are capable of phosphorylating and modulating the function of ligand-gated ion channels (Boxall and Lancaster, 1998). Tyrosine kinase phosphorylation of both β and γ subunits of GABA_A receptors has been reported (Moss et al., 1995; Valenzuela et al., 1995; Wan et al., 1997). Tyrosine phosphorylation of β subunits has been reported to increase the amplitude of whole-cell currents in HEK293 cells (Wan et al., 1997). However, Moss et al. (1995) originally reported that the β subunit phosphorylation was functionally silent and that phosphorylation of $Tyr^{365:367}$ on the $\gamma 2L$ subunit alone was responsible for the increase in whole-cell currents. Valenzuela and colleagues (1995) determined that $\beta 2$ and $\gamma 2$ GABA_A receptor subunits purified from bovine brain could be phosphorylated by Src as well as the intracellular loop of $\beta 1$ and $\gamma 2L$ in vitro. They also observed in Xenopus oocytes expressing $\alpha 1\beta 1$ and $\alpha 1\beta 1\gamma 2L$ receptors an inhibition of GABA currents on bath application of genistein suggesting in this case β 1 phosphorylation had a functional effect (Valenzuela *et al.*, 1995). However, genistein can have a direct effect on GABA_A receptor function (Dunne et al., 1998). Tyrosine phosphorylation may maintain $GABA_A$ receptor activity and a

loss of basal tyrosine kinase activity may be responsible for the phenomenon of "rundown" (Huang and Dillon, 1998).

The α subunit of the GABA_A receptor is not thought to be a good substrate for phosphorylation. There is a lack of suitable consensus sites within the TM3-4 intracellular loop and the α l GST-fusion protein cannot be phosphorylated *in vitro* (Moss *et al.*, 1992a; Moss and Smart, 1996). However, CaMK-II dependent phosphorylation of the α l subunit has been reported using synaptosomal membrane fractions (Churn *et al.*, 2002). In addition, phosphorylation of the α l TM3-4 intracellular loop, that supports GABA_A receptor activity and prevents receptor rundown during electrophysiological recording, has been observed in acutely dissociated rat neocortical neurones (Laschet *et al.*, 2004).

1.3 Phosphorylation of the GABA_A receptor: neuronal systems.

There is a wide range of evidence that phosphorylation of the GABA_A receptor can modulate function in a neurone in a physiologically significant way. However, there has been a wide range of different functional effects reported. Part of this variation in response to phosphorylation between different cell-types and culture conditions may reflect differences in the GABA_A receptor subunits that are expressed. However, even taking this into account there are still discrepancies between reported effects of phosphorylation in recombinant systems and neurones.

1.3.1 PKC dependent effects in neuronal preparations.

The effects of PKA and PKC dependent phosphorylation have been the most widely studied. Activation of PKC in cultured sympathetic neurones results in a decrease in whole-cell currents recorded, similar to that seen in HEK293 cells
expressing $\alpha 1\beta 1\gamma 2$ receptors (Krishek *et al.*, 1994). PKC activity has been reported to have different effects in different cell types. In CA1 pyramidal neurones, PKC was without effect (Poisbeau *et al.*, 1999); however in dentate gyrus granule cells, PKC increased the peak amplitude of spontaneous (s)IPSCs (Poisbeau *et al.*, 1999). Dentate gyrus cells express pre-dominantly $\beta 3$ subunits (Miralles *et al.*, 1999; Pirker *et al.*, 2000) and the increase in amplitude is consistent with $\beta 3$ subunit mediated increases in GABA_A receptor function observed in other neuronal cell types (Jovanovic *et al.*, 2004). However, in cortical neurones, constitutive phosphorylation of $\beta 3$ Ser^{408,409} by PKC has been reported to decrease the amplitude of whole-cell currents (Brandon *et al.*, 2000).

The anchoring protein RACK-1 has been shown to form complexes with PKC and GABA_A receptors in the adult rat brain. In rat superior cervical ganglion neurones activation of PKC led to a decrease in the amplitude of GABA-mediated currents. Inclusion of a peptide that competes with RACK-1 for binding to the β 1 subunit reduced the effect of PKC on these GABA_A receptors (Brandon *et al.*, 2002b). The absence or presence of RACK-1 or other similar proteins within a cell-type could significantly alter the apparent effect of phosphorylation and may explain some of the differences reported for PKC modulation within cells that express similar GABA_A receptor subtypes.

Brain-Derived Neurotrophic factor (BDNF) can modulate GABA_A receptor mediated mIPSCs through the activation of PKC, followed by protein phosphatase (PP2A) mediated dephosphorylation of the receptor. There was differential recruitment of PKC, RACK-1 and PP2A according to the phosphorylation state of the β 3 subunit. Phosphorylation of β 3 at Ser^{408:409} was associated with an increase in miniature (m)IPSC amplitude, with no change in the decay time (Jovanovic *et al.*, 2004).

Recombinant studies highlighted potential phosphorylation sites on the $\gamma 2$ subunit for PKC at Ser³²⁷ and Ser³⁴³ within the eight amino acid insert that forms the $\gamma 2L$ splice variant. Within neurones the role of PKC phosphorylation of Ser³²⁷ is unknown. However, phosphorylation of Ser³⁴³ has been reported in neurones to facilitate the trafficking of $\gamma 2S$ to the synapse (Meier and Grantyn, 2004). Activation of PKC has also been reported to increase receptor internalisation in cultured cortical neurones and promotes receptor endocytosis in HEK293 cells. This appears to occur via a dynamin endocytic pathway that is dependent on a dileucine motif in the $\beta 2$ subunit and also requires the presence of the $\gamma 2$ subunit (Herring *et al.*, 2005).

1.3.2 PKA dependent effects in neuronal preparations.

Modulation of $GABA_A$ receptors by PKA in neurones has also revealed different functional effects in different cell-types (Porter *et al.*, 1990; Kano and Konnerth, 1992; Moss *et al.*, 1992b; Kapur and Macdonald, 1996). Again, some of this may be related to differential effects on different subunits. There is however, some discrepancy between reported effects in a recombinant system and that seen in neurones.

Inclusion of the catalytic domain of PKA in the patch pipette was reported to reduce the amplitude of whole-cell currents recorded from mouse spinal neurones. This reduction was reported to be mediated by a reduction in channel opening frequency (Porter *et al.*, 1990). It has also been reported that in CA1 hippocampal pyramidal cells PKA reduced the amplitudes of IPSCs and altered the fraction of events with a double exponential decay. However, in hippocampal dentate gyrus . granule cells PKA was without effect (Poisbeau et al., 1999). In granule cells of the olfactory bulb which express predominantly the β 3 subunit, constitutively-active PKA resulted in an increase in IPSC amplitudes with no change in decay (Nusser et al., 1999). An increase in amplitude would correlate with the up-regulation of function seen in HEK293 cells expressing β 3 subunit-containing receptors (McDonald et al., 1998). The lack of a reported effect of PKA on dentate gyrus cells is surprising given this cell-type also expresses predominantly β 3 subunit-containing receptors (Miralles et al., 1999; Pirker et al., 2000). However a PKA dependent increase in whole-cell currents recorded from hippocampal dentate granule cells has been reported by another group (Kapur and Macdonald, 1996). Therefore, even within the same cell type, variations in response to phosphorylation occurs possibly as a result of differences in tissue preparation or experimental method of kinase activation. Addition of PKA to cerebellar stellate and basket cells that express predominantly $\beta 2$ subunit-containing receptors prolonged the time-course of IPSCs without changing the peak amplitude (Nusser et al., 1999). This is in contrast to the lack of any functional modulation observed on $\beta 2$ subunit-containing receptors in HEK293 cells (McDonald et al., 1998).

Phosphorylation of β 3 Ser^{408,409} by PKA can be counteracted by protein phosphatase 1 (PP1). This interaction can also be regulated by additional binding proteins. For instance, phospholipase C (PLC)-related inactive protein type 1 (PRIP-1) binds and inactivates protein phosphatase (PP)1 allowing PKA phosphorylation to occur (Terunuma *et al.*, 2004). The <u>A-kinase anchoring protein (AKAP) 79/150 has</u> been identified which allows PKA to be selectively targeted to β 1/3 subunitcontaining receptors. This protein binds to β 1 and β 3 only and is critical for PKA mediated phosphorylation (Brandon *et al.*, 2003). PKA dependent effects on GABA_A receptors in neurones have been observed in cell-types thought to express predominantly $\beta 2$ subunit-containing receptors (Kano and Konnerth, 1992; Nusser *et al.*, 1999). It may be that other, as yet unidentified, kinase anchoring proteins exist that are able to allow PKA dependent phosphorylation of the $\beta 2$ subunit. Or alternatively, the activation of PKA or PKC in neurones might activate secondary pathways downstream or in parallel which influence GABA_A receptor function. PKA or PKC might be able to phosphorylate proteins associated with the GABA_A receptor which may have different effects to direct phosphorylation of the receptor itself. The considerable variation in response to activation of these kinases in neuronal systems implies that phosphorylation dependent modulation of GABA_A receptors is complex and may well involve multiple kinase anchoring proteins and/or different mechanisms of phosphorylation mediated effects.

1.3.3 PKB and Src.

In addition to PKA and PKC the Ser/Thr kinase PKB has also been demonstrated to phosphorylate the GABA_A receptor in neurones at Ser⁴¹⁰ in the β 2 subunit. PKB activity leads to an increase in the number of functional GABA_A receptors at the cell surface, increasing IPSC amplitudes with no change in decay times. The tyrosine kinase Src has also been demonstrated to alter GABA_A receptor function in neurones. Src mediated phosphorylation of Tyr^{365:367} on γ 2L leads to an increase in the whole-cell GABA current amplitude recorded from superior cervical ganglion neurones in culture. Single channel recordings revealed an increase in mean open time and probability of opening of the channel after activation of Src (Moss *et al.*, 1995). Src is capable of interacting with β and γ 2 subunits. The

residues Tyr^{365,367} on γ 2 have also been shown to be phosphorylated in the adult rat brain (Brandon *et al.*, 2001).

1.3.4 Mechanisms of phosphorylation dependent changes in receptor function.

The process of phosphorylation appears to be more complex than a simple covalent modification of receptor structure. There are a number of potential mechanisms for phosphorylation-dependent effects. The simplest mechanism is direct phosphorylation of the receptor followed by a structural change which alters the ion channel properties. For instance tyrosine kinase dependent changes in mean open time and probability of opening (Moss *et al.*, 1995). These types of changes could also alter the amplitude or decay phase of whole-cell currents or IPSCs. One particular mechanism that has been identified is phosphorylation-dependent modulation of desensitisation (Jones and Westbrook, 1996). This could alter the apparent decay times of IPSCs, a phenomenon that has been observed in response to phosphorylation. Phosphorylation dependent changes in desensitisation of the protein phosphatase calcineurin (Jones and Westbrook, 1997).

Another potential mechanism of phosphorylation-dependent changes is through modulation of receptor trafficking which could occur through direct phosphorylation of the receptor or indirectly through phosphorylation of other proteins involved in the correct targeting and insertion of receptors at the synapse. For instance, PKC can alter the trafficking of GABA_A receptors by blocking receptor recycling to the cell surface thereby reducing the number of functional receptors at the plasma membrane (Connolly *et al.*, 1999). It has also been proposed that PKC activity can induce receptor internalisation through a dynamin-dependent endocytic pathway (Herring *et al.*, 2005). In addition, PKC phosphorylation of Ser³⁴³ has been linked to the correct targeting of γ 2L subunits to the synapse (Meier and Grantyn, 2004).

Phosphorylation dependent changes to receptor trafficking may also allow the insertion of new receptors in order to potentiate inhibitory synapses. There is evidence that PKB phosphorylation may function in this way (Wang *et al.*, 2003b). The modulation of excitatory AMPA receptors by phosphorylation has been shown to occur by both mechanisms (Lisman *et al.*, 2002). Phosphorylation can directly change the properties of the AMPA channel (Barria *et al.*, 1997; Benke *et al.*, 1998); however, synaptic plasticity driven by phosphorylation can result in the insertion of new AMPA receptors at the synapse (Hayashi *et al.*, 2000; Malinow and Malenka, 2002; Bredt and Nicoll, 2003; Ju *et al.*, 2004).

1.3.5 Perturbation of normal phosphorylation pathways in disease states.

The abnormal function of protein kinases or perturbation of specific signalling pathways can be linked to particular disease states. Mice lacking the γ isoform of PKC have been shown to exhibit decreased anxiety (Bowers *et al.*, 2000). The mutant tottering (tg) mouse, displays seizures similar to human petit-mal epilepsy. It has been reported that there is an abnormal elevation of PKA activity in these mice and a reduction in GABA_A receptor function, indicating that this may contribute to the induction of seizures (Tehrani and Barnes, 1995). Cell membranes isolated from patients with drug-resistant temporal lobe epilepsy were injected into *Xenopus oocytes* and the response of GABA_A receptors to BDNF was determined. BDNF potentiated the epileptic GABA_A receptors probably through activation of tyrosine kinase receptor B and both PLC and PKC. Recombinant $\alpha l\beta 2\gamma 2$ receptors

and $GABA_A$ receptors transplanted from a non-epileptic brain were unaffected by BDNF (Palma *et al.*, 2005). The onset of epilepsy appears to induce changes in the $GABA_A$ receptor in terms of its response to signalling pathways and phosphorylation.

1.4 Calcium/calmodulin-dependent protein kinase II: structure and function.

Calcium/calmodulin dependent protein kinase II is a serine/threonine 2nd messenger dependent protein kinase that responds to increases in intracellular Ca²⁺ concentration (Fink and Meyer, 2002; Hudmon and Schulman, 2002; Soderling *et al.*, 2002). It is a multifunctional protein that phosphorylates a large number of different substrates and is involved in many different cellular processes (Schulman, 2004; Colbran, 2004b). There are four main families of CaMK-II subunits (α , β , γ , δ) and around 28 different isoforms. The α and β subunits are most prominent in the brain where they are highly expressed (Hudmon and Schulman, 2002). CaMK-II is known to phosphorylate a number of proteins involved in synaptic transmission and is thought to play a key role in synaptic plasticity (Lisman *et al.*, 2002; Colbran and Brown, 2004).

1.4.1 CaMK-II structure and function: autophosphorylation at Thr²⁸⁶.

CaMK-II has a unique complex structure containing self-regulatory properties. Each subunit consists of an N-terminal catalytic domain, a central regulatory domain (containing within it an auto-inhibitory domain), a Ca²⁺/CaM binding motif and a C-terminal association domain (Shen and Meyer, 1998; Hudmon and Schulman, 2002; Lisman *et al.*, 2002). Subunits can associate together in groups

of 10-12 to form a holoenzyme arranged as two stacked rings (See Fig 1.3, 1.4) with the association domains in the centre and the catalytic domains facing outwards (Kolodziej *et al.*, 2000; Hoelz *et al.*, 2003).

In the absence of Ca^{2+}/CaM , CaMK-II remains relatively inactive. The autoinhibitory domain (AID) acts as a pseudo-substrate which binds with the catalytic region of the same subunit acting as a gate that inhibits enzyme activity (Mukherji *et al.*, 1994; Mukherji and Soderling, 1994). The AID makes multiple interactions with the catalytic domain. Part of the AID occupies the substrate binding site or the S-site of the kinase whereas a more N-terminal portion of the AID makes an additional interaction on the C-terminal lobe of the catalytic domain (T-site). The interaction at the T-site is thought to induce conformational changes that interfere with ATP binding to the kinase. In this way the kinase remains inactive in the absence of Ca^{2+}/CaM (Hudmon and Schulman, 2002; Lisman *et al.*, 2002; Colbran, 2004b). The Ca^{2+}/CaM binding motif overlaps the AID, so Ca^{2+}/CaM binding is thought to disrupt the interaction between the AID and the catalytic domain allowing the kinase to become active (See Fig 1.4).

After binding of Ca^{2+}/CaM and subsequent activation of the kinase, autophosphorylation can occur at Thr²⁸⁶ (α -subunit, Thr²⁸⁷ - β -subunit) within the AID. This occurs between adjacent subunits that have bound Ca²⁺/CaM (Hanson *et al.*, 1994). Autophosphorylation causes the dissociation rate for Ca²⁺/CaM to fall significantly so that it is effectively bound to the kinase. Even after Ca²⁺/CaM dissociates the kinase retains partial Ca²⁺/CaM independent activity (Colbran *et al.*, 1988; Fong *et al.*, 1989), until it is dephosphorylated by a protein phosphatase (Hashimoto *et al.*, 1987). Autophosphorylation at Thr²⁸⁶ allows CaMK-II to remain persistently active after the initial stimulus and Ca²⁺ signal has dissipated. Thus, a transient rise in Ca^{2+} can stimulate long-term kinase activation (See Fig 1.3 and 1.4). It also allows CaMK-II to respond to the frequency of Ca^{2+} oscillations (Hanson *et al.*, 1994; De Koninck and Schulman, 1998). The extent of CaMK-II activity is also dependent on the rate at which subunits are de-phosphorylated. Evidence suggests that in the PSD this dephosphorylation is primarily mediated by PP1 (Hudmon and Schulman, 2002; Lisman *et al.*, 2002).



Fig 1.3: A schematic representation of the cycle of activation and deactivation of CaMK-II.

The binding of Ca^{2+}/CaM activates CaMK-II and enables it to phosphorylate substrates. It also allows autophosphorylation to occur at Thr²⁸⁶ between adjacent subunits that have bound Ca²⁺/CaM.

Autophosphorylation reduces the rate of Ca^{2+}/CaM dissociation, trapping Ca^{2+}/CaM to the kinase. When Ca^{2+}/CaM eventually dissociates the kinase retains some Ca^{2+}/CaM independent activity until Thr²⁸⁶ is dephosphorylated by a protein phosphatase and the kinase becomes inactive again.

CaMK-II forms holoenzymes of 10-12 subunits. Each subunit consists of three functional domains, a catalytic domain, an autoinhibitory domain and an association domain. Thr²⁸⁶ autophosphorylation and Ca²⁺/CaM binding occurs within the autoinhibitory domain.

Adapted from Molecular Biology of the Cell (1994), Garland Publishing.



Fig 1.4: Proposed 3D structure of CaMK-II and schematic representation of the interaction between the catalytic and autoinhibitory domains.

(A) Three-dimensional structure of CaMK-II holoenzyme, showing a hexameric ring from a plan-view perspective.

(B) The autoinhibitory domain acts as a gate to regulate catalytic activity. The enzyme is inhibited when the autoinhibitory domain binds at the S and T sites. The binding of Ca^{2+}/CaM disrupts the association and the enzyme becomes active. A site on the NR2B subunit of the NMDA receptor can bind the T site, disrupting the association of the autoinhibitory domain and keeping the enzyme active in the absence of Ca^{2+}/CaM . Autophosphorylation at Thr²⁸⁶ is also sufficient to disrupt the association between the catalytic domain and the autoinhibitory domain and to allow CaMK-II to be active after the dissociation of Ca^{2+}/CaM .

Adapted from Lisman et al. (2002) and Kolodziej et al. (2000).

1.4.2 CaMK-II structure and function: autophosphorylation at Thr^{305/6}.

CaMK-II can also undergo autophosphorylation at other sites. This can occur as a slow intra-subunit autophosphorylation that occurs when CaMK-II is in its basal state leading to phosphorylation of $Thr^{305/6}$. This has the effect of blocking subsequent Ca²⁺/CaM binding and so locks the kinase in an inactive state. Following autophosphorylation at Thr^{286} removal of Ca²⁺/CaM can induce a burst of inhibitory autophosphorylation at Thr^{305} and/or Thr^{306} , either of which is sufficient to block further Ca²⁺/CaM binding. In this case the kinase retains partial activity due to prior phsophorylation at Thr^{286} (Hanson and Schulman, 1992; Colbran, 2004b).

Phosphorylation of these sites is strongly opposed by phosphatase activity and it has been difficult to observe their phosphorylation *in vivo*. However, formation of knock-out mice has revealed that these sites may be important in regulating sub-cellular localisation of CaMK-II as well as regulating CaMK-II activity. Phosphorylation of these sites is elevated in a mouse model of the neurodevelopmental disorder Angelman syndrome (Elgersma *et al.*, 2002; Weeber *et al.*, 2003).

1.4.3 Regulation of CaMK-II by protein phosphatases.

It is important to consider how the activity of protein phosphatases regulates CaMK-II activity in the brain (Colbran, 2004a). It has been known for a number of years that PP1, PP2A and PP2C can dephosphorylate Thr²⁸⁶ and that in some cell types autophosphorylation of CaMK-II is strongly opposed by phosphatase activity (Fukunaga *et al.*, 1989; Molloy and Kennedy, 1991; Fukunaga *et al.*, 1993). It is thought that PP1 acts to dephosphorylate Thr²⁸⁶ at the PSD, whereas PP2A might dephosphorylate CaMK-II in the cytosol (Strack *et al.*, 1997). Therefore the

localisation of the kinase may alter which phosphatase is the more physiologically relevant for dephosphorylation (Strack *et al.*, 1997; Colbran, 2004a). The presence of phosphatases may alter CaMK-II function, as the presence of PP1 results in an ultra-sensitive kinase activation within a tighter range of Ca^{2+} concentrations than that observed for the kinase alone (Bradshaw *et al.*, 2003). The opposing actions of CaMK-II and PP1 have been suggested to be such that the CaMK-II/PP1 system could function as an energy efficient bi-stable switch (Lisman and Zhabotinsky, 2001). Phosphatase activity against CaMK-II has also been shown to be physiologically and pathologically relevant in that PP1 activity modulates LTP induction (Brown *et al.*, 2000) and decreased PP1/PP2A activity has been shown to be involved in the alterations in CaMK-II inhibitory autophosphorylation associated with the mouse model of Angelman syndrome (Weeber *et al.*, 2003; Elgersma *et al.*, 2004).

The binding of Ca^{2+} to CaM and recruitment of the Ca^{2+}/CaM complex to a CaMK-II enzyme is not simply a diffusion of Ca^{2+} and CaM in solution but is a regulated process in itself involving multiple cellular proteins and processes. Ca^{2+} influx to a cell can occur via voltage-gated Ca^{2+} channels or ligand-gated channels (e.g., NMDA receptors). Within the cell release of Ca^{2+} in stores can also provide a stimulus to activate CaMK-II. There are a large number of proteins present in neurones that can bind CaM. Increases in Ca^{2+} cause the release of this stored CaM and phosphorylation of some CaM binding proteins can cause CaM to dissociate, allowing it to participate in other signalling events (Gnegy, 2000; Meldolesi, 2001; Xia and Storm, 2005). As well as activating CaMK-II, CaM is involved in several other cellular signalling cascades and can have direct effects on ligand-gated ion channel function. For example, CaM has been shown to bind to the NR1 subunit of

the NMDA receptor and cause a four-fold reduction in channel open probability (Ehlers *et al.*, 1996; Zhang *et al.*, 1998; Rycroft and Gibb, 2002). It can also activate the phosphatase calcineurin (PP2B). PP2B is not thought to directly dephosphorylate Thr²⁸⁶ like PP1/PP2A but could act indirectly through dephosphorylation of DARPP-32 which inhibits PP1 activity. Therefore PP2B activity against DARPP-32 might indirectly increase PP1 activity against CaMK-II (Kawaguchi and Hirano, 2002; Svenningsson *et al.*, 2004; Colbran, 2004a). In addition, activation of PP2B/calcineurin is thought to be involved in some forms of LTD at both excitatory and inhibitory synapses (Winder and Sweatt, 2001; Wang *et al.*, 2003a).

1.5 CaMK-II localisation and role in synaptic plasticity.

CaMK-II is an abundant enzyme in the nervous system with multiple substrates that responds to changes in intracellular Ca²⁺ (Bayer and Schulman, 2001; Hudmon and Schulman, 2002; Griffith *et al.*, 2003; Colbran, 2004b). Calcium itself is a ubiquitous signalling molecule which plays a role in multiple cellular processes. Ca²⁺ signalling is tightly controlled both temporally and spatially (Berridge *et al.*, 1998). Increases in Ca²⁺ usually occur within discrete microdomains, small local regions of Ca²⁺ influx and localisation of Ca²⁺-sensitive effector proteins to these regions of Ca²⁺ influx is often crucial for the proper function of Ca²⁺-dependent signalling pathways (Franks and Sejnowski, 2002; Augustine *et al.*, 2003). The localisation of CaMK-II to sites of action and an ability to translocate on stimulation to specific sites of action is thought to be crucial for the proper function of the kinase (Bayer and Schulman, 2001; Griffith *et al.*, 2003; Schulman, 2004; Colbran, 2004b).

1.5.1 The role of α and β subunits of CaMK-II.

Examination of different forms of CaMK-II has revealed the importance of the different subunits of CaMK-II. The α and β isoforms can exist as homomers or can form heteromers with varying amounts of α : β subunits. Within the cerebellum there is a high ratio of β -CaMK-II to α -CaMK-II but within the forebrain α -CaMK-II is dominant (Miller and Kennedy, 1985). The physical properties of these two isoforms of CaMK-II are identical in that homomers of α or β subunits possess the same substrate specificity. However, β -CaMK-II appears to be more sensitive to Ca²⁺ and becomes autophosphorylated at Thr²⁸⁷ at much lower Ca²⁺/CaM levels (Miller and Kennedy, 1985). Half-maximal autophosphorylation of the α subunit occurs at 130 nM CaM whereas the same level of autophosphorylation of β subunits occurs at 15 nM CaM (Brocke *et al.*, 1999).

The β subunit is expressed earlier in development, within the embryo, whereas the α subunit is only expressed postnatally and the ratio of α to β increases throughout synaptogenesis (Burgin *et al.*, 1990; Vallano, 1990; Griffith *et al.*, 2003). The localisation of α and β subunits also appears to differ and it is thought that the proportion of α to β subunits within a heteromer may determine kinase localisation. The α subunit is often found to be cytosolic whereas the β subunit is thought to be more closely localised to the cytoskeleton. The β subunit can bind F-actin whereas the α subunit does not and the presence of a small number of β subunits is enough to localise CaMK-II to the cytoskeleton (Shen *et al.*, 1998; Bayer and Schulman, 2001; Fink *et al.*, 2003; Griffith *et al.*, 2003). The mRNA for α -CaMK-II but not β -CaMK-II can be found at dendrites and an increase in mRNA and *de novo* synthesis of this subunit has been reported in response to synaptic plasticity (Ouyang *et al.*, 1997; Ouyang *et al.*, 1999).

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The proportion of α to β subunits is known to change in response to stimulation. The α -subunit increases in expression during increased activity and the β -subunit increases in expression during decreased activity. Over-expression of α -CaMK-II or β -CaMK-II can have opposite effects on synaptic strength. Overexpression of α -CaMK-II leads to an increase in miniature excitatory postsynaptic current (mEPSC) amplitudes and a lengthening of decay times and over-expression of the β -subunit has no effect on amplitude whilst decay time is reduced. Overall, increased expression of α -CaMK-II increased charge transfer of mEPSCs whilst over-expression of β -CaMK-II decreased charge transfer (Thiagarajan et al., 2002). Despite similar substrate specificity (Miller and Kennedy, 1985), α and β isoforms in heteromers may alter cellular localisation and translocation rates (Griffith et al., 2003; Colbran, 2004b) and so differences in the relative access to specific substrates may mediate this differential effect on EPSC synaptic strength. Therefore the ratio of α to β subunits within a cell can play an important role in synaptic transmission and plasticity. In addition, the β subunit appears to be involved in morphological and structural changes associated with synaptogenesis in line with its close association with the cytoskeleton, whereas the α subunit has no morphogenic activity (Fink et al., 2003).

1.5.2 Translocation of CaMK-II to the PSD.

CaMK-II is capable of rapid translocation in response to a stimulus in time scales of seconds / minutes (Gleason *et al.*, 2003). Expression of GFP-tagged α -CaMK-II and β -CaMK-II has revealed that these subunits differ in terms of their translocation in response to a stimulus. NMDA receptor stimulation can cause reversible CaMK-II translocation from an F-actin bound to a postsynaptic density (PSD) bound state. The time taken for translocation to occur is controlled by the ratio of α : β subunits expressed. F-actin dissociation, PSD translocation and localisation at the PSD can be controlled by Ca²⁺/CaM binding and autophosphorylation (Shen and Meyer, 1999). Early reports estimated that CaMK-II constituted 20-50 % of the total protein isolated in PSDs. Subsequently it has become clear that in some cases there can be a post-mortem accumulation of CaMK-II at the PSD (Suzuki *et al.*, 1994; Lengyel *et al.*, 2001). In addition, in certain pathological conditions such as during ischemic episodes CaMK-II can translocate and form clusters of CaMK-II within the cytoplasm (Dosemeci *et al.*, 2000; Tao-Cheng *et al.*, 2002).

Association of CaMK-II with the PSD can be dynamically regulated in an activity-dependent way (Griffith *et al.*, 2003; Schulman, 2004; Colbran, 2004b). The α subunit of CaMK-II appears to be cytosolic but can co-localise with the PSD in puncta within 20 s of glutamate addition (Shen and Meyer, 1999). The β subunit can associate with F-actin under basal conditions, but after stimulation slowly dissociates and accumulates at the PSD (Shen and Meyer, 1999). Autophosphorylation at Thr²⁸⁶ stabilises the association of CaMK-II with the PSD and PP1 activity encourages dissociation (Strack *et al.*, 1997; Shen and Meyer, 1999; Shen *et al.*, 2000). Autophosphorylation at Thr³⁰⁵ and/or Thr³⁰⁶ also modulates the interaction of CaMK-II with the PSD. It appears that phosphorylation of these sites leads to CaMK-II dissociation from the PSD (Shen *et al.*, 2000; Elgersma *et al.*, 2002; Weeber *et al.*, 2003).

1.5.3 CaMK-II targeting and/or anchoring proteins.

The importance of kinase anchoring proteins in localising a kinase to its substrate and controlling phosphorylation-dependent signalling has been revealed in recent years (Pawson and Scott, 1997). As a result there has been a great deal of interest in identifying potential CaMK-II anchoring proteins. A number of proteins located at the synapse have been found to bind CaMK-II and may play a role in localistion of the kinase. In some cases these interactions can alter the function of the kinase itself (Bayer and Schulman, 2001; Colbran, 2004b).

The NMDA receptor is capable of binding to CaMK-II (See Fig 1.4). This association is thought to localise CaMK-II to the site of Ca^{2+} entry. In addition, the association of CaMK-II with the NR2B subunit may alter the function of the kinase itself (See Fig 1.4) acting to disrupt the AID and allow Ca^{2+} -independent activity (Strack *et al.*, 2000a; Bayer *et al.*, 2001).

As well as a known association between F-actin and β -CaMK-II (Shen *et al.*, 1998) both α and β CaMK-II can interact with α -actinin at a region within the catalytic domain (Walikonis *et al.*, 2001; Dhavan *et al.*, 2002). CaMK-II also interacts with another synaptic protein, densin-180, which can also interact with α -actinin, suggesting a possible ternary complex at the synapse (Walikonis *et al.*, 2001). α -actinin can bind a number of receptors and ion channels and so this association may act to localise CaMK-II and target its activities to specific sites. Densin-180 is capable of directing CaMK-II localisation within HEK293 cells and α -CaMK-II and densin-180 can be co-immunoprecipitated from brain extracts (Strack *et al.*, 2000b; Walikonis *et al.*, 2001). As densin-180 can interact with other synaptic proteins it is also a strong candidate for a protein that can determine CaMK-II localisation and therefore function (Colbran, 2004b).

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Another identified CaMK-II binding protein is a membrane associated guanylate kinase (MAGUK) found in Drosophila called Camguk. Camguk is similar to the mammalian CASK protein which is part of a subfamily of MAGUKs which are defined by an N-terminal CaMK-II like domain. In mammalian neurones, CASK has been shown to be localised at synapses both pre- and postsynaptically (Hata et al., 1996). In Drosophila, Camguk, associates with CaMK-II in the absence of Ca²⁺/CaM and Thr²⁸⁶ autophosphorylation and promotes phosphorylation of Thr^{305/6} and subsequent inactivation of the kinase. In the presence of Ca²⁺/CaM, CaMK-II associated with camguk can autophosphorylate at Thr²⁸⁶ and became constitutivelyactive. Camguk in the presence of Ca^{2+}/CaM provides a local source of active CaMK-II but when synaptic activity is low and levels of Ca^{2+}/CaM are low, Camguk promotes autophosphorylation at inhibitory sites that then requires phosphatase activity to dephosphorylate these inhibitory phosphorylation sites and allow subsequent activation of the kinase (Griffith et al., 2003; Lu et al., 2003). This kind of interaction may provide a mechanism for the local control of CaMK-II activity at active and inactive synapses.

1.5.4 CaMK-II and synaptic plasticity.

The most well studied role for CaMK-II at the synapse is in the induction of long-term potentiation (LTP) of AMPA receptor-mediated synaptic currents. This is a process by which a repetitive stimuli at a synapse leads to an increase in the strength of AMPA receptor mediated transmission at that synapse for hours after the initial stimulus has dissipated (Lisman *et al.*, 2002; Rongo, 2002; Collingridge *et al.*, 2004). CaMK-II activity is now known to be crucial for this process to occur. For a number of years the ability of CaMK-II to maintain activity after a Ca²⁺ stimulus

through autophosphorylation at Thr²⁸⁶ led to the hypothesis that CaMK-II was a "memory molecule" capable of maintaining long-term changes in synaptic function. CaMK-II was found to phosphorylate the AMPA receptor directly at Ser⁸³¹ on the GluR1 subunit (Barria *et al.*, 1997; Derkach *et al.*, 1999; Lee *et al.*, 2000). This phosphorylation led to an up-regulation of receptor function through a change in the properties of the channel, by increasing the single channel conductance (Benke *et al.*, 1998; Derkach *et al.*, 1999).

Recently it has been demonstrated that CaMK-II dependent changes associated with LTP can also occur through another separate mechanism. CaMK-II dependent activation of a Ras/Rap GTPase pathway (Zhu *et al.*, 2002) and a subsequent insertion of receptors (Lledo *et al.*, 1998) into the synapse is thought to up-regulate AMPA receptor function during LTP. This process does not require direct phosphorylation of the AMPA receptor (Hayashi *et al.*, 2000). However, receptor insertion is thought to be subunit dependent as GluR1/4 subunits are thought to be inserted into the membrane during LTP and this process is dependent on the GluR1 subunit, whereas the maintenance of receptor number through recycling is thought to involve GluR2/3 subunits (Shi *et al.*, 2001; Malinow and Malenka, 2002; Bredt and Nicoll, 2003; Ju *et al.*, 2004). In this way receptor number is increased and then maintained (See Fig 1.5).

1.5.5 CaMK-II, LTP and behaviour.

It is clear that CaMK-II is crucial for certain forms of LTP and that disruption of CaMK-II and subsequently LTP prevents proper memory formation and alters animal behaviour (Lisman *et al.*, 2002; Elgersma *et al.*, 2004; Ehrlich and Malinow, 2004). Mice devoid of α -CaMK-II have been generated which revealed the importance of this subunit for the generation of NMDA receptor-dependent LTP at the CA1 pyramidal neurone in the hippocampus. In addition these mice exhibited specific spatial learning impairments strengthening the view that LTP forms the cellular basis of spatial learning (Silva *et al.*, 1992a; Silva *et al.*, 1992b). In T286A α -CaMK-II mutant mice, in which the α subunit cannot undergo autophosphorylation and therefore display Ca²⁺-independent activity, CA1-LTP is completely absent (Giese *et al.*, 1998). These mice also displayed deficits in spatial learning. This illustrated the importance of autophosphorylation for LTP and the subsequent effects on physiology and behaviour.

Within mice which contain the α -CaMK-II T305D mutation (mimicking inhibitory autophosphorylation), CaMK-II activation and autophosphorylation at Thr²⁸⁶ is blocked as is subsequent translocation to the PSD (Hanson and Schulman, 1992; Elgersma et al., 2002). In this case a stimulus which has no effect in wild-type mice can induce LTD. Mice which express α -CaMK-II which cannot be phosphorylated at Thr^{305,306}; show a marked LTP in response to the same stimulus demonstrating that these sites are important in the regulation of CaMK-II activity and can control the threshold for LTP induction (Elgersma et al., 2002). Alteration in LTP seems to correlate with cognitive impairment. For instance T305D and T286D mice, which lack any LTP at CA1 synapses and show no evidence of β -CaMK-II compensation, have severely impaired spatial learning (Elgersma et al., 2002). A reduction in the level of CaMK-II at the PSD, and in CaMK-II activity as well as impairment of LTP and deficits in context-dependent learning, have been observed within a mouse model of Angelman syndrome (Weeber et al., 2003). The human form of this disease is characterised by severe mental retardation and epilepsy. This forms a clear link between alterations in the normal function of CaMK-II, changes in hippocampal LTP and subsequent alterations in memory learning and behaviour (Elgersma *et al.*, 2004).

1.6 CaMK-II and GABA_A receptor function.

1.6.1 Ca²⁺-dependent depression of GABA_A receptor function.

There have been a number of studies that have demonstrated Ca^{2+} dependent modulation of GABA_A receptor function. Increases in intracellular calcium levels have been reported to depress GABA currents in sensory neurones of the bullfrog. In this case, Ca^{2+} influx led to a rapid decrease in the affinity of the GABA_A receptor to GABA (Inoue *et al.*, 1986). In CA1 hippocampal pyramidal cells Ca^{2+} entry through NMDA receptors was reported to depress the amplitude of whole-cell currents. This effect was prevented by the addition of a specific protein phosphatase 2B (PP2B) / Calcineurin (CaN) inhibitor (Stelzer and Shi, 1994). Ca^{2+} -dependent phosphatases have been reported to depress GABA_A receptor currents in a number of different celltypes (Akopian *et al.*, 1998; Huang and Dillon, 1998; Mozrzymas and Cherubini, 1998; Wang *et al.*, 2003a; Sanchez *et al.*, 2005). Calcineurin has also been reported to alter desensitisation, acting to alter the decay phase of IPSCs (Martina *et al.*, 1996; Jones and Westbrook, 1997; Mozrzymas and Cherubini, 1998; Sanchez *et al.*, 2005) as well as playing a pivotal role in the process of long-term depression (LTD) of CA1 inhibitory synapses in the hippocampus (Wang *et al.*, 2003a).

1.6.2 Bidirectional plasticity.

At excitatory synapses long-term plasticity is thought to be mediated through phosphorylation (Soderling and Derkach, 2000; Lee *et al.*, 2000; Lisman *et al.*, 2002; Collingridge *et al.*, 2004). The protein phosphatase, CaN, and the protein kinase,

CaMK-II, may act in opposition to each other in response to Ca^{2+} signals that vary in magnitude (Yang *et al.*, 1999; Cormier *et al.*, 2001; Winder and Sweatt, 2001; Collingridge *et al.*, 2004; Xia and Storm, 2005). It is possible that a similar situation exists at inhibitory synapses. Indeed it has been demonstrated that long-term plasticity at inhibitory synapses can be bi-directional (potentiation or depression) and that this is also mediated by Ca^{2+} signals of different magnitudes (McLean *et al.*, 1996; Aizenman *et al.*, 1998).

1.6.3 CaMK-II mediated potentiation of GABA_A receptor function.

As well as depression in response to Ca^{2+} influx there are also a number of studies which have identified up-regulation of GABA_A receptor function which is dependent on Ca^{2+} (See Table 1.2). For instance, Aguayo *et al.* (1998) demonstrated that increases in intracellular Ca^{2+} in mouse cortical neurones could induce a transient increase in the amplitude of whole-cell GABA_A receptor-mediated currents. The addition of CaM and CaMK-II specific inhibitors blocked this potentiating effect suggesting that CaMK-II was a necessary part of the signal cascade.

There are several studies which have reported a CaMK-II dependent modulation of GABA_A receptor function. Wang *et al.* (1995) intracellularly-applied pre-activated (autophosphorylated at Thr²⁸⁶) α -CaMK-II to acutely isolated rat spinal dorsal horn neurones and observed a significant increase in the peak amplitude of whole-cell currents induced by a fixed dose of GABA. They found a similar effect when recording IPSCs from CA1 neurones in a hippocampal slice preparation. As well as an increase in amplitude they also reported a slight change in desensitisation acting to prolong the decay of the currents they recorded. A Ca²⁺ dependent increase in the decay of IPSCs has also been reported in response to amputation of the dendrites of dentate gyrus granule cells in central neurones (Soltesz and Mody, 1995).

Churn *et al.* (1998) examined how CaMK-II activity towards endogenous substrates in a synaptosomal membrane fraction altered GABA_A receptor agonist binding. Activation of endogenous CaMK-II led to an increase in muscimol binding, without altering binding affinity. The results suggested that the increase in binding was due to an increase in the number of functional GABA_A receptors (Churn and DeLorenzo, 1998). In an additional study they used a similar protocol to demonstrate that CaMK-II activity increased allosteric-modulator binding (Churn *et al.*, 2002). Again this was thought to be due to an increase in the apparent number of receptors available for binding. In this study, CaMK-II was reported to phosphorylate the α l subunit (Churn *et al.*, 2002).

Alix *et al.* (2002) characterised a Ca^{2+} sensitive GABA mediated chloride (Cl⁻) current in cockroach dorsal unpaired median neurones. This current could be potentiated by increases in intracellular Ca^{2+} and the potentiation could be blocked by addition of specific CaMK-II and CaM inhibitors (KN-62 and W7 respectively). Insect GABA-stimulated Cl⁻ channels are pharmacologically and structurally distinct from vertebrate GABA_A receptors (Alix *et al.*, 2002), but it appears they may be regulated by phosphorylation in a similar way to the vertebrate GABA_A receptors.

During the development of the hippocampus BDNF has been reported to have a potentiating action on GABA currents recorded from CA1 pyramidal neurones isolated from P6 rats. This was prevented by the addition of the CaMK-II inhibitor KN-62 (Mizoguchi *et al.*, 2003). Potentiation of GABA currents has also been reported after mechanical injury of cortical neurones. This potentiation took the form of an increase in peak current density (pA/pF) that could be blocked by the CaMK-II inhibitor KN-93 (Kao *et al.*, 2004).

It appears possible that $GABA_A$ receptors can be modulated by Ca^{2+} dependent phosphorylation processes and that in some cases Ca^{2+} -dependent activation of CaMK-II results in a potentiation of $GABA_A$ receptor mediated currents. CaMK-II dependent phosphorylation is critical for excitatory synaptic plasticity, like long-term potentiation (LTP). There is also evidence that CaMK-II may also play a role in inhibitory synaptic plasticity.

Table	1.2:
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Cell-Type	Method	Response	Reference
Rat dorsal hom neurones Hippocampal CA1 pyramidal	i/c pre-activated α-CaMK-II	 ↑ Peak w-c amplitude ↓Desensitisation ↑ IPSC amplitude 	Wang <i>et al</i> ., (1995)
Mouse cortical neurones.	† i/c Ca²+	t Peak w-c amplitude Blocked by KN-93	Aguayo <i>et al.</i> , (1998)
Rat forebrain synaptosomal membrane fractions.	i/c Ca²⁺/CaM i/c α-CaMK-II	 Agonist binding Allosteric modulator binding α1 subunit phosphorylation 	Churn and DeLorenzo (1998) Churn <i>et al</i> ., (2002)
Cerebellar purkinje neurones	ti/c Ca ²⁺ via VGCCs i/c pre-activated α-CaMK-II	↑ Peak w-c Amplitude ↑ sIPSCs amplitude No change in decay. (Rebound Potentiation)	Kano <i>et al.</i> , (1996)
Hippocampal CA1 pyramidal neurones	BDNF application	↑ Peak w-c Amplitude (P6 not P14) Blocked by BAPTA and KN-62	Mizoguchi <i>et al.,</i> (2003)
Cockroach dorsal unpaired median neurones	t i/c Ca ²⁺ via depolarisation	t w-c amplitude Biocked by calmodulin blocker (W7) and KN-62	Alix <i>et al.</i> , (2002)
CA1 hippocampal neurones	† Ca ²⁺ /CaM	t IPSC amplitude. Blocked by CaMK autoinhibitory peptide.	Wei <i>et al.,</i> (2004)

Table 1.2: A summary of the published evidence for CaMK-II modulation of the GABA_A receptor.

In a number of neuronal preparations increases in Ca^{2+}/CaM and activation of CaMK-II or inclusion of exogenous pre-activated CaMK-II in the patch pipette has been demonstrated to up-regulate GABA_A receptor function.

i/c = intracellular; w-c = whole-cell GABA current.

1.7 Inhibitory Synaptic Plasticity

A number of Ca^{2+} dependent forms of inhibitory synaptic plasticity have been reported (Komatsu and Iwakiri, 1993; Komatsu, 1994; McLean *et al.*, 1996; Caillard *et al.*, 1999; Gaiarsa *et al.*, 2002; Patenaude *et al.*, 2003; Bauer and LeDoux, 2004). The mechanisms behind plasticity vary with some reports of a presynaptic effect on frequency (Caillard *et al.*, 1999; Ouardouz and Sastry, 2000) and others suggesting a purely postsynaptic effect mediated by an up-regulation of GABA_A receptor function (Patenaude *et al.*, 2003; Bauer and LeDoux, 2004).

1.7.1 Activity and development dependent plasticity.

One of the first forms of inhibitory plasticity to be identified occurs within the visual cortex. High frequency stimulation could induce a long-term increase in the amplitude of IPSPs recorded. This form of LTP of inhibitory synapses shared many of the features of excitatory LTP recorded from CA1 hippocampal neurones (Komatsu and Iwakiri, 1993; Komatsu, 1994). One of the interesting features of this form of plasticity was that LTP was more easily inducible in developing rats rather than mature. A similar form of activity-dependent plasticity of IPSCs mediated by GABA_A receptors has been reported in rat hippocampal slices. Stimulation induced a long-lasting change in both the frequency and amplitude of IPSCs recorded, however, this plasticity could only be induced within a restricted period of development (Gubellini *et al.*, 2001).

Some of these reported forms of inhibitory plasticity are bi-directional (McLean *et al.*, 1996; Aizenman *et al.*, 1998) and rely on differences in magnitude of Ca^{2+} signals in a similar way to the plasticity of excitatory synapses (Yang *et al.*, 1999; Cormier *et al.*, 2001). In the same way that CaMK-II and CaN can bi-

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directionally regulate excitatory synaptic transmission it has been proposed that CaMK-II may potentiate inhibitory synapses and CaN may depress them (Morishita and Sastry, 1996; Wang *et al.*, 2003a; Sanchez *et al.*, 2005). Aizenman *et al.* (1998) found that at GABAergic synapses onto deep cerebellar nuclei neurones, a long-term potentiation of IPSC amplitudes could be induced by a strong stimulation and large Ca^{2+} influx into the postsynaptic cell whereas a more moderate stimulation and Ca^{2+} influx resulted in a long-term depression of the IPSC amplitudes.

In neurones of the deep cerebellar nuclei (DCN) long-term potentiation (LTP) of inhibitory GABA_A receptor mediated currents can be evoked by high frequency stimulation of the white matter surrounding the DCN. This potentiation is dependent on NMDA receptor activation and Ca^{2+} influx and is blocked by an NMDA receptor antagonist (APV) and the Ca^{2+} chelator BAPTA, suggesting that Ca^{2+} influx via NMDA receptors stimulated the potentiation. Intracellular application of tetanus toxin, a blocker of exocytosis, also prevented the induction of LTP of inhibitory currents. Suggesting a role for exocytosis and receptor trafficking in this form of Ca^{2+} stimulated inhibitory plasticity (Ouardouz and Sastry, 2000).

Another form of long-term inhibitory plasticity has been identified at inhibitory synapses onto CA1 hippocampal pyramidal neurones (Wei *et al.*, 2004). In this case a Ca^{2+}/CaM dependent increase in the amplitude of IPSCs was prevented by postsynaptic dialysis of an auto-inhibitory peptide of CaM-dependent kinases, suggesting the enhancement observed was CaMK-II dependent. In addition, inhibition of tubulin or actin polymerization attenuated this effect implying an intact cytoskeleton was required for this to occur.

Modelling and functional studies of GABAergic synapses suggests that if GABA is saturating at a given synapse, the most effective mechanism to potentiate GABA mediated currents would be to increase receptor number at the synapse, as alterations in transmitter concentration and the probability of channel opening would have no impact on the amplitude of the inhibitory currents (Otis *et al.*, 1994; Nusser *et al.*, 1998). In this regard, it is now thought AMPA receptor insertion may be the predominant mechanism operating during excitatory LTP (Hayashi *et al.*, 2000; Zhu *et al.*, 2002).

Experimental models of epilepsy in which chronic epilepsy is induced through a process known as kindling have revealed inhibitory synaptic plasticity in response to the induction of epilepsy. This increase in inhibitory synaptic transmission is thought to be associated with an increase in the number of GABA_A receptors at inhibitory synapses (Otis *et al.*, 1994; Nusser *et al.*, 1998). These kinds of alterations in inhibitory synaptic transmission have been suggested to function as homeostatic plasticity (Turrigiano and Nelson, 2004). Blockade of activity in cortical neuronal cultures by TTX leads to a decrease in the amplitude of IPSCs recorded. This is also accompanied by a reduction in the number of channels at the cell surface (Kilman *et al.*, 2002). A similar treatment caused an increase in the amplitude of mEPSCs, suggesting that the balance between excitatory and inhibitory synaptic currents could be dynamically modulated (Turrigiano *et al.*, 1998).

1.7.2 Rebound potentiation

Kano *et al.* (1992; 1996) characterised a phenomenon known as rebound potentiation in which stimulation of climbing fibre synapses onto cerebellar Purkinje cells could produce a long term potentiation of GABA_A receptor mediated currents (in the same cell). The Ca²⁺ chelator BAPTA can block this rebound potentiation suggesting the process is Ca²⁺ dependent. Calmodulin binding domain (CBD) synthetic peptide inhibits binding of Ca^{2+}/CaM to CaMK-II and blocks the activation of the kinase. Its introduction into Purkinje cells blocked the potentiation of GABAmediated currents. The CaM antagonist, calmidazolium, and the selective CaMK-II antagonist, KN-62, had a similar effect. Moreover, the intracellular introduction of purified pre-activated CaMK-II increased the amplitude of whole-cell GABA currents recorded. Addition of CaMK-II inhibitors five minutes after the initial stimulus (Ca²⁺ rise) failed to block the effect, suggesting that CaMK-II activation is involved in the induction not the maintenance of rebound potentiation.

Further investigation has elucidated some of the molecular pathways involved in rebound potentiation. It has been demonstrated that multiple kinase pathways interact to modulate GABA_A receptors in Purkinje cells. Kawaguchi and Hirano (2002) confirmed that inhibition of CaMK-II reduces or blocks rebound potentiation and observed that inhibition of PKA has a similar effect. Inhibition of PP1 or CaN impairs the suppression of potentiation caused by PKA inhibition, but not that caused by CaMK-II inhibition. The substrate of PKA and CaN, DARPP-32, inhibits PP1 when phosphorylated by PKA (Svenningsson *et al.*, 2004). PP1 inhibits CaMK-II, which alters GABA_A receptor function more directly by a mechanism that is yet to be determined (Fig 1.5B).

In ionotropic glutamate receptor subunit $\delta 2$ (GluR $\delta 2$) knockout mice GABAergic synaptic transmission is enhanced and it is not possible to stimulate rebound potentiation suggesting that rebound potentiation occurs *in vivo*, potentially as a form of homeostatic plasticity (Ohtsuki *et al.*, 2004). Stimulation of metabotropic P2Y purinoceptors and subsequent G-Protein-coupled stimulation of Ca²⁺ release from intracellular stores can induce a long-term enhancement of GABA_A receptor mediated currents recorded from Purkinje cells. This process

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appears similar to rebound potentiation in that it is dependent on postsynaptic calcium increases. However, the time course is much slower and does not require Ca^{2+} influx through voltage gated calcium channels indicating that different pathways may be involved (Saitow *et al.*, 2005).

There are a number of similarities between long-term plasticity of excitatory and inhibitory synaptic transmission in that both kinds of synapse can be modified bi-directionally, dependent upon the magnitude of the Ca^{2+} signal. Under certain circumstances Ca^{2+} -dependent potentiation of GABA_A receptor function has been linked to activation of CaMK-II. It appears, therefore, that CaMK-II may be capable of mediating some forms of long-term inhibitory plasticity and as a result CaMK-II dependent modulation of the GABA_A receptor may have important functional consequences for understanding network behaviour.

Fig 1.5:





Fig 1.5: Schematic representation of signalling pathways involved in excitatory and inhibitory forms of long-term synaptic plasticity.

(A) Schematic representation of the signalling pathways thought to be involved in excitatory LTP. CaMK-II is able to directly phosphorylate the AMPA receptor and can also drive AMPA receptor insertion into the plasma membrane. GluR1/4 receptors are inserted into the synapse during LTP. GluR2/3 receptors are continuously recycled to and from the plasma membrane and maintain a constant level of receptors at the surface. LTD may involve Ca²⁺-dependent phosphatase activity directly against the AMPA receptor but may also involve removal of receptors from the plasma membrane. Plasticity is bi-directional in response to Ca²⁺ signals of differing magnitude.

Adapted from Zhu et al. (2002).

(B) Schematic representation of the signalling pathways involved in rebound potentiation, a form of inhibitory long-term potentiation.

In a similar way to excitatory LTP pathways, activation of CaMK-II is thought to induce a long-term potentiation of $GABA_A$ receptor function, and activation of a protein phosphatase (CaN) is thought to counteract that process. PKA is able to induce rebound potentiation through an interaction with DARPP-32 and a subsequent downstream activation of CaMK-II through inhibition of PP1.

The process or events downstream of CaMK-II activation that result in modification of GABA_A receptor function, are at present unknown.

Adapted from Kawaguchi and Hirano (2002).

1.8 CONCLUSIONS

The process of phosphorylation is known to regulate GABA_A receptor function through direct changes to the properties of the channel and through changes in cell surface trafficking in ways that have been shown to be physiologically relevant (Moss and Smart, 1996; Brandon *et al.*, 2002a; Kittler and Moss, 2003; Lüscher and Keller, 2004). CaMK-II can phosphorylate GST-fusion proteins of the TM3-4 intracellular loop of β and γ subunits (McDonald and Moss, 1994; McDonald and Moss, 1997). CaMK-II has also been shown to potentiate GABA_A receptor mediated currents in several neuronal cell types and has been implicated in the induction of certain forms of inhibitory synaptic plasticity (Wang *et al.*, 1995; Kano *et al.*, 1996; Wei *et al.*, 2004). It is important to determine the mechanism of CaMK-II dependent modulation of GABA_A receptors as the processes involved may play important physiological roles. For instance, up-regulation of inhibitory synaptic strength may be important in forms of homeostatic plasticity and in the proper formation of inhibitory networks during development.

At present it is unclear whether CaMK-II dependent modulation of GABA_A receptors occurs through direct phosphorylation or through indirect downstream processes. In addition, given the heterogeneity of GABA_A receptors and the differential effects of other kinases on different GABA_A receptor subtypes, it is unknown whether CaMK-II shows any subunit-dependent effects on GABA_A receptors.

1.9 AIMS

- To establish whether CaMK-II dependent modulation of GABA_A receptors occurs through direct phosphorylation of the receptor.
- To determine if CaMK-II dependent modulation of the GABA_A receptor has differential effects on different GABA_A receptor subtypes.
- To establish which specific subunits are required for CaMK-II modulation, and if possible, which specific residues might be phosphorylated.
- To understand more about the possible mechanism(s) of CaMK-II dependent functional modulation of the GABA_A receptor.

<u>Chapter Two:</u> <u>Materials and Methods</u>

2.1 MOLECULAR BIOLOGY

All molecular biology was carried out in collaboration with Dr A.M Hosie and Ms H.M.A da Silva. In brief the methods were as follows:

2.1.1 Site-directed mutageneis.

Murine $\alpha 1$, $\beta 2/3$, $\gamma 2S$ GABA_A receptor cDNAs were cloned into the plasmid vector pRK5 (Moss *et al.*, 1991; Connolly *et al.*, 1996). Expression of heterologous subunits in this vector is under the control of the cytomegalovirus promoter. Site-directed mutagenesis was carried out using the Quikchange method (Stratagene). This involved:

- The design of appropriate oligonucleotides containing the desired mutation flanked by an unmodified nucleotide sequence; to be used as mutagenic primers.
- The amplification of the relevant product by polymerase <u>chain reaction</u> (PCR); electrophoresis to check for amplification of the desired product and digestion of unmutated DNA.
- Transformation into *E.Coli*.
- Purification and sequencing of plasmid DNA.

Complementary olignucleotides containing the desired mutation were used as primers for PCR (See Appendix A1.1). The PCR reaction contained 5 μ l of 10X reaction buffer, 10-100 ng of dsDNA template (cloned into pRK5), 125 ng primer 1 (Invitrogen) and 125 ng of primer 2 (Invitrogen), 1 μ l of deoxynucleotide triphosphate (dNTP) mix (10 mM of each dNTP), 2.5U of *Pfu turbo* DNA polymerase and double distilled water to a final volume of 50 μ l.

The PCR reaction mix underwent thermocycling with the following parameters: 99°C for 30 s, followed by 18 cycles of 95°C for 30 s (denaturation), 55 to 65°C for 30 s to 60 s (annealing), and 72°C for 2 min/kb of plasmid length (extension). The PCR mix was then electrophoresed on a 1 % w/v agarose gel, to check for the formation of a product. The PCR product was then digested with *Dpn I* restriction enzyme (10U-New England Biolabs) to cut the methylated template DNA, leaving only mutated PCR product DNA (1 hr at 37°C).

After digestion, the PCR product was transformed into *E-Coli* using TOP10 one shot Kit (Invitrogen – See Appendix A1.2). Top10 competent cells (Invitrogen) were grown on LB Agar (Sigma) containing 50 μ g/ml ampicillin. Bacterial colonies containing the mutated DNA were selected and grown overnight in Lennox L broth base (LB) medium (Sigma) to allow the transformed bacteria to multiply. The plasmid was then purified using the QIAprep Spin Miniprep Kit (Qiagen – See Appendix A1.3).

The sequence of the entire coding region of the cDNA was then determined to ensure the mutation had been correctly incorporated into the template DNA sequence without any ectopic mutations (DNA sequencing service – The Wolfson Institute for Biomedical Research, Cruciform Building, UCL). The analysis of the base sequence was performed by Sequencher 3.1.2 software. If the mutation had been successfully incorporated into the base sequence the bacterial stock was grown to increase the amount of the mutated cDNA using a large volume of LB media (100 ml midi prep; 250 ml, maxi prep Qiagen; grown overnight). As before, cDNA was

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purified but this time at higher concentrations $(1 \ \mu g/\mu l)$ that could then be used for transfection of cell lines or neurones.

2.2 CELL CULTURE

2.2.1 HEK293 cells.

HEK293 cells (Graham *et al.*, 1977) were grown in 10 cm diameter culture dishes (Nunc) in media containing <u>D</u>ulbecco's <u>m</u>odified <u>E</u>agle's <u>m</u>edium (DMEM -Gibco) supplemented with 10 % v/v foetal calf serum (FCS), 2 mM glutamine, 100 U/ml penicillin G, 100 μ g/ml streptomycin and incubated at 37°C, in 95 % air / 5 % CO₂. Cells were passaged when they reached approximately 70 % confluency, usually every 2-3 days. Cells requiring passaging were washed once with 5 ml <u>H</u>ank's <u>b</u>alanced <u>s</u>alt <u>s</u>olution (HBSS) and then incubated in 2 ml trypsin (0.25 % w/v) until the cells floated free off the dish. DMEM (10 ml) was then added to quench the trypsin and the cells were transferred to a 50 ml conical tube for centrifugation at 1000 rpm for 2 min. The supernatant was then removed and cells were re-suspended in 5 ml of media and re-plated at a dilution factor of 1:10 / 1:20 cell-containing media to fresh media (Krishek *et al.*, 1994; Dunne *et al.*, 1998; Thomas and Smart, 2002; Thomas and Smart, 2005). Cells used for transfection by electroporation or calcium phosphate were subsequently triturated to ensure a singlecell suspension and were re-suspended in OPTIMEM (Gibco) at this point.

2.2.2 NG108-15 cells.

NG108-15 cells (Klee and Nirenberg, 1974; Daniels and Hamprecht, 1974; Docherty *et al.*, 1991) were grown in 50 ml culture flasks (Nunc) in DMEM (4.5 g/L glucose - Gibco) supplemented with 10 % v/v FCS, 2 mM glutamine, 100 μ M

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hypoxanthine, 0.4 μ M aminopterin, 16 μ M thymidine (Sigma) and 20 U/ml penicillin G and 20 mg/ml streptomycin and incubated at 37°C, 90 % air /10 % CO₂. Cells were passaged when they reached approximately 70 % confluency every two-three days at a ratio of 1:3 or 1:6 cell-containing media to fresh media. Cells were detached from the bottom of the flask by a gentle wash and spun at 900 rpm for 5 min; they were then re-suspended in fresh media and re-plated to continue the culture. At this point cells for transfection were further diluted (1:2-3) and drops of cell-containing media were dropped onto pre-prepared (2.2.3) glass coverslips in 35 mm dishes to allow the cells to adhere (Nunc, See Fig 2.1A).

2.2.3 Primary cell culture – rat/mouse cerebellar granule cells.

The cerebella of postnatal day 1 (P1) Sprague-Dawley rats were removed after decapitation and placed in ice-cold dissection media (DM = HBSS – without Ca^{2+}/Mg^{2+} (Gibco) + Gentomycin 10 µg/ml (Sigma). The meninges were then carefully removed and the cerebella transferred to DM that had been heated to 37°C and CO₂ saturated in the incubator. They were then centrifuged at 1000 rpm for 1.5 min; the supernatant was removed and another 10 ml DM added and the tissue centrifuged again. The supernatant was removed and the cerebella were then incubated in trypsin (0.1 % w/v, Sigma) at 37°C for 10 min. The trypsin was removed and the cerebella washed with 10 ml DM. They were centrifuged again for 3 min at 1000 rpm followed by a second wash in DM and centrifuged at 1000 rpm for 1.5 min.

The DM was removed virtually completely using a 200 μ l Gilson pipette. Then, 1 ml DNase (0.05 % w/v in 12 mM MgSO₄, Sigma) was added and the cerebella were triturated with two glass pipettes that had been fire-polished to small
bore diameters. The final trituration step was carried out with a 200 μ l pipette tip until a single-cell suspension had formed. Another 9 ml DM was added and the mixture centrifuged at 1200 rpm for 5 min. Care was taken to remove all the supernatant at this point with a 200 μ l pipette and the cells were re-suspended in 1 ml of seeding media (<u>Basal Medium Eagle</u> – BME (Gibco) + 10 % v/v FCS). An aliquot (15 μ l) of the suspension were then transferred to a haemocytometer and an estimate of cell number made. Cells were diluted to give approximately 5 x 10⁶ cells/ml; 90 μ l of this suspension was then placed onto the pre-prepared glass coverslips coated in poly-L-ornithine (Sigma). Coverslips were washed in 100 % v/v ethanol and flamed, twice and coated in 500 μ g/ml poly-L-ornithine in borate buffer (50 mM Boric acid, 12.5 mM sodium tetraborate), 0.5 ml per coverslip, overnight at 4°C, then washed three times with sterile water and finally left to air dry before use. Dishes were left in the incubator for 3 hrs to allow cells to adhere to the glass after which the appropriate growth media was added.

[1] Serum free media – 20 mM KCl.

BME (Gibco) supplemented with 0.5 % w/v glucose, 5 mg/L insulin, 5 mg/L transferrin, 5 mg/L selenium (Sigma), 20 U/ml pencillin-G and 20 μ g/ml streptomycin, 0.14 mM glutamine, 1.4 mM NaCl, 0.01 mg/ml <u>B</u>ovine <u>Serum Albumin (BSA, Sigma) and 20 mM KCl (See Fig 2.1B).</u>

[2] Serum-containing media – 5 mM KCl.

BME (containing 5mM KCl) supplemented with 0.5 % w/v glucose, 20 U/ml penicillin-G, 20 μ g/ml streptomycin, 0.2 mM glutamine, 5 % v/v FCS, 1.2 mM NaCl and 5 mg/ml insulin, 5 mg/ml transferrin and 5 mg/ml selenium (Sigma, See Fig 2.1B).

Cells maintained in serum-containing media were treated with 1-2 μ M cytosine arabinoside (Sigma) at 4-5 DIV to control the growth of non-neuronal cells, which otherwise threatened to overwhelm the culture. Serum-free media maintained cells had 1 ml of media replaced every 7-10 days. Cells for transfection were chosen and transfected at 3-4 DIV (Furuya *et al.*, 1998; Leao *et al.*, 2000; Tabata *et al.*, 2000; Ives *et al.*, 2002).

2.2.4 Embryonic cortical neuronal culture.

The cortical neuronal cultures (See Chapter 4, Fig 4.1) were prepared by Dr J.N Jovanovic and Ms K McAinsh as follows:

Cortices were dissected from embryonic day 19 rats and the tissue incubated in 0.25 % w/v trypsin in HBSS for 15 min followed by 3 x 5 min washes in HBSS. The tissue was then dissociated by titration with a fire-polished pipette. Cells were plated on 0.5 mg/ml poly-L-ornithine-coated glass coverslips at a density of 10^5 cells/cm² and incubated at 37°C, 95 % air / 5 % CO₂ for 6 days before use (Brandon *et al.*, 1999).

Fig 2.1:





Fig 2.1: Images of NG108-15 cells and cerebellar granule cells in culture.

(A) NG108-15 cells transfected with cDNAs for $\alpha 1\beta 3\gamma 2S$ GABA_A receptor subunits and EGFP (Enhanced Green Florescent Protein). Cells were then stained with a $\beta 3$ specific antibody (red fluorescence - See table 2.1) 48 hrs after transfection.

(B) Differential interference contrast (DIC) images of cerebellar granule cells maintained in culture in 20 mM KCl and 5 mM KCl. Cells in 5 mM KCl tended to form clusters whereas cells in 20 mM KCl had a more uniform distribution.

Images acquired on a Zeiss laser scanning confocal microscope (LSM 510 Meta; See 2.6.2).

2.3 TRANSFECTION

2.3.1 Electroporation – HEK293 cells.

For the production of a large number of dishes expressing the same receptor subunits the following method was used (Wooltorton *et al.*, 1997; Dunne *et al.*, 1998). A 10 cm dish of HEK293 cells at 70 % confluency was used for each transfection. Cells were dissociated from the dish with trypsin (See 2.2.1) and resuspended in 25 ml OPTIMEM in a 50 ml conical tube and centrifuged for 2 min at 1000 rpm. The supernatant was removed and the cells re-suspended in 0.5 ml OPTIMEM for electroporation and 0.3 ml for continuation of the culture. The cells were triturated with a flame-polished pipette to form a single-cell suspension. Electroporation cuvettes were washed twice with sterile water and twice with OPTIMEM before use. The appropriate cDNA suspension was then placed in the bottom of the cuvette, e.g, $\alpha 1:\beta 3:\gamma 2S:EGFP$ at a ratio of 1:1:1:1 (Stock solutions of DNA at 1 $\mu g/\mu l$, 1 μl of each cDNA was used per dish).

Then, 0.5 ml of the cell suspension was added to the cDNA in the cuvette and mixed gently. The cuvette contents were mixed before, in-between and after electroporation with the Biorad Gene pulser II. Settings were "infinite resistance", voltage 0.4 kV and capacitance 0.125 mF. The time constant for the electric field used for electroporation generally fell between 3-4 ms.

Cells were allowed to stand for 10 min after electroporation. Then, 8-10 ml of media (See 2.2.1) was then placed in a 10 ml tube and with a glass pipette the contents of the cuvette were added and the suspension inverted to mix well. After checking the cell density, 0.5 ml of media was placed onto pre-prepared glass coverslips in 35 mm dishes (Nunc) and a suitable number of drops of the cell suspension were added to this. Coverslips were prepared by washing in 100 % v/v

ethanol and flaming, then adding 0.5 ml of poly-L-lysine (100 μ g/ml) for 1 hour, followed by three washes with sterile water and then left to dry before use.

Coverslips with cells added were left in the incubator for 10 min at 37°C to allow the cells to adhere to the substratum. Then 1-2 ml of media (DMEM) was added to each dish and they were placed into the incubator $(37^{\circ}C / 95 \% \text{ air} / 5 \% CO_2)$ for at least 24 hrs. These cells could be recorded from for 24-48 hrs after transfection.

2.3.2 Calcium phosphate – HEK293 and NG108-15 cells.

This procedure was used when a number of different cDNAs were to be transfected into different dishes for recording on the same day (Krishek *et al.*, 1994). To 1 μ l of cDNA stock solution (1000 ng/ μ l) per dish per subunit (+ EGFP), 340 mM CaCl₂ (20 μ l per dish to be transfected) was added. This was left for 20 min to allow a precipitate to form. This suspension was then mixed with double-strength HBSS (24 μ l per dish to be transfected; 2 x HBSS = 280 mM NaCl , 2.8 mM Na₂HPO₄, 50 mM HEPES, NaOH to pH 7.2). Once mixed, 45 μ l was then applied to each 35 mm dish of cells. Dishes of cells were prepared by dissociating cells from the stock dish they were growing in (when at 70 % confluency) and re-suspending them in their normal media (See 2.2.1 and 2.2.2). Pre-prepared coverslips in 35 mm dishes (Nunc), already coated with poly-L-lysine for HEK293 cells (See 2.3.1) and poly-L-ornithine (See 2.2.3, 1 hr incubation only) for NG108-15 cells, were coated with 0.5 ml of normal media, prior to seeding with drops of cell suspension to obtain the appropriate cell density for recording.

After seeding the cells, the coverslips were left for 10-15 min to allow the cells to adhere. After this, 1.5 ml of fresh media was applied to each dish. At this

point the cells were ready for calcium phosphate transfection. After transfection cells were left overnight and the media changed in the morning. Cells could be recorded from 24 hr after transfection (See Fig 2.1A).

2.3.3 Effectene transfection - NG108-15 cells and cerebellar granule cells.

Both NG108-15 and cerebellar granule cell cultures were transfected using the Effectene transfection reagent (Qiagen, See Fig 2.1A and Fig 7.6-7).

On the day of transfection, NG108-15 cells were seeded onto glass coverslips (See 2.3.2) and left for 10 min in the incubator for cells to adhere. Cerebellar granule cell cultures were transfected at 3-4 DIV. A total of 0.4 μ g DNA per dish (0.3 μ g for granule cells) was prepared (containing an equal ratio of each subunit and enhanced GFP, 1:1:1:1) and added to EC Buffer (DNA-condensation buffer, 75 μ l per 35 mm dish – supplied with reagent). Then, 8 μ l Enhancer (supplied with reagent) was then added per 1 μ g of total DNA. The mixture was then vortexed and allowed to stand for 5 min. Then, 10 μ l of Effectene reagent was added per dish for the transfection and the mixture vortexed and allowed to stand for a further 10 min to allow the DNA transfection-complex to form. Finally 0.5 ml of the appropriate media was added for each dish and mixed. The DNA suspension was then added immediately to the previously prepared cells. Cells were left in the transfection reagent containing media for 3-4 hrs and then the media was replaced. NG108-15 cells could be recorded from 24 hrs later (See Fig 2.1A). Cerebellar granule cells were left for a further 3-5 days and recorded from at 7-12 DIV.

2.4 ELECTROPHYSIOLOGY

2.4.1 The equipment.

The cells to be recorded from were identified under a Nikon Eclipse TE300 microscope at 200-400X magnification, with additional X5 magnification, as required sometimes for cerebellar granule cells. Transfected NG108-15 and HEK293 cells could be identified by GFP fluorescence observed using the Nikon epifluorescence unit (See Fig 2.2). Drugs were applied to cells using a fast perfusion system based on a modified Y-tube (Wooltorton *et al.*, 1997) controlled with solenoid valves (Lee products ltd, Buckinghamshire) operated electronically (See Fig 2.4). Cells were voltage-clamped using an Axopatch 1C patch clamp amplifier (Axon Instruments) with a CV-3 headstage (Axon Instruments) at -50mV (HEK293 or NG108-15 cells) or -60mV (cerebellar granule cells). Membrane currents were filtered at 3kHz (8 pole Bessel filter) and digitised via a Digidata 1200 (Axon Instruments) and viewed with a PC (Dell – Optiplex GX260) using the software programme Clampex 8.2 (Axon Instruments - See Fig 2.3).

2.4.2 Whole-cell patch clamp recording.

Cells were cultured on glass coverslips that could be transferred to the recording chamber and anchored in place in the bath. Cells were continuously perfused with Krebs solution at room temperature containing: 140 mM NaCl, 4.7 mM KCl, 1.2 mM MgCl₂, 2.52 mM CaCl₂, 11 mM glucose and 5 mM HEPES, pH adjusted to 7.4 with NaOH and filtered through Whatman paper. Osmolarity was confirmed as 300 ± 10 mOsm with an Osmometer (Automatic – CAMLAB).

Patch pipettes were pulled from thick-walled borosilicate glass (Harvard, GC150-10) and filled with an internal patch pipette solution containing: 120 mM

KCl, 1 mM MgCl₂, 11 mM EGTA, 10 mM HEPES, 1mM CaCl₂ and 4mM Na₂ATP, pH adjusted to 7.2 with NaOH (osmolarity confirmed as 280-300 mOsm). This solution was used to record from HEK293 cells in Figs 3.2-4. At all other times the patch pipette solution contained: 150 mM CsCl, 1 mM MgCl₂, 10 mM HEPES, 4 mM Na₂ATP, 0.1 mM CaCl₂ and 1.1 mM EGTA, pH adjusted to 7.2 with CsOH (osmolarity confirmed as 280-300 mOsm). Free [Ca²⁺] \leq 100nM.¹

The resistance of patch electrodes was monitored using the seal-test option in Clampex 8.2. The resistance of electrodes used to record from HEK293 and NG108-15 cells was 5-6 M Ω , and for cerebellar granule cells, 8-9 M Ω . Electrodes were manoeuvred into position onto the surface of the cell using a Sutter MP-285 micromanipulator.

The seal-test option and the membrane test option in Clampex 8.2 were used to monitor the fidelity of the whole-cell recording (See Fig 2.3B and C). The seal test measured the seal resistance between pipette and ground by repetitive application of -5mV voltage pulses and measuring the current response. This was then used to set the capacitance compensation on the patch clamp amplifier. Series resistance and whole-cell capacitance were optimised before the start of experiments using the shape of the waveform in the seal-test as a guide (See Fig 2.3C).

During patch clamp recording a proportion of the command potential is expected to be lost as it passes across the patch electrode and cell membrane in series (series resistance). Voltage clamp errors are expected to be low if the cell is healthy and the membrane seal around the electrode is high (G Ω). Voltage clamp errors were further reduced by compensating for series resistance by using the patch clamp amplifier to inject current into the cell to compensate what is lost due to series resistance. Voltage clamp errors unavoidably increase for larger membrane currents

^{1.} Calculated using Mathcad 2000i.

(Thomas and Smart, 2002). Thus as a result, membrane currents of over 4 nA, usually recorded from larger NG108-15 cells, were discarded from the analysis.

The stability of the connection between cell and electrode was monitored throughout the course of the experiment using the seal-test and membrane test options to continuously monitor seal resistance, access resistance (Ra) and series resistance (Rs). Series resistance can also vary as access resistance alters which tends to occur as the electrode tip becomes occluded with cellular contents. Cells showing changes in access resistance or series resistance of over 15 % were discarded from the analysis.

The cell's current response after application of GABA was recorded, in the gap-free mode (Clampex 8.2) and stored on the PC for further analysis. Typical values for series resistance were $<1M\Omega$, and for capacitance, 21.7 ± 3.6 pF (n = 6, HEK293), 56.5 ± 7.6 pF (n = 6, NG108-15), 11.1 ± 0.6 pF (n = 15, CGCs – 20mM KCl) and 8.3 ± 0.5 pF (n = 15, CGCs – 5mM KCl). The resting membrane potentials of cells immediately after achieving the whole-cell configuration were recorded as a good approximation of cell health. HEK293 cells and NG108-15 cells generally had typical values -33.5 ± 5.6 mV (n = 6, HEK293 cells), -39.5 ± 4 mV (n = 6, NG108-15). CGCs maintained in depolarising conditions had slightly lower resting potentials on average (-39.2 ± 2.5 mV) as compared to CGCs maintained under physiological conditions (-47.5 ± 5.9 mV; comparable to that reported by Mellor *et al.*, 1998).



Fig 2.2: Photographs of the electrophysiology rig.



A



pClamp gap-free recording



Fig 2.3: Software used for data acquisition: pClamp 8.2.

(A) Whole cell currents were recorded in the gap-free mode.

(B/C) Membrane test and seal test in pClamp were used throughout the experiment to monitor Ra and Rs. The seal test waveform was used as a guide for compensation of series resistance on the amplifier.

2.5 DRUG APPLICATION AND SOLUTIONS

2.5.1 Drug application.

Drugs were rapidly-applied to cells via a Y-tube positioned approximately 400 μ m away from the recorded cell. Cells were continuously superfused by a gravity-fed Krebs solution (See Fig 2.4) whilst solenoid 1 was open. Water (to save drug solutions) was perfused around the Y-tube whilst solenoid 2 was open, powered by the vacuum pump. Just before the drug was added water was replaced by the appropriate drug solution dissolved in Krebs solution. Both solenoids were closed simultaneously allowing the drug solution to flow rapidly over the cell whilst stopping the normal Krebs from flowing. When both solenoids were opened again (after a fixed amount of time), the normal Krebs could wash-off the drug solution from the exposed cell (Wooltorton *et al.*, 1997).





Fig 2.4: Schematic representation of the rapid perfusion Y-tube system.

The flow of a drug was controlled by two solenoids which acted as gates to allow or stop drug application and wash. The solenoids were controlled electronically. The typical rise time (10-90 %) of the 10 μ M GABA response in NG108-15 cells was 276.2 ± 28.9 ms (n = 6).

2.5.2 Drug solutions (intracellular ligands).

- KN-93 (Sigma), an inhibitor of CaMK-II activity (Sumi *et al.*, 1991) was dissolved in DMSO to form a stock concentration of 2.4 mM. KN-93 was added to the pipette solution at a dilution factor of 1 in 4800 to ensure DMSO did not interfere with the stability of recordings.
- KN-92 (Sigma), an inactive analogue of KN-93 was dissolved in DMSO to form a stock concentration of 1 mM and was dissolved in the pipette solution at a dilution of 1 in 2000.
- Genistein a specific inhibitor of tyrosine kinase activity (Akiyama *et al.*, 1987) was dissolved in DMSO to give a final stock solution of 100 mM and diluted 1 in 1000 in the patch pipette solution.
- Okadaic acid sodium salt (Sigma), an inhibitor of protein phosphatase activity (Bialojan and Takai, 1988) was dissolved in internal pipette solution to form a stock solution of 100-120 µM. This was further diluted at a ratio of 1:100-120 (1 µM final concentration) or 1:10-12 (10 µM final concentration) okadaic acid-containing pipette solution to normal pipette solution.

2.5.3 Pre-activation of CaMK-II (Upstate) and *in vitro* phosphorylation assay.

Purified rat forebrain CaMK-II (50 ng/µl in 50 mM HEPES, pH 7.5, 50 % v/v glycerol, 10 % v/v ethylene glycol, 2.5 mM EDTA and 1 mM dithiothreitol) consisted of a mixture of CaMK-II isoforms (α , β , γ , δ), 50-60 kDa/isoform and was pre-activated by incubation with 10 mM MgCl₂, 1 µM CaM, 0.2 mM EGTA, 0.3 mM CaCl₂ for 15 min at 25°C. The pre-activation buffer containing CaMK-II was then diluted in the patch pipette solution to give a final concentration of 60-100 nM CaMK-II.

In vitro phosphorylation of β 3 TM3-4 loop GST-fusion proteins by CaMK-II (See Fig 3.4) was carried out in collaboration with Dr J.N Jovanovic as follows:

Purified CaMK-II (Upstate – 10 ng/µl final concentration in pre-activation buffer), was pre-activated (10 min) and incubated with GST- β 3 (1.6 µg/µl) and ³²P-ATP (0.4mM) at 25°C or 5°C, with or without 150 mM KCl. The reaction was stopped (t = 10 min) by addition of an equal volume of SDS-PAGE (sodium dodecyl sulphate-polyacrylamide gel electrophoresis) sample buffer. Phosphorylation was analysed by SDS-PAGE and visualized by autoradiography. KCl (150 mM) was added to the pre-activation/autophosphorylation buffer to simulate addition to the patch pipette solution. CaMK-II was pre-activated by incubation in 10 mM MgCl₂, 1 µM CaM, 0.2 mM EGTA, 0.3 mM CaCl₂ (McDonald and Moss, 1994; McDonald and Moss, 1997).

2.5.4 Pre-activation of recombinant α-CaMK-II (New England Biolabs).

In all other experiments a truncated monomer of rat α -CaMK-II (1-325 amino acid residues) with a molecular weight of 33 kDa (New England Biolabs) obtained from *Spodoptera frugiperda* (Sf9) cells infected with recombinant baculovirus carrying the recombinant α -CaMK-II cDNA (Brickey *et al.*, 1990; Suzuki-Takeuchi *et al.*, 1992) was used. This had a specific activity of \approx 5,000,000 units/mg, supplied at 500 units/µl (100 ng/µl). One unit is defined as the amount of activated CaMK-II required to catalyze the transfer of 1 pmol of phosphate to Autocamtide-2 (CaMK-II peptide substrate) KKALRRQETVDAL (NEB – 50 µM) in 1 min at 30°C in a total reaction volume of 30 µl.

 α -CaMK-II was supplied in 100 mM NaCl , 50 mM Tris-HCl (pH 7.5 @ 25°C), 0.1 mM Na₂EDTA, 1 mM dithiothreitol, 0.02 % v/v Tween-20 and 50 % v/v

glycerol. Glycerol at this concentration was detrimental to the health of the cells so α -CaMK-II was dialysed with the Slide-A-Lyzer MINI Dialysis unit to reduce the glycerol concentration. Dialysis units were soaked in de-ionised water for 15 min. The dialysis unit was then placed in the microtube supplied, filled with 1.3 ml of dialysate containing 100 mM NaCl, 50 mM Tris-HCl (pH 7.5 @ 25°C), 0.1 mM EDTA and 1 mM dithiothreitol. The α -CaMK-II as supplied was then added just above the membrane and allowed to sink. The dialysis unit was then capped and placed on a shaker at 4°C. CaMK-II and the microtube were kept on ice throughout the procedure. After 1-2 hrs the sample was collected from the dialysis unit and aliquots ready for use in patch clamp recordings were frozen and kept at -70°C. The dilution factor generally fell between 2-2.5 fold giving a final concentration of 1200-1500 nM (40-50 ng/ μ l). Stocks were defrosted just before use and incubated in a pre-activation/autophosphorylation buffer as suggested by the manufacturers: 8-10 µl CaMK-II stock (40-50 ng/ μ l = 1200-1500 nM) diluted into CaMK-II buffer as supplied (50 mM Tris-HCl, 10 mM MgCl₂, 2 mM dithiothreitol, 0.1 mM Na₂EDTA, pH 7.5 @ 25°C) in a total volume of 16-20 µl supplemented with 1.2 µM calmodulin (CaM), 1.5 mM CaCl₂, 0.4 mM ATP- γ -S incubated at 25°C for 15 min (McGlade-McCulloh et al., 1993).

The CaMK-II pre-activation buffer was then diluted into the patch pipette solution (McGlade-McCulloh *et al.*, 1993; Lledo *et al.*, 1995; Barria *et al.*, 1997; Nelson *et al.*, 2005) to give a final CaMK-II concentration of 35 nM (1.16 ng/µl), 60 nM (2 ng/µl), 85 nM (2.85 ng/µl) or 100 nM (3.33 ng/µl). At the highest concentration the dilution factor was approximately 1:12 α -CaMK-II stock solution to patch pipette solution. Control recordings were made with the pre-activation mixture without CaMK-II diluted into the patch pipette solution at the same ratio, in

order to ensure there were no non-specific effects on the stability of recordings caused by residual CaM and CaCl₂ from the pre-activation buffer or that the buffer diluted into the patch pipette solution did not activate endogenous Ca^{2+}/CaM dependent kinases or phosphatases.

2.6 IMMUNOCYTOCHEMISTRY

2.6.1 Procedure for immunolabelling of cell cultures.

Cells cultured on glass coverslips (See 2.2.3) were washed gently twice with ice-cold phosphate buffered saline (PBS). They were then fixed in 4 % paraformaldehyde (Sigma) in PBS, for 15 min at room temperature. The fixative was then washed with PBS twice and quenched with 50 mM NH₄Cl (in PBS) twice for 5 min. Cells were then washed again with PBS (twice). At this point the fixed cells were kept until required at 4°C.

If the desired antigen was intracellular (anti- β 3) the cells were then permeabilised by incubation at room temperature with 0.05 % v/v Triton-X-100, 10 % v/v FCS, 0.5 % w/v BSA (in PBS) for 10 min. Followed by a further two washes in PBS. If the desired antigen was extracellular (anti-myc) this stage was omitted as only cell-surface labelling was required.

Prior to antibody incubation, cells were washed in 10 % v/v FCS, 0.5 % w/v BSA (in PBS). The appropriate antibody was then diluted (See Table 2.1) in the same buffer. Then, 50 μ l of the diluted antibody was placed onto a parafilm segment, the coverslip was removed from solution and the edge briefly dried on a tissue before inverting over the drop. Coverslips were covered with the dish lid to prevent drying and left for 45 min at room temperature.

After the primary antibody incubation cells were placed back into 35 mm dishes and washed three times with 10 % v/v FCS, 0.5 % w/v BSA in PBS. The secondary antibody was diluted into the same buffer at the appropriate concentration (See Table 2.1) and again coverslips were inverted over 50 μ l of the diluted antibody mixture dropped onto parafilm. Coverslips were covered with the lid of the dish to prevent drying and covered with foil to protect from the light. Cells were incubated in the secondary antibody for 45 min at room temperature.

After incubation, the coverslips were returned to the 35 mm dishes and washed three times in 10 % v/v FCS, 0.5 % w/v BSA (in PBS) and then three times in PBS. Coverslips were fixed to microscope slides using glycerol. The coverslips were lifted from the dish and excess fluid was removed by dipping the edge on a tissue. A drop of (pre-heated) glycerol (30 μ l) was placed on the slide and the coverslip was inverted onto it and left to dry (Duguid and Smart, 2004).

Antibody	Species	Epitope	Dilution	Source
Anti-myc (1°)	Polyclonal Raised in Rabbit	C-teminus of c-Myc of human origin.	1:250	Santa Cruz Biotechnology
Anti-β3 (1°)	Polyclonal Raised in Rabbit	TM3-4 Loop of the β3 subunit.	1:200	Gift from Prof. W. Sieghart (Vienna)
Anti-rabbit TRITC (2°)	Raised in Goat	lgG pooled rabbit serum	1:250	Sigma

Table 2.1:

Table 2.1: Primary and secondary antibodies used in immunocytochemistry.

2.6.2 Image acquisition.

Images were acquired using a Zeiss laser-scanning confocal microscope (LSM 510 Meta). An oil-immersion objective lens (x 40 Plan-Neofluar or x 63 Plan-Apochromat) was used to visualise individual cells. Multi-track images were obtained with an Argon laser (488 nM) to visualise EGFP fluorescence and a Krypton (568 nM) laser to visualise tetramethyl rhodamine isothiocyanate (TRITC) fluorescence. Optical sections (z series) of intervals of 0.3-0.4 μ m were taken through cells and 8-10 images were reconstructed to yield the final projection image. The precise settings can be seen in Fig 2.5.

Fig 2.5:



Fig 2.5: Settings of Zeiss LSM 510 Meta laser scanning confocal microscope.

Optimal settings are shown. Detector gain and amplifier offset can be altered to increase the signal to noise ratio for each channel. The pinhole size was set at 1 Airy unit with an optical slice of $<1 \mu m$. A series of optical slices (Z series) can be taken through the cell to form a projected image.

2.7 KNOCKOUT MICE

The β 2 knockout mouse strain (from Merck Sharp and Dohme, UK) was created by deleting exons 6 and 7 of the β 2 subunit gene by homologous recombination (Sur *et al.*, 2001). Heterozygous F3 generation mice were cross-bred to produce β 2 -/- and wild-type (wt) strains. The offspring of breeding pairs of the F5-6 generation of wt and β 2 -/- mice were taken at P0-2 to prepare cerebellar granule cell cultures (See 2.2.3).

The tails of offspring were removed after dissection for genotyping to confirm that the $\beta 2$ subunit was missing from the $\beta 2$ -/- cultures. DNA was extracted from the mouse tails of every pup (See appendix A1.4) in a litter used for each culture and this genomic DNA was included in a PCR reaction using primers complimentary to the $\beta 2$ subunit gene and a primer complimentary to the neomycin cassette, to act as a positive control for homologous recombination in $\beta 2$ -/- cultures (Sur *et al.*, 2001).

2.8 ANALYSIS

2.8.1 Whole-cell currents.

Whole-cell currents were analysed in Clampfit 8.2 (See Fig 2.3A) by measuring the peak amplitude and subtracting the baseline. Cells were chosen for analysis if access resistance and series resistance remained stable throughout the period of recording (See 2.5.2). If traces had deflections in their rising phase that current was discarded from the analysis. The peak amplitude recorded at 2 min intervals was normalised to the peak amplitude of the response obtained 3-4 min after achieving the whole-cell configuration.

The peak amplitude of the response to different concentrations of GABA was used to construct a concentration-response curve (See Fig 4.2). The peak amplitudes were normalised to the capacitance of the cell (pA/pF). These values were then plotted against GABA concentration. At least 5 different GABA concentrations were used to define the threshold or heel of the curve, the linear portion, and the clear saturating maximum. These data were then fitted by the Hill equation (shown below) within the program Origin 6.0 (Microcal):

$$I = 1 / (1 + (EC_{50}/[GABA]^{n}))$$

where, I = the current activated by GABA (normalised to membrane capacitance). EC₅₀ = the concentration of GABA to activate a current that is 50 % of the amplitude of current induced by a maximal dose of GABA and **n** = the Hill coefficient.

2.8.2 MiniAnalysis

The software program MiniAnalysis (Synaptosoft) was used to analyse synaptic currents. Recordings were captured in Clampex 8.2 over 2 min or 5-6 min epochs starting 4-5 min after achieving the whole-cell configuration. Within MiniAnalyis a period of recording could be displayed and synaptic currents were detected automatically with an amplitude threshold of 6-10 pA, depending on the level of baseline noise, which was typically around 5 pA. After automatic detection, each event was manually checked and verified by eye. The program could then analyse and display a mean value for the frequency, the peak amplitude and the half-

width (the time taken to decay to 50 % of the peak amplitude) over that period of recording (See Fig 2.6A).

Events could also be grouped manually and these groups could be used to analyse the decay time constants of each period of recording. Synaptic events were chosen for analysis if there were no deflections in the rising or decaying phases and they decayed back to the baseline. Events of low amplitude (<5 pA) were discarded from this analysis as they were thought to represent events occurring at the most distant synapses and therefore more affected by dendritic filtering or cabling. All the events chosen were grouped together for a given recording and aligned by their mean rise-times (50 %) for analyses of decay time constants. The program calculated an average current from the group. From this the 10-90 % rise-time could be calculated. The average current can then be fitted with a sum of one, two or three exponentials. The best fit could be identified visually but the program also allowed the best fit to be determined by regression analysis.

The resulting best fit curve was described by the following equation:

$$y = A1 \exp(-x/\tau 1) + A2 \exp(-x/\tau 2) + baseline,$$

where, A1 and A2 are the fractions of the fast (1) and slow (2) components and $\tau 1$ and $\tau 2$ are their respective decay time constants.

A decay time constant is defined as the time taken for each component to decay to 37 % of its original amplitude. From this, values for the decay time constant of each component could be recorded as well as the relative area of each

component. When several cells had been analysed in this way, mean values could then be determined (See Fig 2.6B).



Fig 2.6: Software used for analysis of synaptic currents: MiniAnalysis.

(A) In the main display window the green x indicates the location of the baseline. The yellow dot (\bullet) indicates the starting time of the rising phase of the peak. The red x indicates the maximum point of the peak. The purple dot (\bullet) indicates the decay of the peak (50 %).

(B) 10-90 % rise time is indicated by two yellow dots (\bullet). 90-37 % decay is shown by two purple dots (\bullet , indicated by the arrows). The red x is where the peak is located. The green x and green horizontal line are where the baseline average is located. The peak amplitude is calculated as peak minus baseline average. The area under the curve is calculated from the whole trace with respect to the green line. The equation describing the curve is shown in 2.8.2.

2.8.3 Gaussian fits

Raw current data could be transferred from MiniAnalysis into Origin 6.0 for analysis of distributions. If the data was formed from populations described by a Gaussian distribution this program fitted the data (with multiple Gaussian distributions if necessary) to identify separate populations. The best fit was determined visually.

2.8.4 Statistics

For comparison of two values, statistical analyses were carried out using the unpaired Student's t-test (with a Welch correction if the difference in standard deviations between groups was large). When analysing multiple means, statistical analyses were carried out using ANOVA, with a Bonferroni post-test to compare selected groups. With all statistical tests, two means were considered significantly different if the P value was less than 0.05. All statistical tests were carried out using the software program GraphPad Instat 3.01 (GraphPad, USA).

Chapter Three:

<u>CaMK-II modulation of recombinant GABA_A Receptors</u> <u>expressed in HEK293 cells.</u>

3.1 INTRODUCTION

The HEK293 cell line is endothelial in origin so is generally not thought to express any functional neuronal ligand-gated ion channels (Graham *et al.*, 1977). These cells are suitable for whole-cell patch clamp, which allows electrophysiological recordings to be made of membrane currents following activation of recombinant ion channels that are expressed after transfection. In this way the exact isoform/s or combinations of subunits making up a given receptor/ion channel can be largely controlled, allowing functional dissection of the roles of different subunit isoforms, and the signalling processes that alter their function.

Expression of GABA_A receptor subunits in HEK293 cells has been used previously to investigate the role of PKA-induced phosphorylation, revealing functional heterogeneity between different β subunits (McDonald *et al.*, 1998). Similar approaches have been used to investigate PKC-dependent phosphorylation of GABA_A receptors (Brandon *et al.*, 2002b) and tyrosine kinase-Src dependent modulation (Moss *et al.*, 1995).

It has been shown that CaMK-II can phosphorylate GST-fusion proteins formed from the intracellular loop between transmembrane (TM) domains 3 and 4 of β and γ subunits of GABA_A receptors *in vitro* (Machu *et al.*, 1993; McDonald and Moss, 1994; McDonald and Moss, 1997). In neuronal preparations intracellular addition of pre-activated CaMK-II, through inclusion in the patch pipette, led to a significant increase in the peak amplitude of GABA_A mediated currents (Wang *et al.*, 1995; Kano *et al.*, 1996). Synaptosomal membrane fractions have also been used to show CaMK-II dependent modulation of agonist binding to GABA_A receptors and alterations in benzodiazepine binding to these receptors possibly through phosphorylation of the α l subunit (Churn and DeLorenzo, 1998; Churn *et al.*, 2002).

However, as yet, no attempt has been made to study the functional effects of CaMK-II dependent phosphorylation of GABA_A receptors in a controlled environment such as the HEK293 cell. Such HEK293 cells have been used to successfully demonstrate CaMK-II dependent functional modulation of AMPA receptors (Barria et al., 1997; Derkach et al., 1999). The advantages of using such a system are that any potential differences in effect caused by phosphorylation on different GABA_A receptor subunits may be determined. This is especially important in the case of GABAergic signalling mechanisms as the GABA_A receptor displays considerable heterogeneity in subunit combinations and variable functional effects depending on the different subunits present in a particular recombinant system or neuronal cell type (Moss and Smart, 1996; Nusser et al., 1999; Brandon et al., 2002a). It is also possible, through site-directed mutagenesis, to disrupt phosphorylation of specific residues within subunits and determine the specific sites of phosphorylation and their relative importance. Using these advantages, the initial aim of these experiments was to express different subunit combinations of the GABA_A receptor in HEK293 cells and to induce CaMK-II dependent phosphorylation to study the functional consequences (See Fig 3.1).



Fig 3.1: Studying CaMK-II dependent effects on GABA_A receptors using HEK293 cells.

The cDNAs of various $GABA_A$ receptor subunits are transfected into HEK293 cells by electroporation (See Chapter 2) along with the plasmid encoding for enhanced GFP to allow identification of transfected cells.

The cells are left for 24 hours to allow expression of the recombinant $GABA_A$ receptors at the cell surface.

Whole-cell currents can then be recorded in response to applied GABA, over time, as pre-activated CaMK-II within the patch pipette dialyses into the cell and modulates the response (dotted lines).

3.2 **RESULTS**

In order to characterise the effect of CaMK-II on GABA_A receptor mediated currents recombinant GABA_A receptors were expressed in HEK293 cells. The cells were transfected by electroporation with cDNAs for specific GABA_A receptor subunits in equal ratio (1:1:1) and enhanced GFP to allow easy identification of transfected cells (See Chapter 2). The GABA_A receptor α and β subunits, when expressed together, are known to form robust functional channels that respond to GABA (Levitan *et al.*, 1988a; Levitan *et al.*, 1988b; Sigel *et al.*, 1990; Connolly *et al.*, 1996). To simplify the search for the effect(s) of CaMK-II on GABA_A receptors, initial experiments involved HEK cells expressing $\alpha 1\beta 2$ and $\alpha 1\beta 3$ alone, as the $\gamma 2$ subunit also possesses potential phosphorylation sites for this kinase (Machu *et al.*, 1993; McDonald and Moss, 1994).

Whole-cell patch clamp recordings were conducted to measure GABA mediated currents. To assess the stability of these currents over time, a fixed concentration of GABA (10 μ M) was applied every 2 min for 20-30 min. The peak amplitude was then normalised to the peak amplitude recorded 3-4 min after achieving the whole cell configuration. Under control conditions, the peak amplitude of the GABA-activated current remained at a relatively constant level for the time of recording with only a slight decrease of about 10-20 % apparent over 30 min (Fig 3.2-3.4). This "run-down" of GABA_A receptor currents has been observed in neurones and in recombinant expression systems and is thought to be associated with a disruption of the phosphorylation state(s) of the receptor. This is possibly through depletion of ATP or an alteration in the concentration of unbuffered Ca²⁺ caused by differences between the internal pipette solution and the normal intracellular milieu

(Stelzer *et al.*, 1988; Chen *et al.*, 1990; Huang and Dillon, 1998; Laschet *et al.*, 2004).

3.2.1 Recombinant $\alpha 1\beta 2$ receptors and α -CaMK-II.

Purified recombinant α-CaMK-II (Brickey *et al.*, 1990; Suzuki-Takeuchi *et al.*, 1992) was pre-incubated in 1.2 μM Calmodulin (CaM), 10 mM MgCl₂, 2 mM CaCl₂ and 0.4 mM ATP- γ -S to promote autophosphorylation at Thr²⁸⁶ (McGlade-McCulloh *et al.*, 1993; Lledo *et al.*, 1995; Wang *et al.*, 1995; Barria *et al.*, 1997; Nelson *et al.*, 2005). This recombinant form of α-CaMK-II represents a structurally simplified form of the kinase that cannot form a holoenzyme but which is capable of phosphorylating the same substrates as the full length form of the kinase (Suzuki-Takeuchi *et al.*, 1992). This recombinant form of α-CaMK-II can be extracted and purified after being expressed at high levels in an appropriate cell line, avoiding the difficulties associated with obtaining purified CaMK-II from whole-brain extracts. In addition this monomeric form of the kinase has a molecular weight of 33 kDa as opposed to the full holoenzyme with a molecular weight of 600 kDa thereby avoiding problems associated with diffusion of such a large molecule out of the patch pipette and into the cell.

ATP- γ -S was used rather than ATP in the pre-activation/autophosphorylation reaction, as thiophosphorylated proteins are less subject to dephosphorylation by the endogenous protein phosphatases that might be present in the HEK cell. The α -CaMK-II and the pre-activation buffer were then diluted into the pipette solution immediately before recording and then maintained on ice (See Chapter 2). To determine that the pre-activation buffer had no effect on GABA mediated currents,

all controls included a mock pre-activation buffer (lacking α -CaMK-II) diluted into the pipette solution at the same ratio.

In previous experiments using a similar form of recombinant α -CaMK-II, 200 nM α -CaMK-II significantly potentiated GABA mediated currents in rat dorsal horn and hippocampal neurones (Wang *et al.*, 1995). Since HEK293 cells are more compact without the large processes of neurones to diffuse through, the concentration of α -CaMK-II was reduced to 100 nM (3.33 ng/µl, See Chapter 2). This concentration should be sufficient to allow phosphorylation of the GABA_A receptor without compromising the quality of recordings. In HEK cells expressing $\alpha 1\beta 2$ subunits, the addition of 100 nM pre-activated α -CaMK-II in the patch pipette induced no significant differences from control GABA currents over a recording period of 30 min. Control currents recorded in response to application of 10 µM GABA at t = 10 min was 88.5 ± 2.3 % (Fig 3.2A, n = 13) and in the presence of α -CaMK-II at t = 10 min was 87.5 ± 3.1 % (Fig 3.2A, n = 5).

In addition, purified rat forebrain CaMK-II (full-length, consisting of a mixture of α , β , γ and δ subunits) was also used and pre-activated in a similar way (co-incubation with 10 mM MgCl₂, 1 μ M CaM, 0.2 mM EGTA, 0.3 mM CaCl₂, 0.4 mM ATP- γ -S; See Chapter 2). This form of CaMK-II (60-100 nM) also had no significant effects on GABA mediated currents which attained 87.8 ± 1.8 % at t = 10 min (n = 4, Fig 3.4A) as compared to control (88.5 ± 2.3 %, n = 13; Fig 3.2A). This form of CaMK-II was tested biochemically and shown to be capable of phosphorylating the β 3 subunit intracellular domain *in vitro*, after the pre-activation reaction and in the presence of 150 mM KCl, used to simulate conditions in the patch pipette solution (in collaboration with Dr. J. Jovanovic; Fig 3.4B).

As an alternative method, HEK cells were transfected with $\alpha 1\beta 2$ cDNA and with cDNA for α -CaMK-II in equal ratio. Inclusion of a relatively high CaM (600 nM) concentration in an internal solution with a higher free Ca²⁺ (10 μ M) concentration may be expected to activate α -CaMK-II expressed in the HEK cells and thereby modulate GABA mediated currents. However no significant differences were seen between control recordings without CaMK-II co-transfection, CaMK-II co-transfection with normal patch pipette solution, and CaMK-II co-transfection with a high CaM and high free Ca²⁺ patch pipette solution. The free Ca²⁺ concentration in the pipette solution was altered by changing the ratio between EGTA : CaCl₂ from 10:1 (10 nM free Ca²⁺) in the normal pipette solution to 1:1 (10 μ M free Ca²⁺ (Aguayo *et al.*, 1998). After 10 minutes recording, GABA current amplitudes were: Control, 88.5 ± 2.3 % (n = 13); CaMK-II co-transfection, 92.9 ± 5 % (n = 5); and CaMK-II co-transfection with additional Ca²⁺/CaM, 96.4 ± 6.2 % (n = 9, Fig 3.2B).

Previously, it has been demonstrated that CaM itself can have a functional effect on some ion channels including ligand-gated ion channels like GABA_A receptors. For instance it can have a direct inhibitory effect on NMDA receptor function (Ehlers *et al.*, 1996; Zhang *et al.*, 1998; Rycroft and Gibb, 2002). It can also activate CaM dependent phosphatases such as PP2B/Calcineurin, which may have a direct effect on the GABA_A receptor (Jones and Westbrook, 1997; Wang *et al.*, 2003). In order to determine whether CaM itself was modulating GABA_A receptor function and occluding CaMK-II dependent effects, GABA currents were recorded in the presence of higher concentrations of CaM (1.5 μ M) alone. However there were no significant differences observed between control GABA current amplitudes, 88.5 ± 2.3 % (Fig 3.2C, n = 13), and those in the presence of CaM, 90.5 $\% \pm 4.2$ % (Fig 3.2C, n = 6) at t = 10 min.

Taken together it appeared that CaMK-II had no functional effect on $\alpha 1\beta^2$ subunit-containing GABA_A receptors expressed in HEK293 cells. Interestingly, cAMP-dependent kinase (PKA) was also unable to functionally modulate $\alpha 1\beta^2$ subunit-containing receptors in HEK cells but had a significant effect on whole-cell currents recorded from $\alpha 1\beta^3$ subunit-containing receptors (McDonald *et al.*, 1998). Therefore the same series of experiments with CaMK-II were repeated in HEK293 cells expressing $\alpha 1\beta^3$ subunits.



Fig 3.2: α -CaMK-II modulation of whole-cell currents recorded from HEK293 cells expressing $\alpha 1\beta 2$ subunits.

Peak amplitudes of GABA-mediated currents recorded following the application of 10 μ M GABA to transfected HEK293 cells expressing $\alpha 1\beta 2$ GABA_A receptors. GABA was re-applied at 2 min intervals for up to 30 min. All currents were normalised to the peak amplitude of the response measured during the first 3-4 min (= 100 %) after achieving the whole-cell configuration (t = 0 min). Control recordings were made with normal patch pipette solution supplemented with the pre-activation buffer without α -CaMK-II.

(A) GABA currents recorded in the absence (Control, n = 13) and presence of 100 nM pre-activated α -CaMK-II, which was introduced into the cells via the patch pipette solution (n = 5).

(B) Using HEK293 cells now also transfected with α -CaMK-II cDNA, GABA currents were recorded using two different patch pipette solutions, containing: normal Ca²⁺ buffered patch pipette solution (EGTA 10:1 CaCl₂, free Ca²⁺ 10 nM; n = 5); and a solution supplemented with CaM (600 nM) and high Ca²⁺ (EGTA 1:1 CaCl₂, free Ca²⁺ 10 μ M; n = 9). The control cells were transfected with cDNA for $\alpha l\beta 2$ only (Taken from 3.2A, n = 13).

(C) GABA currents recorded from HEK293 cells expressing only $\alpha 1\beta 2$ GABA_A receptors using either a normal patch pipette solution (Taken from 3.2A, Control, n = 13) or one supplemented with 1.5 μ M CaM (n = 6).

(D) Representative 10 μ M GABA-activated currents recorded at t = 2 and 10 min following internal dialysis with 100 nM α -CaMK-II in the patch pipette solution.

All points represent the mean \pm s.e.

In A, B and C, the exposure to α -CaMK-II or CaM did not significantly affect the GABA currents recorded.

P < 0.05, t-test, ANOVA.

Recombinant $\alpha 1\beta 3$ GABA_A receptors and CaMK-II 3.2.2

The addition of autophosphorylated α -CaMK-II to the pipette solution also had no significant effects on GABA mediated currents compared to control in cells expressing $\alpha 1\beta 3$ (Control = 92 ± 3.3 %, n = 11; + CaMK-II = 81.7 ± 4.5 %, n = 6; Fig 3.3A, t = 10 min). Furthermore, the co-transfection with α -CaMK-II cDNA and the addition of CaM to a high Ca²⁺ containing patch pipette solution had no significant effects on GABA (10 μ M) currents compared to control (Control = 92 ± 3.3 %, n = 11, α -CaMK-II transfection + CaM = 85.2 ± 2.9 %, n = 4, t = 10 min, Fig. 3.3A). Finally the addition of 1.5 μ M CaM alone to the pipette solution also had no significant effects on the $\alpha 1\beta 3$ GABA_A receptors (Control = 92 ± 3.3 %, n = 11, + $CaM = 92.2 \pm 3.5$ %, n = 6, t = 10 min, Fig 3.3C).

Pre-activated CaMK-II purified from rat forebrain also had no significant effect on GABA mediated currents recorded from HEK293 cells expressing $\alpha 1\beta 3$ $(84.4 \pm 11.3 \%, n = 5, t = 10 \text{ min}, \text{Fig 3.4A})$. Taken overall CaMK-II was unable to modulate $\alpha 1\beta 3$ subunit-containing receptors in HEK cells.







Fig 3.3: α -CaMK-II modulation of whole-cell currents recorded from HEK293 cells expressing $\alpha 1\beta 3$ subunits.

Peak amplitudes of GABA-mediated currents recorded following the application of 10 μ M GABA to transfected HEK293 cells expressing $\alpha 1\beta 3$ GABA_A receptors. GABA was re-applied at 2 min intervals for up to 30 min. All currents were normalised to the peak amplitude of the response measured during the first 3-4 min (= 100 %) after achieving the whole-cell configuration (t = 0 min). Control recordings were made with normal patch pipette solution supplemented with the pre-activation buffer without α -CaMK-II.

(A) GABA currents recorded in the absence (Control, n = 11) and presence of 100 nM pre-activated α -CaMK-II, which was introduced into the cells via the patch pipette solution (n = 6).

(B) Using HEK293 cells now also transfected with α -CaMK-II cDNA, GABA currents were recorded using normal patch pipette solution supplemented with CaM (600 nM) and high Ca²⁺ (EGTA 1:1 CaCl₂: free Ca²⁺ \approx 10 μ M; n = 4). The control cells were transfected with GABA_A receptor cDNAs only (Taken from 3.3A, Control, n = 11).

(C) GABA currents recorded from HEK293 cells expressing only $\alpha 1\beta 3$ GABA_A receptors using either a normal patch pipette solution (Taken from 3.3A, Control, n = 11) or one supplemented with 1.5 μ M CaM (n = 6).

All points = mean \pm s.e.

In A, B and C, the exposure to α -CaMK-II or CaM did not significantly affect the GABA currents recorded.

P < 0.05, t-test.



Fig 3.4: Purified rat forebrain CaMK-II phosphorylation of the β 3 subunit *in vitro* and modulation of GABA currents in HEK293 cells.

(A) GABA currents recorded with patch pipette solution supplemented with purified pre-activated CaMK-II (60-100 nM) from HEK293 cells expressing $\alpha 1\beta 2$ (n = 4) and $\alpha 1\beta 3$ (n = 5). All points are mean \pm s.e.

(B) Purified pre-activated CaMK-II incubated with GST- β 3 and ³²P-ATP at 25°C or 5°C with or without 150 mM KCl, visualized by autoradiography.

Phosphorylation reaction was carried out in collaboration with Dr J. Jovanovic (See Chapter 2).

3.2.3 Recombinant $\alpha 1\beta 2/3\gamma 2$ GABA_A receptors and CaMK-II.

The $\gamma 2$ subunit also contains potential phosphorylation sites for CaMK-II as identified by studies involving GST-fusion proteins in vitro (Machu et al., 1993; McDonald and Moss, 1994). It is also thought that $GABA_A$ receptors require a γ subunit to form functional synaptic receptors (Essrich et al., 1998; Brickley et al., 1999; Studler et al., 2002). Therefore the possibility remained that CaMK-II dependent modulation may be solely dependent upon the γ^2 subunit, which would explain why the modulation was not observed on $\alpha\beta$ heteromers. Recordings were made from HEK cells expressing $\alpha 1\beta 2\gamma 2S$ and $\alpha 1\beta 3\gamma 2S$ with 100 nM α -CaMK-II in the patch pipette. As for the $\alpha\beta$ heteromers again no significant differences in the 10 µM GABA currents were observed from control in either case (Fig 3.5A, 3.5B). Given that the proposed stoichiometry for an $\alpha\beta\gamma$ subunit-containing receptor is thought to be 2:2:1 (Chang et al., 1996; Tretter et al., 1997) transfection of HEK cells with cDNA for each subunit in a ratio of 1:1:1 should ensure an overabundance of $\gamma 2$ cDNA and a preference for $\alpha\beta\gamma$ subunit-containing receptors at the cell surface. There remains a possibility that some $\alpha\beta$ subunit-containing GABA_A receptors may have been present and there is evidence that this can occur in HEK cells (Boileau et al., 2003). There is also, however, evidence that when $\alpha\beta$ and γ subunits are expressed together within a cell line $\alpha\beta\gamma$ channels are formed preferentially (Farrar et al., 1999) and that the ratio of cDNA has to be increased to $\alpha\beta\gamma$ 1:4:1 before $\alpha\beta$ receptors can be detected alongside $\alpha\beta\gamma$ (Angelotti et al., 1993; Angelotti and Macdonald, 1993). Angelotti et al., (2004) also observed that incorporation of the $\gamma 2$ subunit increased the peak amplitudes of the currents recorded. This was also observed in the current study (Fig 3.2D, Fig 3.5C). For the purposes of this experiment there should have been sufficient expression of $\gamma 2S$ subunit-containing
receptors at the cell surface to determine if CaMK-II can functionally modulate these receptors.

Fig 3.5:







Fig 3.5: α -CaMK-II modulation of whole-cell currents recorded from HEK293 cells expressing $\alpha 1\beta 2\gamma 2S$ and $\alpha 1\beta 3\gamma 2S$ subunits.

Peak amplitudes of GABA-mediated currents recorded following the application of 10 μ M GABA to transfected HEK293 cells expressing $\alpha 1\beta 2\gamma 2S$ and $\alpha 1\beta 3\gamma 2S$ GABA_A receptors. GABA was re-applied at 2 min intervals for up to 30 min. All currents were normalised to the peak amplitude of the response measured during the first 3-4 min (= 100 %) after achieving the whole-cell configuration (t = 0 min). Control recordings were made with normal patch pipette solution supplemented with the pre-activation buffer without α -CaMK-II.

(A) GABA currents recorded from HEK293 cells expressing $\alpha 1\beta 2\gamma 2S$ in the absence (Control, n = 6) or presence of 100 nM α -CaMK-II (n = 6).

(B) GABA currents recorded from HEK293 cells expressing $\alpha 1\beta 3\gamma 2S$ in the absence (Control, n = 5) or presence of 100 nM CaMK-II (n = 8).

(C) Representative 10 μ M GABA-activated currents recorded from HEK293 cells expressing $\alpha 1\beta 2\gamma 2S$ receptors a t = 2 and 10 min following internal dialysis with 100 nM α -CaMK-II in the patch pipette solution. Note: Increased current amplitude compared to Fig 3.2C.

All points = mean \pm s.e.

In A, B and C, the exposure to α -CaMK-II or CaM did not significantly affect the GABA currents recorded.

P < 0.05, t-test.

3.3 DISCUSSION

CaMK-II has previously been shown to modulate GABA_A receptor mediated currents in neurones that are known to express predominantly $\beta 2/3$ and $\gamma 2$ subunits (Wang *et al.*, 1995; Kano *et al.*, 1996; Wei *et al.*, 2004). Biochemical analysis of CaMK-II dependent phosphorylation of fusion proteins formed from the intracellular loops of different subunits, indicates that the β and γ subunits are the main candidates for phosphorylation (McDonald and Moss, 1994; Connolly *et al.*, 1996; McDonald and Moss, 1997). However, the present study shows that the expression of receptors containing $\beta 2/3$ and $\gamma 2S$ in HEK cells are not functionally modulated by CaMK-II.

It is possible that other subunits or subunit combinations that were not tested here are responsible for the CaMK-II mediated effect observed in neurones. For instance there is a potential site for phosphorylation in the 8 amino acid insert in the γ 2L splice variant. There are also potential sites for CaMK-II phosphorylation on the β 1 subunit (See Chapter 1, Table 1.1). The majority of β subunits expressed throughout the brain are β 2 followed by lower levels of β 3 and β 1 (Benke *et al.*, 1994). The expression of γ 2 is widespread with relatively low expression levels of other γ isoforms in a few selected cell types (Persohn *et al.*, 1992; Wisden *et al.*, 1992; Laurie *et al.*, 1992a; Laurie *et al.*, 1992b; Pirker *et al.*, 2000). CaMK-II dependent effects GABA_A receptors have been observed in hippocampal CA1 neurones, which are thought to express β 2/3/ γ 2 subunits. CaMK-II mediated effects have also been observed in cerebellar Purkinje cells that are thought to express predominantly α 1 β 2 γ 2 receptor subunits (Persohn *et al.*, 1992; Laurie *et al.*, 1992a) and so it would be predicted therefore, that GABA_A receptors containing β 2/3/ γ 2 are good candidates for CaMK-II dependent modulation. More recently, it has been suggested that the GABA_A receptor α l subunit may also be phosphorylated by CaMK-II and that this phosphorylation modulates agonist and allosteric-modulator binding (Churn *et al.*, 2002). This subunit was included in all of the combinations tested and there was no evidence of altered GABA_A receptor function. The majority of biochemical studies have identified β and γ and not α as the main sites of phosphorylation suggesting the α subunit is a poor substrate for phosphorylation (Moss *et al.*, 1992; McDonald and Moss, 1994; Moss and Smart, 1996).

Given the contradictory nature of the literature and the present results it is a more likely possibility that CaMK-II dependent modulation of GABAA receptor mediated currents is not possible in HEK293 cells because of an inherent difference between a neuronal environment and that of a HEK cell which is of epithelial origin. For instance the background of protein kinase and protein phosphatase activity within the intracellular milieu may be quite different. A very high level of phosphatase activity in HEK cells may act to prevent any modulation by CaMK-II by acting to dephosphorylate the GABA_A receptor itself. Recombinant GABA_A receptors expressed within different cell lines have been shown to display different functional properties. Comparison of $\alpha 1\beta 2\gamma 2$ mediated currents recorded in HEK293 cells and QT6 cells reveals that the deactivation kinetics of the receptor can be modulated by the cell type used for expression. Possibly, this may result because of differences in endogenous kinases and phosphatases, or in the proteins linking the receptors to the cytoskeleton (Mercik et al., 2003). In support of this there is evidence that the scaffolding protein GABARAP, which is thought to promote clustering of GABA_A receptors, also affects the channel properties of the receptor

when GABARAP and GABA_A receptors are co-expressed in L929 cells (Everitt *et al.*, 2004).

In addition, there is some evidence that HEK293 cells can express endogenous GABA_A receptor subunits that can co-assemble with subunits transfected into the cell altering the observed properties of the GABA_A receptor mediated currents recorded from HEK293 cells (Ueno *et al.*, 1996). However, this scenario has not been reported by all laboratories using HEK293 cells (Pritchett *et al.*, 1988) and has not been observed using the batches of HEK293 cells used in the current experiments (Connolly *et al.*, 1996; Thomas and Smart, 2005).

Despite these potential reservations, it is also clear that HEK cells have been used quite successfully in the study of PKA, PKC and the tyrosine kinase Src on GABA_A receptor function (Moss *et al.*, 1995; McDonald *et al.*, 1998; Brandon *et al.*, 2002b). The apparent inability of CaMK-II to modulate receptor function in HEK293 cells may simply reflect the inability of CaMK-II to associate with the GABA_A receptor in these cells.

There is a possibility that the diffusion of CaMK-II into the cell was too slow to reach full concentration in the time allowed. However, co-transfection of cDNA for CaMK-II would have produced a large amount of CaMK-II within the HEK cell before electrophysiological recording. The addition of CaM and an increase in free Ca^{2+} still produced no obvious modulation of GABA_A receptors. A recombinant form of α -CaMK-II, identical to the one used in these experiments, has been shown to have a diffusion coefficient of 4-5µm²/s within rat basophilic leukemia cells (RBL 2H3) (Shen and Meyer, 1998). This should be sufficient to allow diffusion throughout the relatively compact HEK cell within the time allowed. Examination of CaMK-II effects on AMPA receptors within HEK cells using a similar form of recombinant, truncated CaMK-II, activated by a similar autophosphorylation protocol, showed a significant modulation of currents within 10 min (Barria *et al.*, 1997). Another experiment with the same form of CaMK-II and using the *in vitro* autophosphorylation procedure as described in this study also demonstrated a functional effect on excitatory post synaptic currents (EPSCs) and whole-cell responses to AMPA in hippocampal pyramidal cells within 10-20 min (Lledo *et al.*, 1995).

There is also a possibility that CaMK-II activity was low when in the patch pipette solution either due to problems with the pre-activation reaction or due to deactivation within the patch pipette solution. This could be tested functionally by applying the same preparation to a neurone to establish whether neuronal GABA_A receptors could be modulated by CaMK-II. Under such conditions, the same recombinant α -CaMK-II preparation as that used to record from HEK293 cells could functionally modulate GABA mediated currents recorded from cerebellar granule cells (See Chapter 4).

In addition to recombinant α -CaMK-II a second preparation of purified rat forebrain CaMK-II consisting of a mixture of different isoforms of CaMK-II was also pre-activated and added to the patch pipette solution. This preparation of CaMK-II was tested for its ability to phosphorylate a GST-fusion protein of the intracellular loop of the β 3 subunit (See Fig 3.4B). This form of CaMK-II was shown to be able to phosphorylate the β 3 subunit TM3-4 intracellular loop *in vitro* (McDonald and Moss, 1997) but the same preparation also failed to significantly modulate GABA currents recorded from HEK293 cells expressing α 1 β 2/3 after being included in the patch pipette in the same way as the recombinant α -CaMK-II (See Fig 3.4). The evidence that CaMK-II can directly phosphorylate GABA_A receptors comes from analysis of GST-fusion proteins (McDonald and Moss, 1994; McDonald and Moss, 1997). The results indicated that CaMK-II could phosphorylate $\beta 1/2$ and 3 receptor subunits and $\gamma 2S/L$ at previously identified consensus sites on the intracellular loop between TM 3 and 4 (See Chapter 1, Table 1.1). It should be noted however that fusion proteins are not directly comparable to full-length, native receptor subunits. For example the $\beta 2$ subunit cannot be functionally modulated by PKA within HEK293 cells although it can be phosphorylated *in vitro* (McDonald *et al.*, 1998). It therefore needs to be confirmed that full-length native GABA_A receptors can be phosphorylated by CaMK-II. It is possible that within neurones CaMK-II does not phosphorylate the GABA_A receptor directly, acting instead, through another kinase or signalling pathway.

The signalling cascade involved in forms of excitatory plasticity such as LTP requires CaMK-II for induction. It has been revealed there are two mechanisms of action involved in CaMK-II modulation of AMPA receptors (Poncer *et al.*, 2002). One involves a direct phosphorylation of the GluR1 subunit at Ser⁸³¹ (Barria *et al.*, 1997) and the other involving an insertion of AMPA receptors into the postsynaptic membrane. This insertion of receptors does not occur through direct phosphorylation but through CaMK-II dependent activation of a downstream-signalling cascade involving Ras/Rap GTPases (Hayashi *et al.*, 2000; Zhu *et al.*, 2002). So CaMK-II can act both directly and indirectly to modulate AMPA receptor function. It is interesting, however, that CaMK-II modulation of AMPA receptors can be observed in HEK cells (Barria *et al.*, 1997), suggesting there is a key difference between CaMK-II modulation of AMPA and GABA_A receptors.

CaMK-II is found in high levels throughout the brain and has a large number of potential substrates (Hudmon and Schulman, 2002). It may be that in order to phosphorylate the GABA_A receptor or other targets successfully, it has to be targeted to the correct location. Anchoring proteins can restrict the subcellular location, at which a signalling molecule like CaMK-II exerts its effect, controlling which substrates are phosphorylated, by the location of the kinase (Bayer and Schulman, 2001; Schulman, 2004; Colbran, 2004). Such anchoring proteins can also have a direct effect on the activity of the kinase. The interaction between CaMK-II and the NMDA receptor is one example of this. Binding of CaMK-II to the NMDA receptor acts in the same way as Ca^{2+}/CaM binding, allowing CaMK-II to be active when bound to the NMDA receptor. In this way CaMK-II is localised to a specific location and its activity is altered through an interaction with a binding partner (Bayer *et al.*, 2001).

A number of potential CaMK-II anchoring proteins have been identified which localise the kinase to targets of phosphorylation (See Chapter 1). E.g. CaMK-II binding to the NMDA receptor (Strack *et al.*, 2000; Bayer *et al.*, 2001) localising the kinase at the excitatory post-synaptic density (PSD). In addition a non-catalytic subunit of CaMK-II has been identified which can localise CaMK-II to the sarcoplasmic reticulum in muscle cells (Bayer *et al.*, 1998). It is thought that there are probably a number of other proteins with a similar function that have yet to be identified (Bayer and Schulman, 2001; Griffith *et al.*, 2003; Colbran, 2004).

Other serine/threonine kinases like PKC are known to interact with anchoring proteins at the GABA_A receptor, e.g. RACK-1, which binds to the GABA_A receptor enhancing PKC phosphorylation of Ser⁴⁰⁹ (Brandon *et al.*, 2002b). Furthermore the anchoring protein, AKAP150 can also directly bind the GABA_A receptor and has

been shown to be critical for PKA mediated phosphorylation of the β 3 subunit (Brandon *et al.*, 2003). Currently relatively little is known about the scaffolding and signalling complexes that exist for the GABA_A receptor at inhibitory synapses. It may be that in order for CaMK-II to exert its effect, an additional anchoring protein may be required which is not present in HEK cells or alternatively, that CaMK-II phosphorylates another substrate closely linked to the GABA_A receptor which then goes on to modulate GABA_A receptor function.

It may be possible to determine in a neuronal preparation which subunits are important for CaMK-II modulation through transfection of neurones or the use of knock-out animals lacking specific subunits of the GABA_A receptor. It seems a neuronal system or neuronal cell-type environment may be crucial to reliably determine the signalling pathways and mechanisms of action involved in CaMK-II dependent modulation of GABA_A receptors. It is also important to try to determine if CaMK-II directly phosphorylates the GABA_A receptor, or if it acts through another kinase or signalling pathway.

3.4 CONCLUSIONS

- Intracellular dialysis of two different preparations of pre-activated CaMK-II failed to modulate $\alpha 1\beta 2/3\gamma 2$ GABA_A receptors.
- Co-transfection with α -CaMK-II cDNA and intracellular dialysis of CaM and Ca²⁺ fails to modulate GABA_A receptors expressed in HEK293 cells.
- Intracellular dialysis of CaM alone also had no effect on the amplitude of GABA mediated currents.
- Recombinant GABA_A receptors expressed in HEK293 cells cannot be modulated by CaMK-II.

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Chapter Four:

<u>CaMK-II dependent modulation of GABA_A receptors in</u> <u>cerebellar granule cells.</u>

4.1 INTRODUCTION

CaMK-II has previously been shown to modulate GABA_A mediated currents in a number of neuronal preparations (Kano *et al.*, 1992; Wang *et al.*, 1995; Kano *et al.*, 1996; Aguayo *et al.*, 1998; Kawaguchi and Hirano, 2002; Mizoguchi *et al.*, 2003; Wei *et al.*, 2004). In order to confirm these results it was necessary to repeat the experiments of the previous chapter in a neuronal cell type, given the lack of significant modulation observed in HEK293 cells. The same protocols that were applied in chapter three were adapted for cerebellar granule cells maintained in culture. The aim was to determine if CaMK-II can modulate GABA_A receptors in this cell type and if so, to characterise this modulation.

Intracellular dialysis of pre-activated α -CaMK-II has been previously shown to increase the peak amplitude of whole-cell GABA currents and IPSCs recorded in rat spinal dorsal horn neurones, hippocampal pyramidal and cerebellar Purkinje cells (Wang *et al.*, 1995; Kano *et al.*, 1996). Calcium dependent increases in the peak amplitude of GABA_A mediated currents that could be blocked by CaMK-II inhibitors have also been observed in mouse cortical neurones and hippocampal pyramidal neurones (Aguayo *et al.*, 1998; Mizoguchi *et al.*, 2003; Wei *et al.*, 2004). As CaMK-II dependent modulation of GABA_A occurs in a range of different neuronal cell types, it seemed reasonable to assume that CaMK-II dependent modulation of GABA_A receptors might be observed in cerebellar granule cells (CGCs). CGCs are thought to express a wide variety of different GABA_A receptor subunits (Wisden *et* *al.*, 1996) so if only certain subunits of GABA_A receptors could be modulated by CaMK-II there was a high probability that they would exist in CGCs. This array of subunits could be important, since some GABA_A receptors are differentially sensitive to phosphorylation. For example, the expression of different β subunits has been shown using heterologous expression systems and in neuronal preparations to alter the response to PKA phosphorylation (McDonald *et al.*, 1998; Nusser *et al.*, 1999).

Equally important in determining the phosphorylation state of GABA_A receptors, and therefore potential changes in function, are protein phosphatases, which act to dephosphorylate the GABA_A receptor itself and can also regulate CaMK-II activity. Wang and colleagues (1995) demonstrated that in isolated spinal dorsal horn neurones the addition of calyculin A, an inhibitor of protein phosphatase 1 (PP1) and protein phosphatase 2A (PP2A), caused an increase in the peak amplitude of whole-cell GABA currents recorded similar to that observed after the addition of pre-activated CaMK-II. Although PP1, PP2A and protein phosphatase 2B (PP2B)/calcineurin have been shown to directly modulate GABA_A receptor mediated currents (Wang *et al.*, 2003; Jovanovic *et al.*, 2004; Terunuma *et al.*, 2004; Sanchez *et al.*, 2005), protein phosphatases can also directly modulate CaMK-II activity. For example in neurones, phosphatases can alter the levels of Thr²⁸⁶ autophosphorylation and also the inhibitory autophosphorylation sites Thr^{305/6} (Fukunaga *et al.*, 1989; Fukunaga *et al.*, 1993; Kasahara *et al.*, 1999; Colbran, 2004), which will have indirect consequences for GABA_A receptor function.

The dynamic balance between phosphatase and kinase activities will determine the overall phosphorylation state of the receptor and result in altered function accordingly. Therefore the phosphatase inhibitor okadaic acid (OA) was used to determine the role of various phosphatases in cerebellar granule cells in altering GABA_A receptor function. At different concentrations, OA inhibits various protein phosphatases (Bialojan and Takai, 1988) and so the relative contribution of different protein phosphatases in modulating GABA_A receptor function could be determined. The relative balance between CaMK-II activity and phosphatase activity, under basal conditions, could then be determined in CGCs under these particular culture conditions.

4.2 **RESULTS**

Rat cerebellar granule cells were cultured in a 20 mM KCl supplemented, serum free medium and recorded from after 10-14 days in vitro (DIV, See Fig 2.1B). A recombinant form of α -CaMK-II, (Brickey et al., 1990; Suzuki-Takeuchi et al., 1992) 1200-1500 nM (40-50 ng/µl), was pre-incubated in 2 mM CaCl₂, 1.2 µM CaM, 0.4 mM ATP-y-S, 10 mM MgCl₂ in a total volume of 16-20 µl (See Chapter 2 and 3.2.1). This was then transferred to the patch pipette solution at a dilution of at least 1:12 and maintained on ice throughout the recording (Brickey et al., 1990; Suzuki-Takeuchi et al., 1992; McGlade-McCulloh et al., 1993). Controls consisted of using the same patch pipette solution containing pre-activation buffer diluted at the same ratio but without α -CaMK-II present or with heat-inactivated α -CaMK-II (95°C for 15 min). Whole-cell patch clamp recordings were then made in response to 10 μ M GABA applied intermittently for a period of 15-20 min. Cerebellar granule cells Their relatively compact morphology results in have very small cell bodies. recordings with very low noise and high fidelity for voltage clamp; however, due to their size, recordings of over 20 min were often difficult to maintain.

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4.2.1 Cerebellar granule cells in culture and modulation by α-CaMK-II

The inclusion of 100 nM α -CaMK-II in the patch pipette solution and subsequent dialysis into the cell resulted in a significant increase in the peak amplitude of GABA currents to 204.9 ± 24.7 % (n = 7) as compared to control currents, 96.5 ± 4.2 % (n = 7, Fig 4.1A) 10 min into the recording. A reduction in the concentration of α -CaMK-II to 35 nM resulted in a smaller, but still statistically significant, increase in the peak amplitude of GABA currents recorded to 127.3 ± 4.7 % (n = 7, Fig 4.1A) at t = 10 min. It appeared therefore that the effect of α -CaMK-II was dose-dependent and reached a plateau within 10 min. As a second control, α -CaMK-II was heat-inactivated and applied in the same way (100 nM). In this case no increase was observed compared to control currents (93 ± 7.3 %, n = 7, Fig 4.1A) at t = 10 min.

To confirm that this effect is present in different neuronal cell types, 85 nM pre-activated α -CaMK-II was added to the patch pipette solution and whole-cell currents were recorded from cortical pyramidal neurones in culture (DIV6-8). Reducing the concentration to 85 nM improved the quality and duration of recordings whilst producing an easily observed effect. Intracellular addition of 85 nM α -CaMK-II led to a significant increase in the peak amplitude of GABA currents recorded to 122.3 ± 5.3 % (n = 14, Fig 4.1C) at t = 10 min as compared to the equivalent time in control, 90.2 ± 4.1 % (n = 9, Fig 4.1C). The increase in this case appeared to be proportionately smaller than in CGCs and diminished to control levels within 25 min. In CGCs the effect remained stable and did not decay, although recordings of over 20 min were not routinely possible.

In order to further characterise the nature of this up-regulation of $GABA_A$ receptor function the dose-response relationship to GABA was determined under control conditions at t = 10 min and in the presence of 85 nM α -CaMK-II. The EC₅₀ value for GABA in the presence of α -CaMK-II was 6.3 ± 0.5 μ M (n = 13, Fig 4.2), and in control conditions, it was not significantly different, at 5.9 ± 0.7 μ M (n = 12, Fig 4.2). Indicating there was no change in the potency of GABA at these receptors. There was also no significant difference in the Hill slope (Control = 0.85 ± 1.7, CaMK-II 0.93 ± 0.6). However the maximal response to GABA appeared to have significantly increased from 127.4 ± 15.1 pA/pF to 212.1 ± 26.1 pA/pF.





Fig 4.1: α-CaMK-II modulation of whole-cell GABA currents recorded from cerebellar granule and cortical pyramidal cells in culture.

Peak amplitudes of GABA-mediated currents were recorded after application of 10 μ M GABA to cerebellar granule or cortical pyramidal neurones in culture. GABA was re-applied at 2 min intervals for up to 30 min. All currents were normalised to the peak response of the GABA current measured during the first 3-4 min (= 100 %) after achieving the whole-cell configuration (t = 0 min). Control recordings were made with normal patch pipette solution supplemented with the pre-activation buffer without α -CaMK-II.

(A) GABA currents recorded from cerebellar granule neurones using patch pipette solution supplemented with 100 nM pre-activated α -CaMK-II (n = 7), 35 nM α -CaMK-II (n = 7), 100 nM heat-inactivated (HI) α -CaMK-II (n = 7) or in the absence of CaMK-II (Control, n = 6).

(B) Representative 10 μ M GABA-activated currents recorded from a cerebellar granule cell at t = 2 and 10 min following internal dialysis with 100 nM α -CaMK in the patch pipette solution

(C) GABA currents recorded from cortical pyramidal neurones in the absence (Control, n = 14) or presence of 85 nM α -CaMK-II introduced into the cells via the patch pipette solution (n = 9).

All data points = mean \pm s.e.

Significance is indicated by the symbol * (100 nM) * (35 nM), CaMK-II compared to control and *, CaMK-II (35 nM and 100 nM) compared to HI CaMK-II. P < 0.05, t-test, ANOVA.



Fig 4.2: The concentration response relationship of GABA in the absence and presence of α -CaMK-II.

Peak amplitudes GABA-mediated currents were recorded in response to various concentrations of GABA normalised to the cerebellar granule cell's membrane capacitance (pA/pF).

GABA currents were recorded with a patch pipette solution supplemented with 85 nM pre-activated α -CaMK-II (n = 13) and normal patch pipette solution supplemented with pre-activation buffer without CaMK-II (Control, n = 12). Currents were recorded at t = 10 min after the effect of α -CaMK-II had reached a plateau.

All data points = mean \pm s.e. Significance is indicated by the symbol *, CaMK-II compared to control. P < 0.05, t-test.

4.2.2 Modulation of GABA_A receptors in cerebellar granule cells by endogenous CaMK-II.

CaMK-II is activated by the binding of Ca²⁺/CaM, which allows substrate access to the catalytic site of the enzyme. Autophosphorylation at Thr²⁸⁶ can then occur between adjacent subunits in the holoenzyme that have bound Ca²⁺/CaM (Hudmon and Schulman, 2002). The formation of a holoenzyme is critical for this autophosphorylation to occur within the cell (Hanson *et al.*, 1994; Mukherji and Soderling, 1994; Shen and Meyer, 1998). The recombinant form of CaMK-II used in this study lacks the association domain of the full-length α subunit and so it cannot form a holoenzyme. However it has the same catalytic properties as native CaMK-II and can phosphorylate the same substrates (Suzuki-Takeuchi *et al.*, 1992; Lledo *et al.*, 1995). Incubation of Ca²⁺/CaM and ATP- γ -S *in vitro*, in a small volume, enables autophosphorylation to occur.

The CaMK-II inhibitor, KN-93, prevents Ca²⁺/CaM binding and so in this way prevents Ca²⁺/CaM from increasing CaMK-II activity (Sumi *et al.*, 1991). Application of KN-93 to the pre-activation buffer (15-20 μ M) should prevent Ca²⁺/CaM binding and autophosphorylation at Thr²⁸⁶ from occurring and therefore prevent CaMK-II activation. When α -CaMK-II and the pre-activation buffer are then diluted into the patch pipette solution the α -CaMK-II that then dialyses into the cell will be in its basal state and therefore inactive. Endogenous Ca²⁺/CaM within the cell is unlikely to activate the recombinant α -CaMK-II because it cannot form a holoenzyme and in addition KN-93 will still be present in the patch pipette solution at a concentration of 500 nM.

The addition of the pre-activation buffer containing KN-93 (500 nM in patch pipette solution) and α -CaMK-II (35 nM) to the patch pipette solution resulted in a

significant depression of the peak amplitude of whole-cell GABA currents recorded to 71.9 ± 4.6 % at t = 10 min (n = 10, Fig 4.3A), compared to controls of, 96.5 ± 5.1 % (n = 6, Fig 4.3A). This suggested that endogenous CaMK-II is active and maintaining GABA_A receptor function under basal conditions. The addition of KN-93 alone led to a similar significant decrease in the amplitude of currents recorded to 69.3 ± 7.6 % at t = 10 min. The fact that the depression with or without α -CaMK-II present was not significantly different suggests that pre-incubation in KN-93 was able to prevent the activation of exogenously applied, as well as endogenous α -CaMK-II.

As a control the inactive analogue of KN-93, KN-92 was added at the same concentration (500 nM) to the patch pipette. In this case there was no significant difference from controls over 16 min (91.2 \pm 8.5 % at t = 10 min). Taken together, this suggests there is a basal level of Ca²⁺/CaM dependent activity within these cells that can be inhibited by KN-93. Inhibition of CaMK-II activity by KN-93 results in a depression of GABA currents, presumably due to a loss of CaMK-II dependent upregulation of GABA_A receptor function.





Fig 4.3: Modulation of whole-cell GABA currents recorded from rat cerebellar granule cells in culture by KN-93.

Peak amplitudes of GABA-mediated currents were recorded after application of 10 μ M GABA to cerebellar granule cells in culture. GABA was re-applied at 2 min intervals for up to 20 min. All currents were normalised to the peak response of the GABA current measured during the first 3-4 min (= 100 %) after achieving the whole-cell configuration (t = 0 min). Control recordings were made with normal patch pipette solution supplemented with the pre-activation buffer without α -CaMK-II.

(A) GABA currents recorded with: normal patch pipette solution (Control, n = 6); patch pipette solution supplemented with 500 nM KN-93 (n = 10); and patch pipette solution supplemented with 35 nM CaMK-II and 500 nM KN-93, pre-incubated during the pre-activation reaction (n = 10).

(B) GABA currents recorded with a patch pipette solution supplemented with 500 nM KN-92 (n = 6) compared to normal patch pipette solution (Control, n = 6) and patch pipette solution with 500 nM KN-93 (n = 10, taken from A).

All data points = Mean \pm s.e.

Significance is indicated by the symbol *, KN-93 compared to control and the symbol *, KN-92 compared to KN-93.

P < 0.05, t-test, ANOVA.

4.2.3 Protein phosphatase activity in cerebellar granule cells in culture.

Regulation of ion channel function by phosphorylation is often bi-directional involving protein kinases and phosphatases. It is the balance of activity between the two that ultimately controls the functional effect. In this case it appeared that α -CaMK-II was able to functionally modulate GABA_A receptors in granule cells under basal conditions in this culture. In order to see if this modulation was bi-directional the phosphatase inhibitor okadaic acid (OA) was added to the patch pipette. Okadaic acid is a general phosphatase inhibitor but at different concentrations affects a different range of phosphatases. At 10 µM, OA blocks the activity of PP1, PP2A and PP2B/calcineurin, but at 1 µM, only the activity of PP1 and PP2A would be affected (Bialojan and Takai, 1988). In this way the relative contribution of different phosphatases to the regulation of GABA_A receptor function could be determined in this cell type (Lieberman and Mody, 1994; Kurz *et al.*, 2001). In addition, inclusion of KN-93 could be used to determine the contribution of CaMK-II activity to the effect of blocking various protein phosphatases.

Including 10 μ M OA in the patch pipette resulted in a significant increase in the peak amplitude of whole-cell GABA currents recorded within 4 min to 126 ± 11.5 % as compared to control, 95.6 ± 3.3 % (n = 7, Fig 4.4A). This increase was relatively short-lived and there was no significant difference from control after 15 min. Lowering the concentration of okadaic acid to 1 μ M appeared to have no significant effect on the increase in amplitude, 117.1 ± 4.3 % at t = 4 min (n = 6, Fig 4.4A). This suggested that PP2B/calcineuin activity played no significant role in the modulation of GABA currents observed in these cells under basal conditions.

This increase in amplitude could have been caused by a release of protein kinase activity from inhibition by protein phosphatases. A number of different kinases may be involved. As it had already been established that CaMK-II might be involved in altering GABA_A receptor function in these cells, the calcium chelator 1,2-bis(2-aminophenoxy)ethane-N,N,N',N'-tetraacetic acid (BAPTA) was added to the patch pipette to see what proportion of the effect was related to Ca²⁺ sensitive kinases. Inclusion of 20 mM BAPTA to the patch pipette with 5 μ M OA completely blocked the rise in amplitude and resulted in a significant depression to 75.8 ± 5.7 % (n = 5, Fig 4.4B). This indicated that all of the effect produced by blocking phosphatase activity was mediated through Ca²⁺ sensitive processes and confirmed that some of this activity maintained GABA_A receptor function under basal conditions.

In order to determine the role of CaMK-II in increasing GABA_A receptor function after block of phosphatase activity, the CaMK-II inhibitor KN-93 (500 nM) was also added to the patch pipette containing 1 µM of okadaic acid. This led to a decrease in the effect of OA, $99.3 \pm 3.5\%$ (n = 6, Fig 4.4C) at t = 4 min so that there was no significant difference from control, 95.6 ± 3.3 % (n = 6, Fig 4.4C). The addition of 500 nM KN-93 previously led to a significant depression of amplitude. This did not occur with OA included in the patch pipette. This suggests that another kinase may also have been involved in increasing GABA_A receptor activity when protein phosphatase activity was blocked. As the addition of 20 mM BAPTA led to depression of currents recorded this might suggest the additional а kinase/phosphatase is Ca^{2+} sensitive. As an additional control, recordings were made with 1 µM OA and the inactive analogue of KN-93, KN-92. In this case the effect of 1 μ M OA was not significantly different from 1 μ M OA alone, 125 \pm 6.7 % at t = 4 $\min(n = 6, Fig 4.4C).$

The increase in amplitude caused by OA addition in all cases was rapid but transient. There were no significant differences from control after 15 min. This is possibly an indication that other OA insensitive phosphatases were present that counteracted the effect of blocking PP1/2A but on a slower time-scale.



Fig 4.4: Modulation of whole-cell GABA currents recorded from cerebellar granule cells by okadaic acid.

Peak amplitudes of GABA-mediated currents were recorded after application of 10 μ M GABA to cerebellar granule in culture. GABA was re-applied at 2 min intervals for up to 20 min. All currents were normalised to the peak response of the GABA current measured during the first 3-4 min (= 100 %) after achieving the whole-cell configuration (t = 0 min). Control recordings were made with normal patch pipette solution supplemented with the pre-activation buffer without α -CaMK-II.

(A) GABA currents recorded with normal patch pipette solution (Control, n = 6); a patch pipette solution supplemented with 10 μ M okadaic acid (n = 7); and with a patch pipette solution supplemented with 1 μ M okadaic acid (n = 6).

(B) GABA currents recorded with a patch pipette solution supplemented with 20 mM BAPTA and 5 μ M okadaic acid (n = 5) compared to control (n = 6).

(C) GABA currents recorded with a patch pipette solution supplemented with 1 μ M okadaic acid and 500 nM KN-93 (n = 6) and with a patch pipette solution with 1 μ M okadaic acid and 500 nM KN-92 (n = 6) compared to control (n = 6).

All data points = Mean \pm s.e.

Significance is indicated by the symbol *. Each group (indicated by colour) is compared to control, P < 0.05, t-test, ANOVA.

4.2.4 Depolarisation of cerebellar granule cells

CaMK-II activity is increased in response to a Ca²⁺ signal. In order to determine if endogenous CaMK-II in CGCs could be modulated by an increase in internal Ca²⁺ concentration, CGCs were depolarised to 0 mV for 2 s to allow Ca²⁺ entry through voltage-gated calcium channels (VGCCs). In this case depolarisation resulted in a slight but significant depression of currents recorded over time to $80 \pm 4.5 \%$ (n = 9, Fig 4.5) at t = 10 min. It is possible that this was due to activation of Ca²⁺ sensitive kinases/phosphatases or signalling pathways other than CaMK-II (Wang *et al.*, 2003).

Calcium signalling is complex, controlling many cellular signalling complexes/pathways. Often slight changes in the amplitude and/or localisation of a calcium signal can alter significantly the target-signalling molecule that responds (Franks and Sejnowski, 2002; Augustine *et al.*, 2003). It appeared that depolarisation was not sufficient to increase endogenous CaMK-II activity or that any increase was occluded by activation of other Ca^{2+} sensitive signalling molecules.



Fig 4.5: Depolarisation of cerebellar granule cells.

Peak amplitudes GABA-mediated currents were recorded after application of 10 μ M GABA to cerebellar granule cells in culture. GABA was re-applied at 2 min intervals for up to 20 min. All currents were normalised to the peak response of the GABA current measured during the first 3-4 min (= 100 %) after achieving the whole-cell configuration (t = 0 min). Control recordings were made with normal patch pipette solution supplemented with the pre-activation buffer without α -CaMK-II.

GABA currents were recorded from cerebellar granule cells with normal patch pipette solution (Control, n = 6). This is compared to cells that were depolarised to 0 mV for 2 s (indicated by the arrow, n = 9).

All data points = Mean \pm s.e. Significance is indicated the symbol *, depolarisation compared to control. P < 0.05, t-test.

4.3 **DISCUSSION**

CaMK-II dependent modulation of GABA_A receptors has been previously observed in a number of neuronal cell types (Wang *et al.*, 1995; Kano *et al.*, 1996; Aguayo *et al.*, 1998; Kawaguchi and Hirano, 2002; Mizoguchi *et al.*, 2003; Wei *et al.*, 2004). In HEK293 cells it was not possible to observe any functional modification, however when recording GABA mediated currents from cerebellar granule cells the activity of CaMK-II had a significant effect on GABA_A receptor function. CaMK-II appeared to up-regulate GABA_A receptor function causing a significant increase in the peak amplitude of currents recorded. The lack of functional effect observed in HEK293 cells suggested that this modulation by CaMK-II required a "neuronal environment" to be operative.

Intracellular dialysis of pre-activated α -CaMK-II (100 nM) to cerebellar granule cells resulted in a significant potentiation of whole-cell currents recorded. This effect appeared to be concentration-dependent. Heat-inactivated α -CaMK-II had no significant effects compared to control suggesting it was CaMK-II activity that was responsible for the effect and not another non-specific mechanism (See Fig 4.1A). Intracellular dialysis of α -CaMK-II into cortical pyramidal neurones in culture also induced a significant increase in GABA mediated currents (See Fig 4.1C). This effect was comparatively smaller than that observed in granule cells suggesting a difference in sensitivity between the two cell types. This potentially could have been mediated by differences in GABA_A receptor subunits present or differences in the background levels of kinase and phosphatase activities within the two cell types.

Construction of GABA dose-response relationships in control conditions and in the presence of CaMK-II revealed a difference in the maximal response without an alteration in the GABA EC_{50} (See Fig 4.2). A change in the number of functional receptors at the cell surface, an increase in channel conductance or a change in the probability of opening could mediate this increase in the maximal response. Aguayo and colleagues (1998) studied Ca²⁺ dependent potentiation of GABA_A currents in mouse cortical neurones. This modulation could be blocked by addition of the CaMK inhibitor KN-62 and the CaM inhibitor W-7, indicating the likely They examined the GABA concentration-response involvement of CaMK-II. relationship in the presence of high Ca^{2+} (10 μ M) and saw no change in the GABA Using a saturating concentration of GABA they still observed a Ca²⁺ EC₅₀. dependent potentiation, indicating that an increase in the maximal GABAA receptor function had occurred. In cultured rat cortical neurones it has also been observed that mild stretch injury resulted in a potentiation of the maximal GABA current density (pA/pF - with no change in EC_{50}) that could be blocked by KN-93 (Kao et al., 2004). An increase in the overall charge transfer on activation of GABAA receptors could have significant effects on neuronal excitability. Modulation of the maximal response to GABA with no change to EC_{50} appears to be a common feature of phosphorylation dependent modulation of GABA_A receptor function. For example PKC dependent down-regulation of receptor function results in a depression of the maximal response to GABA without alteration of the EC₅₀ (Krishek et al., 1994; Brandon et al., 2000).

In the present study intracellular dialysis of the CaMK inhibitor KN-93 led to a significant decrease in the amplitude of currents recorded indicating that there is a certain level of Ca^{2+}/CaM dependent CaMK-II activity under basal conditions that acts to enhance GABA_A receptor function. Cerebellar granule cells in culture are often maintained in depolarising conditions (20 mM KCl) to increase CGC survival (Galli *et al.*, 1995; Borodinsky *et al.*, 2002; Borodinsky *et al.*, 2003; Linseman *et al.*, 2003). There is some controversy over how depolarisation increases cell survival but it seems that maintenance in chronically depolarising conditions significantly alters the development of these neurones. It would appear that basal calcium levels and calcium dynamics are also altered by such culture conditions (Moulder *et al.*, 2003).

Borodinsky *et al.* (2002; 2003) observed after addition of KN-93 that CaMK-II activity was involved in neuritogenesis and neuronal survival of CGCs grown under depolarising conditions. Linesman *et al.* (2003) also identified basal levels of CaMK-II activity in CGCs in culture under depolarising conditions that were responsible for increased neuronal survival through activation of myocyte enhancer factor 2 (MEF2) transcription factors. MEF2 transcription factors are thought to be involved in activity dependent neuronal survival (McKinsey *et al.*, 2002). CGCs cultured under depolarising conditions may therefore have significant levels of basal Ca²⁺/CaM dependent CaMK-II activity and this activity may mediate part of the observed increase in neuronal survival seen under these conditions (Hack *et al.*, 2004).

CaMK-II within the brain consists of two main isoforms, α/β . These isoforms are able to phosphorylate the same substrates but they differ in their localisation and sensitivity to Ca²⁺/CaM (Hudmon and Schulman, 2002). CaMK-II isolated from the cerebellum is thought to contain a higher proportion of β subunits compared to that isolated from the forebrain. Cerebellar CaMK-II has a two-fold higher affinity to Ca²⁺ and associates differently with subcellular structures (Miller

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and Kennedy, 1985). Under non-saturating levels of CaM, β -CaMK-II has a higher sensitivity to lower levels of Ca²⁺ (Brocke *et al.*, 1999).

Evidence suggests that the ratio of $\alpha:\beta$ CaMK-II may be important for CaMK-II modulation of AMPA receptors in terms of localisation and translocation (Shen *et al.*, 1998; Shen and Meyer, 1999; Thiagarajan *et al.*, 2002) even though both subunits can phosphorylate the same substrates. The role of the different isoforms of CaMK-II in modulation of inhibitory synaptic transmission has yet to be determined. Previous experiments have involved introduction of pre-activated α -CaMK-II (Wang *et al.*, 1995; Kano *et al.*, 1996). This along with the current study suggests α -CaMK-II in when applied to the cell through intracellular dialysis is capable of modulating GABA_A receptor function. However, it is thought that CGCs express a high level of β -CaMK-II and little or no α -CaMK-II. This would suggest that the basal activity observed in cerebellar granule cells on application of KN-93 is mediated through β -CaMK-II. Since β -CaMK-II has a higher affinity for Ca²⁺ it is possible it may be basely active within CGCs which is in line with experiments that suggested basal CaMK-II activity is involved in CGC survival when cultured under depolarising conditions (Borodinsky *et al.*, 2002; Linseman *et al.*, 2003).

It can be predicted that the role of the α and β subunits would have little significance on the effect observed in the current study as this involved preactivating CaMK-II *in vitro*. Pre-activated recombinant β -CaMK-II applied to the cell through the patch pipette would be predicted to behave in exactly the same way as pre-activated α -CaMK-II. The role of the α : β subunits within a CaMK-II heteromer would be significant in a physiological situation if CaMK-II is activated by an endogenous Ca²⁺ signal. The evidence suggests that basal CaMK-II activity within CGC cultures under depolarising conditions can modulate GABA_A receptor function. Analysis of CaMK-II activity by Fukanaga *et al.* (1989) in CGCs through examination of the proportion of CaMK activity that is Ca^{2+} independent (presumed to be autophosphorylated at Thr²⁸⁶), revealed that 4-5 % of CaMK-II in CGCs was Ca^{2+} independent. This level could be reduced to 1-2 % (reduction of 50 %) by removal of extracellular Ca^{2+} suggesting basal intracellular Ca^{2+} levels must allow a certain level of CaMK-II autophosphorylation at Thr²⁸⁶.

They also provided evidence for high levels of phosphatase activity within these cells, which could reduce Ca^{2+} independent activity rapidly after an initial stimulus. Application of OA increased the amount of Ca^{2+} independent activity over basal levels but this was not sustained and returned to baseline within 10-15 min. They proposed that PP1/PP2A strongly regulated CaMK-II activity in CGCs and that another OA insensitive protein phosphatase (PP2C) was responsible for the slow return of CaMK-II activity to baseline (Fukunaga *et al.*, 1989; Fukunaga *et al.*, 1993).

Analysis of CaMK-II activity in organotypic hippocampal slices also revealed a significant level of autophosphorylation under basal Ca^{2+} conditions that was reduced by the removal of extracellular Ca^{2+} and increased by treatment with OA. This suggested that the relatively high level of CaMK-II activity in basal conditions could allow for a bi-directional regulation of CaMK-II activity (Molloy and Kennedy, 1991).

Both Fukunaga *et al.* (1989) and Molloy and Kennedy (1991) looked at the relative proportions of Ca^{2+} -independent activity as a measure of CaMK-II activity. This reveals levels of CaMK-II autophosphorylation at Thr²⁸⁶ but does not reveal the

absolute levels of CaMK-II activity in the cell which would include Ca^{2+} independent and Ca^{2+} -dependent activity. The levels of Ca^{2+} -independent activity reported in these studies are low (<10%) however, even maximally autophosphorylated homogenates display only 45 % Ca^{2+} -independent activity using the methods used in these studies (Molloy and Kennedy, 1991). It is interesting though that both of these studies suggest a certain level of basal Ca^{2+} -independent activity (autophosphorylation at Thr²⁸⁶) which must reflect a basal level of CaMK-II activity which can be up or down-regulated by the activity of protein phosphatases.

Applying these details to CaMK-II modulation of CGC GABA currents, it can be speculated that endogenous CaMK-II within cerebellar granule cells, which predominantly express the β isoform of CaMK-II, can be basally active in a Ca²⁺/CaM-dependent manner and that this basal activity acts to up-regulate GABA_A receptor function.

The amplitude of GABA mediated currents within CGCs in this study was altered by the phosphatase inhibitor OA. Different concentrations of OA were used to determine which protein phosphatases regulated GABA_A receptor function. Inclusion of OA in the patch pipette resulted in an increase in the amplitude of GABA mediated currents that was the same regardless of the concentration of OA used; suggesting that PP1/PP2A modulated GABA_A receptor function but that PP2B did not. The increased amplitude returned to baseline within 15 min indicating that another OA insensitive protein phosphatase may also be involved. This time-course and sensitivity to OA of this effect on GABA_A receptors was very similar to the observed effects of OA on Ca²⁺-independent CaMK-II activity levels (Fukunaga *et al.*, 1989).

The increased amplitude caused by the block of these phosphatases was removed by the addition of BAPTA. Indeed, there was a decrease similar to that observed with KN-93, suggesting Ca^{2+} sensitive kinase/phosphatase activities mediated the increase in GABA_A receptor function observed on addition of OA. Addition of KN-93 and OA resulted in no significant increase or decrease in amplitude. Suggesting CaMK activity was partly responsible for an increase in amplitude initiated by protein phosphatase blockade but that other Ca²⁺ sensitive kinases/phosphatases were also involved. It appears that GABA_A receptor function can be bi-directionally altered by CaMK-II activity and protein phosphatase activity.

It may be that for CGCs maintained under depolarising conditions, a relatively high basal Ca²⁺ level and/or a proportionately high level of β -CaMK-II, may act to allow a significant level of basal CaMK-II activity, which can modulate GABA_A receptor function. Blocking the activity of protein phosphatases may act to increase the level of CaMK-II activity, e.g, PP1 is known to dephosphorylate CaMK-II (Colbran, 2004), or may act directly on the GABA_A receptor itself. The presence of PP1 has been shown to alter the sensitivity of CaMK-II to Ca²⁺ levels (Bradshaw *et al.*, 2003). Inhibition of PP1 has been linked to increased CaMK-II activity in a number of systems (Brown *et al.*, 2000; Kawaguchi and Hirano, 2002; Colbran, 2004). PP1 is also thought to modulate GABA_A receptor function directly (Terunuma *et al.*, 2004). Moreover, PP2A has also been suggested to modulate CaMK-II activity (Fukunaga *et al.*, 1989; Colbran, 2004) and GABA_A receptor activity directly (Jovanovic *et al.*, 2004).

Basal GABA_A receptor phosphorylation by CaMK-II would require CaMK-II to be localised to inhibitory synapses. Attempts to identify the location of CaMK-II subunits have suggested that the α subunit can be localised to excitatory but not

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inhibitory synapses within the CA1 region of the hippocampus, the thalamus and the cerebral cortex (Liu and Jones, 1996; Liu and Jones, 1997). However, in the rat basolateral amygdala α-CaMK-II has been identified postsynaptically at GABAergic synapses (McDonald et al., 2002). Differences between papers may be due to the different antibodies used and slight variations in immunocytochemical techniques. In addition, recent evidence suggests that CaMK-II is capable of very rapid translocation (Shen and Meyer, 1999; Gleason et al., 2003). It has also been suggested CaMK-II can alter localisation and form clusters in response to ischaemic stress and that the process of preparing tissue for staining may alter CaMK-II localisation (Dosemeci et al., 2000; Lengyel et al., 2001). Depolarisation and excitotoxic stress appears to increase the apparent accumulation of α -CaMK-II at the PSD (Tao-Cheng et al., 2002). Early on in development the β -CaMK-II is more prevalent with an increase in α subunit occurring in the rat in the 2nd and 3rd postnatal week (Burgin *et al.*, 1990). There is also variation in the relative proportion of $\alpha:\beta$ in different tissues (Miller and Kennedy, 1985; Vallano, 1990). Since the ratio of $\alpha:\beta$ subunits can alter the localisation of the kinase, it may be that at different developmental time points the distribution of CaMK-II observed may also be different.

Maintenance of CGC cultures in depolarising conditions not only alters CaMK-II activity; it can also alter the expression of ligand-gated ion channels (Vallano *et al.*, 1996; Mellor *et al.*, 1998; Ives *et al.*, 2002). Ives *et al.* (2002) studied mRNA levels and protein levels with immunocytochemistry of different GABA_A subunits in mouse cerebellar granule cells cultured in 25 mM KCl, or 5 mM KCl. They reported that under depolarising conditions, αl , $\alpha \delta$ and $\beta 2$ expression was greatly reduced and they failed to detect any $\beta 2$ subunit protein. In addition there was an increase in $\alpha 3$ and $\beta 3$ subunit expression with no difference in $\alpha 2$. Given that $\beta 2$ shows a preference to assembly with $\alpha 1$ and $\beta 3$ with $\alpha 2/3$ (Benke *et al.*, 1994) it could be predicted that CGCs grown in 20 mM KCl medium, may express predominantly $\alpha 2/3\beta 3$ subunit-containing receptors. In more physiological conditions (5 mM KCl) there may be a mixture of $\alpha 1/\beta 2$ and $\alpha 2/3\beta 3$ subunit-containing receptors.

The β 3 subunit is, in many parts of the brain, expressed earlier in development with $\beta 2$ expression increasing at later developmental stages (Laurie et al., 1992). In support of a crucial role for β 3 in early development β 3 knock-out mice display severe developmental impairments (Homanics et al., 1997; DeLorey et al., 1998). Ives et al. (2002) suggested that CGCs maintained under depolarising conditions are "arrested" in their development and that they represent an immature stage in CGC development. This is supported by the fact that these cells are generally electrically silent, characteristic of a lack of fully functional synapses. In further support of this idea, depolarisation appears to prevent of subunit expression in mouse cerebellar granule cultures which is known to be expressed in mature CGCs (Mellor et al., 1998). It should be noted though that ∞ expression has been reported in rat CGC cultures under depolarising conditions (Thompson and Stephenson, 1994). Rat CGC cultures appear to behave slightly differently and so the alteration in GABA_A subunit expression may not be as pronounced as in mouse cultures. However rat CGCs are similar to mouse CGCs in that they are electrically silent and so they may also represent an earlier stage in development and express different levels of GABA_A receptor β subunits accordingly.

The age of the animal used for culture will also have an impact on the GABA_A subunits present. In the current study P0 rats were used. As $\alpha 6/\delta$

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expression does not occur until the end of the 1st postnatal week (Wisden *et al.*, 1996) the cultures used in these experiments would probably not have substantially expressed these subunits. CaMK-II has been shown to phosphorylate β and γ subunits *in vitro* (Machu *et al.*, 1993; McDonald and Moss, 1994; McDonald and Moss, 1997). It is reasonable to suggest that the cultures used in the current experiments may have expressed predominantly $\alpha 2/3\beta 3\gamma 2$ subunit-containing receptors and so this suggests that $\beta 3$ and $\gamma 2$ may be good candidates for CaMK-II modulation in this cell type.

In the current study, an attempt was also made to depolarise CGCs to encourage Ca²⁺ entry via VGCCs to see if endogenous CaMK-II activity could be increased thereby increasing the amplitude of GABAA currents. However, depolarisation led to a small decrease in amplitude. There are several possible reasons why depolarisation did not appear to activate endogenous CaMK-II. Calcium signalling is complex, Ca^{2+} signals are often under tight spatial and temporal control to allow a neurone to use multiple Ca²⁺ sensitive signalling pathways (Augustine *et al.*, 2003). The source and localisation of Ca^{2+} entry can have a profound effect on CaMK-II activation (Franks and Sejnowski, 2002). CaMK-II can respond with differing sensitivities to Ca²⁺ signals depending on the proportion of $\alpha:\beta$ subunits within the heteromer. In addition CaMK-II has been shown to respond not only to the amplitude of a Ca^{2+} signal but also to its frequency (De Koninck and Schulman, 1998). Complex processes within the cell can also regulate the availability of CaM (Gnegy, 2000). Therefore the link between depolarisation and increases in the intracellular Ca^{2+} concentration is not always straightforward and Ca²⁺ entry, in itself, is not a guarantee that CaMK-II will be activated.

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Global Ca^{2+} entry may have activated a number of different Ca^{2+} sensitive processes that may have had direct or indirect effects on GABA_A receptor function. Using different concentrations of OA it appeared that PP2B/Calcineurin had little basal effect on GABA_A receptor function. However, Ca^{2+} entry may have activated this phosphatase. Calcineurin has been shown to decrease the amplitude of IPSCs through an interaction with the $\gamma 2$ subunit (Wang *et al.*, 2003). Previously the alteration of intracellular Ca^{2+} levels has been associated with both increased (Kano *et al.*, 1992; Aguayo *et al.*, 1998) and decreased GABA_A receptor function (Martina *et al.*, 1994; Stelzer and Shi, 1994). This may be due to GABA_A receptor heterogeneity, but also to activation of different downstream signalling pathways. In addition the Ca^{2+} sensitive kinase PKC has also been shown to up or down-regulate GABA_A receptor function depending on the cell type and the GABA_A receptor subtypes present (Krishek *et al.*, 1994; Lin *et al.*, 1994; Lin *et al.*, 1996; Brandon *et al.*, 2000; Jovanovic *et al.*, 2004).

In addition to the problems associated with the complexity of Ca^{2+} signalling it has been reported by Moulder *et al.* (2003) that CGCs maintained in depolarising conditions have a reduced calcium channel density. Ca^{2+} entry on depolarisation may be significantly reduced after maintenance in these conditions. The authors also suggested that reduced Ca^{2+} entry, presynaptically, might be responsible, in part, for the electrical silence of these cells.

In summary it appears that CaMK-II is capable of potentiating GABA currents recorded from cerebellar granule cells. There also appears to be a certain level of basal CaMK-II activity in these cells which acts to enhance GABA_A receptor function under resting conditions. In order for CaMK-II modulation of GABA_A receptors to play a physiological role, CaMK-II activity must be able to significantly
modulate synaptic function. Since CGCs are electrically silent and morphologically immature when maintained under depolarising conditions it was necessary to alter the CGC culture conditions to those that are more physiologically representative (5 mM KCl) in order to determine if CaMK-II activity can indeed modulate synaptic GABA_A receptors.

4.4 CONCLUSIONS

- α-CaMK-II activity was capable of potentiating GABA_A receptor function in cerebellar granule and cortical pyramidal neurones.
- Within CGCs, CaMK-II activity appeared to play a basal role in maintaining GABA_A receptor function.
- Protein phosphatase activity, probably involving PP1/PP2A, modulated CaMK-II activity towards GABA_A receptors either acting on CaMK-II or the GABA_A receptor directly.

<u>Chapter Five:</u> <u>Modulation of inhibitory synaptic transmission by</u> <u>*α*-CaMK-II</u>

5.1 INTRODUCTION

GABA_A receptors mediate the majority of fast synaptic inhibition in the brain. Their activity can shape the output of a given neurone and the general excitability of a network and their function is thought to be crucial for the generation of network oscillations relevant to behaviour (Gaiarsa et al., 2002). Inhibitory synaptic transmission has also been shown to be altered in certain pathological conditions such as epilepsy (Macdonald et al., 2004). At the receptor level, covalent modification by phosphorylation has been demonstrated to modulate GABAA receptor mediated synaptic transmission (Wang et al., 1995; Wang et al., 2003b; Wei et al., 2004; Jovanovic et al., 2004), although there is controversy over the exact effects observed in different cell types (Nusser et al., 1999; Poisbeau et al., 1999). Phosphorylation appears to play an important role in long-term plasticity of both excitatory and inhibitory synapses (Soderling and Derkach, 2000; Kittler and Moss, 2003; Colbran and Brown, 2004; Lüscher and Keller, 2004). For CaMK-II modulation of GABA_A receptors to have any role in shaping the activity of a neurone or network it is important to establish if CaMK-II can modulate inhibitory synaptic transmission. Therefore this chapter addressed this issue by examining CaMK-II dependent effects on GABA_A receptor mediated synaptic transmission in neurones.

There are a number of examples of kinase and phosphatase activities altering fast synaptic inhibition. PKC activity has been shown to increase miniature inhibitory postsynaptic current (mIPSC) amplitudes and protein phosphatase 2A

(PP2A) has been shown to subsequently reverse this effect (Jovanovic *et al.*, 2004), without any changes to mIPSC decay time constants. Using recombinant GABA_A receptors expressed in *Xenopus oocytes*, PKC activity can lead to a depression of GABA currents (Krishek *et al.*, 1994) with no change in the time constants for decay. More recent evidence suggests PKC can have different functional effects in different neuronal cell types (Poisbeau *et al.*, 1999). PKA activity can also have different functional effects in different cell types and has been shown to lead to an increase, no change, or a depression of IPSC amplitudes. The decay time course of IPSCs was in some cases prolonged, but in others remaining unaffected (Nusser *et al.*, 1999).

Alteration in the decay phase of an IPSC appears to be a common mechanism for phosphorylation-dependent modulation of inhibitory synaptic transmission (Nusser *et al.*, 1999; Poisbeau *et al.*, 1999; Hinkle and Macdonald, 2003). In addition to the lengthening of IPSC decay times by PKC and PKA, the phosphatase calcineurin (CaN)/PP2B has also been shown to regulate IPSC decay times (Martina *et al.*, 1996; Jones and Westbrook, 1997; Sanchez *et al.*, 2005). However under certain conditions CaN/PP2B causes a reduction in mIPSC amplitude without affecting decay times (Wang *et al.*, 2003a). As well as Ser/Thr kinases, tyrosine kinase phosphorylation has been shown to up-regulate GABA_A receptor function (Moss *et al.*, 1995) and can increase mIPSC amplitude without apparent changes in their decay times (Boxall, 2000; Mizoguchi *et al.*, 2003; Ma *et al.*, 2003).

Similarly CaMK-II activity has been shown to alter fast synaptic inhibition. Wang and colleagues (1995) applied pre-activated α -CaMK-II to hippocampal CA1 neurones through inclusion in the patch pipette and observed an increase in evoked IPSP amplitudes. They also observed a reduction in desensitisation, a possible mechanism for alteration of IPSC decay (Jones and Westbrook, 1996). Kano *et al.* (1996) observed increases in the amplitude of mIPSCs recorded from Purkinje neurones without a change in the decay time constant after application of preactivated α -CaMK-II. In addition, Wei and colleagues (2004) observed a calciumdependent increase in spontaneous IPSC (sIPSC) amplitudes that could be blocked by CaMK-II inhibitors.

The evidence suggests that CaMK-II is capable of altering GABA_A receptor mediated synaptic transmission through changes in amplitude and possibly the rate of desensitisation of IPSCs. However phosphorylation-dependent effects on inhibitory synaptic transmission mediated by other kinases/phosphatases have suggested considerable variation in the effect observed amongst different cell types.

Cerebellar granule cells (CGCs) maintained in depolarising conditions do not display synaptic activity and so these cells were cultured in more physiological conditions in order to examine synaptic transmission. Mixed cerebellar cultures maintained in physiological (5 mM KCl) conditions are able to form inhibitory synapses. *In vivo* cerebellar granule cells receive inhibitory inputs from Golgi cells and Purkinje cells receive inhibitory input from stellate and basket cells (Wisden *et al.*, 1996; Nakanishi, 2005). In culture it is possible that GABAergic synapses were formed by all of these interneurones onto the granule cell. Recordings were made from rat cerebellar granule cells with pre-activated α -CaMK-II in the patch pipette using the same protocol as described in the previous chapters (See Chapter 2 and 3.2.2) to determine if there was a CaMK-II dependent effect on inhibitory synaptic transmission, and if so, what form that modulation took.

5.2 **RESULTS**

Cerebellar granule cells maintained in physiological levels of KCl (5 mM) for 7-14 days (See Chapter 2) had a slightly altered morphology to cells maintained under depolarising conditions (Moulder et al., 2003). Generally they had slightly smaller cell bodies and correspondingly lower cell capacitance $(8.3 \pm 0.5 \text{ pF}, n = 15)$ compared to cells maintained in depolarising conditions $(11.1 \pm 0.6 \text{ pF}, n = 15)$. In 5 mM KCl, CGCs tended to form small clusters of cells (See Fig 2.1B) rather than the uniform distribution of cells maintained in 20 mM KCl (Ives et al., 2002). Most CGCs within clusters displayed spontaneous synaptic activity, although a proportion displayed no activity at all (15-20 %). Both excitatory and inhibitory postsynaptic currents could be observed. Excitatory input from surrounding granule cells could be observed as spontaneous excitatory postsynaptic currents (sEPSCs). Application of the NMDA receptor blocker AP-5 (20 μ M) and AMPA receptor blocker CNQX (10 μ M), revealed GABAergic currents (IPSCs) that could be blocked by the GABA_A receptor antagonist bicuculline (20 µM; See Fig 5.1). Bath application of tetrodotoxin (TTX - 500 nM) almost completely abolished all IPSCs. This is in line with the observation that most CGC IPSCs in culture and in thin brain slices are action potential dependent (Puia et al., 1994; Leao et al., 2000; Farrant and Brickley, 2003). The Na²⁺ channel blocker OX-314 (5 mM) was included within the patch pipette to prevent action potentials from occurring within the cell being recorded from whilst allowing the activity of the rest of the network to remain relatively unaltered. In this way action potential driven sIPSCs could be recorded.

A tonic GABA_A receptor mediated current has been observed in cerebellar granule cells in culture and in slices, presumed to be mediated by the basal GABA concentration or by GABA spillover acting on $\alpha 6/\delta$ subunit containing extrasynaptic

receptors (Brickley *et al.*, 1996; Leao *et al.*, 2000; Wall, 2002). On application of bicuculline the tonic noise was altered in only a small subset of the cells recorded from these cultures (1 in 20). This is possibly due to the immature nature of the cultures and a possible lack of δ subunit expression, or perhaps due to a reduced number/density of GABA releasing cells in culture. Expression of δ subunits is thought to be negligible until postnatal day 6-7 and does not reach adult levels until the third postnatal week (Wisden *et al.*, 1996). As these cultures were prepared from P0 rats it may be that δ subunit expression never reached high enough levels to support a tonic GABA current.

Fig 5.1:

AP-5 + CNQX



Fig 5.1: sIPSCs recorded from rat cerebellar granule cells in culture.

Representative synaptic currents recorded from rat cerebellar granule cells grown and maintained in a physiological culture medium containing (5 mM) KCl and 5 % FCS.

Application of bicuculline (20 μ M) reveals EPSCs. Application of AP-5 (20 μ M), CNQX (10 μ M) and bicuculline (20 μ M) removes all events. Application of AP-5 and CNQX reveals sIPSCs.

5.2.1 CaMK-II modulation of sIPSC amplitude.

Recordings were made with either 85 nM pre-activated α -CaMK-II (See Chapter 2 and 3) within the patch pipette or normal patch pipette solution containing the pre-activation buffer without α -CaMK-II as a control. This was to ensure that the pre-activation buffer (although diluted in the patch pipette solution) did not activate endogenous CaMK-II or other Ca²⁺ sensitive kinases or phosphatases. Continuous recordings were made in 2 min periods starting 4-5 min after achieving the whole-cell configuration (t = 0 min), for a total period of 20 min. Analysis of IPSC kinetics required continuous recordings to be made from t = 6 min to t = 12 min to ensure an adequate number of events for analysis.

The addition of 85 nM α -CaMK-II led to a significant increase in the mean peak amplitude of sIPSCs to 129.7 ± 6.1 % (n = 7) as compared to control, 93.7 ± 7.8 % (n = 9, Fig 5.2A) at t = 6-8 min. This effect reached a steady-state within 6 min. The observed increase in the mean sIPSC amplitude did not necessarily indicate a uniform increase in amplitude of all events within the population. In order to further evaluate the effects of CaMK-II on sIPSC amplitude, an amplitude distribution histogram was constructed. Under control conditions the population was best fitted by three Gaussian populations (n = 7, 7000 events). The presence of more than one population has been previously observed in rat cerebellar cultures (Leao *et al.*, 2000) and is presumed to correspond to action potential dependent (sIPSCs) and action potential independent (mIPSC) forms of GABA release. However, the addition of TTX abolishes almost all IPSCs in this culture so it is difficult to confirm that large amplitude events are action potential driven as there are not enough mIPSCs for reliable analysis. The distribution of amplitudes appeared to be skewed towards higher amplitude events as has also been described previously (De Koninck and Mody, 1994; Nusser et al., 1997).

The addition of α -CaMK-II led to the appearance of a group of large amplitude IPSCs of over 400 pA (n = 4, 7000 events, Fig 5.2C). There was a large drop in the number of events of around 200 pA and an increase in the number of smaller amplitude events. The population was now composed of four Gaussian populations. The appearance of larger amplitude events appears to have been at the expense of the larger amplitude populations of events seen in controls (around 200 pA). The smallest amplitude population seen in controls shifts slightly to the right (to give rise to a fourth population, around 70 pA) and the number of the smallest events (<70 pA) is increased. This could be partly due to smaller amplitude events that were previously undetectable, increasing in amplitude, and appearing out of the noise. This might be expected to be reflected by an increase in frequency, however, the frequency of events was not significantly altered over the time of recording (Fig 5.2D) or in the presence of α -CaMK-II (n = 7, 2.2 ± 0.6 Hz) as compared to control (n = 8, 2.8 ± 1.2 Hz). It appears though that the effect of CaMK-II is not uniform across the whole population.



Fig 5.2: α -CaMK-II modulation of sIPSCs recorded from rat cerebellar granule cells.

sIPSCs were recorded from cerebellar granule cells in culture (DIV10-14) maintained in 5 mM KCl and 5 % FCS. Recordings were made in 2 min periods with the first starting 4-5 min after achieving the whole-cell configuration (t = 0-2 min) or in 5-6 min periods starting 6 min after achieving the whole-cell configuration. Control recordings were made using normal patch pipette solution supplemented with preactivation buffer without α -CaMK-II.

(A) Mean sIPSC peak amplitudes of events within a 2 min recording period were normalised to the mean peak amplitude at t = 0-2 min, in the absence (Control, n = 9) and presence of 85 nM α -CaMK-II (n = 7). All points are mean \pm s.e.

(B) Representative recordings of sIPSCs at t = 2 min and t = 8 min with 85 nM α -CaMK-II present within the patch pipette solution.

(C) Amplitude distribution histogram of sIPSCs in the absence (Control, n = 7,7000 events) and in the presence of 85 nM α -CaMK-II (n = 4,7000 events) at t = 6-12 min. (D) Bar chart showing the frequency at t = 6-8 min normalised to the frequency at t = 0-2 min in control conditions (n = 8) and in the presence of 85 nM α -CaMK-II (n = 7).

Significance is indicated by the symbol *, P < 0.05, t-test.

5.2.2 CaMK-II modulation of sIPSC decay.

Previous studies have shown that there is considerable variation in the decay of IPSCs between different cell types, cells of the same type, and even within the same cell (Auger and Marty, 1997; Bacci *et al.*, 2003; Mody and Pearce, 2004). Control recordings from rat CGCs in physiological KCl (5 mM) revealed that there were three distinct Gaussian populations of decay times, as determined by half-width (time taken to decay to 50 % of peak amplitude; n = 6, 2600 events, Fig 5.3A).

The variation in decay times has been observed in other neuronal preparations and is thought to reflect the expression of different GABA_A receptor subtypes within the cell (Puia *et al.*, 1994; Koulen *et al.*, 1996; Bacci *et al.*, 2003) and different phosphorylation states (Jones and Westbrook, 1997), but may also reflect variations in GABA concentration within the synaptic cleft of different synapses (Nusser *et al.*, 2001; Mody and Pearce, 2004). A typical example trace taken from a single CGC recording illustrates events with similar rise times but different decay profiles (Fig 5.3A).

The distribution of half-widths of sIPSCs recorded in the presence of α -CaMK-II displays a shift to the right towards a slower decay time (n = 7, 2600 events, Fig 5.3A). There is an increase in the number of events decaying within the 25-45 ms range and a drop in the number of events within the 10-25 ms range.

At t = 6-12 min, sIPSC events (50-100) were averaged from control cells and from those in the presence of α -CaMK-II. Average traces were then used to calculate decay time constants (See Chapter 2, 2.7.2). Events recorded in the absence and in the presence of α -CaMK-II were best fitted by a bi-exponential decay. The decay constant τ_1 significantly increased from 10 ± 0.9 ms (n = 7) to 22.7 ± 3.2 ms in the presence of α -CaMK-II (n = 8, Fig 5.3C). The decay constant τ_2 also significantly increased from 46.7 \pm 6 ms (n = 7) to 78 \pm 10 ms in the presence of α -CaMK-II (n = 8, Fig 5.3C). The proportion of area $\tau_{1:}\tau_{2}$ was not significantly altered (A1 Control = 53.6 \pm 4.7 %, n = 7, A1 + CaMK-II = 51.7 \pm 3.6 %, n = 7, Fig 5.3D) by the presence of α -CaMK-II.

Variation in IPSC decay times could also result from dendritic filtering, however, CGCs are electrotonically very compact and so the effect of dendritic filtering should be comparatively low (Gabbiani *et al.*, 1994; Rossi and Hamann, 1998). To confirm this, scatter plots of IPSC rise-time against IPSC amplitude were constructed for each cell to check there was no linear relationship between these two variables. Only cells lacking any clear correlation between IPSC rise-time and amplitude were used in the analysis of IPSC decay times (r = 0.15, Fig 5.4). This suggests that variation in decay times in the majority of cases is related to genuine differences in the properties of GABA_A receptors at different synapses and not due to dendritic cabling. Moreover, α -CaMK-II did not affect the mean rise-time of the IPSCs, from 10-90 % of peak amplitude (Control, 1.2 ± 0.2 ms, n = 8, + α -CaMK-II, 1.1 ± 0.1 ms, n = 7, Fig 5.4B).





Fig 5.3: Modulation of sIPSC decay times by α-CaMK-II.

(A) Frequency distribution histogram of sIPSC half-widths (τ_{50}) at t = 6-8 min in the absence (Control, n = 6, 2600 events) and the presence of α -CaMK-II (n = 7, 2600 events). Inset is a representative trace illustrating variation in the decay times under control conditions.

(B) Superimposed scaled averages of 80 sIPSCs recorded from one cell in the presence of α -CaMK-II at t = 0-2 min and t = 6-8 min and re-scaled to the same amplitude.

(C) Bar charts of the mean values for the exponential decay time constants, $\tau 1$ and $\tau 2$ at t = 6-12 min in the absence (Control, n = 8, minimum 50 events per cell) and the presence of α -CaMK-II (n = 9, minimum 50 events per cell).

(D) Bar chart showing the relative contribution of $\tau 1$ (A1) and $\tau 2$ (A2) at t = 6-12 min in the absence (Control, n = 8) and presence of CaMK-II (n = 7).

Mean \pm s.e. Significance is indicated by the symbol *, CaMK-II compared to control.

P < 0.05, t-test.





Fig 5.4: sIPSC rise-times in the absence and presence of α-CaMK-II.

(A) Example scatter plot of sIPSC amplitudes against sIPSC rise-times in the absence (Control, n = 3, 780 events) of α -CaMK-II. Note the lack of a linear relationship (r = 0.15).

(B) Bar chart showing the mean sIPSC rise-time in the absence (n = 8) and the presence of 85 nM α -CaMK-II (n = 7). For each cell the mean rise-time was calculated from 50-100 events averaged from recordings, t = 6-12 min.

5.2.3 CaMK-II modulation of the whole population of sIPSCs.

Currently, little is known about CaMK-II dependent modulation of GABA_A receptors, so the possibility remains that CaMK-II preferentially modulates certain GABA_A receptor subtypes or can differentially modulate certain subtypes in different ways. The latter may involve either direct phosphorylation, indirect phosphorylation of GABA_A receptor-associated proteins or possibly may act through downstream signalling pathways. Scatter plots of sIPSC amplitudes against half-widths recorded at t = 6-12 min in the absence (Control, n = 7, 7000 events) and in the presence of 85 nM α -CaMK-II (n = 4, 7000 events, Fig 5.5A) were constructed to examine the effects of α -CaMK-II on the population as a whole. Addition of α -CaMK-II results in a shift of a sub-population of IPSCs to much larger amplitudes with relatively little change in decay. In addition another population of events appears to increase in half-width with relatively little change in amplitude. There is also another sub-population of events that are not significantly altered by α -CaMK-II at all.

It is possible this is a reflection of a variation in response to α -CaMK-II, with one population responding to α -CaMK-II with a large increase in amplitude and another separate population increasing in decay time. It certainly appears that the effect of α -CaMK-II was not uniform and consistent across the whole population of IPSCs.

Increases in sIPSC amplitude and decay would increase the area of each IPSC and therefore the overall charge transfer. The amplitude distribution histogram of sIPSC areas in the absence (n = 6, 2600 events) and the presence of α -CaMK-II (n = 7, 2600 events, Fig 5.4B) at t = 6-12 min displays a significant shift to events with much greater area. This would be consistent with a significant increase in the overall charge transfer. In addition the mean area of sIPSCs significantly increases from



 1931 ± 265 fC (n = 14) to 4440 ± 714 fC (n = 9, Fig 5.5C) in the presence of 85nM

Fig 5.5: CaMK-II modulation of sIPSCs as a population and changes in synaptic charge transfer.

(A) Scatter plots of sIPSC amplitude against half-width at t = 6-12 min in the absence (left, Control, n = 7, 7000 events) and presence (right) of 85 nM α -CaMK-II (n = 4, 7000 events).

Different populations are highlighted by the circles.

(B) Frequency distribution histogram of the area of each IPSC (pC) at t = 6-12 min in control conditions (n = 6, 2600 events) and in the presence of 85 nM α -CaMK-II (n = 7, 2600 events). (1nA . ms = 1pC; 1C = 1 coulomb of charge) (C) Bar chart showing the mean area of IPSCs (fC) in the absence (Control, n = 14)

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or presence of 85 nM α -CaMK-II (n = 9) at t = 6-12 min and t = 6-8 min.

Data points = mean \pm s.e. Significance is indicated by the symbol *, CaMK-II compared to control. P < 0.05, t-test.

5.3 **DISCUSSION**

The application of pre-activated α -CaMK-II to CGCs appears to enhance inhibitory synaptic transmission. This occurs through an increase in the mean peak amplitude and decay time constants of sIPSCs. These changes are consistent with other studies on phosphorylation-dependent effects on IPSCs with both PKA/PKC being shown to alter IPSC amplitude and/or decay in different preparations (Nusser *et al.*, 1999; Poisbeau *et al.*, 1999; Jovanovic *et al.*, 2004). Such changes potentially have significant implications for the modulation of inhibition and subsequently, neuronal excitability.

The dose-response relationship in the presence of α -CaMK-II (See Chapter 4, Fig 4.2) revealed an increase in the maximal response to GABA. This implies an increase in the number of functional receptors at the synapse, a change in

conductance or a change in the probability of ion channel opening may be responsible for the up-regulation of $GABA_A$ receptor function observed. A change in the number of receptors at a synapse has been proposed to be the most efficient way of enhancing GABAergic transmission given that GABA is presumably saturating at those synapses (Nusser *et al.*, 1998).

There is some evidence that at certain synapses in certain cell types, GABA is indeed saturating (De Koninck and Mody, 1994; Nusser *et al.*, 1997; Hájos *et al.*, 2000). Non-stationary noise analysis has allowed the numbers of functional receptors at synapses to be estimated (Traynelis and Jaramillo, 1998) and has shown that in some cases up-regulation of GABA_A receptor function is attributable to increases in receptor number (Otis *et al.*, 1994; De Koninck and Mody, 1994; Mizoguchi *et al.*, 2003). These increases are usually not associated with a change in decay. Recently Wang *et al.* (2003) demonstrated that PKB phosphorylation can increase the number of receptors at the cell surface via phosphorylation of the β 2 subunit at Ser⁴¹⁰, which results in an increase in mIPSC amplitude without affecting IPSC decay.

Recent evidence, however, has suggested that GABA is not saturating at all synapses in the CNS (Perrais and Ropert, 1999) and that receptor occupancy can vary between cell types and synapses within the same cell (Nusser *et al.*, 1997; Hájos *et al.*, 2000). Application of benzodiazepines (BDZ's) will increase the amplitude and the decay time constant of IPSCs, but only the decay time constant, if GABA is saturating. A number of studies have shown BDZs can increase IPSC amplitude in a number of cell types at different synapses. This is true for multi-quantal release (sIPSCs) as well as spontaneous miniature release (mIPSCs) of GABA (Rumpel and Behrends, 2000).

In the current study it is not yet apparent whether some or all of the synapses formed in these cultures are saturating. However there is a possibility that, as in other cell types, receptor occupancy is variable (Hájos *et al.*, 2000). Nusser and colleagues (1997) observed that a quantum of GABA results in incomplete receptor occupancy for a sub-population of events recorded from cerebellar stellate cells whilst others were thought to be saturated. Hajos *et al.* (2000) also observed differential receptor occupancy in olfactory bulb granule cells. They observed two populations of events with different decay time constants only one of which appeared to be saturated by a quantum of GABA. In the current study, α -CaMK-II appeared to have different effects on different populations of events; one possible explanation for this could be differences in the level of receptor occupancy after GABA release at different synapses within these cells.

An alteration in the probability of channel opening and the conductance of a channel could account for the apparent changes in amplitude observed (Moss *et al.*, 1995). An increase in the probability of channel opening could increase the apparent amplitude of whole-cell GABA currents and sIPSCs but would only produce large effects if GABA were non-saturating (Nusser *et al.*, 1998). Probability of opening may also affect the decay phase of IPSCs. These parameters could be changed by the covalent addition of a phosphate group (phosphorylation) altering conformation of the receptor, resulting in an alteration of single channel properties (Soltesz and Mody, 1995). Tyrosine kinase phosphorylation of the γ 2 subunit has been shown to increase the mean open time and the probability of opening of the GABA_A receptor (Moss *et al.*, 1995) corresponding to an increase in the amplitude of whole-cell currents recorded.

Another potential mechanism for the increase of IPSC amplitude and decay times observed could involve changes in the desensitisation of the receptor. Alterations in desensitisation have been shown occur through to phosphorylation/dephosphorylation (Jones and Westbrook, 1997). This may be a relatively common mechanism to modulate fast synaptic transmission (Jones and Westbrook, 1996). The bi-exponential decay of most IPSCs is thought to reflect the movement of the GABA_A receptor through desensitised states (representing the slower phase of decay). Although no or limited current flows through the desensitised channel, entering into a desensitised state prolongs the amount of time the channel can be active after a transient exposure to GABA.

The number of IPSCs with a bi-exponential decay and the relative contribution of each phase has been shown to be modulated by phosphorylation (Nusser *et al.*, 1999; Poisbeau *et al.*, 1999). The current study reveals an increase in amplitude and a slowing of the decay phase without alteration in the relative contribution of each component, resulting in a significant increase in synaptic charge transfer. Soltesz and Mody (1995) identified a calcium-dependent alteration in decay from a single exponential decay to a prolonged double exponential after amputation of dentate gyrus granule cell dendrites (>66 % of the dendritic tree – dendrotomy), which resulted in a significant increase in synaptic charge transfer. α -CaMK-II has been observed to alter desensitisation in rat spinal dorsal horn neurons acting to prolong decay times (Wang *et al.*, 1995). In addition the Ca²⁺ sensitive phosphatase calcineurin (CaN) has been reported to modulate desensitisation acting to lengthen and in some cases to shorten decay times (Martina *et al.*, 1996; Jones and Westbrook, 1997; Mozrzymas and Cherubini, 1998).

IPSCs in general display considerable variation in decay times between different cell types and even within the same cell (Banks et al., 1998; Bacci et al., 2003). The different populations of decay times seen in the present study were entirely consistent with this. Such variation may have significant functional effects with different GABA_A receptor subtypes and their distinct kinetic properties being incorporated into discrete synapses to perform particular roles in controlling neuronal excitability (Jones and Westbrook, 1996; Somogyi, 2005). Variation can occur as a result of several different reasons. Different GABA_A receptor subunit combinations can produce receptors with different kinetic properties (Puia et al., 1994; Gingrich et al., 1995; Haas and Macdonald, 1999; Bacci et al., 2003). Often these subunit combinations and their corresponding decay time constants can alter during development (Tia et al., 1996; Hollrigel and Soltesz, 1997; Hutcheon et al., 2000; Vicini et al., 2001). For example, $\alpha 2/3\beta 3$ receptors have a slower apparent IPSC decay time and are often expressed earlier in development being replaced by faster decaying $\alpha l \beta 2$ receptors as synapses reach maturity (Vicini et al., 2001; Jüttner et al., 2001). The GABA_A receptor β 3 subunit, knock-out mouse displays CGC IPSCs that decay faster than their wild-type counterparts, which may be a result of an associated down-regulation of $\alpha 2/3$ expression (Ramadan et al., 2003). There is evidence that different synapses within the same cell may express different GABA_A receptor subunits and that accordingly, decay time constants at different synapses within the same cell are different (Koulen et al., 1996; Banks et al., 1998; Brünig et al., 2002). This possibly suggests that the slower decaying events in the current study may correspond to $\alpha 2/3\beta 3$ subunit-containing receptors.

Variation in the rate of decay can also result from the activation of perisynaptic receptors, activated by the spillover of GABA from the synaptic cleft

(Wall, 2002; Wei *et al.*, 2003). This phenomenon has been observed at Golgigranule cell synapses in the cerebellum (Rossi and Hamann, 1998; Hamann *et al.*, 2002). A preferential up-regulation of these perisynaptic receptors by CaMK-II dependent phosphorylation could theoretically increase the apparent decay of synaptic events without involving phosphorylation of the synaptic GABA_A receptors themselves. However, this would affect the rise-times of IPSCs as well as the decay times which was not observed in the current study.

Variations in IPSC decay times can also result at non-saturating synapses as a result of variations in GABA concentration within the synaptic cleft (Nusser *et al.*, 2001). Clustering of receptors and associated scaffolding proteins at the synapse can also alter decay. Co-expression of recombinant $\alpha 1\beta 2\gamma 2L$ receptors with GABARAP in QT6 quail fibroblasts promoted clustering. These clustered receptors deactivated faster and densensitised slower (Chen *et al.*, 2000).

Alteration in the level of variation around the mean (IPSC amplitude and/or decay) can in itself alter neuronal excitability. Increasing event to event variance of peri-somatically injected IPSCs caused either a decrease, an increase or no change in the firing rate of CA1 hippocampal pyramidal cells, depending on the mean around which the scatter was introduced and the degree of scatter (Aradi *et al.*, 2002; Aradi *et al.*, 2004). Alterations in the decays of IPSCs can have significant effects on synaptic inhibition and the excitability of a cell (Jones and Westbrook, 1996). Changes in decay have also been linked to pathological states, for example, receptor mutations that alter the decay phase of IPSCs have been linked to familial epilepsy and insomnia (Bianchi *et al.*, 2002; Buhr *et al.*, 2002; Fisher, 2004).

It is possible that in the current study some of the variation in IPSC decay times observed occurred as a result of dendritic filtering. An inadequate voltage

clamp may alter the profile of IPSCs occurring at distant synaptic sites. It is unlikely this effect is significant in these cells as they are electrotonically compact (Gabbiani *et al.*, 1994; Rossi and Hamann, 1998), so the effect of dendritic filtering should be minimal. To confirm this, scatter plots of sIPSC rise-times were plotted against amplitudes and it was confirmed there was no linear relationship between the two.

Given the large number of potential mechanisms that could result in the changes in sIPSC amplitude and decay times observed in the present study it is not possible to say precisely how α -CaMK-II mediates its effects on IPSCs. It is also not clear whether this effect is through direct phosphorylation or whether modulation occurs indirectly. There is some evidence to suggest that CaMK-II can bind to the β subunit TM3-4 intracellular loop and that phosphorylation of the β 3 subunit occurs within embryonic cortical neurones in culture in response to CaMK-II activation (McAinsh K, Jovanovic JN and Moss SJ, unpublished observations). However, the data in the current study suggests that the effect of α -CaMK-II is not uniform across all IPSC events. It may be that more than one process is involved and/or that some GABA_A receptor subunit combinations are functionally modulated in different ways by α -CaMK-II. In addition, variation in synapses such as receptor occupancy, receptor number and GABA concentration within the synaptic cleft following action potential driven release could alter the way an IPSC is modulated by CaMK-II phosphorylation.

It is interesting that α -CaMK-II appears to potentiate GABA_A receptor function when CaMK-II is also involved in potentiation of AMPA receptor function (Lisman *et al.*, 2002; Colbran and Brown, 2004). There are a number of parallels that exist between excitatory and inhibitory long-term synaptic plasticity, in that both respond bi-directionally to Ca²⁺ signals (Aizenman *et al.*, 1998; Yang *et al.*, 1999; Cormier *et al.*, 2001). These Ca²⁺ signals, at excitatory synapses, are thought to preferentially activate either calcineurin or CaMK-II (Xia and Storm, 2005). GABAergic synapses have also been shown to be depressed by calcineurin and potentiated by CaMK-II (Kano *et al.*, 1996; Wang *et al.*, 2003a). It may be that this form of potentiation of GABA_A receptor synapses occurs as a homeostatic mechanism in response to large increases in excitatory drive (Turrigiano and Nelson, 2004). Long lasting changes in inhibitory synaptic transmission can as a consequence alter the thresholds for inducing excitatory LTP, a process known as metaplasticity (Abraham and Tate, 1997; Chevaleyre and Castillo, 2004).

Kano et al. (1992; 1996) identified a phenomenon termed rebound potentiation, involving an α -CaMK-II dependent increase in amplitude (with no change in the decay times) of sIPSCs in cerebellar Purkinje neurones. Purkinje cells are thought to express predominantly $\alpha 1\beta 2\gamma 2$ receptors (Persohn et al., 1992). The CGCs in the current study are thought to express $\alpha 1/2/3/6$, $\beta 2/3$, $\gamma 2$ and possibly δ subunit-containing receptors. Maintaining cells in depolarising conditions may shift the expression of GABA_A receptor subunits towards $\alpha 2/3$, $\beta 3$ subunit-containing receptors (Ives et al., 2002) but in low KCl (5 mM), CGC development may be expected to follow the same pattern of development as in vivo and as a result express an increased number of different GABA_A receptor subunits (Wisden et al., 1996). The differential effect of α -CaMK-II on different populations of IPSCs may reflect different GABA_A receptor subtypes at different synapses. As modulation of IPSCs in Purkinje cells by CaMK-II involves an increase in amplitude with no change in decay the sub-population of events in the current study that increases in amplitude with little change in decay times could be proposed to be mediated by the same GABA_A receptor subtype as that in Purkinje cells. The alteration in decay times seen

in the current study may reflect α -CaMK-II action on another GABA_A receptor subtype other than that seen in Purkinje cells.

In order to understand the variations in response to α -CaMK-II observed in the present study and within the literature it is necessary to determine the effect of CaMK-II on defined receptor populations. This has proved effective in investigating the variation in response to PKA dependent phosphorylation of GABA_A receptors (McDonald *et al.*, 1998). The lack of response to α -CaMK-II in HEK293 cells suggests a neuronal environment may be needed to observe any effects. However, it is difficult to control the expression of GABA_A receptor subunits expressed in neuronal cells. A cell line of neuronal origin may allow CaMK-II dependent modulation of the GABA_A receptor to be observed whilst allowing more control over the GABA_A receptor subunits that are expressed.

5.4 CONCLUSIONS

- Application of α-CaMK-II results in a significant potentiation of the amplitudes and an increase in the decay times of sIPSCs recorded from cerebellar granule cells.
- The effect of α-CaMK-II is not uniform across all events in the population suggesting possible differential effects of α-CaMK-II at different synapses possibly expressing different GABA_A receptor subtypes.

Chapter Six:

<u>CaMK-II modulation of recombinant GABA_A receptors</u> <u>expressed in NG108-15 cells.</u>

6.1 INTRODUCTION

CaMK-II is capable of modulating GABA_A receptor mediated currents in a number of neuronal cell types (Wang *et al.*, 1995; Kano *et al.*, 1996; Wei *et al.*, 2004) including cerebellar granule cells (Chapters 4-5). However it appears unable to modulate recombinant GABA_A receptors expressed in HEK293 cells. This is presumed to be because the intracellular environment of a HEK293 cell is unable to support the CaMK-II modulation of GABA_A receptors. However, expression of recombinant receptors in a cell line that does not express endogenous GABA_A receptors is desirable because it allows the exact subunit combination present to be controlled (McDonald *et al.*, 1998). For this reason, and to find an environment that would support CaMK-II modulation, GABA_A receptor subunits were expressed in NG108-15 cells, a cell line of neuronal and glial origin (Klee and Nirenberg, 1974; Daniels and Hamprecht, 1974). In this way, CaMK-II modulation of recombinant GABA_A receptors could be investigated and used to determine the relative effects of the kinase on different subunit types.

The NG108-15 clonal cell line is a hybrid cell line formed from the fusion of mouse neuroblastoma N18TG-2 cells and rat glioma C6BU-1 cells (Klee and Nirenberg, 1974; Daniels and Hamprecht, 1974). Since the formation of this cell line in the 1970s these cells have been used extensively to study neuronal differentiation (Tojima and Ito, 2004). Under basic culture conditions NG108-15 cells proliferate, but after treatment to increase internal cAMP levels, they stop proliferating and

produce extended processes consistent with a neuronal cell-type morphology (Docherty *et al.*, 1991; Wu *et al.*, 1998; Tojima and Ito, 2004). This property has been used as a model for the study of neuritogenesis and synaptogenesis (Nirenberg *et al.*, 1983; Han *et al.*, 1991; Chen *et al.*, 2001; Tojima *et al.*, 2003a; Tojima *et al.*, 2003b).

These cells display a wide range of voltage-dependent currents comparable to those observed in neurones (Docherty et al., 1991; Lukyanetz, 1998; Bowden et al., 1999). They are thought to be comparable to sympathetic neurones of the autonomic nervous system and are known to express choline acetyltransferase (ChAT; Daniels and Hamprecht, 1974). In co-culture with myotubes they form synapses and miniature endplate potentials (mEPPS) can be recorded (Nelson et al., 1976; Christian et al., 1977; Chen et al., 2001). For this reason they have also been used as a basic model of the neuromuscular junction. Furthermore, under certain culture conditions, they can express a number of ligand-gated ion channels. For example, they have been reported to express 5HT₃ receptors (Chan et al., 1994; Bartrup and Newberry, 1996; Boddeke et al., 1996), δ opioid receptors (Beczkowska et al., 1997; Buzas et al., 1998), NMDA receptors (Ohkuma et al., 1994; Beczkowska et al., 1997) and muscarinic acetylcholine receptors (Fukuda et al., 1988). It should be noted that although this cell line is termed clonal, this refers to the fact that they are derived from a common ancestor and does not indicate that all cells are genetically identical. Over time in culture they can undergo selection depending on the specific culture conditions within a given laboratory. For this reason not all NG108-15s will display the same characteristics and may not express the same neuronal receptors (Docherty et al., 1991).

NG108-15 cells have even been reported to display spontaneous synaptic currents when cultured alone (Tojima *et al.*, 2000a; Tojima *et al.*, 2003a). In general they are able, given the appropriate culture conditions, to form morphological characteristics of neurones as well as certain pre- and post-synaptic structures.

Under non-differentiating conditions when they are allowed to grow freely, they can display some neuronal-type morphological characteristics with some small processes present (Docherty *et al.*, 1991); however the expression of neuronal markers (ChAT, synaptophysin, syntaxin-1), the formation of large processes, and the establishment of synaptic connections, tends to occur after alteration of culture conditions to increase intracellular cAMP levels (Tojima *et al.*, 2000b). Under non-differentiating conditions they can express some cytoskeletal components associated with neurones, such as certain neurofilaments (e.g. NF200), and the actin depolymerising protein, cofilin (Tojima *et al.*, 2000b; Tojima *et al.*, 2003b). Under both proliferating and differentiated conditions they are not thought to express GABA_A receptors (Searles and Singer, 1988; Ishizawa *et al.*, 1997) although they have been reported to express glutamic acid decarboxylase (GAD) and possibly GABA (Searles and Singer, 1988).

NG108-15 cells have been used to study the properties of recombinant ion channels in the same way as HEK293 cells have by transfection with the relevant cDNAs. They have been stably transfected with dopamine D_3 receptors which have been shown to be functionally coupled to G-proteins. Dopamine D_3 receptors only couple weakly with G-proteins in Chinese Hamster Ovary cells, GH_4C_1 pituitary cells and non-differentiated NG108-15 cells (Pilon *et al.*, 1994; Seabrook *et al.*, 1994). NG108-15 cells have also been successfully transfected with GABA_B receptors to study GABA_B receptor function. This approach revealed possible

endogenous but non-functional $GABA_B$ receptor subunits within these cells (Easter and Spruce, 2002).

NG108-15 cells have also been used to investigate Ca²⁺ dependent signalling pathways involving Ca²⁺/CaM dependent phosphatase and kinase activity. Calcineurin activity has been shown to regulate the function of voltage-operated Ca²⁺ channels and 5-HT₃ receptors in NG108-15 cells (Boddeke *et al.*, 1996; Lukyanetz *et al.*, 1998). It has also been shown that calcineurin and possibly CaMK activity can modulate δ opioid mRNA expression in these cells (Buzas *et al.*, 1998). CaMK-II and CaMK-IV activity has been detected in NG108-15 cells after stimulation by direct depolarisation or following application of bradykinin (Yamakawa *et al.*, 1992; Enslen *et al.*, 1996). Endogenous CaMK-II within NG108-15s exists at a low level but is significantly increased by treatment that induces cellular differentiation (Vallano and Beaman-Hall, 1989). Activation of endogenous CaMK-II has been linked to peripheral actin polymerization and filopodia formation (Chen *et al.*, 2003) and overexpression of α -CaMK-II and autophosphorylation of α -CaMK-II have been shown to alter growth cone motility and neurite outgrowth in this cell line (Goshima *et al.*, 1993).

It seemed, therefore, that because of its neuronal origin and evidence of endogenous Ca^{2+}/CaM signalling pathways, that CaMK-II may be able to modulate the function of recombinant GABA_A receptors expressed within this cell line. In order to eliminate the possibility of endogenous GABA_{A/B} receptors being present or other ligand-gated ion channels, and to limit the contribution of endogenous CaMK-II; proliferating, undifferentiated NG108-15s were transfected with various GABA_A receptor subunit cDNAs. In this way NG108-15s provided a neuronal-like environment where α -CaMK-II modulation of recombinant receptors could be

observed and the relative contribution of different receptor subunits to the action of CaMK-II could be determined.

6.2 **RESULTS**

Using a similar protocol to that established in the previous chapters, preactivated recombinant α -CaMK-II (Suzuki-Takeuchi *et al.*, 1992) was included in the patch pipette solution (McGlade-McCulloh *et al.*, 1993) and whole-cell GABA currents were recorded from transfected NG108-15 cells. This was compared to controls which included either pre-activation buffer without α -CaMK-II or heatinactivated α -CaMK-II (pre-incubated in activation buffer) in the patch pipette. The peak amplitude of GABA currents were recorded over time as α -CaMK-II dialysed into the cell and normalised to the peak GABA current amplitude obtained 3-4 min after achieving the whole-cell configuration.

As NG108-15 cells in their undifferentiated state have a more compact round morphology (Docherty *et al.*, 1991) compared to cerebellar granule cells, and as in previous experiments 85-100 nM α -CaMK-II produced large increases in amplitude, the concentration was reduced to 60 nM α -CaMK-II to produce an easily observable effect whilst maintaining stable recordings. After stimulation to undergo differentiation NG108-15 cells can form large processes that make it difficult to establish an ideal voltage clamp of membrane currents (Docherty *et al.*, 1991). Using undifferentiated cells therefore, had the added advantage of avoiding problems of inadequate space clamp, as in this state, cellular processes are generally small.

NG108-15 cells were transfected with an equal ratio of $\alpha:\beta:\gamma$ subunit and enhanced GFP cDNAs using the transfection reagent Effectene (See Chapter 2). Transfected cells were subsequently identified by their fluorescence. As NG108-15

cells tended to form aggregates, they were plated at a low density following transfection and recorded from 24-72 hrs later. Cells were selected for electrophysiology if they were isolated from other cells, had only a few small processes and displayed low levels of fluorescence. As the cells were large and generally transfection efficiency was high, some GABA currents recorded were over 4000pA in amplitude even with a 10 μ M GABA application (EC₅₀). These currents were discarded from further analysis due to the potential for insufficient voltage clamp and series resistance problems.

6.2.1 NG108-15 cells did not express functional endogenous GABA_A receptors.

It is thought that NG108-15 cells do not express GABA_A receptors, however as these cells differ in their characteristics depending on the culture conditions and there was no systematic or thorough demonstration of this lack of expression in the literature, cells were transfected with cDNAs for GFP alone or for GFP and αl subunits alone, and GABA was applied to untransfected and transfected cells. As can be seen in Fig 6.1 no response to GABA was seen in any control situation (Untransfected, n = 9, GFP only, n = 5, αl subunit only, n = 4). This confirmed that there were no functional GABA_A receptors expressed at the cell surface of untransfected cells. As αl subunit transfected cells also failed to respond to GABA this confirmed that there were no endogenous β subunits present as if there had been, it would be predicted that the co-expression of αl would allow $\alpha l\beta$ receptors to be formed as has been observed in some cases with HEK293 cells (Ueno *et al.*, 1996; Davies *et al.*, 2000). As transfection with cDNAs for $\alpha\beta/\gamma$ subunits produced large currents in response to GABA it appeared that recombinant GABA_A receptors were expressed with high efficiency. It was thought, therefore, that recombinant subunits would dominate any endogenous protein that might be present (Davies *et al.*, 2000).

Fig 6.1:

10µM GABA

 NG108-15 GFP

 500pA

 NG108-15 GFP +

 2s

 NG108-15 GFP + $\alpha 1$

 NG108-15 GFP + $\alpha 1$

Fig 6.1: Expression of GABA_A receptors in NG108-15 cells.

Representative membrane currents showing response to applied GABA (10 μ M) recorded from:

(A) Untransfected NG108-15 cell (n = 9).

(B) Enhanced GFP (EGFP) positive NG108-15 cell transfected with EGFP alone (n = 5).

(C) EGFP positive NG108-15 cell transfected with the EGFP and the α 1 subunit only (n = 5).

(D) EGFP positive NG108-15 cell transfected with enhanced EGFP and $\alpha 1\beta 3$.

6.2.2 Recombinant $\alpha\beta$ receptors and CaMK-II.

Application of 60 nM α -CaMK-II to cells transfected with $\alpha 1\beta 1$ produced no significant effects (107.4 ± 7.3 %, n = 7) as compared to control (93.8 ± 3.7 %, n = 3, Fig 6.2A) at t = 6 min. Although a slight increase seemed apparent in the first few minutes this was not statistically significant and was indistinguishable from control at t = 8 min. Application of 60 nM CaMK-II to $\alpha 1\beta 2$ receptors also produced no significant differences (85.5 ± 4.2 %, n = 5) from control GABA currents (91 ± 6.3 %, n = 4, Fig 6.2B) at t = 6 min.

However, application of 60 nM α -CaMK-II to $\alpha 1\beta 3$ receptors produced a statistically significant increase in the peak amplitude of currents recorded to 118 ± 4.8 % (n = 7) as compared to control, 96.4 ± 4.6 % (n = 7, Fig 6.2C) at t = 6 min which reached a plateau within 6-8 min.

6.2.3 Recombinant $\alpha\beta\gamma$ receptors and CaMK-II.

Application of 60 nM α -CaMK-II to $\alpha 1\beta 1\gamma 2S$ receptors expressed in NG108-15 cells resulted in a significant potentiation of the whole cell currents recorded to 125.6 ± 6.4 % (n = 11) as compared to control, 97.1 ± 3.9 % (n = 5) at t = 6 min. The inclusion of heat inactivated α -CaMK-II though, had no significant effects (89.2 ± 3.1 % at t = 6 min, n = 7, Fig 6.3A).

As for the $\alpha 1\beta 2$ receptor, application of 60 nM α -CaMK-II to $\alpha 1\beta 2\gamma 2S$ expressing NG108-15 cells had no significant effect (102.5 ± 5.3 %, n = 11) as compared to control (105.1 ± 6.7 %, n = 6, Fig 6.3B). However, application of 60 nM α -CaMK-II to $\alpha 1\beta 3\gamma 2S$ receptors significantly increased the peak amplitude of whole-cell currents to 145.8 ± 10.8 % (n = 7) at t = 6 min as compared to control, 102.8 ± 5 % (n = 4). Heat inactivated α -CaMK-II again had no significant effect $(90.5 \pm 11.6, n = 5, Fig 6.3C)$ as compared to control, but was significantly different from addition of pre-activated CaMK-II, at t = 6 min. The response to α -CaMK-II for $\alpha 1\beta 3$ receptors was significantly increased by the inclusion of the $\gamma 2S$ subunit.

Transfection with cDNA for $\alpha\beta\gamma$ subunits in equal ratio should ensure that the majority of receptors at the surface of the cell are $\alpha\beta\gamma$ containing receptors as opposed to $\alpha\beta$ (Angelotti *et al.*, 1993; Angelotti and Macdonald, 1993). In agreement with this the amplitudes of currents recorded from $\alpha1\beta3\gamma2S$ receptors were larger than those recorded from $\alpha\beta$ receptors (Angelotti *et al.*, 1993). In addition, application of 10 μ M diazepam (Herring *et al.*, 2003) routinely increased the amplitude of the currents recorded indicating the presence of the $\gamma2$ subunit (Fig 6.3D).





Fig 6.2: Whole-cell currents recorded from NG108-15 cells expressing $\alpha 1\beta 1-3$ subunits were differentially modulated by α -CaMK-II.

Peak amplitude of GABA mediated currents recorded following the application of 10 μ M GABA to transfected NG108-15 cells expressing $\alpha 1\beta 1$, $\alpha 1\beta 2$ and $\alpha 1\beta 3$ GABA_A receptors. GABA was re-applied at 2 min intervals for up to 20 min. All currents were normalised to the peak amplitude of the response measured during the first 3-4 min (= 100 %) after achieving the whole-cell configuration (t = 0 min). Control recordings were made with normal patch pipette solution supplemented with the pre-activation buffer without α -CaMK-II.

(A) GABA currents recorded from NG108-15 cells transfected with $\alpha 1\beta 1$ subunits in the absence (Control, n = 3) and presence of 60 nM pre-activated α -CaMK-II, which was introduced into the cells via the patch pipette solution (n = 7).

(B) GABA currents recorded from NG108-15 cells transfected with $\alpha 1\beta 2$ subunits in the absence (Control, n = 4) and the presence of 60 nM α -CaMK-II (n = 5).

(C) GABA currents recorded from NG108-15 cells transfected with $\alpha 1\beta 3$ subunits in the absence (Control, n = 7) and the presence of 60nM α -CaMK-II (n = 7).

All points = mean \pm s.e. Significance is indicated by the symbol *, CaMK-II compared to control. P < 0.05, t-test.



Fig 6.3: Whole-cell currents recorded from NG108-15 cells expressing $\alpha 1\beta 1$ -3 $\gamma 2S$ subunits were differentially modulated by α -CaMK-II.

Peak amplitude of GABA mediated currents recorded following the application of 10 μ M GABA to transfected NG108-15 cells expressing $\alpha 1\beta 1\gamma 2S$, $\alpha 1\beta 2\gamma 2S$ and $\alpha 1\beta 3\gamma 2S$ GABA_A receptors. GABA was re-applied at 2 min intervals for up to 20 min. All currents were normalised to the peak amplitude of the response measured during the first 3-4 min (= 100 %) after achieving the whole-cell configuration (t = 0 min). Control recordings were made with normal patch pipette solution supplemented with the pre-activation buffer without α -CaMK-II.

(A) GABA currents recorded from NG108-15 cells transfected with $\alpha 1\beta 1\gamma 2S$ subunits in the absence (Control, n = 5) and presence of 60 nM pre-activated α -CaMK-II (n = 11) and with 60 nM heat inactivated (HI) α -CaMK-II (n = 7).

(B) GABA currents recorded from NG108-15 cells transfected with $\alpha 1\beta 2\gamma 2S$ subunits in the absence (Control, n = 6) and the presence of 60 nM α -CaMK-II (n = 11).

(C) GABA currents recorded from NG108-15 cells transfected with $\alpha 1\beta 3\gamma 2S$ subunits in the absence (Control, n = 4) and the presence of 60 nM α -CaMK-II (n = 7) and in the presence of 60 nM heat inactivated (HI) α -CaMK-II (n = 5).

(D) Representative membrane currents recorded from NG108-15 cells expressing $\alpha 1\beta 3\gamma 2S$ receptors before and after the bath application of 10 μ M diazepam.

All points = mean \pm s.e.

Significance is indicated by the symbol *, CaMK-II compared to control and the symbol *, CaMK-II compared to HI CaMK-II. P < 0.05, t-test, ANOVA.
6.3 **DISCUSSION**

Recombinant GABA_A receptors expressed in NG108-15 cells can be functionally modulated by α -CaMK-II in a manner similar to that observed in cerebellar granule cells. However, control over the subunits that are expressed revealed different CaMK-II dependent effects depending on the precise subunit combination present. The only $\alpha\beta$ heteromer that could be modulated by α -CaMK-II was $\alpha1\beta3$ suggesting a significant role for the $\beta3$ subunit in the α -CaMK-II modulation of GABA_A receptors. Addition of the $\gamma2S$ subunit increased this level of modulation significantly and revealed a lower, but still significant, level of modulation of $\alpha1\beta1$ subunit-containing receptors. However, both $\alpha1\beta2$ and $\alpha1\beta2\gamma2S$ receptors failed to be modulated by α -CaMK-II. It appears, therefore, that CaMK-II has a differential effect on GABA_A receptors depending on the subunits present, a phenomena that has been observed in relation to other phosphorylation-dependent effects on GABA_A receptor function (McDonald *et al.*, 1998; Nusser *et al.*, 1999).

The different responses to CaMK-II mediated by different β subunits shows some similarity to the effects seen on GABA_A receptors mediated by PKA (McDonald *et al.*, 1998). In this case PKA phosphorylation of β 3 is thought to increase the peak amplitude of GABA currents and have no effect on β 2 subunitcontaining receptors. However, phosphorylation of the β 1 subunit by PKA caused a depression of receptor function. In addition PKC is thought to have different effects on different GABA_A receptor subtypes. Again phosphorylation of β 3 subunitcontaining receptors is thought to increase GABA_A receptor function (Jovanovic *et al.*, 2004) whilst PKC dependent phosphorylation of β 1 subunit-containing receptors depresses GABA_A receptor function (Leidenheimer *et al.*, 1992; Krishek *et al.*, 1994; Brandon *et al.*, 2002b). Although there is some discrepancy in the functional effect observed (Lin *et al.*, 1994; Lin *et al.*, 1996).

The variation in response of native GABA_A receptors to activation of these kinases in neurones is thought, in part, to be mediated by these differential effects of kinases on different receptor subunit combinations. PKA and PKC have both been shown in neurones to have slightly different effects on GABA_A receptor mediated mIPSCs (Nusser *et al.*, 1999; Poisbeau *et al.*, 1999). However the response in neurones does not always correlate to that seen in recombinant systems. For instance, PKA appears able to modulate IPSCs in cells that predominantly express the β 2 subunit (Nusser *et al.*, 1999) despite a lack of response of the β 2 subunit-containing GABA_A receptors in HEK293 cells to PKA (McDonald *et al.*, 1998).

In different recombinant systems there is also reported variation in the effects of phosphorylation. For example PKC activity has been reported to increase the amplitude of GABA currents mediated by $\alpha 1\beta 1\gamma 2L$ receptors expressed in L929 fibroblasts (Lin *et al.*, 1996) although in *Xenopus oocytes* and HEK293 cells this has been reported to result in a depression of receptor function (Krishek *et al.*, 1994). In addition, examination of desensitisation rates of $\alpha 1\beta 1\gamma 2L$ and $\alpha 1\beta 3\gamma 2L$ receptors expressed in HEK293 cells in response to fast application of GABA revealed identical effects of PKA phosphorylation on macroscopic desensitisation rates (Hinkle and Macdonald, 2003) regardless of the β subunit present.

An important observation in the current study is that the addition of $\gamma 2S$ to the $\alpha 1\beta 3$ heteromer increased the response to α -CaMK-II. It appeared that both $\beta 3$ and $\gamma 2S$ subunits might be important for CaMK-II modulation and that both subunits are required for the full effect to be observed. It is thought that $\alpha\beta\gamma$ receptors form the majority of GABA_A receptors in neurones (Benke *et al.*, 1994) and the $\gamma 2S$ subunit is thought to be required for GABA_A receptors to be correctly targeted to the synapse (Essrich *et al.*, 1998). It is significant, therefore, that α -CaMK-II modulates γ 2S subunit-containing receptors to a greater degree than $\alpha\beta$ heteromers, as these are the synaptic and therefore more physiologically relevant receptor subtype. This finding also supports data from the previous chapter that α -CaMK-II is capable of significant modulation of synaptic GABA currents in cerebellar granule cells.

Modulation of GABA_A receptors by PKA and PKC has also been suggested to be dependent on both β and γ subunits, with both being required for the full effect to be observed (Krishek *et al.*, 1994). Ser⁴⁰⁹ on the β 1 subunit with Ser^{327/343} on the γ 2L were found to be important for PKC modulation in L929 fibroblasts (Krishek *et al.*, 1994; Lin *et al.*, 1996). Ser⁴¹⁰ and Ser³²⁷ were also found to be important when looking at PKC modulation of $\alpha 1\beta 2\gamma 2S$ receptors in *Xenopus oocytes* (Kellenberger *et al.*, 1992). In both cases it was suggested that phosphorylation of both β and γ subunits were required for the full PKC mediated effect to be apparent.

The β and γ subunits are thought to be the main targets of phosphorylation of GABA_A receptors. It may be that phosphorylation of both subunits is required in some cases to mediate a functional effect. Little is currently known about how the intracellular loops of different subunits might interact. There is evidence that γ 2 intracellular loops can interact with β subunit intracellular loops (Nymann-Andersen *et al.*, 2002). It is also thought intracellular loops between neighbouring receptors may be able to interact as the γ 2 TM3-4 intracellular loop can also associate with other γ 2 TM3-4 intracellular loops expressed as GST fusion proteins and this may play a role in receptor clustering (Nymann-Andersen *et al.*, 2002). It has become apparent that phosphorylation can be regulated by the targeting of kinases to specific sites and that anchoring proteins can significantly alter how a kinase behaves

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(Brandon *et al.*, 2002a). It may be that the phosphorylation of β and γ subunits is intricately linked within the receptor complex.

In the current study, $\beta 2$ subunit-containing receptors are not modulated by α -CaMK-II (similar to PKA). The presence of $\gamma 2$, therefore, is not enough on its own to ensure CaMK-II dependent modulation. In the case of $\beta 1$ subunit-containing receptors CaMK-II modulation is only observed with $\gamma 2S$ present. This implies that individual subunits cannot be regarded in isolation, but that the receptor as a whole and all its subunits together, determine the effects of phosphorylation. It might be that the β subunit acts in a permissive manner to allow $\gamma 2S$ related phosphorylation.

Cerebellar granule cells maintained in depolarising or more physiological conditions may both express $\beta 2/\beta 3$ subunits. In depolarising conditions it has been suggested there may be a higher level of $\beta 3$ subunit expression (Ives *et al.*, 2002a) possibly reflecting the immature nature of these cultures. This would be consistent with the large response to α -CaMK-II in these cultures and the large response to α -CaMK-II seen in NG108-15 cells expressing $\alpha 1\beta 3\gamma 2S$. In more physiological conditions, it is likely that the expression of GABA_A receptor subunits more closely follows that seen *in vivo* (Laurie *et al.*, 1992a; Wisden *et al.*, 1996). As a result $\beta 2$ and $\beta 3$ expression may both be significant, although $\beta 1$ levels would be expected to be low (Persohn *et al.*, 1992; Laurie *et al.*, 1992b). IPSCs in CGCs were modulated by α -CaMK-II but there was evidence for some variation in the response to α -CaMK-II of $\beta 3$ versus $\beta 2$ subunit-containing receptors.

Between subunit isoforms of the GABA_A receptor, there is typically around 70-90 % sequence homology (Moss and Smart, 1996). The β subunits are similar in structure with the main differences between isoforms related to the TM 3-4

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intracellular domain (Moss and Smart, 1996). It is known that the β 2 subunit is the most prevalent throughout the CNS (Benke *et al.*, 1994), but the β 3 subunit is thought to be expressed more predominantly at early stages in development in certain brain regions (Laurie *et al.*, 1992b). This is possibly significant given that β 3 appears to be strongly regulated by phosphorylation by a number of kinases. In the adult brain there are clear differences in the pattern of β 2 and β 3 subunit expression (Miralles *et al.*, 1999). As another example of the differences between these two subunits, β 2 subunit knock-out mice are apparently normal in development and behaviour (Sur *et al.*, 2001) whereas β 3 knock-out mice are severely developmentally impaired (Homanics *et al.*, 1997; DeLorey *et al.*, 1998).

It is surprising that recombinant receptors containing $\beta 2$ do not appear to be modulated by PKA (McDonald *et al.*, 1998) or CaMK-II given the large number of these receptors within the brain. There are also surprisingly few examples of $\beta 2$ subunit dependent phosphorylation in the literature, however, phosphorylation of Ser⁴¹⁰ by PKB causing an up-regulation of receptor number has been reported (Wang *et al.*, 2003). Ser⁴¹⁰ appears, so far, to be the main identified Ser/Thr kinase site of phosphorylation on this subunit (McDonald and Moss, 1997). Interestingly when $\beta 2$ Ser⁴¹⁰ is phosphorylated by PKC it appears to result in a down-regulation in receptor function (Kellenberger *et al.*, 1992) when receptors are expressed in *Xenopus oocytes*. Chapell *et al.* (1998) also observed this down-regulation and related it to a loss of receptors from the cell surface, however, in this case, mutation of Ser⁴¹⁰ had no effect on PKC modulation. Both of these studies involved activation of PKC via addition of phorbol 12-myristate 13-acetate (PMA). Differences may reflect variations in experimental protocol (application of PMA) or may of course reflect another example of the cellular environment, in this case *Xenopus oocytes*, affecting the functional consequences of phosphorylation.

The trafficking of receptors to the synapse can also be altered by phosphorylation and alterations in the number of receptors at the synapse have been suggested to mediate changes in synaptic strength (Kittler and Moss, 2003; Lüscher and Keller, 2004). PKC dependent phosphorylation of $\beta 2$ subunit-containing GABA_A receptors has been linked to alterations in the cell surface stability of the receptors (Chapell et al., 1998; Connolly et al., 1999). PKB dependent phosphorylation at Ser⁴¹⁰ has also been suggested to alter receptor number at the cell surface. PKA activity has been shown to differentially regulate trafficking and cell surface expression of $\beta 2$ and $\beta 3$ subunit-containing receptors in CGCs (Ives *et al.*, 2002b). Moreover, there is evidence that $\beta 2$ and $\beta 3$ subunits can be targeted differently within cells to different locations (Connolly et al., 1996). Interestingly, β^2 and $\beta^{1/3}$ subunits are reported to require different assembly signals within α^1 and γ^2 to be correctly assembled and trafficked to the cell surface (Bollan *et al.*, 2003). It is possible that phosphorylation may differentially affect the trafficking of these subunits and some of the variation in response to phosphorylation, observed in neurones, may be due to different levels of activation of mechanisms that alter the number of receptors at the synapse through changes in cell surface stability, or insertion, and those that alter the channel properties of the receptors directly. This type of model has also been proposed for modulation of AMPA receptors by CaMK-II (Hayashi et al., 2000; Shi et al., 2001; Malinow and Malenka, 2002).

The phenomena of rebound potentiation of $GABA_A$ receptor function that occurs in Purkinje cells of the cerebellum, is also CaMK-II dependent (Kano *et al.*, 1992; Kano *et al.*, 1996; Kawaguchi and Hirano, 2002). This cell type is thought to

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express predominantly $\alpha l\beta 2\gamma 2$ receptors (Persohn *et al.*, 1992; Laurie *et al.*, 1992b), however, this subunit combination, in a recombinant system, is not modulated by α -CaMK-II. As yet there is no apparent explanation for this; however, the lack of response to phosphorylation seen in recombinant systems expressing $\beta 2$ subunitcontaining receptors may reflect unique functional properties of such receptors. A possible explanation is the need for additional co-factors scaffolding/anchoring proteins for $\beta 2$ subunit phosphorylation to occur, or that phosphorylation dependent effects on $\beta 2$ subunit-containing receptors are indirect, occurring through downstream pathways or phosphorylation of associated proteins. Interestingly the insertion of AMPA receptors stimulated by CaMK-II activity is dependent on GluR1 receptors but not by their direct phosphorylation (Hayashi *et al.*, 2000). It could be speculated that modulation of $\beta 2$ subunit-containing receptors by CaMK-II is similar in that modulation of function occurs via downstream pathways / indirect effects and that this is difficult to observe in recombinant systems.

Expression of recombinant receptors in NG108-15 cells suggests that CaMK-II modulation of GABA_A receptors is mediated mainly through $\alpha 1\beta 3\gamma 2S$ receptors and partially through $\alpha 1\beta 1\gamma 2S$. Within the $\beta 1/\beta 3$ and $\gamma 2S$ subunits there are a number of potential sites for phosphorylation by α -CaMK-II (McDonald and Moss, 1994; McDonald and Moss, 1997). Mutation of these residues so they cannot be phosphorylated will allow determination of which sites are the most important on each subunit and also if CaMK-II mediates its effects through direct phosphorylation or whether it can act indirectly through other downstream signalling pathways.

6.4 CONCLUSIONS

- Recombinant $\alpha 1\beta 3$ receptors expressed in NG108-15 cells can be functionally up-regulated by α -CaMK-II.
- Recombinant $\alpha 1\beta 1$ and $\alpha 1\beta 2$ receptors are not functionally modulated by α -CaMK-II.
- Inclusion of the $\gamma 2S$ subunit to $\alpha 1\beta 3$ subunit-containing receptors enhances the modulation by α -CaMK-II.
- Inclusion of the $\gamma 2S$ subunit to $\alpha 1\beta 1$ subunit-containing receptors reveals a functional modulation by α -CaMK-II that is smaller than the effect observed for $\alpha 1\beta 3\gamma 2S$ receptors.
- Inclusion of the $\gamma 2S$ subunit to $\alpha 1\beta 2$ subunit-containing receptors has no effect. GABA_A receptors containing the $\beta 2$ subunit do not appear to be regulated by α -CaMK-II in a recombinant system.

<u>Chapter Seven:</u> <u>Sites of CaMK-II phosphorylation on recombinant GABA_A</u> <u>receptors expressed in NG108-15 cells and cerebellar</u> <u>granule cells.</u>

7.1 INTRODUCTION

Modulation of the GABA_A receptor by α -CaMK-II was observed in cerebellar granule cells and modulation of recombinant GABAA receptor subunits was also observed in the neuronal cell line NG108-15 (See Chapter 6). α -CaMK-II was capable of potentiating GABA currents recorded from NG108-15 cells expressing $\alpha 1\beta 3 / \alpha 1\beta 1\gamma 2S$ and $\alpha 1\beta 3\gamma 2S$ receptors implying that the $\beta 1/3$ and $\gamma 2S$ subunits were required for the effect. It was, however, unclear if α -CaMK-II mediated its effect through: direct phosphorylation of these subunits; phosphorylation of accessory proteins associated with them; or through activation of further downstream signalling pathways. In order to address these questions, potential phosphorylation sites on the β and $\gamma 2$ subunits were altered through sitedirected mutagenesis to prevent phosphorylation. These mutated recombinant receptors were then expressed in NG108-15 cells, which do not express endogenous GABA_A receptors, and the effect on α -CaMK-II modulation was determined. To confirm the importance of potential sites of phosphorylation recombinant receptors were also expressed in cerebellar granule cells in culture.

The use of site-directed mutagenesis and expression of recombinant receptors in a cell line with no endogenous receptors has been used successfully to determine the sites of phosphorylation important for modulation of GABA_A receptors by

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PKC/PKA and the tyrosine kinase, Src (Kellenberger *et al.*, 1992; Moss *et al.*, 1995; Lin *et al.*, 1996; McDonald *et al.*, 1998; Chapell *et al.*, 1998). In addition, GST fusion proteins, formed from the TM3-4 intracellular loop of different subunits, have been used to identify potential sites of phosphorylation by different kinases on GABA_A receptors (Machu *et al.*, 1993; McDonald and Moss, 1994; McDonald and Moss, 1997). The fusion proteins formed from the β 1 subunit could be phosphorylated by PKG, PKA, PKC and CaMK-II on Ser⁴⁰⁹, and on Ser³⁸⁴ by CaMK-II only (Moss *et al.*, 1992; McDonald and Moss, 1994). The Ser⁴⁰⁹ site was found to be important for PKC dependent functional modulation of GABA_A receptors (Krishek *et al.*, 1994; Lin *et al.*, 1996; McDonald *et al.*, 1998; Brandon *et al.*, 2002b; Hinkle and Macdonald, 2003).

The β 2 subunit has one main site of phosphorylation at Ser⁴¹⁰, (the equivalent of Ser⁴⁰⁹ in β 1/3). PKG, PKC, PKA and CaMK-II were all capable of phosphorylating this site on a GST fusion protein of the TM3-4 intacellular loop of the β 2 subunit (McDonald and Moss, 1997). This site has been shown to be important in PKC dependent modulation of whole-cell currents recorded from recombinant α 1 β 2 γ 2S receptors expressed in *Xenopus oocytes* (Kellenberger *et al.*, 1992). It has also been shown to be required for PKB mediated increases in the number of β 2 subunit-containing receptors at the cell surface of HEK293 cells and hippocampal neurones (Wang *et al.*, 2003a).

The use of GST-fusion proteins has also identified Ser⁴⁰⁹ as the major site of phosphorylation in the β 3 subunit. Using such constructs PKC, PKA, PKG and CaMK-II all phosphorylated this site. Moreover, Ser⁴⁰⁸ was additionally phosphorylated by PKC, and Ser³⁸³ was also subject to phosphorylation, but only by CaMK-II (See Table 1.1). In a recombinant system, Ser⁴⁰⁸ and Ser⁴⁰⁹

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phosphorylation of β 3 by PKA was required for an up-regulation of function and mutation of Ser⁴⁰⁸ converted the up-regulation into a down-regulation similar to that observed for PKA modulation of β 1 subunit-containing receptors (McDonald *et al.*, 1998). It appears, therefore, that there are three potential sites of CaMK-II phosphorylation on the β 3 subunit, Ser^{383,408,409}.

The $\gamma 2S$ subunit intracellular loop, expressed as a fusion protein, can be phosphorylated by PKC at Ser³²⁷, and at Ser³⁴⁸ and Thr³⁵⁰ by CaMK-II (Moss *et al.*, 1992; McDonald and Moss, 1994). Both PKC and CaMK-II were capable of phosphorylating Ser³⁴³ within the 8 amino acid insert that forms the $\gamma 2L$ splice variant (Moss *et al.*, 1992; Machu *et al.*, 1993; McDonald and Moss, 1994). PKA did not phosphorylate the $\gamma 2S$ subunit intracellular loop fusion protein (Moss *et al.*, 1992). Phosphorylation of Ser³²⁷ and Ser³⁴³ in $\gamma 2L$ by PKC have both been shown to be functionally important for GABA_A receptors expressed in heterologous systems (Kellenberger *et al.*, 1992; Krishek *et al.*, 1994; Lin *et al.*, 1996).

As well as Ser/Thr kinases, tyrosine kinases have also been shown to phosphorylate the GABA_A receptor and modulate function in both heterologous expression (Moss *et al.*, 1995; Wan *et al.*, 1997) and neuronal systems (Moss *et al.*, 1995; Valenzuela *et al.*, 1995). Tyrosine kinase activity has been shown to potentiate GABA currents (Moss *et al.*, 1995; Huang and Dillon, 1998) including IPSC amplitudes (Boxall, 2000; Ma *et al.*, 2003). It has been reported that application of pp60^{c-src} to HEK293 cells resulted in phosphorylation of $\beta 2/3$ subunits (Wan *et al.*, 1997). Modulation of $\alpha 1\beta 1$ receptors expressed in *Xenopus oocytes* by tyrosine kinase activity has also been reported (Valenzuela *et al.*, 1995). In addition phosphorylation of tyrosines 384 and 386 within the $\beta 1$ subunit has also been reported, but in this case, phosphorylation appeared to be functionally silent (Moss *et* *al.*, 1995). Mutation of the tyrosine residues at 365 and 367 on the γ 2L subunit (corresponding to Y357 and Y359 on γ 2S) was found to block functional modulation by Src of GABA_A receptors in HEK293 cells (Moss *et al.*, 1995). It appears therefore that these two residues may be the most functionally relevant sites of phosphorylation on the γ 2 subunit. These sites have been shown to be phosphorylated within the brain and can be dynamically regulated by tyrosine phosphatase activity in cultured cortical neurones (Brandon *et al.*, 2001).

Thus, there are a number of sites identified on GABA_A receptors that may be directly phosphorylated by CaMK-II. Expression of $\alpha\beta$ heteromers in NG108-15 cells suggests that β 3 is the only β subunit that could be directly phosphorylated by α -CaMK-II. In addition, γ 2S subunits appeared to play a role in α -CaMK-II modulation of $\alpha 1\beta 1\gamma 2S$ and $\alpha 1\beta 3\gamma 2S$ receptors. A series of mutant receptor subunits were prepared, with potential sites of phosphorylation ablated so they could not be phosphorylated, using site-specific mutagenesis (See Chapter 2.1). These mutant receptors were expressed in NG108-15 cells and cerebellar granule cells to determine the functional effect on α -CaMK-II modulation.

7.2 RESULTS

In each case, NG108-15 cells and cerebellar granule cells were transfected with $\alpha\beta$ or $\alpha\beta\gamma$ subunit cDNA combinations, in equal ratios, along with enhanced GFP for identification of transfected cells. 60 nM α -CaMK-II (pre-activated) was introduced into the cells via the patch pipette solution (See Chapter 2 and 3.2.1). Control recordings consisted of a normal patch pipette solution containing the preactivation buffer without α -CaMK-II. Whole-cell currents were recorded in response to GABA and the percentage change in the peak current amplitude was recorded over time. Current amplitudes were then normalised to the peak amplitude recorded 3-4 min after first achieving the whole-cell configuration.

7.2.1 α -CaMK-II modulation of GABA_A receptor mutant $\alpha\beta$ heteromers.

The $\alpha 1\beta 3$ receptor was selected as this exhibited the most significant modulation by α -CaMK-II (See Fig 6.2). All three potential phosphorylation sites, Ser^{383,408,409} were converted to alanines within the $\beta 3$ subunit. It was presumed that this would remove all the potential sites of CaMK-II phosphorylation within this subunit. On the sequential application of 10 μ M GABA over 20 min in control conditions, the amplitude of GABA currents recorded was quite stable with little rundown.

On application of 60 nM α -CaMK-II, no significant effect was seen on the amplitude of the currents recorded (93.2 ± 7.6 %, n = 8) as compared to the controls (101.3 ± 4.6 %, n = 5 at t = 6 min, Fig 7.1A) indicating that phosphorylation of at least one of these residues was capable of mediating the α -CaMK-II modulation of the $\alpha 1\beta$ 3 receptor. PKA phosphorylation of β 3 has been reported to increase the amplitude of GABA_A receptor mediated currents through phosphorylation of Ser^{408,409}. However, the application of 60 nM α -CaMK-II to $\alpha 1\beta 3^{S408A,S409A}$ receptors resulted in a significant potentiation of the amplitude of currents recorded to 120.6 ± 6.2 % (n = 7) as compared to control, 93.9 ± 3.8 % (n = 4) at t = 6 min, which was not significantly different from the effect of 60 nM α -CaMK-II on $\alpha 1\beta 3$ wild-type receptors (118 ± 4.8 %, n = 7, Fig 7.1B). This effect of α -CaMK-II had a similar time-course to the effect on wild-type receptors reaching a steady-state within 6 min.

Although Ser^{408,409} appeared unimportant for the α -CaMK-II effect, application of 60 nM α -CaMK-II to $\alpha 1\beta 3^{S383A}$ receptors resulted in no significant increase in amplitude (94 ± 6.8 %, n = 5) as compared to control (101.7 ± 1.5 %, n = 6 at t = 6 min, Fig 7.1C). This indicates that α -CaMK-II modulation of the $\alpha 1\beta 3$ receptor is mediated solely through phosphorylation of Ser³⁸³.

Fig 7.1:



Fig 7.1: α -CaMK-II modulation of $\alpha 1\beta 3$ heteromers is mediated through phosphorylation of Ser³⁸³.

Peak amplitudes of GABA-mediated currents recorded following the application of 10 μ M GABA to transfected NG108-15 cells expressing: $\alpha 1\beta 3^{S383A,S408A,S409A}$; $\alpha 1\beta 3^{S408A,S409A}$ or $\alpha 1\beta 3^{S383A}$ GABA_A receptors. GABA was re-applied at 2 min intervals for up to 20 min. All currents were normalised to the peak amplitude of the response measured during the first 3-4 min (= 100 %) after achieving the whole-cell configuration (t = 0 min). Control recordings were made with a normal patch pipette solution supplemented with the pre-activation buffer without α -CaMK-II.

(A) GABA currents recorded from NG108-15 cells transfected with $\alpha 1\beta 3^{S383A,S408A,S409A}$ subunits in the absence (Control, n = 5) and presence of internal 60 nM pre-activated α -CaMK-II (n = 8). In this figure, $\alpha 1\beta 3$ wild-type responses to 60 nM CaMK-II (taken from Fig 6.2C) are shown for comparison (n = 7).

(B) GABA currents recorded from NG108-15 cells transfected with $\alpha 1\beta 3^{S408A,S409A}$ subunits in the absence (Control, n = 4) and the presence of 60 nM α -CaMK-II (n = 7).

(C) GABA currents recorded from NG108-15 cells transfected with $\alpha 1\beta 3^{S383A}$ subunits in the absence (Control, n = 6) and the presence of 60 nM α -CaMK-II (n = 5).

All points = mean \pm s.e.

Significance is indicated by the symbol *, CaMK-II compared to control and *, CaMK-II wild-type compared to CaMK-II mutant.

P < 0.05, t-test, ANOVA.

7.2.2 α -CaMK-II modulation of mutant $\alpha 1\beta 3\gamma 2S$ subunit-containing receptors.

CaMK-II modulation of $\alpha 1\beta 3\gamma 2S$ receptors appeared significantly larger when compared to the modulation of $\alpha 1\beta 3$ receptors (Figs 6.2C, 6.3C) indicating that α -CaMK-II modulation may be partly mediated through the $\gamma 2S$ subunit. Application of 60 nM α -CaMK-II to $\alpha 1\beta 3^{S383A,S408A,S409A}\gamma 2S$ receptors still resulted in a significant potentiation of the amplitude of GABA (10 μ M) currents recorded to 117.8 ± 5 % (n = 8) as compared to control, 94.9 ± 4 % (n = 4, Fig 7.2A) at t = 6 min. However, this effect was significantly smaller (107.5 ± 3.5 %, n = 8) than the effect of α -CaMK-II on $\alpha 1\beta 3\gamma 2S$ wild-type receptors (145.8 ± 10.8 %, n = 7) at t = 4 min. This confirmed the likelihood that α -CaMK-II mediated part of its functional modulation via the $\gamma 2S$ subunit and that mutation of all known sites of phosphorylation in the $\beta 3$ subunit did not remove the CaMK-II modulation. The time-course of the residual response appeared slightly slower than for wild-type receptors and the effect was also transient with amplitudes returning to baseline levels within 20 min.

Analysis of the potential sites for CaMK-II phosphorylation on the $\gamma 2S$ subunit, using fusion proteins, suggested that the main sites involved were Ser³⁴⁸ and Thr³⁵⁰. Application of 60 nM α -CaMK-II to $\alpha 1\beta 3\gamma 2S^{S348A,T350A}$ receptors resulted in a significant potentiation of GABA currents recorded to 142.7 ± 11.2 % (n = 8) as compared to control, 101.3 ± 6.7 % (n = 5, Fig 7.2B) at t = 6 min. Interestingly, this was not significantly different from the potentiation seen when CaMK-II was applied to wild-type $\alpha 1\beta 3\gamma 2S$ receptors indicating that these residues are not functionally important in modulating the amplitude of $\alpha 1\beta 3\gamma 2S$ receptor mediated currents via CaMK-II phosphorylation.

The application of 60 nM α -CaMK-II to the combined mutant receptor $\alpha 1\beta 3^{S383A,S408A,S409A}\gamma 2S^{S348A,T350A}$ also resulted in a significant potentiation to 127.4 ± **9** % (n = 8) as compared to control, 104.2 ± 2.8 % (n = 4, Fig 7.2C) at t = 6 min. This was not significantly different from that seen on $\alpha 1\beta 3^{S383A,S408A,S409A}\gamma 2S$ receptors, further indicating that phosphorylation of Ser³⁴⁸ and Thr³⁵⁰, in this case, does not mediate the effect of α -CaMK-II. Again, the effect was transient and by t = 10 min the effect of α -CaMK on $\alpha 1\beta 3^{S383A,S408A,S409A}\gamma 2S^{S348A,T350A}$ receptors was significantly different (114.2 ± 6.4 %, n = 8) from the effect on wild-type (148.5 ± 12.2 %, n = 7) but was not significantly different from control (102.7 ± 5.8 %, n = 4, Fig7.2C).

Application of 10 μ M diazepam to $\alpha 1\beta 3\gamma 2S^{S348A,T350A}$ receptors resulted in potentiation of GABA currents, indicating that the $\gamma 2S^{S348A,T350A}$ subunit was expressed at the cell surface (Fig 7.2D). In addition, if this subunit was not expressed, it would be predicted that α -CaMK-II would be without effect as previous results indicated α -CaMK-II could not functionally modulate $\alpha 1\beta 3^{S383A,S408A,S409A}$ receptors (Fig 7.1A), however, a significant increase in amplitude was observed on α -CaMK-II application to $\alpha 1\beta 3^{S383A:S408A:S409A}\gamma 2S^{S348A,T350A}$ receptors suggesting an $\alpha\beta\gamma$ heteromer was correctly assembled and expressed (Fig 7.2C).



Fig 7.2: CaMK-II modulation of $\gamma 2S$ subunit-containing receptors is not mediated through phosphorylation of GABA_A Ser³⁴⁸ and Thr³⁵⁰.

Peak amplitude GABA mediated currents recorded following the application of 10 μ M GABA to transfected NG108-15 cells expressing: $\alpha 1\beta 3^{S383A,S408A,S409A}\gamma 2S$; $\alpha 1\beta 3\gamma 2S^{S348A:T350A}$ or $\alpha 1\beta 3^{S383A,S408A,S409A}\gamma 2S^{S348A,T350A}$ GABA_A receptors. GABA was re-applied at 2 min intervals for up to 20 min. All currents were normalised to the peak amplitude of the response measured during the first 3-4 min (= 100 %) after achieving the whole-cell configuration (t = 0 min). Control recordings were made with normal patch pipette solution supplemented with the pre-activation buffer without α -CaMK-II.

(A) GABA currents recorded from NG108-15 cells transfected with $\alpha 1\beta 3^{S383A,S408A,S409A}\gamma 2S$ subunits in the absence (Control, n = 4) and presence of 60 nM pre-activated α -CaMK-II (n = 8). In this figure, $\alpha 1\beta 3\gamma 2S$ wild-type responses to 60 nM CaMK-II (taken from Fig 6.3C) are shown for comparison (n = 7).

(B) GABA currents recorded from NG108-15 cells transfected with $\alpha 1\beta 3\gamma 2S^{S348A,T350A}$ subunits in the absence (Control, n = 5) and presence of 60 nM α -CaMK-II (n = 8).

(C) GABA currents recorded from NG108-15 cells transfected with $\alpha 1\beta 3^{S383A,S408A,S409A}\gamma 2S^{S348,T350A}$ subunits in the absence (n = 4) and the presence of 60 nM α -CaMK-II (n = 8).

(D) Representative currents recorded from NG108-15 cells expressing $\alpha 1\beta 3\gamma 2S^{S348A,T350A}$ receptors, before and after bath application of 10 μ M diazepam.

All points = mean \pm s.e.

Significance is indicated by the symbol *, CaMK-II compared to control and the symbol *, CaMK-II wild-type compared to CaMK-II mutant. P < 0.05, t-test, ANOVA.

7.2.3 Cross-talk between CaMK-II and tyrosine kinase phosphorylation of the $\alpha 1\beta 3\gamma 2S$ receptor.

Analysis of fusion proteins in vitro suggested that Ser³⁴⁸ and Thr³⁵⁰ were the main sites of CaMK-II phosphorylation on the y2S subunit however mutation of these sites to alanines clearly had no effect on α -CaMK-II modulation of $\alpha 1\beta 3\gamma 2S$ receptors. This left three possibilities, either CaMK-II was capable of phosphorylating y2S at another site in an NG108-15 cell that it could not phosphorylate in vitro; or CaMK-II was acting on the $\gamma 2S$ subunit through phosphorylation of an accessory protein; or CaMK-II was acting through the subsequent activation of another downstream kinase. It has previously been shown that phosphorylation of two tyrosine residues in the $\gamma 2S$ subunit can result in an increase in whole-cell current amplitudes similar in magnitude to that observed following the action of α -CaMK-II on γ 2S subunit-containing receptors (Moss *et al.*, 1995). Therefore, 100 µM genistein an inhibitor of tyrosine kinase activity (Akiyama et al., 1987), was co-applied in the patch pipette solution with 60 nM α -CaMK-II. It was not possible to pre-apply genistein in the bath, because if applied extracellularly, genistein is capable of directly inhibiting the GABAA receptor (Dunne et al., 1998). Application of 60 nM CaMK-II and 100 µM genistein to $\alpha 1\beta 3\gamma 2S$ receptors resulted in a significant increase in the amplitude of currents recorded to 120.3 ± 7.6 % (n = 7) as compared to control, 100.2 ± 2.3 % (n = 4, Fig. 7.3A) at t = 4 min. This potentiation however was significantly smaller than that seen on application of α -CaMK-II alone (145.8 \pm 10.8 %, n = 7). It should also be noted in this case that application of genistein alone had no effect on the amplitude of currents recorded over 20 min. This indicated that, in this cell type, the γ 2S subunit is not subject to significant basal phosphorylation by tyrosine kinases. Application

of 60 nM α -CaMK-II and 100 μ M genistein to $\alpha 1\beta 3^{S383A}\gamma 2S$ receptors resulted in no significant changes in amplitude (103.6 ± 2.8 %, n= 7) over 20 min as compared to control (97 ± 7.1 %, n = 4 at t = 6 min, Fig 7.3B). This suggested that phosphorylation at Ser³⁸³ in the $\beta 3$ subunit and α -CaMK-II dependent activation of tyrosine kinase activity was fully responsible for the modulation of GABA_A receptors by α -CaMK-II.

The application of 60 nM α -CaMK-II to $\alpha 1\beta 3\gamma 2S^{Y357F,Y359F}$ receptors resulted in a significant potentiation of the amplitude of currents recorded to 126.2 ± 4 % (n = 7) as compared to control, 107.6 ± 6.5 % (n = 4, Fig 7.3C) at t = 4 min. However, as before, this potentiation was significantly less than the effect of α -CaMK-II on $\alpha 1\beta 3\gamma 2S$ receptors (145.7 ± 10.2 %, n = 7) at t = 6 min. However after 8-10 min of recording, there was no significant difference in response to α -CaMK-II as compared to wild-type. This coincides with the observation that the response to α -CaMK-II mediated by the $\gamma 2S$ subunit is transient and is no longer evident after 10-15 min.

Finally, application of 60 nM α -CaMK-II to $\alpha 1\beta 3^{S383A}\gamma 2S^{Y357F,Y359F}$ receptors abolished the response to α -CaMK-II with no significant effect on the amplitude of currents recorded over 20 min (103.3 ± 3.4 %, n = 6, at t = 6 min) as compared to control (108.4 ± 4.3 %, n = 6, at t = 6 min, Fig 7.3D). This indicated that CaMK-II dependent phosphorylation of $\beta 3$ Ser³⁸³ and $\gamma 2S$ Tyr^{357,359} mediates α -CaMK-II dependent modulation of $\alpha 1\beta 3\gamma 2S$ receptors. This suggested a possible direct site of phosphorylation by α -CaMK-II and the activation of a second signalling pathway, acting through tyrosine kinase phosphorylation of the $\gamma 2S$ subunit.

Application of 10 μ M diazepam to $\alpha 1\beta 3\gamma 2S^{Y357F,Y359F}$ receptors resulted in a potentiation of the current recorded indicating that the $\gamma 2S^{Y357F,Y359F}$ subunit was expressed and correctly assembled at the cell surface (Fig 7.4A). It has been

observed that the addition of the γ subunit to $\alpha\beta$ heteromers increases the amplitudes of the GABA currents recorded (Angelotti *et al.*, 1993). The current density recorded from NG108-15 cells expressing $\alpha1\beta3$ (20.2 ± 11.5 pA/pF, n = 6) was significantly increased for $\alpha1\beta3\gamma2S$ receptors (94.1 ± 19 pA/pF, n = 6, Fig 7.4B). There were no significant differences in current density between $\alpha1\beta3\gamma2S$, $\alpha1\beta3\gamma2S^{S348A,T350A}$ (87.1 ± 21.3 pA/pF, n=7) and $\alpha1\beta3\gamma2S^{Y357F,Y359F}$ (82.1 ± 15.6 pA/pF, n = 6, Fig 7.4B) receptors, indicating that site-specific mutation of phosphorylation sites in the $\gamma2S$ subunit was unlikely to interfere with subunit expression or trafficking to the cell surface.

Fig 7.3:



Fig 7.3: α -CaMK-II modulation of $\alpha 1\beta 3\gamma 2S$ subunit-containing GABA_A receptors is mediated partly through tyrosine kinase phosphorylation of Tyr^{357,359}.

Peak amplitudes of GABA mediated currents recorded following the application of 10 μ M GABA to transfected NG108-15 cells expressing: $\alpha 1\beta 3\gamma 2S$; $\alpha 1\beta 3^{S383A}\gamma 2S$; $\alpha 1\beta 3^{\gamma} 2S^{\gamma} 57F, \gamma 359F}$ or $\alpha 1\beta 3^{S383A}\gamma 2S^{\gamma} 57F, \gamma 359F}$ GABA_A receptors. GABA was reapplied at 2 min intervals for up to 20 min. All currents were normalised to the peak amplitude of the response measured during the first 3-4 min (= 100 %) after achieving the whole-cell configuration (t = 0 min). Control recordings were made with normal patch pipette solution supplemented with the pre-activation buffer without α -CaMK-II.

(A) GABA currents recorded from NG108-15 cells transfected with $\alpha 1\beta 3\gamma 2S$ subunits in the absence (Control, n = 4) and presence of 60 nM pre-activated α -CaMK-II (n = 7) which also contained 100 μ M genistein. In this panel and C $\alpha 1\beta 3\gamma 2S$ wild-type responses to 60 nM CaMK-II (taken from Fig 6.3C) are shown for comparison (n = 7).

(B) GABA currents recorded from NG108-15 cells transfected with $\alpha 1\beta 3^{S383A}\gamma 2S$ subunits in the absence (Control, n = 4) and the presence of internal 60 nM α -CaMK-II (n = 7) but always including 100 μ M genistein.

(C) GABA currents recorded from NG108-15 cells transfected with $\alpha 1\beta 3\gamma 2S^{Y357F,Y359F}$ subunits in the absence (Control, n = 4) and the presence of 60 nM α -CaMK-II (n = 7).

(D) GABA currents recorded from NG108-15 cells transfected with $\alpha 1\beta 3^{S383A}\gamma 2S^{Y357F,Y359F}$ subunits in the absence (Control, n = 6) and the presence of 60 nM α -CaMK-II (n = 6).

All points = mean \pm s.e.

Significance is indicated by the symbol *, CaMK-II compared to control, the symbol *, CaMK-II wild-type compared to CaMK-II mutant and the symbol *, CaMK-II and inhibitor compared to control.

P < 0.05, t-test, ANOVA.





Fig 7.4: $\gamma 2S^{S348A,T350A}$ and $\gamma 2S^{Y357F,Y359F}$ receptor subunits are expressed at the cell surface.

(A) Representative GABA currents recorded from NG108-15 cells expressing $\alpha 1\beta 3\gamma 2S^{Y357F,Y359F}$ receptors before and after bath application of 10 μ M diazepam.

(B) Bar chart showing the mean current density (pA/pF) of NG108-15 cells expressing $\alpha 1\beta 3$ (n = 6), $\alpha 1\beta 3\gamma 2S$ (n = 6), $\alpha 1\beta 3\gamma 2S^{S348A,T350A}$ (n = 7) and $\alpha 1\beta 3^{S383A}\gamma 2S^{Y357F,Y359F}$ (n = 6) GABA receptors after application of 10 μ M GABA at t = 0 min in control conditions.

Significance is indicated by the symbol *, $\alpha 1\beta 3$ compared to $\alpha 1\beta 3\gamma 2S$. P < 0.05, t-test.

7.2.4 Cross-talk between CaMK-II and tyrosine kinases acting on $\alpha 1\beta 1\gamma 2S$ receptors.

 α -CaMK-II was capable of modulating $\alpha 1\beta 1\gamma 2S$ receptors, but not $\alpha 1\beta 1$ receptors, suggesting that modulation of $\alpha 1\beta 1\gamma 2S$ receptors occurred primarily through the $\gamma 2S$ subunit. In order to confirm if the same tyrosine kinase dependent mechanism revealed for $\alpha 1\beta 3\gamma 2S$ receptors, was also involved, 60 nM α -CaMK-II and 100 μ M genistein were internally applied to $\alpha 1\beta 1\gamma 2S$ receptors. In this case there was no significant effect on the amplitude of currents recorded (98.5 ± 5 %, n = 6) as compared to control (105.1 ± 2.5 %, n = 4, Fig 7.5A) at t = 6 min. The application of genistein in control conditions again had no significant effect on the amplitude of the GABA currents recorded, suggesting a lack of any basal tyrosine kinase phosphorylation of these receptors in NG108-15 cells.

Application of 60 nM α -CaMK-II to $\alpha 1\beta 1\gamma S^{Y3576F,Y359F}$ receptors abolished the effect of α -CaMK-II, with no significant difference in the amplitude of GABA currents (99.7 ± 5.5 %, n = 7) compared to control (102.1 ± 3.5 %, n = 4, Fig 7.5B) at t = 6 min. This indicated that the effect of α -CaMK-II on $\alpha 1\beta 1\gamma 2S$ receptors was entirely mediated by CaMK-II dependent activation of tyrosine kinase activity.

The current density of $\alpha 1\beta 1\gamma 2S$ receptors (87.9 ± 16.3 pA/pF, n = 6) was significantly higher than the current density of $\alpha 1\beta 1$ receptors (19.5 ± 4.8 pA/pF, n = 6) and the current density of $\alpha 1\beta 1\gamma 2S^{Y357F,Y359F}$ receptors was not significantly different (75.2 ± 32 pA/pF, n = 6, Fig 7.5C) from $\alpha 1\beta 1\gamma 2S$. This indicated that the $\gamma 2s^{Y357F,Y359F}$ subunit was most likely expressed at the cell surface and assembled correctly with $\alpha 1\beta 1$.



Fig 7.5: CaMK-II modulation of $\alpha 1\beta 1\gamma 2S$ subunit-containing GABA_A receptors is mediated through tyrosine kinase phosphorylation of Tyr^{357,359}.

Peak amplitudes of GABA-mediated currents recorded following the application of 10 μ M GABA to transfected NG108-15 cells expressing $\alpha 1\beta 1\gamma 2S$ and $\alpha 1\beta 1\gamma 2S^{Y357F,Y359F}$ GABA_A receptors. GABA was re-applied at 2 min intervals for up to 20 min. All currents were normalised to the peak amplitude of the response measured during the first 3-4 min (= 100 %) after achieving the whole-cell configuration (t = 0 min). Control recordings were made with normal patch pipette solution supplemented with the pre-activation buffer without α -CaMK-II.

(A) GABA currents recorded from NG108-15 cells transfected with $\alpha 1\beta 1\gamma 2S$ subunits in the absence (Control, n = 4) and presence of 60 nM pre-activated α -CaMK-II with 100 μ M genistein present throughout (n = 6). Data for $\alpha 1\beta 1\gamma 2S$ wild-type responses to 60 nM CaMK-II (taken from Fig 6.3A) are shown for comparison (n = 11).

(B) GABA currents recorded from NG108-15 cells transfected with $\alpha 1\beta 1\gamma 2S^{Y357F,Y359F}$ subunits in the absence (Control, n = 4) and the presence of 60 nM α -CaMK-II (n = 7).

(C) Bar chart showing the mean current densities (pA/pF) of NG108-15 cells expressing: $\alpha l\beta l$ (n = 6), $\alpha l\beta l\gamma 2S$ (n = 6), $\alpha l\beta l\gamma 2S^{Y357,Y367F}$ (n = 6) GABA_A receptors after application of 10 μ M GABA at t = 0 min in control conditions.

All points = mean \pm s.e.

Significance is indicated by the symbol *, CaMK-II wild-type compared to CaMK-II mutant or CaMK-II and inhibitor.

P < 0.05, t-test, ANOVA.

7.2.5 Recombinant GABA_A receptors expressed in cerebellar granule cells are expressed at the cell surface.

There is a certain amount of discrepancy between the reported effects of phosphorylation in recombinant systems and that observed in neuronal systems (See Chapter 1). In addition, it has been suggested that the expression of recombinant receptors in different cell lines can alter the properties of the receptors expressed (Ueno *et al.*, 1996; Mercik *et al.*, 2003; Everitt *et al.*, 2004) as has also been demonstrated in the current study (See Chapter 3). In order to confirm the importance of the sites of phosphorylation identified in NG108-15 cells in a neuronal system, recombinant GABA_A receptors were expressed in cerebellar granule cells in culture.

A number of studies have reported that expression of single GABA_A receptor subunits in neurones allows expression of these recombinant receptors at the surface through assembly with endogenous subunits (Okada *et al.*, 2000; Cheng *et al.*, 2001). However a number of other studies have reported that recombinant single subunits cannot be expressed at the cell surface without co-transfection of the other recombinant GABA_A receptor subunits required for cell surface expression (Gorrie *et al.*, 1997; Kittler *et al.*, 2000).

Cerebellar granule cells maintained in depolarising conditions were transfected using Effectene (See Chapter 2) with α l GABA_A receptor subunit cDNA co-expressed with β 2 subunit cDNA containing a functionally silent extracellular myc epitope (See Chapter 2) inserted at the N terminus (Connolly *et al.*, 1996a; Connolly *et al.*, 1996b; Farrar *et al.*, 1999). The presence of recombinant receptors at the cell surface was determined by immunocytochemistry. Cultures were immunolabeled with a primary antibody directed against the myc epitope. As these

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cells were not permeabilised and the myc epitope was inserted in a part of the subunit expected to be extracellular, the presence of fluorescence would indicate cell surface expression of the recombinant subunit.

In order to observe strong cell surface expression in the current study it was observed that β subunits had to be co-expressed with a recombinant α subunit. Coexpression of α l with β 2 and α l with β 3 resulted in strong cell surface expression suggesting that these granule cells now expressed a large number of $\alpha\beta$ heteromers at the cell surface most likely in excess of endogenous receptors given the strong viral promoter within the plasmid cDNA (See Fig 7.6A and B).

Co-expression of $\alpha 1$ with $\beta 3^{S408A,S409A}$ and $\beta 3^{S383A}$ revealed that despite the disruption to the phosphorylation sites in these recombinant subunits they were still strongly expressed at the cell surface of cerebellar granule cells after transfection (See Fig 7.6 C and D). Co-expression of $\gamma 2S^{myc}$ with $\alpha 1$ and $\beta 2$ or $\beta 3$ resulted in strong surface expression suggesting that these cells now predominantly express recombinant $\alpha\beta\gamma$ GABA_A receptors (See Fig 7.7 A and B). Co-expression of $\alpha 1$ and $\beta 3$ with $\gamma 2S^{Y357F,Y359F,myc}$ also resulted in expression of the recombinant $\alpha 1\beta 3\gamma 2^{Y357F,Y359F,myc}$ receptors at the cell surface (See Fig 7.7C).













α1β3^{S383A,myc}

D

в







Fig 7.6:

С

Fig 7.6: Expression of recombinant $\alpha\beta$ subunit-containing GABA_A receptors at the cell surface of transfected cerebellar granule cells:

Cultures of rat cerebellar granule cells maintained in depolarising conditions were transfected with $\alpha 1\beta 2^{myc}$, $\alpha 1\beta 3^{myc}$, $\alpha 1\beta 3^{S408A,S409A,myc}$ and $\alpha 1\beta 3^{S383A,myc}$ GABA_A receptor subunits along with enhanced GFP for identification of transfected cells. Cells were immunolabelled with an anti-myc epitope antibody without permeabilisation so that only subunits at the cell-surface were detected. Images were obtained on a Zeiss Meta 510 laser scanning confocal microscope. Unless otherwise stated each image consists of a Z series of 8-10 images at intervals of 0.3-0.4 μ m (See Chapter 2). Each image is representative of at least five different transfected cells (GFP expressing). All transfected cells observed showed cell surface expression of recombinant receptors.

(A) Cerebellar granule cell transfected with $\alpha 1\beta 2^{myc}$ GABA_A receptor subunits. First panel shows GFP fluorescence indicating transfection (Green), the second panel shows anti-myc immunolabelling (red) and the third panel is an overlay of the two showing co-localisation (yellow).

(B) Cerebellar granule cell transfected with $\alpha 1\beta 3^{myc}$ GABA_A receptor subunits.

(C) Cerebellar granule cell transfected with $\alpha 1\beta 3^{S408A,S409A,myc}$ GABA_A receptor subunits.

(**D**) Cerebellar granule cell transfected with $\alpha 1\beta 3^{S383A,myc}$ GABA_A receptor subunits. (Calibration bar = 10 μ m)

Fig 7.7:



Fig 7.7: Expression of recombinant $\alpha\beta\gamma$ subunit-containing GABA_A receptors at the cell surface of transfected cerebellar granule cells:

Cultures of rat cerebellar granule cells maintained in depolarising conditions were transfected with $\alpha 1\beta 3\gamma 2S^{myc}$, $\alpha 1\beta 2\gamma 2S^{myc}$ and $\alpha 1\beta 3\gamma 2S^{Y357F,Y359F,myc}$ GABA_A receptor subunits along with enhanced GFP for identification of transfected cells. Cells were immunolabelled with an anti-myc epitope antibody without permeabilisation so that only subunits at the cell-surface were detected (See Chapter 2). Each image is representative of at least five different transfected cells (GFP expressing). All transfected cells observed showed cell surface expression of recombinant receptors.

(A) Cerebellar granule cell transfected with $\alpha 1\beta 3\gamma 2S^{myc}$ GABA_A receptor subunits. (B) Cerebellar granule cell transfected with $\alpha 1\beta 2\gamma 2S^{myc}$ GABA_A receptor subunits. (C) Cerebellar granule cell transfected with $\alpha 1\beta 3\gamma 2S^{Y357F,Y359F,myc}$ GABA_A receptor subunits. (Calibration bar = 10 μ m)

7.2.6 α -CaMK-II modulation of recombinant $\alpha\beta$ heteromers expressed in cerebellar granule cells.

Cerebellar granule cells transfected with GFP only were recorded from with normal patch pipette solution supplemented with pre-activation buffer without CaMK-II. The whole-cell GABA currents recorded remained stable throughout 16 minutes of recording (See Fig 7.8). The resting potential of transfected cerebellar granule cells was recorded as an indication that transfection did not significantly alter cell physiology. The resting potential of transfected granule cells was not significantly different (-43.9 \pm 2.8 mV, n = 8) to untransfected granule cells -39.2 \pm 2.5 mV (n = 6).

Application of 60 nM α -CaMK-II to granule cells expressing $\alpha 1\beta 3$ heteromers resulted in a significant potentiation of the amplitude of whole-cell currents to 130.6 ± 6.3 % at t = 4 min (n = 7) as compared to 60 nM α -CaMK-II applied to granule cells expressing $\alpha 1\beta 2$ heteromers (105.6 ± 2.6 %, n = 7, at t = 4 min, Fig 7.8A). Application of 60 nM α -CaMK-II to $\alpha 1\beta 3^{S408A,S409A}$ expressing granule cells resulted in a significant increase in amplitude to 127.1 ± 8 % (n = 4) at t = 6 min as compared to application of α -CaMK-II to $\alpha 1\beta 3^{S383A}$ expressing granule cells (104.2 ± 2.6 %, n = 4, at t = 6 min, Fig 7.8B), which for $\alpha 1\beta 3^{S408A,S409A}$, was not significantly different from the application of α -CaMK-II to wild-type $\alpha 1\beta 3$ heteromers.

The response of cerebellar granule cells expressing recombinant GABA_A receptors to α -CaMK-II was similar to that observed in NG108-15 cells (See Fig 7.1). Again, the results indicated that $\alpha 1\beta 2$ heteromers were not modulated by α -CaMK-II in this system and that Ser³⁸³ on the $\beta 3$ subunit appeared to mediate the α -CaMK-II dependent effect on $\alpha 1\beta 3$ heteromers. The extent of α -CaMK-II

modulation was also similar to that observed in NG108-15 cells, suggesting that in granule cells the $\gamma 2$ subunit may also mediate part of the response to CaMK-II. It also suggested that the expressed recombinant $\alpha\beta$ heteromers dominated over endogenous receptors in granule cells which would be expected to consist of $\alpha\beta\gamma$ receptors.

7.2.7 α -CaMK-II modulation of $\alpha\beta\gamma$ receptors expressed in cerebellar granule cells.

Application of 60 nM α -CaMK-II to granule cells expressing $\alpha l\beta 3\gamma 2S$ recombinant receptors resulted in a significant potentiation of the amplitude of whole-cell currents to 167.8 ± 13.7 % (n = 4) at t = 6 min, as compared to granule cells expressing $\alpha l\beta 2\gamma 2S$ recombinant receptors (112.8 ± 8 %, n = 7, at t = 6 min, Fig 7.9A). In addition this increase was significantly greater than the effect of α -CaMK-II on $\alpha l\beta 3$ expressing granule cells at t = 6 min (127.1 ± 8 %, n = 7, Fig 7.8A). This indicated that similar to NG108-15 cells, the $\gamma 2S$ subunit mediated part of the GABA_A receptor response to α -CaMK-II. In contrast to NG108-15 cells (Fig 6.3), application of α -CaMK-II to $\alpha l\beta 2\gamma 2S$ expressing granule cells resulted in a significant but transient potentiation of whole cell currents to 115.6 ± 4.2 % (n = 7) at t = 4 min as compared to control (96.4 ± 6.1 %, n =7, at t = 4 min, Fig 7.8A). This was possibly as a result of CaMK-II modulation of a population of endogenous GABA_A receptors. Again, this suggests that the recombinant receptor expression dominates over the endogenous receptors when measuring whole-cell currents from transfected cells.

Application of 60 nM α -CaMK-II to granule cells expressing $\alpha 1\beta 3\gamma 2S^{Y357F,Y359F}$ resulted in a significant increase of the amplitude of whole-cell

currents measured to 120.4 ± 8 % (n = 7) at t = 6 min as compared to control (92.3 ± 5.8 %, n = 7, at t = 6 min, Fig 7.9B). This increase, however, was significantly smaller than the increase observed after application of α -CaMK-II to $\alpha 1\beta 3\gamma 2S$ expressing granule cells (167.8 \pm 13.7 %, n = 4, at t = 6 min, Fig 7.9B). This indicated that, as in NG108-15 cells (See Fig 7.3), a proportion of the effect of α -CaMK-II on $\alpha 1\beta 3\gamma 2S$ receptors was mediated through tyrosine kinase phosphorylation of the $\gamma 2S$ subunit. In addition, similar to that observed in NG108-15 cells (See Fig 7.3C), this proportion of the effect was transient as there was no significant difference between $\alpha 1\beta 3\gamma 2S$ and $\alpha 1\beta 3\gamma 2S^{Y357F,Y359F}$ in the presence of α -CaMK-II after 10 min. The overall similarity between results obtained in NG108-15 cells and cerebellar granule cells confirms that NG108-15 cells are an appropriate model for the study of CaMK-II phosphorylation of GABA_A receptors. The mechanisms and signalling pathways involved in CaMK-II modulation of GABAA receptors in NG108-15 cells must, therefore, be comparable to that observed in cerebellar granule cells (maintained in depolarising conditions).



Fig 7.8: CaMK-II dependent modulation of whole-cell currents recorded from transfected cerebellar granule cells.

Peak amplitude of GABA mediated currents recorded following the application of 10 μ M GABA to cerebellar granule cells transfected with $\alpha 1\beta 2$, $\alpha 1\beta 3$, $\alpha 1\beta 3^{S408A,S409A}$ and $\alpha 1\beta 3^{S383A}$ GABA_A receptor subunits. GABA was re-applied at 2 min intervals for up to 16 min. All currents were normalised to the peak amplitude of the response measured during the first 3-4 min (= 100 %) after achieving the whole-cell configuration (t = 0 min). Control recordings were made with normal patch pipette solution supplemented with the pre-activation buffer without α -CaMK-II.

(A) GABA currents recorded from cerebellar granule cells transfected with GFP only in the absence of α -CaMK-II (Control, n = 7) and transfected with $\alpha 1\beta 3$ (n = 7) and $\alpha 1\beta 2$ (n = 7) GABA_A receptor subunits in the presence of 60 nM α -CaMK-II.

(B) GABA currents recorded from cerebellar granule cells transfected with GFP only in the absence of α -CaMK-II (Control, n = 7, taken from A) and transfected with $\alpha l\beta 3^{S408A,S409A}$ (n = 4), $\alpha l\beta 3^{S383A}$ (n = 4) and $\alpha l\beta 3$ (n = 7, taken from A) GABA_A receptor subunits in the presence of 60 nM α -CaMK-II.

All data points mean \pm s.e.

Significance is indicated by the symbol *, comparing different subunit combinations and mutants in the presence of CaMK-II.

P<0.05%, t-test, ANOVA.




Fig 7.9: CaMK-II dependent modulation of whole-cell currents recorded from cerebellar granule cells transfected with recombinant $\alpha\beta\gamma$ receptors.

Peak amplitude of GABA mediated currents recorded following the application of 10 μ M GABA to cerebellar granule cells transfected with $\alpha 1\beta 2\gamma 2S$, $\alpha 1\beta 3\gamma 2S$ and $\alpha 1\beta 3\gamma 2s^{Y365F,Y367F}$ GABA_A receptor subunits. GABA was re-applied at 2 min intervals for up to 16 min. All currents were normalised to the peak amplitude of the response measured during the first 3-4 min (= 100 %) after achieving the whole-cell configuration (t = 0 min). Control recordings were made with normal patch pipette solution supplemented with the pre-activation buffer without α -CaMK-II.

(A) GABA currents recorded from cerebellar granule cells transfected with GFP only in the absence of α -CaMK-II (Control, n = 7) and transfected with $\alpha 1\beta 2\gamma 2S$ (n = 7), $\alpha 1\beta 3\gamma 2S$ (n = 6) and $\alpha 1\beta 3$ (n = 7, taken from A) GABA_A receptor subunits in the presence of 60 nM α -CaMK-II.

(B) GABA currents recorded from cerebellar granule cells transfected with GFP only in the absence of α -CaMK-II (Control, n = 7) and transfected with $\alpha 1\beta 3\gamma 2S$ (n = 8, taken from A) and $\alpha 1\beta 3\gamma 2S^{Y357F,Y359F}$ (n =7) GABA_A receptor subunits in the presence of 60 nM α -CaMK-II.

All data points mean \pm s.e.

Significance is indicated by the symbol *, comparing $\alpha 1\beta 3\gamma 2S$ to $\alpha 1\beta 2\gamma 2S$ and $\alpha 1\beta 3\gamma 2S^{Y357F,Y359F}$ in the presence of α -CaMK-II. The symbol * indicates $\alpha 1\beta 3\gamma 2S$ compared to $\alpha 1\beta 3$ and the symbol *, indicates $\alpha 1\beta 2\gamma 2S$ and $\alpha 1\beta 3\gamma 2S^{Y357F,Y359F} \alpha$ -CaMK-II compared to control.

P < 0.05, t-test, ANOVA.

7.3 DISCUSSION

Using site-directed mutagenesis and the expression of recombinant GABA_A receptors in NG108-15 cells and cerebellar granule cells, it was possible to identify the sites of phosphorylation that are important for α -CaMK-II dependent modulation of $\alpha 1\beta 1\gamma 2S$ and $\alpha 1\beta 3\gamma 2S$ GABA_A receptors. It appeared that phosphorylation of Ser³⁸³ on the $\beta 3$ subunit mediated part of this response to α -CaMK-II through direct phosphorylation of the GABA_A receptor. In addition, phosphorylation of the $\gamma 2S$ subunit contributed to the effect, but in this case via CaMK-II dependent activation of tyrosine kinase activity and phosphorylation of Tyr^{357;359} in the $\gamma 2S$ subunit.

To date, the Ser³⁸³ site on the β 3 subunit is the only site reported to be phosphorylated by α -CaMK-II and no other Ser/Thr kinases when the TM3-4 intracellular loop is expressed as a fusion protein (McDonald and Moss, 1997). The same comparable site, Ser³⁸⁴, can be phosphorylated by CaMK-II on the β 1 subunit, when expressed as a fusion protein (McDonald and Moss, 1994), but phosphorylation of this site does not seem to be involved in mediating the functional effect of α -CaMK-II on $\alpha 1\beta 1\gamma 2S$ receptors, as $\alpha 1\beta 1$ heteromers do not respond to α -CaMK-II. It appears that the tyrosine kinase-dependent phosphorylation of the $\gamma 2S$ subunit must also be dependent on the β subunit present in the receptor complex, as $\alpha 1\beta 2\gamma 2S$ receptors are not modulated by α -CaMK-II at all in NG108-15 cells, or in transfected cerebellar granule cells.

This may be an emerging theme in phosphorylation dependent modulation of GABA_A receptors as differences between $\beta 1/3$ and $\beta 2$ subunits have previously been demonstrated. PKA appears able to phosphorylate $\beta 1/3$ in a recombinant system whereas $\beta 2$ is not (McDonald *et al.*, 1998), even though in a neuronal environment PKA can modulate mIPSCs thought to be mediated by $\alpha 1\beta 2\gamma 2S$ receptors (Kano and

Konnerth, 1992; Nusser *et al.*, 1999). This is comparable to the fact that recombinant $\alpha 1\beta 2\gamma 2S$ receptors cannot be modulated by CaMK-II in NG108-15 cells or cerebellar granule cells, but CaMK-II dependent modulation of GABA_A receptors involved in rebound potentiation (Kano *et al.*, 1996; Kawaguchi and Hirano, 2002) may be mediated by $\alpha 1\beta 2\gamma 2$ receptors (Persohn *et al.*, 1992). Despite the high level of homology in the β subunit family, it has been demonstrated that $\beta 1/3$ and $\beta 2$ subunits require different assembly signals on the α / γ subunits for correct assembly and trafficking to the cell surface (Bollan *et al.*, 2003). In addition, different β subunits appear to be targeted to different specific regions of a cell (Connolly *et al.*, 1996b). This implies that GABA_A receptors containing different β subunits are assembled, trafficked and targeted differentially to different synaptic sites. It is not surprising therefore that they might also be modulated by phosphorylation in different ways.

It is also becoming increasingly clear that for many multi-functional kinases with multiple substrates to function efficiently, correct targeting within the cell or localization to points of action is necessary (Pawson and Scott, 1997; Brandon *et al.*, 2002a). The action of kinase anchoring proteins that can bind both kinase and receptor substrate may provide explanations for some of the conflicting reports regarding the effects of phosphorylation on GABA_A receptors. For example, the kinase anchoring protein RACK-1, which potentiates PKC dependent phosphorylation of the β subunit of the GABA_A receptor (Brandon *et al.*, 2002b), has been shown to associate with β 1/3 subunits (Brandon *et al.*, 1999; Brandon *et al.*, 2002b). Interestingly, RACK-1 is able to bind tyrosine kinases as well, in particular the cytosolic Src family (Yaka *et al.*, 2002). The action of PKA has also been reported to be dependent on AKAP 79/150 which only binds β 1 and β 3 and not β 2

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subunits (Brandon *et al.*, 2003) giving a possible mechanism for the observed lack of functional modulation of $\beta 2$ subunit-containing GABA_A receptors in HEK293 cells (McDonald *et al.*, 1998). It is possible that another, as yet unidentified, anchoring protein mediates PKA dependent effects on $\alpha 1\beta 2\gamma 2$ receptors observed in some neuronal cell-types (Kano and Konnerth, 1992; Nusser *et al.*, 1999). The differential effect of α -CaMK-II on $\alpha 1\beta 1/3\gamma 2S$ and $\alpha 1\beta 2\gamma 2S$ receptors may be mediated by an anchoring protein with differential binding affinities for the different subunits. This model of phosphorylation may also explain why α -CaMK-II modulation was not observed at all in HEK293 cells, if this cell line lacks the relevant anchoring protein(s).

The fact that $\gamma 2S^{Y357F,Y359F}$ mutant subunit-containing GABA_A receptors had their response to α -CaMK-II reduced or abolished, suggests that α -CaMK-II is capable of directly or indirectly activating a tyrosine kinase which can then phosphorylate the $\gamma 2S$ subunit. As there did not appear to be any basal tyrosine phosphorylation of GABA_A receptors in this cell line, as has been reported elsewhere (Moss *et al.*, 1995; Brandon *et al.*, 2001), phosphorylation of these residues would be expected to result in an increase in the whole-cell GABA current amplitude. Moss *et al.* (1995) determined that phosphorylation of these residues led to an increase in whole-cell GABA currents primarily due to an increase in the single GABA channel mean open time and probability of opening. They also suggested that in neurones, these residues are basally phosphorylated. Using phospho-specific antibodies directed against Y365 and Y367 on the $\gamma 2L$ subunit, Brandon *et al.* (2001) demonstrated that phosphorylation of phosphorylation of these sites through tyrosine kinase / phosphatase activities in cortical neurones in culture.

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In cells where tyrosine phosphorylation has already occurred under basal conditions, this may be expected to reduce the effect of CaMK-II modulation. Interestingly, in cortical neuronal cultures, addition of α -CaMK-II had a less profound effect on GABA currents than that observed in cerebellar granule cell cultures (See Chapter 4, Fig 4.1). Cortical neuronal cultures of this type have been reported to have a very high level of tyrosine phosphatase activity and subsequently low levels of basal phosphorylation of Y365 and Y367 even after stimulation to activate tyrosine kinases (Brandon *et al.*, 2001). This may also explain why the response of GABA_A receptors to α -CaMK-II in cortical neurones (Chapter 4, Fig 4.1) appeared comparably lower. The importance of these sites of tyrosine kinase phosphorylation on the γ 2 subunit has been recently demonstrated by the formation of a transgenic mouse where these tyrosine residues have been mutated to prevent tyrosine kinase phosphorylation of the γ 2 subunit. This mutation is fatal with -/- mice not surviving beyond birth (V. Tretter, C. Houston, T. Smart and S. Moss, unpublished observations).

It is also possible that mutation of these tyrosine residues to phenylalanines altered the normal behaviour of the receptor such that it appeared to reduce the response to α -CaMK-II, but in fact, was simply an indirect effect, altering the receptors basal phosphorylation state. In addition it is always possible that in different neuronal cell types phosphorylation of other residues that have not yet been identified might also be important. However, these hypotheses are not the simplest ones to adopt. Within the γ 2S subunit the only residue that was not tested was Ser³²⁷ the residue thought to be partially responsible for a PKC dependent depression (Kellenberger *et al.*, 1992; Krishek *et al.*, 1994) and a potentiation of $\alpha l\beta l/2\gamma 2S$ containing receptors (Lin *et al.*, 1996). In addition a reduction in the level of

phosphorylation of this residue was reported after induction of a calcineurindependent form of LTD of inhibitory synapses (Wang et al., 2003). Given that CaMK-II appeared unable to phosphorylate this residue in vitro (fusion proteins) it seems α -CaMK-II action on the γ 2S subunit is unlikely to be through direct phosphorylation of this residue. Downstream activation by CaMK-II of PKA and/or PKC is also possible but Ser^{408,409} mutants in β 3 subunits seemed to have no effect, arguing against this. The residues Ser³⁴⁸ and Thr³⁵⁰, identified in fusion proteins, were thought to be main the main sites of phosphorylation by CaMK-II in $\gamma 2S$ subunits (McDonald and Moss, 1994). Mutation of these residues to alanine abolished CaMK-II dependent phosphorylation in vitro. However, mutation of these residues had no functional effect on α -CaMK-II modulation of $\alpha 1\beta 3\gamma 2S$ GABA_A receptors in NG108-15 cells. The evidence therefore suggests that CaMK-II modulation of GABA_A receptors is dependent on both the β and γ 2 subunits, but not through direct phosphorylation by CaMK-II of the γ 2S subunit. The depression of CaMK-II mediated effects by tyrosine kinase inhibitors and mutation of sites of tyrosine kinase phosphorylation on the γ 2S subunit both suggest that CaMK-II may activate a tyrosine kinase which in turn phosphorylates the $\gamma 2S$ subunit and upregulates receptor function.

An important question relating to the above hypothesis is can CaMK-II activate a tyrosine kinase? There is some evidence that Ca^{2+} sensitive tyrosine kinases exist (Rusanescu *et al.*, 1995; Boxall and Lancaster, 1998). In addition, it has been determined that PKC can influence the activity of the non-receptor tyrosine kinase Src, suggesting that an interaction between Ser/Thr kinases and the Src family of tyrosine kinases is possible (Lu *et al.*, 1999; Ali and Salter, 2001). It is generally thought that activation of Src through PKC is mediated by activation of PYK2 (Dikic

et al., 1996), a member of the focal adhesion kinase family highly expressed in the nervous system (Sasaki et al., 1995; Lev et al., 1995), which responds to calcium (Lev et al., 1995). Activation of PYK2 leads to autophosphorylation which creates an SH2 ligand that can bind the SH2 domain of Src, thereby activating it (Dikic et al., 1996; Ali and Salter, 2001).

Activation of PYK2, and subsequently, Src has been demonstrated in some cases to be $Ca^{2+}/Calmodulin$ dependent, but not PKC dependent (Della Rocca *et al.*, 1997; Heidinger *et al.*, 2002). Indeed, there is evidence for CaMK-II dependent activation of Src in glomerular mesangial cells (Wang *et al.*, 2003b) and HEK293 cells (Lo and Wong, 2004). There is also evidence for CaMK-II dependent activation of PYK2 and Src in vascular smooth muscle (Ginnan and Singer, 2002; Ginnan *et al.*, 2004), PC12 cells (Zwick *et al.*, 1999) and in hippocampal neurones following ischemia (Guo *et al.*, 2004). It is unknown if NG108-15 cells express PYK2, but given that this protein is abundant in the CNS (Sasaki *et al.*, 1995; Lev *et al.*, 1995) and the neuronal origin of the cell line, it seems a plausible possibility especially considering how NG108-15 cells are also used as a model of neuronal differentiation and neuritogenesis.

Src activation has been implicated in the regulation of excitatory synaptic plasticity through the action of Src on the NMDA receptor (Lu *et al.*, 1999; Huang and Hsu, 1999; Salter and Kalia, 2004). PYK2 induced Src activation is thought to be necessary for certain forms of excitatory LTP (Huang *et al.*, 2001). At the NMDA receptor, Src kinase acts as a point of convergence for a number of different signalling pathways (Salter and Kalia, 2004).

For Src and also other tyrosine kinases, binding to the NMDA receptor is thought to occur via association with PSD-95, or possibly another adaptor protein rather than directly interacting with the receptor itself (Tezuka *et al.*, 1999; Gingrich *et al.*, 2004). For the present study, it may be that binding of the PYK2:Src complex, either directly or through an intermediate protein to the GABA_A receptor, may be dependent on the subunits present. Binding may be $\beta 1/3$ subunit dependent and unable to occur if $\beta 2$ is present within the receptor complex. This would explain the lack of a tyrosine kinase dependent effect seen on $\alpha 1\beta 2\gamma 2S$ receptors.

The actions of Src on GABA_A receptors in neurones suggest that Src can increase whole-cell current amplitudes (Moss *et al.*, 1995; Wan *et al.*, 1997). There is also evidence for a Src mediated increase in the amplitude of IPSCs without a change in their decays (Boxall, 2000). This was considered to be in accord with the findings of Moss *et al.* (1995) and the observed increase in mean open time and probability of channel opening. Activation of PYK2 and Src in prefrontal cortical neurones was also shown to increase the amplitude of IPSCs without changing their decays (Ma *et al.*, 2003). The known association between Ca²⁺ and PYK2 (Lev *et al.*, 1995) and the activation of Src (Rusanescu *et al.*, 1995; Dikic *et al.*, 1996), together with the evidence for CaMK-II dependent activation of Src (Wang *et al.*, 2003b; Lo and Wong, 2004; Ginnan *et al.*, 2004) possibly through PYK2 (Zwick *et al.*, 1999; Ginnan and Singer, 2002), links several signal transduction pathways. It seems that a CaMK-II dependent activation of PYK2 and subsequent activation of Src tyrosine kinase is the most plausible, potential signalling pathway for CaMK-II dependent, tyrosine kinase dependent, phosphorylation of the GABA_A receptor.

The application of α -CaMK-II to cerebellar granule cells (maintained in physiological conditions) led to an increase in IPSC amplitude and a change in decay. It appeared that CaMK-II did not have a uniform effect across all populations of synaptic events. It is possible that the analysis of the effect of CaMK-II on sIPSCs

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is further complicated by the fact that there may be two different effects being mediated by phosphorylation of the β 3 subunit and the γ 2S subunit.

In conclusion, the evidence suggests that α -CaMK-II mediates its effects on $\alpha 1\beta 1\gamma 2S$ and $\alpha 1\beta 3\gamma 2S$ receptors in NG108-15 cells and cerebellar granule cells through a direct phosphorylation of the $\beta 3$ subunit at Ser³⁸³ and through CaMK-II activation of a tyrosine kinase that phosphorylates $\gamma 2S$ at Tyr^{357,359}. This tyrosine kinase dependent phosphorylation is partially dependent on the identity of the β subunit, as it is only observed if the $\beta 1/3$ subunit is present and not when $\beta 2$ is present. It is assumed that the modulation of GABA_A receptors expressed in NG108-15 cells is comparable to that of native receptors in neurones; however, there remain unanswered questions about the role of the $\beta 2$ subunit-containing GABA_A receptors.

7.4 CONCLUSIONS

- The modulation of $\alpha 1\beta 3$ heteromers expressed in NG108-15 cells and cerebellar granule cells by α -CaMK-II is mediated through phosphorylation of the Ser³⁸³ residue.
- Modulation of $\alpha 1\beta 3\gamma 2S$ receptors expressed in NG108-15 cells and cerebellar granule cells is mediated through phosphorylation of $\beta 3$ Ser³⁸³ and through tyrosine kinase mediated phosphorylation of Tyr^{357,359} on the $\gamma 2S$ subunit.
- Modulation of $\alpha 1\beta 1\gamma 2S$ receptors expressed in NG108-15 cells is mediated through tyrosine kinase phosphorylation of Tyr^{357,359}.
- CaMK-II modulation of the GABA_A receptor can occur through direct phosphorylation of the receptor and through activation of a second downstream kinase signalling pathway.

<u>Chapter Eight:</u> <u>CaMK-II modulation of sIPSCs in cerebellar granule cells</u> <u>from *β*2 knockout mice.</u>

8.1 INTRODUCTION

Analysis of the effects of α -CaMK-II on recombinant GABA_A receptors expressed in NG108-15 cells and cerebellar granule cells suggested that there was a differential effect of CaMK-II on GABA_A receptors depending on the combination of β and γ subunits assembled within the receptor. The evidence suggested that β 2 subunit-containing receptors could not be functionally modulated by α -CaMK-II in a recombinant system or in neurones expressing recombinant receptors but that β 1/3 subunit-containing recombinant receptors can be up-regulated by a direct phosphorylation of the β 3 subunit and a downstream activation of a tyrosine kinase and subsequent phosphorylation of the γ 2S subunit. However, a lack of PKA dependent modulation of recombinant receptors containing the β 2 subunit has been observed in HEK293 cells (McDonald *et al.*, 1998), but PKA activity in neurones can alter IPSCs within cells that express predominantly α 1 β 2 γ 2 receptors (Kano and Konnerth, 1992; Nusser *et al.*, 1999). A lack of functional modulation of β 2 subunitcontaining receptors in a recombinant system is therefore not necessarily an indication that modulation cannot occur for native GABA_A receptors at the synapse.

In order to determine more clearly how CaMK-II might modulate β^2 versus β^3 subunit-containing receptors in a neuronal environment it was necessary to make use of a GABA_A receptor β^2 subunit knockout mouse (Sur *et al.*, 2001). In any neuronal preparation it is difficult to study the effects of phosphorylation on a discrete receptor population because of heterogeneity. However, cultures prepared from β^2 knockout mice will always lack this subunit and so after comparison with

wild-type mice it may be possible to determine if there are any differences between modulation of receptors that contain different β subunits.

Knockout mice have been used to study the relative importance and contribution of different GABA_A receptor subunits to synaptic transmission. At present knockout mice lacking the $\alpha 1$, $\alpha 5$, $\alpha 6$, $\beta 2$, $\beta 3$, δ and $\gamma 2$ S/L subunits have been generated (Vicini and Ortinski, 2004; Rudolph and Möhler, 2004). The $\gamma 2$ subunit knockout mouse revealed the importance of this subunit in receptor clustering at the synapse (Essrich *et al.*, 1998) and for survival as homozygotes have a high level of morbidity. By contrast, the $\alpha 1$ knockout mouse has provided insight into the observed reduction in decay time of IPSCs during development observed in some cell types (Vicini *et al.*, 2001; Jüttner *et al.*, 2001; Goldstein *et al.*, 2002; Ortinski *et al.*, 2004). This is thought to be a result of increased $\alpha 1$ expression later in development (Ortinski *et al.*, 2004). The $\alpha 1$ knockout mouse phenotype is relatively normal; except for an occasional handling induced tremor, these mice are indistinguishable from wild-type (Sur *et al.*, 2001) in many behavioural tests.

The β 3 subunit knockout mouse suffers from severe neurological impairments including epilepsy (Homanics *et al.*, 1997) and has many of the phenotypic characteristics of the human neurodevelopmental disorder, Angelman syndrome (DeLorey *et al.*, 1998). Interestingly Angelman syndrome has also been linked to deficits in normal CaMK-II signalling pathways (Weeber *et al.*, 2003).

The removal of the β 3 subunit reduces IPSC duration in β 3 -/- mouse cortical neurones in culture and also decreases expression of the α 2/3 subunits (Ramadan *et al.*, 2003). This supports the idea that there is a preferential association of α 2/3 with β 3 and α 1 with β 2. Disruption of β 3 has been shown in β 3 -/- mice to alter network oscillations in the olfactory bulb (Nusser *et al.*, 2001)

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demonstrating that specific β subunit-containing receptors can play specific physiological roles.

The $\beta 2$ knockout mouse unlike the $\beta 3$ knockout mouse is behaviourally normal with no evidence of spontaneous seizures. The only distinguishable difference from wild-type is a slight increase in locomotor activity (Sur et al., 2001). The difference between the $\beta 2$ and $\beta 3$ knockout mice is striking given the comparable loss in receptor density across the brain seen in the two lines of mice (Homanics et al., 1997; Sur et al., 2001; Vicini and Ortinski, 2004). There was no observed compensatory increase in $\beta 1/3$ subunits observed in the $\beta 2$ knockout mouse but there was a decrease in expression of all six α subunits (Sur *et al.*, 2001). The generation of knockout mice lacking subunits of the GABA_A receptor has started to reveal differences in the function of different GABA_A receptor subtypes. Although compensatory increased expression of subunits can sometimes mask the role(s) of particular receptor subtypes, it appears that $\beta 2$ and $\beta 3$ subunit-containing receptors must play very different roles in development (Homanics et al., 1997; DeLorey et al., 1998; Sur et al., 2001; Sinkkonen et al., 2003) and potentially in pathological states such as epilepsy (Brooks-Kayal et al., 1998; DeLorey et al., 1998; Sur et al., 2001; Jones-Davis and Macdonald, 2003). Knock-in mice which lack the developmental changes that can occur in whole-subunit knockouts have revealed that different effects of anaesthetics are mediated through specific $\beta 2$ or $\beta 3$ subunit-containing GABA_A receptors (Jurd et al., 2003; Reynolds et al., 2003; Rudolph and Möhler, 2004). Using the β 2 knockout mouse, changes in the response of GABA_A receptors to phosphorylation may reveal how distinct receptor populations are modulated by protein kinases.

Application of α -CaMK-II to rat cerebellar granule cells in culture under depolarising conditions (20 mM KCl) led to a significant increase in the whole-cell currents recorded (see Chapter 4). Application of α -CaMK-II to cerebellar granule cells cultured under more physiological conditions (5 mM KCl) led to a significant modulation of the amplitude and decay times of sIPSCs. This modulation of sIPSCs was not consistent across the whole population suggesting that there were possibly different effects of phosphorylation at different synapses (See Chapter 5). This could be related to differential effects of α -CaMK-II on different GABA_A receptor subunit combinations.

Cerebellar granule cells are thought to express a number of different GABA_A receptor subunits. Including $\alpha 1/2/3/6$, $\beta 2/3$, $\gamma 2$ and δ subunits (Persohn *et al.*, 1992; Laurie *et al.*, 1992b; Wisden *et al.*, 1996; Pirker *et al.*, 2000). They are thought to express very low levels of the $\beta 1$ subunit (Persohn *et al.*, 1992; Laurie *et al.*, 1992b). The $\beta 2$ subunit knockout mouse causes a concurrent decrease in the expression of all six α subunits but there has been no reported compensatory increase in the expression of different β subunits (Sur *et al.*, 2001). Therefore cultures obtained from $\beta 2$ knockout mice may be expected to consist predominantly of $\beta 3$ subunitcontaining receptors as compared to wild-type mice which may be expected to express both $\beta 2$ and $\beta 3$.

Mouse cerebellar cultures maintained under depolarising conditions have been reported to express high levels of β 3 subunits and a decreased expression of β 2 subunits as compared to cultures maintained under more physiological conditions (Ives *et al.*, 2002a). The α -CaMK-II dependent effect observed on whole-cell currents under depolarising conditions therefore may reflect CaMK-II action on β 3 subunit-containing receptors. Cells maintained under physiological conditions are

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thought to express both $\beta 2$ and $\beta 3$ subunits (Laurie *et al.*, 1992b; Ives *et al.*, 2002a; Ortinski *et al.*, 2004) so the effect of CaMK-II on sIPSCs could potentially be mediated by receptors containing either or both of these subunits. Analysis of the effects of α -CaMK-II cultures prepared from $\beta 2$ knockout mice maintained under both conditions should allow the determination of the relative contribution of the $\beta 2$ subunit to GABA currents and reveal whether α -CaMK-II can modulate $\beta 2$ subunitcontaining receptors at the inhibitory synapse.

8.2 **RESULTS**

Cerebellar cultures were prepared from β 2 knockout and wild-type mice at P0-2 using the same protocol as that used in the culture of rat cerebellar granule cells (See methods). The β 2 knockout mouse was generated by deletion of exons 6 and 7 of the β 2 subunit gene by homologous recombination (Sur *et al.*, 2001). Heterozygous F3 generation mice were bred to produce β 2 -/- and wild-type strains of mice of the same genetic background (50 % C57BL6, 50 % 129SvEv; Sur *et al.*, 2001). Litters of mice were taken at P0-2 from F5-7 generation wild-type (wt) or β 2 knockout (-/-) mice to produce cerebellar cultures. Tissue (in the form of tail snips) was taken from each mouse pup used for culturing and PCR was used for genotyping (See Chapter 2).

Recombinant α -CaMK-II (Suzuki-Takeuchi *et al.*, 1992) pre-activated by incubation with CaM, CaCl₂, and ATP- γ -S (McGlade-McCulloh *et al.*, 1993) was applied to cerebellar granule cells through inclusion in the patch pipette solution (See Chapter 2 and 3.2.1). The pre-activated α -CaMK-II was diluted into the internal solution (85 nM) and maintained on ice prior to recording. Control recordings were made by supplementing normal patch pipette solution with the pre-activation buffer but without α -CaMK-II.

Whole-cell currents were recorded from cerebellar granule cells maintained in depolarising conditions in response to application of 10 µM GABA every 2 min for 20 min (See Chapter 4). Recordings of sIPSCs were made in the presence of 10 µM CNQX, 20 µM AP-5 to block excitatory postsynaptic currents (EPSCs). All the remaining synaptic currents could be blocked by 20 µM bicuculline indicating they were GABA_A receptor mediated. In a small proportion of cells (15 %) there was evidence of a small tonic current (Fig 8.1) revealed by a shift in baseline current after the application of bicuculline (Brickley et al., 1996; Leao et al., 2000). Application of 500 nM TTX abolished nearly all IPSCs recorded indicating that the majority of IPSCs in this cell-type are action potential driven as has been previously described (Puia et al., 1994; Leao et al., 2000; Farrant and Brickley, 2003). Recordings were made with 5 mM QX-314 in the patch pipette solution to allow recordings of action potential driven sIPSCs whilst suppressing action potentials within the cell being recorded from (See Chapter 5). Continuous 5 min recordings were made starting 5 min after achieving the whole-cell configuration. Cerebellar granule cells were identified by their rounded cell body morphology measuring $< 8 \,\mu m$ in diameter (See Fig 2.1 and Fig 7.6).

Fig 8.1:



Fig 8.1: sIPSCs recorded from mouse cerebellar granule cells in wt cultures.

Representative synaptic currents recorded from mouse cerebellar granule cells grown and maintained in a physiological culture medium (5 mM KCl). Neurones were cultured from P0-2 wt mice.

Recordings were made with 5 mM QX-314 within the patch pipette solution. Bath applied AP-5 (20 μ M) and CNQX (10 μ M) reveals sIPSCs. Additional application of bicuculline (20 μ M) reveals a tonic GABA_A receptor-mediated current.

8.2.1 CaMK-II modulation of whole-cell currents recorded from wt and $\beta 2$ -/mice.

Application of 85 nM α -CaMK-II to cerebellar granule cells grown in depolarising conditions (20 mM KCl) resulted in a significant potentiation of the amplitude of whole-cell currents recorded from wt cultures to 146 ± 4.6 % (n = 8) as compared to control, 95.4 ± 4 % (n = 6, Fig 8.2A) at t = 8 min. Application of 85 nM α -CaMK-II was also significantly potentiated in β 2 -/- cultures to the same extent, where the amplitude was potentiated to 142.8 ± 9 % (n = 10) as compared to control, 100.6 ± 5.7 % (n = 9, Fig 8.2B) at t = 8 min. This indicated that either the β 2 subunit-containing receptors are not functionally modulated by α -CaMK-II in cerebellar granule cells under these culture conditions or that there was a low level of β 2 subunits present even in wild-type cultures. There was no significant difference in the current density of GABA currents recorded in response to 10 μ M GABA in either culture (wt, n = 8, β 2 -/-, n = 10, Fig 8.2C) suggesting that there was a comparable number of functional GABA_A receptors in wt and β 2 -/- cultures.

8.2.2 CaMK-II modulation of sIPSCs.

Application of 85 nM α -CaMK-II to wt cerebellar granule cells grown in physiological (5 mM KCl) conditions resulted in a significant increase in the mean peak amplitude of sIPSCs from 44.9 ± 7.6 pA (n = 9) to 95.4 ± 22.6 pA (n = 12, Fig 8.3A). Application of 85 nM α -CaMK-II to β 2 -/- cerebellar granule cells resulted in no significant change in mean peak amplitude (Control, 47.1 ± 8 pA, n = 8, + CaMK-II, 46.7 ± 5.6 pA, n= 9, Fig 8.3B and Table 8.1). There was also no significant difference between the mean peak amplitude of wt and β 2 -/- mice in control conditions indicating that there was relatively little β 2 subunit expression even under physiological culture conditions (5 mM KCl) or that there was a compensatory increase in the expression of $\beta 1/3$ subunit-containing receptors.

Application of α -CaMK-II significantly increased the half-width (time taken to decay to 50 % amplitude) of sIPSCs recorded from wt cultures from 9.8 ± 1.4 ms (n = 9) to 17.7 ± 2.9 ms (n = 12, Fig 8.3B). Application of α -CaMK-II to β 2 -/cultures also resulted in a significant increase in the half-width of sIPSCs from 10.4 ± 1.3 ms (n = 8) to 17.8 ± 3 ms (n = 9, Fig 8.3B and Table 8.1). The half-width was not significantly different from wt and β 2 -/- cultures in control conditions. There was also no significant difference between half-widths after application of α -CaMK-II in either culture, indicating that removal of the β 2 subunit had little effect on the lengthening of decay by α -CaMK-II. Unlike α 1 knockout mice (Vicini *et al.*, 2001; Goldstein *et al.*, 2002; Ortinski *et al.*, 2004) removal of β 2 does not significantly alter the mean decay time of IPSCs in control conditions.

Application of α -CaMK-II to wt cultures had no significant effect on sIPSC frequency (Control, 3.5 ± 0.8 Hz, n = 9, + CaMK-II, 2.4 ± 0.8 Hz, n = 12). There was no significant difference between wt and $\beta 2$ -/- cultures in control conditions and there was no significant effect on $\beta 2$ -/- cultures after application of α -CaMK-II (Control, 2.6 ± 1 Hz, n = 8, + CaMK-II -/-, 2.5 ± 0.7 Hz, n = 9, Fig 8.3C).



Fig 8.2: Modulation of whole-cell currents by α -CaMK-II recorded from cerebellar granule cells in culture (20 mM KCl) from wild-type (wt) and β 2 knockout mice (-/-).

Peak amplitudes of GABA-mediated currents recorded after application of 10 μ M GABA to mouse cerebellar granule neurones in culture (20 mM KCl) prepared from wild-type (wt) or β 2 knockout mice (-/-). GABA was re-applied at 2 min intervals for up to 20 min. All currents were normalised to the peak response of the GABA current measured during the first 3-4 min (= 100 %) after achieving the whole-cell configuration (t = 0 min). Control recordings were made with normal patch pipette solution supplemented with the pre-activation buffer without α -CaMK-II.

(A) GABA currents recorded from cerebellar granule cells in culture prepared from wt cultures in the absence (Control, n = 8) and presence of 85 nM pre-activated α -CaMK-II (n=6) applied through inclusion in the patch pipette solution.

(B) GABA currents recorded from cerebellar granule cells in culture prepared from $\beta 2$ -/- cultures in the absence (Control, n = 10) and presence of 85 nM α -CaMK-II (n = 9).

(C) A bar chart showing 10 μ M GABA current density (pA/pF) recorded from wt (n = 8) and $\beta 2$ -/- (n = 10) mouse cultures at t = 0 min in control conditions.

All points = mean \pm s.e.

Significance is indicated by the symbol *, CaMK-II compared to control. P < 0.05, t-test.

Fig 8.3:



Fig 8.3: Mean amplitude, half-width and frequency of sIPSCs recorded from cerebellar granule cells in culture (5 mM KCl) from wt and $\beta 2$ -/- knockout mice.

Continuous recordings were made from cerebellar granule cells in culture (5 mM KCl) from t = 5-10 min after achieving the whole-cell configuration in the presence of 10 μ M CNQX, 20 μ M AP-5 and 5 mM QX-314. Control recordings were made with normal patch pipette solution supplemented with the pre-activation buffer without α -CaMK-II.

(A) Bar chart showing the mean peak amplitude of sIPSCs in the absence and presence of 85 nM α -CaMK-II recorded from cerebellar granule cells in culture prepared from wt (Control, n = 9, CaMK-II, n = 12) and $\beta 2$ -/- (Control, n = 8, CaMK-II, n = 9) mouse cultures.

(B) Bar chart showing the mean half-width (τ_{50}) of sIPSCs in the absence and presence of 85 nM α -CaMK-II recorded from cerebellar granule cells in culture prepared from wt (Control, n = 9, CaMK-II, n = 12) and $\beta 2$ -/- (Control, n = 8, CaMK-II, n = 9) mouse cultures.

(C) Bar chart showing the mean frequency of sIPSCs in the absence and presence of 85 nM α -CaMK-II recorded from cerebellar granule cells in culture prepared from wt (Control, n = 9, CaMK-II, n = 12) and $\beta 2$ -/- (Control, n = 8, CaMK-II, n = 9) mouse cultures.

Mean \pm s.e. Significance is indicated by the symbol *, CaMK-II compared to control.

P < 0.05, t-test.

8.2.3 CaMK-II modulation of decay times.

To look at decay times more carefully groups of 50-100 events were selected from each cell that had no deflections in either the rising or decay phases, and which decayed back to the baseline. These average traces were best fitted by a biexponential decay (See Chapter 2, 2.7.2). The mean $\tau 1$ (fast component of decay) of sIPSCs was significantly increased to 29.5 ± 5.3 ms (n = 8) as compared to control, 16.1 ± 2 ms (n = 8) on application of α -CaMK-II to wt cultures. Application of α -CaMK-II to $\beta 2$ -/- cultures also resulted in a significant increase in $\tau 1$ to 22.6 ± 2.2 ms (n = 7) as compared to control, 11.7 ± 1.3 ms (n = 7, Fig 8.4A,E and Table 8.1). There was no significant difference between $\tau 1$ in control conditions in wt or $\beta 2$ -/cultures, or after the application of α -CaMK-II (Fig 8.4B).

Analysis of $\tau 2$ values for wt cultures also showed a significant increase on application of α -CaMK-II to 109 ± 15.9 ms (n = 8) as compared to control, 55.2 ± 7 ms (n = 8). Application of α -CaMK-II to $\beta 2$ -/- cultures also resulted in a significant increase in $\tau 2$ to 78.8 ± 5.1 ms (n = 7) as compared to control 55.2 ± 4.3 ms (n = 7, Fig 8.4B and Table 8.1). There was no significant difference between $\tau 2$ in control conditions in wt or $\beta 2$ -/- cultures or on application of α -CaMK-II between wt and $\beta 2$ -/- cultures (Fig 8.4B).

Phosphorylation has been reported to alter the relative area or contribution of fast and slow decay time constants. However, the proportional area of $\tau 1$ in wt cultures was unaffected by the application of α -CaMK-II (69.5 ± 1.8 %, n =8, as compared to control; 58.7 ± 5.1 %, n = 8). Application of α -CaMK-II to $\beta 2$ -/- cultures also did not affect the relative areas ($\tau 1$ area control, 45.5 ± 7.3 %, n = 7; CaMK-II, 59.5 ± 3.1 %, n = 7, Fig 8.4C,E). The data obtained from rat cerebellar cultures also showed no observable change in the proportional area of $\tau 1$ on

application of α -CaMK-II (Fig 5.3D). In addition there was no significant change between the proportional area of $\tau 1$ from wt cultures in control conditions (58.7 ± 5.1 %, n = 8) and that for $\beta 2$ -/- cultures (45.5 ± 7.3 %, n = 7, Fig 8.4C and Table 8.1) in control conditions. However, the relative contribution of $\tau 2$ on application of α -CaMK-II to $\beta 2$ -/- cultures was significantly greater than when α -CaMK-II was applied to wt cultures (Fig 8.4C).

Changes in IPSC decay in control conditions can also result from dendritic filtering. In order to avoid problems associated with dendritic filtering when analysing decay constants only events with fast rise-times (<3 ms) and clear peak amplitudes were used in the analyses. As cerebellar granule cells are electrotonically compact the effect of filtering should have been relatively small (Gabbiani et al., 1994; Rossi and Hamann, 1998). Synaptic events for analysis were only chosen if there were no deflections in the rising or decaying phase from cells and there was no correlation between rise-times and amplitudes (See Chapter 5, Fig 5.4). The mean rise-times calculated from the grouped average traces used to analyse kinetics showed no significant differences between application of α -CaMK-II (0.76 ± 0.06 ms, n = 8) and control (0.73 \pm 0.08 ms, n = 8) in wt cultures. In β 2 -/- cultures there was no change in the rise-times on application of α -CaMK-II (0.75 ± 0.06 ms, n = 7) as compared to control (0.59 \pm 0.06 ms, n = 7, Fig 8.4D,E) and there was no significant difference between rise-times after application of α -CaMK-II when comparing wt and $\beta 2$ -/- cultures.



Fig 8.4: Modulation of sIPSC decay times by α -CaMK-II in wt and β 2 -/- mouse cultures.

Continuous recordings were made from t = 5-10 min after achieving the whole-cell configuration in the presence of 10 μ M CNQX, 20 μ M AP-5 and 5 mM QX-314. 50-100 IPSCs were grouped together from each cell that had no deflections in either their rising or decaying phases. These average traces were then used to calculate 10-90 % rise-times and decay time constants for each cell. The mean of several cells are displayed as a bar chart. Control recordings were made with normal patch pipette solution supplemented with the pre-activation buffer without α -CaMK-II.

(A) A bar chart showing the mean $\tau 1$ in the absence (Control, n = 8) and the presence (n = 8) of 85 nM α -CaMK-II of sIPSCs recorded from wt and $\beta 2$ -/- (Control, n = 7,CaMK, n = 7) mouse cultures.

(B) A bar chart showing the mean $\tau 2$ in the absence (Control, n = 8) and the presence (n = 8) of 85 nM α -CaMK-II of sIPSCs recorded from wt and $\beta 2$ -/- (Control, n = 7, CaMK-II, n = 7) mouse cultures.

(C) A bar chart showing the relative contributions of the area of $\tau 1$ and $\tau 2$ to the total area of decay in the absence (Control, n = 8) and presence (n = 8) of 85 nM α -CaMK-II of sIPSCs recorded from wt and $\beta 2$ -/- (Control, n = 7, CaMK-II, n = 7) mouse cultures.

(D) A bar chart showing the mean 10-90 % rise-time of sIPSCs in the absence (Control, n = 8) and the presence (n = 8) of 85 nM α -CaMK-II recorded from wt and $\beta 2$ -/- (Control, n = 7, CaMK-II, n = 7) mouse cultures.

(E) Representative average traces of 50-100 IPSCs from a single cell in the absence and the presence of 85 nM α -CaMK-II recorded from wt and β 2 -/- mouse cultures re-scaled to the same amplitude.

Mean \pm s.e. Significance is indicated by the symbol *, CaMK-II compared to Control and the symbol *, wt compared to $\beta 2$ -/-. P < 0.05, t-test.

Table 8.1:

	Amplitude (pA)	H-W (ms)	τ1 (ms)	τ2 (ms)	A1 (%)
Wt Control	44.9 ± 7.6	9.8 ± 1.4	16.1 ± 2	55.2 ± 7	58.7 ± 5.1
Wt CaMK-II	9 5.4 ± 22.6	* 17.7 ± 2.9	* 29.5 ± 5.3	* 109 ± 15.9	69.5 ± 1.8
β2 -/- Control	47.1 ± 8	10.4 ± 1.3	11.7 ± 1.3	55.2 ± 4.3	45.5 ± 7.3
β2 -/- CaMK-II	46.7 ± 5.6	17.8 ± 3	* 22.6 ± 2.2	* 78.8 ± 5.1	* 59.5 ± 3.1

Table 8.1: A table summarising sIPSC characteristics.

The mean values of peak amplitude, half-width (H-W), time constants $\tau 1$ and $\tau 2$ and the proportional area of $\tau 1$, in control conditions and in the presence of 85 nM α -CaMK-II from wt and $\beta 2$ -/- cultures. Significance is indicated by the symbol *, CaMK-II compared to control and the symbol *, Wt compared to $\beta 2$ -/- in the presence of CaMK-II. P < 0.05, t-test.

8.2.4 CaMK-II amplitude and decay time changes across the whole population of IPSCs.

As previously demonstrated, in rat cerebellar cultures the population of sIPSCs did not respond uniformly to α -CaMK-II (See Chapter 5) it was therefore possible that different changes in different sub-populations of events between wt and $\beta 2$ -/- in control conditions or on application of α -CaMK-II, might mask or alter the apparent effect of α -CaMK-II. Therefore, a more beneficial approach would be to look at the whole population of IPSC events.

An IPSC half-width histogram was constructed for wt and β 2 -/- cultures for control conditions and following the response to application of 85 nM α -CaMK-II. In wt cultures there did appear to be a slight shift in half-width values from under 15 ms to over 15 ms (Control, n = 8, 6600 events, CaMK-II, n = 11, 6300 events, Fig 8.5A). The histogram constructed from $\beta 2$ -/- cultures also showed a shift to higher values of half-width (over 15 ms) but in this case it appeared much more pronounced (Control, n = 9, 5500 events, CaMK-II, n = 10, 5500 events, Fig 8.5B).

The more pronounced effect on addition of α -CaMK-II could indicate that the increase in half-width observed in wt cultures is partly masked by the presence of the β 2 subunits which are not affected by CaMK-II. There is also the indication that the loss of the β 2 subunit also creates a small group of events with large half-widths of over 30 ms which are not present in the wild-type, under control conditions. It should be noted from the previous analysis though, that this slight shift in the population was not enough to alter the mean half-width. This population of IPSCs with increased half-widths in $\beta 2$ -/- cultures could possibly be a result of a loss of $\alpha 1$ (faster decaying) subunit-containing receptors alongside $\beta 2$ in favour of $\alpha 2/3/6$ containing receptors which are slower decaying (Gingrich et al., 1995; Tia et al., 1996; Ortinski et al., 2004). It is important to note that analysis of decay time constants and proportional area's required selection of events with no deflections during the rise or decay phase. Slower decaying events are more likely to be superimposed by other events so these slower events may have been underrepresented in the group averages used to calculate τ values and proportional areas.

Amplitude histograms were constructed of sIPSCs recorded from wt cultures in control conditions and after application of α -CaMK-II. In control conditions (n = 8, 6500 events) there appeared to be two populations of events: one under 100 pA and the other of 100-300 pA. On application of α -CaMK-II (n = 11, 6300 events, Fig 8.6A) both populations shifted slightly to the right with an increase in the number of events at around 100 pA and the appearance of events over 300 pA in amplitude (Fig 8.6A, arrow). The amplitude histogram of sIPSCs constructed from events recorded from $\beta 2$ -/- cultures showed a slightly different pattern. In control conditions (n = 9, 5500 events) the distribution of amplitudes was similar to that seen in wild-type with two populations of under 100 pA and between 100-300 pA. On application of α -CaMK-II (n = 10, 5500 events, Fig 8.6B) there was an increase in the number of events around 100 pA; however, in this case, there was no appearance of larger IPSCs around 400pA. This suggested that a sub-population of events did increase in amplitude but that the removal of the $\beta 2$ subunit prevented the appearance of another sub-population of much larger amplitude IPSCs. This is in contrast to the more pronounced effect of α -CaMK-II on half-widths observed in $\beta 2$ -/- cultures. Despite the increase in the number of IPSCs around 100pA in $\beta 2$ -/- cultures this was not enough to significantly alter the mean peak amplitude. The loss of the larger amplitude events is probably responsible for the lack of apparent effect on the mean peak amplitude after α -CaMK-II application to $\beta 2$ -/- cultures.





Continuous recordings were made from t = 5-10 min after achieving the whole-cell configuration in the presence of 10 μ M CNQX, 20 μ M AP-5 and 5 mM QX-314. All the half-widths of sIPSCs from 10-11 cells were grouped together and frequency histograms were constructed. Control recordings were made with normal patch pipette solution supplemented with the pre-activation buffer without α -CaMK-II.

(A) Half-width distribution histogram of sIPSCs in the absence (Control, n = 8, 6600 events) and the presence (n = 11, 6300 events) of 85 nM α -CaMK-II recorded from wt mouse cultures.

(B) Half-width distribution histogram of sIPSCs in the absence (Control, n = 9, 5500 events) and the presence (n = 10, 5500 events) of 85nM α -CaMK-II recorded from $\beta 2$ -/- mouse cultures.



Fig 8.6: Frequency histograms of sIPSC amplitudes recorded from wt and $\beta 2$ -/mouse cultures.

Continuous recordings were made from t = 5-10 min after achieving the whole-cell configuration in the presence of 10 μ M CNQX, 20 μ M AP-5 and 5 mM QX-314. All the peak amplitudes of sIPSCs from a series of cells were grouped together and frequency histograms were constructed. Control recordings were made with normal patch pipette solution supplemented with the pre-activation buffer without α -CaMK-II.

(A) Amplitude distribution histogram of sIPSCs in the absence (Control, n = 8, 6600 events) and the presence (n = 11, 6300 events) of 85 nM α -CaMK-II recorded from wt mouse cultures.

(B) Amplitude distribution histograms of sIPSCs in the absence (Control, n = 9, 5500 events) and the presence (n = 10, 5500 events) of 85 nM α -CaMK-II recorded from β 2 -/- mouse cultures. The red arrow depicts large amplitude events of over 300 pA.

8.2.5 Scatter-plots of sIPSC half-widths and amplitudes.

In order to try and reconcile these different responses when examining sIPSC half-widths and amplitudes separately, scatter-plots of half-width against amplitude were constructed similar to those constructed for the IPSCs measured in rat cultures (Fig 5.5A). The scatter plot of sIPSCs recorded from wt cultures in control conditions (n = 8, 6600 events, Fig 8.7A) reveals at least two populations of events. One population appears to consist of sIPSCs of under 100 pA with a broad range of decay times. There is another distinct population of sIPSCs of 100-300 pA amplitude (as seen in the amplitude histogram) with a half-width of 10-30 ms. In the scatter-plot of sIPSCs from $\beta 2$ -/- cultures in control conditions (n = 9, 5500 events, Fig 8.7C) this larger amplitude population has dispersed with larger amplitude events occurring with varying half-widths mostly over 30 ms. This could be interpreted as a shift from $\alpha 1\beta 2\gamma 2$ subunit-containing events at these synapses to $\alpha 2/3\beta 3\gamma 2$ subunit-containing events with a slower decay.

On application of 85 nM α -CaMK-II to wt cultures (n = 11, 6300 events, Fig 8.7B) there appears to be a shift of sIPSCs to greater half-widths and amplitudes. The population of sIPSCs of 100-300 pA shifts to even higher amplitudes with little effect on half-width. If these events can indeed be interpreted as β 2 subunit-containing events this implies an α -CaMK-II dependent effect on β 2 subunit-containing receptors that is only revealed in a native environment. The addition of α -CaMK-II to rat cerebellar granule cultures revealed a similar effect on the population of sIPSCs to that observed in wt mouse cultures (See Fig 5.5A).

On addition of α -CaMK-II to $\beta 2$ -/- cultures (n = 10, 5500 events, Fig 8.7D) there is a shift of events, or the appearance of another population of events, at 75-200 pA amplitude and 15-35 ms that was not apparent in control conditions. There is,

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however, no shift in amplitude of the larger amplitude events and there are no sIPSCs of over 300 pA amplitude. Again this could be interpreted as a result of the loss of $\beta 2$ subunit-containing events. It cannot be concluded with certainty that this is a specific $\beta 2$ subunit dependent effect due to the possible changes in the expression levels of other subunits on removal of the $\beta 2$ subunit however, it does appear that in this neuronal culture, synaptic β^2 subunit-containing receptors may be modulated by α -CaMK-II which contrasts with the lack of response to α -CaMK-II of $\alpha 1\beta 2\gamma 2S$ receptors expressed in NG108-15 cells.



Control

Fig 8.7: Scatter plots of sIPSC amplitude against half-width recorded from wt and $\beta 2$ -/- mouse cultures.

Continuous recordings were made from t = 5-10 min after achieving the whole-cell configuration in the presence of 10 μ M CNQX, 20 μ M AP-5 and 5 mM QX-314. All the peak amplitudes and half-widths of sIPSCs from a series of cells were grouped together and scatter-plots of the amplitudes and half-widths for the IPSCs were constructed. Control recordings were made with normal patch pipette solution supplemented with the pre-activation buffer without α -CaMK-II.

(A) A scatter-plot of sIPSC amplitudes and half-widths in the absence (Control, n = 8,6600 events) of 85 nM α -CaMK-II recorded from wt mouse cultures.

(B) A scatter-plot of sIPSC amplitudes and half-widths in the presence (n = 11, 6300 events) of 85 nM α -CaMK-II recorded from wt mouse cultures.

(C) A scatter-plot of sIPSC amplitudes and half-widths in the absence (Control, n = 9,5500 events) of 85 nM α -CaMK-II recorded from $\beta 2$ -/- mouse cultures.

(D) A scatter-plot of sIPSC amplitudes and half-widths in the presence (n = 9, 5500 events) of 85 nM α -CaMK-II recorded from $\beta 2$ -/- mouse cultures.

IPSCs potentially mediated by $\beta 2$ subunit-containing receptors are indicated by the dark blue line.

The green line indicates IPSCs potentially mediated by β 3 subunit-containing receptors.

8.3 **DISCUSSION**

Spontaneous IPSCs recorded from cerebellar granule cells lacking the $\beta 2$ subunit display slightly different properties compared to sIPSCs in wild-type cells. The population of sIPSCs from $\beta 2$ -/- cultures has more events with slower decay times although this is not enough to significantly alter the mean half-width. There is however a significant difference in the relative contribution of $\tau 2$ on application of α -CaMK-II between wt and $\beta 2$ -/- cultures indicating a more pronounced effect on decay times which can be observed in the half-width distribution histogram.

On application of α -CaMK-II to wild-type cultures there appears to be two broad trends. An increase in decay time for one population of IPSCs and an increase in amplitude with little change in decay times for another set of IPSCs. By comparing populations of sIPSCs in control conditions for wild-type and $\beta 2$ -/- cells it can be predicted that the larger amplitude sIPSCs with faster decay times that change in amplitude, but not decay, on α -CaMK-II application, may be $\beta 2$ subunitcontaining GABA_A receptor mediated.

This raises an interesting possibility, that $\beta 2$ subunit-containing GABA_A receptors at the synapse can be modulated by α -CaMK-II in a native neuronal environment. It appeared that expression of recombinant receptors in a neuronal cell line or in native neurones was not sufficient to allow α -CaMK-II modulation of $\alpha 1\beta 2\gamma 2$ receptors to occur. This may be because recombinant receptors expressed in neurones are not targeted to the correct subcellular location. For instance $\alpha\beta$ heteromers would not be expected to be targeted to the synapse without the $\gamma 2$ subunit (Essrich *et al.*, 1998). It is also possible that recombinant receptors do not bind to the relevant kinase anchoring proteins and this disrupts their modulation by phosphorylation. The current study seems to suggest that $\alpha 1\beta 2\gamma 2$ receptors need to

be expressed in their native neuronal environment at the synapse in order for α -CaMK-II modulation of GABA_A receptors to be observed. It should be noted that recombinant $\alpha 1\beta 2\gamma 2S$ receptors were transfected into cerebellar granule cells maintained in depolarising conditions, which display no synaptic activity and are thought to be representative of immature neurones (Ives *et al.*, 2002a). It could therefore be that expression of a mature functional synapse and the associated scaffolding and kinase anchoring proteins is critical for modulation of $\alpha 1\beta 2\gamma 2$ GABA_A receptors by CaMK-II.

The functional effect of α -CaMK-II modulation appeared to be different for $\beta 2$ and $\beta 3$ subunit-containing receptors with the $\beta 3$ subunit-containing receptors changing in decay time and $\beta 2$ subunit-containing receptors increasing in amplitude with no change in decay time. It should be noted that a sub-population of IPSCs in $\beta 2$ -/- cultures also increased in amplitude as can be seen from the amplitude distribution (Fig 8.6) but this shift was not enough to alter the mean IPSC amplitude (Fig 8.3). There was also a slight reduction in the number of large amplitude events on application of α -CaMK-II to $\beta 2$ -/- cultures which may have acted to mask this change in amplitude.

It appeared from the study of recombinant receptors expressed in NG108-15 cells and cerebellar granule cells that in looking at α -CaMK-II dependent modulation it is necessary to consider the whole receptor and all subunits together. For instance it appeared that β and γ subunits interacted in different ways to modulate the response to CaMK-II. In a neuronal situation it appears that the loss of the β 2 subunit may alter α subunit expression. A number of studies have suggested that α 1 may preferentially co-assemble with β 2, over α 2/3 (Benke *et al.*, 1994; Ramadan *et al.*, 2003; Ortinski *et al.*, 2004). The loss of faster decaying large amplitude events
in $\beta 2$ -/- cultures may reflect not only the loss of $\beta 2$ but also a concurrent loss of $\alpha 1$, so it is difficult to say with absolute certainty that the $\beta 2$ subunit is modulated directly by CaMK-II as it is possible $\alpha 1$ may be involved as well. Indeed, CaMK-II has been reported to phosphorylate $\alpha 1$ subunits (Churn *et al.*, 2002) although generally, they are thought to be poor substrates for phosphorylation (Moss *et al.*, 1992; Moss and Smart, 1996). The increase in the number of slower decaying events in control conditions in $\beta 2$ -/- cultures may reflect a shift from $\alpha 1$ to $\alpha 2/3$ after a loss of $\alpha 1$ together with $\beta 2$. In support of this, in $\beta 3$ knockout mice, there is a decrease in $\alpha 2/3$ subunit expression and a reduction in IPSC decay times (Ramadan *et al.*, 2003).

It appears that $\beta 2$ subunit-containing receptors predicted to consist of $\alpha 1\beta 2\gamma 2$ subunits may be preferentially targeted to large synapses. The size of the amplitudes of these IPSCs suggests that they were produced by multi-quantal release of GABA. One other possibility is that at these large amplitude synapses, $\beta 2/3$ subunits are coassembled in the same receptor complex and that $\beta 3$ still mediates the CaMK-II effect. However if direct phosphorylation of $\beta 3$ causes an increase in decay time this should have been observable in GABA_A receptors containing both $\beta 2$ and $\beta 3$ subunits. The lack of effect on decay time of the large amplitude IPSCs suggests this effect was not mediated by $\beta 3$.

It is always difficult when interpreting data from knockout mice to say for certain if a loss or change in synaptic transmission is due to the loss of the subunit removed or unforeseen indirect compensatory mechanisms. Despite these potential problems the evidence does suggest that it is possible that β 2 subunit-containing receptors may be modulated at the synapse by α -CaMK-II. Modulation of β 2 subunit-containing receptors by α -CaMK-II in a recombinant system is clearly not

possible (See Chapter 6) suggesting the mechanisms for modulation of both subtypes could be dependent on cell type.

Evidence in the literature suggests that phosphorylation in neurones can have distinctly different functional effects depending on the receptor subtypes present in a cell (Nusser *et al.*, 1999; Poisbeau *et al.*, 1999). There is evidence for PKC dependent down-regulation of receptor function via phosphorylation of $\beta 2$ Ser⁴¹⁰ in a recombinant system (Kellenberger *et al.*, 1992). A similar modulation by PKC has also been reported to be independent of phosphorylation of this subunit (Chapell *et al.*, 1998). In this case the down-regulation observed was thought to be associated with a loss of receptors from the cell surface and phosphorylation of receptor associated proteins rather than the $\beta 2$ subunit directly. PKC has also been reported to modulate receptor endocytosis in HEK293 cells and cortical neurones in culture. This process involved a dynamin endocytic pathway and was dependent on a dileucine motif present on the $\beta 2$ subunit (Herring *et al.*, 2005).

In a neuronal environment, PKA can modulate IPSCs recorded from cerebellar stellate, basket and Purkinje neurones that are thought to predominantly express the β 2 subunit (Persohn *et al.*, 1992; Kano and Konnerth, 1992; Nusser *et al.*, 1999). In addition, CaMK-II is thought to play a role in rebound potentiation (Kano *et al.*, 1992; Kano *et al.*, 1996; Kawaguchi and Hirano, 2002), a phenomenon that results in an increase in the amplitude of IPSCs without a change in decay times. This occurs in Purkinje cells of the cerebellum that are thought to express predominantly $\alpha 1\beta 2\gamma 2$ subunits (Persohn *et al.*, 1992; Laurie *et al.*, 1992a). Wang and colleagues (2003) have demonstrated an increase in the number of receptors at the cell surface related to PKB phosphorylation of β 2 Ser⁴¹⁰ in HEK293 cells and in neurones using immunocytochemistry. Thus, functional changes due to

phosphorylation of $\beta 2$ by PKB can be detected in HEK293 cells, in contrast to the present study with CaMK-II, and in contrast to the lack of PKA dependent functional modulation of GABA_A receptors in HEK293 cells (McDonald *et al.*, 1998). Therefore, phosphorylation of receptors may not only be dependent on cell type and receptor subunit composition, but also on the kinase under investigation.

In neurones the effect of PKB induced an increase in the amplitude of IPSCs without a change in their decay, which was related to an increase in $\beta 2$ subunitcontaining receptors at the synapse (Wang *et al.*, 2003). This is comparable to the $\beta 2$ subunit-containing population of sIPSCs in the current study increasing in amplitude without a change in decay time after exposure to α -CaMK-II. This raises the possibility that $\beta 2$ subunit mediated effects may involve insertion of new receptors rather than a change in their single channel properties. It may be that a lack of functional effect on $\beta 2$ subunit-containing receptors in recombinant systems is related to a lack of the appropriate cellular machinery needed to insert new receptors. Direct modification of channel properties by phosphorylation may be easier to observe and measure in a recombinant environment. However, insertion of receptors into the cell membrane has been observed in recombinant systems in response to phosphorylation (Wan *et al.*, 1997; Wang *et al.*, 2003).

It might simply be that $\beta 2$ subunit phosphorylation is under much tighter control than that of $\beta 1/3$, requiring multiple kinase anchoring proteins or GABA_A receptor scaffolding proteins to occur. There is evidence for differences in $\beta 2$ and $\beta 3$ binding to kinase anchoring proteins as well as differences in the trafficking and targeting of these receptors to the synapse. For example AKAP 79/150 has been shown to directly bind the $\beta 1/3$ subunits but not $\beta 2$ and is critical for PKA-mediated phosphorylation of the $\beta 3$ subunit (Brandon *et al.*, 2003) and RACK-1 has been associated with binding to and facilitation of phosphorylation of $\beta 1/3$ subunits (Brandon *et al.*, 2002; Jovanovic *et al.*, 2004). The GABA_A receptor-associated protein (GRIF-1) has been identified which binds to the $\beta 2$ subunit but not $\beta 1/3$, although the function of this protein is unknown, it has been proposed as another subunit trafficking factor (Beck *et al.*, 2002).

There is some evidence that the trafficking and targeting of $\beta 2$ and $\beta 3$ subunit-containing receptors to the cell surface are different (Connolly *et al.*, 1996) and that these processes can be differentially affected by phosphorylation (Ives *et al.*, 2002b). The $\beta 2$ and $\beta 3$ subunits require different processes for correct assembly of a receptor and trafficking to the cell surface (Bollan *et al.*, 2003). In addition, $\beta 3$ subunits alone can form functional homomeric channels, whereas $\beta 2$ subunits cannot (Connolly *et al.*, 1996; Wooltorton *et al.*, 1997). Moreover, PKC phosphorylation can modulate receptor endocytosis of $\alpha 1\beta 2\gamma 2$ receptors in HEK293 cells and in neurones (Connolly *et al.*, 1999; Herring *et al.*, 2005) as well as directly affecting the properties of the ion channel (Krishek *et al.*, 1994; Lin *et al.*, 1996).

The activity of CaMK-II itself is thought to be controlled through its localisation to targets where it exerts an effect on proteins (Bayer and Schulman, 2001; Schulman, 2004; Colbran, 2004). For instance, it can bind to the NMDA receptor to maintain the activity of the kinase in a calcium-dependent manner (Bayer *et al.*, 2001). A CaMK-II anchoring protein, formed by alternative splicing of the α -CaMK-II gene that is stably integrated into the CaMK-II holoenzyme and influences CaMK-II localisation and function, has been identified in skeletal muscle (Bayer *et al.*, 1998; Bayer and Schulman, 2001) although not yet in neurones. Although a few potential CaMK-II anchoring proteins have been characterised (See Chapter 1) there are probably others that are yet to be found (Colbran, 2004). It is probable, therefore,

that CaMK-II dependent modulation of GABA_A receptors may also be dependent on, or altered by, appropriate anchoring proteins and that some of the differences seen in the present study between recombinant and neuronal cultures may be due to the presence or absence of critical CaMK-II anchoring proteins.

It is possible to interpret the effects of α -CaMK-II in the current study on sIPSCs as a direct β 3 phosphorylation and possibly an indirect tyrosine kinase phosphorylation of γ 2S (See Chapter 7) leading to a possible increase in IPSC amplitude and a change in decay time seen in both wt and β 2 -/- cultures. This would also explain the strong effect of α -CaMK-II on whole-cell currents recorded from cerebellar granule cells cultured in depolarising conditions when cells are thought to express predominantly β 3 subunit-containing receptors (Ives *et al.*, 2002a) and would also be in accord with the modulation of recombinant $\alpha 1\beta 3\gamma 2$ S receptors seen in NG108-15 cells. The lack of a significant change in the extent of the effect of CaMK-II on β 2 -/- cerebellar cultures, maintained in depolarising conditions, could indicate that there is little β 2 in these cells (Ives *et al.*, 2002a), or that β 2 modulation by α -CaMK-II is only possible at a functional synapse since these cultures generally display no synaptic activity (Mellor *et al.*, 1998).

On application of exogenous GABA, the time to reach the peak amplitude of whole-cell current is relatively slow (See Chapter 2, 200-300ms). Direct changes on GABA_A channel function altering probability of opening or the extent of desensitisation could increase the apparent amplitude of whole-cell currents. It has been demonstrated that phosphorylation of the γ 2 subunit by tyrosine kinases alters the mean open time and probability of opening of the GABA_A receptor (Moss *et al.*, 1995) and that this tyrosine kinase phosphorylation of γ 2S can increase whole-cell currents recorded from HEK293 cells and neurones (Moss *et al.*, 1995).

Tyrosine kinase phosphorylation of γ^2 has been reported to increase the amplitude of IPSCs, although not the decay time (Boxall, 2000; Ma *et al.*, 2003). So it could be speculated that Ser³⁸³ phosphorylation of the β^3 subunit and phosphorylation of γ^2 S by down-stream activation of a tyrosine kinase together, alters directly the properties of the channel in such a way as to lengthen the decay times and increase the amplitudes of a sub-population of IPSCs. The β^2 subunit mediated effect may be related to increased numbers of functional receptors and therefore results in an increase in amplitude with no change in decay times. The β^2 subunit-containing receptors appear to be located at synapses responsible for much larger amplitude IPSCs in the present study (See Fig 8.7), possibly reflecting the fact that they are situated near to the soma or that they are located at specific, large, multi-quantal synapses.

There is evidence that the extent of receptor occupancy on release of GABA varies at different synapses within the same cell (Hájos *et al.*, 2000). Changes in channel properties would be expected to increase decay times and amplitudes of sIPSCs if GABA is not saturating, but would have no effect on amplitude at synapses where GABA is saturating. This may mean that phosphorylation dependent changes in channel function have slightly different effects depending on the degree of receptor occupancy at a given synapse. This may explain why α -CaMK-II appears to increase the amplitude of some IPSCs in $\beta 2$ -/- cultures (See Fig 8.6), but has little effect on the mean IPSC amplitude (See Fig 8.3), as some IPSCs increase in amplitude and decay time and others only increase in decay time dependent on the level of receptor occupancy by GABA. At synapses where GABA is saturating an increase in receptor number is predicted to be the most effective way of up-regulating inhibitory transmission (Otis *et al.*, 1994; De Koninck and Mody, 1994;

Nusser *et al.*, 1998). It is possible therefore that different receptor subtypes are modulated by phosphorylation in different ways. This may allow the most appropriate form of modulation to occur at different synapses.

Changes in the decay times of IPSCs can be the result of changes in desensitisation of the receptor (Jones and Westbrook, 1996). The bi-exponential decay observed at many GABAergic synapses has been proposed to be related to desensitisation. The slower component of decay may relate to GABA_A receptors that have bound GABA that are briefly desensitised before re-opening, prolonging the decay phase of the IPSC (Jones and Westbrook, 1996). The alteration in the apparent contribution of $\tau 1$ and $\tau 2$ between wt and $\beta 2$ -/- cultures on application of α -CaMK-II may suggest that CaMK-II dependent phosphorylation can alter the desensitisation of the receptor. In support of this, the calcium-dependent phosphatase calcineurin has been proposed to alter the desensitisation of the GABA_A receptor (Jones and Westbrook, 1997).

The presence of perisynaptic receptors can alter the apparent rise-time and decay time of IPSCs (Mody and Pearce, 2004). Spillover-mediated transmission has been identified at golgi cell to granule cell synapses in brain slice preparations (Rossi and Hamann, 1998; Wall, 2002; Hamann *et al.*, 2002). This is thought to be related to the presence of high affinity α 6 subunits in these cells which have been identified at cerebellar granule cell synapses (Nusser *et al.*, 1996; Ortinski *et al.*, 2004). Within rat cultures, under physiological conditions, there was little evidence of a tonic current, however, in these mouse wild-type and $\beta 2$ -/- cultures, a proportion of cells did express a tonic current indicating the likely presence of α 6 subunit-containing receptors. Up-regulation of the function of perisynaptic receptors could alter the apparent decay times and rise-times of IPSCs. In the current study, a slight tendancy

to change the rise-time was observed in cells from $\beta 2$ -/- cultures on application of α -CaMK-II which could be related to an effect on perisynaptic receptors, however, this effect did not reach statistical significance (P < 0.1, Fig 8.4D).

In conclusion the analysis of knockout mouse data can be complicated by compensatory changes and any differences observed could potentially be non-specific. But the evidence does indicate that an α -CaMK-II dependent effect occurs on a population of events that disappear in the β 2 knockout mouse. The most parsimonious explanation is that α -CaMK-II can modulate β 2 subunit-containing receptors, but only in the native neuronal synaptic environment. It is not clear why this does not occur in a recombinant system, but it could be because specific kinase anchoring proteins, synaptic scaffolding proteins and/or receptor trafficking machinery is needed for α -CaMK-II to modulate these receptors. This has important implications for the functional role of phosphorylation of different GABA_A receptors at inhibitory synapses.

8.4 CONCLUSIONS:

- There was no significant difference in the modulation of whole-cell GABA currents by α-CaMK-II recorded from cerebellar granule cells maintained in depolarising conditions lacking the β2 subunit. Suggesting a lack of β2 in wild-type cultures or a lack of an α-CaMK-II dependent effect on β2 subunit-containing receptors under these culture conditions.
- Application of α-CaMK-II had a different effect on two populations of sIPSCs recorded from cerebellar granule cells from wt mice. One population of sIPSCs increased in decay time and another population of larger amplitude sIPSCs increased significantly in amplitude with no change in decay.

- The larger amplitude population of sIPSCs was no longer present in β 2 -/cultures, suggesting that they were mediated by β 2 subunit-containing receptors and that these sIPSCs could be modulated by α -CaMK-II in the wild-type cultures.
- The loss of the $\beta 2$ subunit resulted in subtle changes to the population of sIPSCs observed in control conditions. There was an apparent loss of large amplitude fast decaying events which were replaced by large amplitude events with slower decay times. This may coincide with a loss of $\alpha 1$ subunit-containing receptors alongside the loss of $\beta 2$.

<u>Chapter Nine:</u> <u>Physiological significance of CaMK-II modulation of</u> <u>GABA_A receptors</u>

9.1 INTRODUCTION

Modulation of GABA_A receptor mediated currents through activation of CaMK-II has been reported in a number of neuronal cell types (Kano et al., 1992; Wang et al., 1995; Kano et al., 1996; Kawaguchi and Hirano, 2002; Wei et al., 2004). The current study provides evidence for the potential mechanisms behind this Use of NG108-15 cells allowed sites of CaMK-II dependent modulation. phosphorylation to be determined revealing that Ser^{383} on the $\beta 3$ subunit is likely to be a direct site of CaMK-II phosphorylation on the GABA_A receptor. In addition, a CaMK-II dependent phosphorylation of tyrosine residues 357 and 359 on the $\gamma 2S$ subunit was also observed (See Fig 9.1). The evidence suggested that CaMK-II can activate a down-stream tyrosine kinase which goes on to phosphorylate $\beta 1$ and $\beta 3$ subunit-containing GABA_A receptors. However, $\beta 2$ subunit-containing receptors are not modulated at all by CaMK-II suggesting a specific kinase anchoring protein may be required for any effect on $\beta 2$ subunit-containing receptors to be observed (See Chapters 6 and 7). In addition, in the HEK293 cell line, it was not possible to identify any functional modulation of GABAA receptors by CaMK-II (See Chapter 3), again suggesting kinase anchoring proteins may be important in regulating CaMK-II dependent phosphorylation of GABA_A receptors. The importance of these sites of phosphorylation was confirmed by expression of recombinant receptors in cerebellar granule cells maintained in depolarising conditions. However it was still not possible to observe any α -CaMK-II dependent modulation of $\alpha 1\beta 2\gamma 2S$ receptors.

CaMK-II is capable of increasing whole-cell current amplitudes in a neuronal culture preparation (cerebellar granule cells in depolarising conditions) that is predicted to express predominantly β 3 subunit-containing receptors (See Chapter 4). Analysis of synaptic currents recorded from cerebellar granule cell cultures predicted to express both β 2 and β 3 subunits (physiological conditions) confirms that CaMK-II can significantly increase the amplitude and decay times of IPSCs. However, these effects are not uniform across the whole population of IPSCs recorded. One population of IPSCs significantly increased in decay time and another population of IPSCs significantly increased in amplitude with no affect on decay time (See Chapter 5).

Analysis of synaptic currents in a culture preparation lacking $\beta 2$ subunitcontaining receptors inferred that CaMK-II can indeed modulate $\beta 2$ subunitcontaining GABA_A receptor mediated IPSCs. The effect of CaMK-II on $\beta 2$ subunitcontaining receptors involves an increase in IPSC peak amplitudes with no change in decay times (See Chapter 8). It is proposed that the mechanism of CaMK-II action on $\beta 2$ subunit-containing receptors may involve insertion of receptors into the plasma membrane rather than a direct effect on channel properties, consistent with other evidence in the literature regarding $\beta 2$ subunit phosphorylation. Whereas, CaMK-II dependent effects on $\beta 3$ subunit-containing receptors may involve direct phosphorylation and alteration of channel properties which is consistent with the lengthened decay times of IPSCs.

Fig 9.1:



Fig 9.1: Summary of the proposed action of CaMK-II on different GABA_A receptor subtypes.

CaMK-II can modulate GABA_A receptor function through direct phosphorylation of the β 3 subunit at Ser³⁸³ and through activation of a tyrosine kinase that can phosphorylate the γ 2S subunit at Tyr^{357, 359}. PYK2 and Src tyrosine kinase (the most likely candidates) have been proposed to be activated by Ca²⁺ dependent signalling pathways and in some cases CaMK-II in other cell types.

CaMK-II modulation of the $\beta 2$ subunit-containing receptors could occur through direct phosphorylation of the subunit or through an indirect mechanism. It is proposed that CaMK-II modulation of $\beta 2$ subunit-containing receptors may occur through insertion of receptors into the plasma membrane.

Arrows in black indicate potential sites of direct phosphorylation. Block arrows indicate sites of phosphorylation mediated by a downstream kinase. Dashed line arrows indicate pathways involving changes to receptor trafficking.

9.2 Limitations in studying phosphorylation.

9.2.1 Discrepancies in the literature.

The investigation of phosphorylation-dependent modulation of GABA_A receptors in both recombinant and neuronal preparations has revealed a considerable amount of diversity in the functional response observed (Moss and Smart, 1996; Brandon *et al.*, 2002a; Lüscher and Keller, 2004). In some cases this may be due to differential effects of phosphorylation by the same kinase on different GABA_A receptor subtypes (McDonald *et al.*, 1998; Nusser *et al.*, 1999). However, this cannot explain all of the differences in the functional effects observed. The importance of kinase anchoring proteins has been revealed by a number of experiments in recent years, and different kinase anchoring proteins in different cell types, which may localise the kinase in question to different sites or enable binding to different GABA_A receptor subtypes (Brandon *et al.*, 2002b; Brandon *et al.*, 2003; Terunuma *et al.*, 2004).

The mechanisms of functional modification by phosphorylation may also vary depending on the kinase involved or again the target GABA_A receptor subtype. Covalent modification of receptor structure and alteration of channel properties, such as desensitisation (Jones and Westbrook, 1997) or probability of channel opening (Moss *et al.*, 1995), have been shown to modulate GABA_A receptor function. In addition, phosphorylation can alter the trafficking of GABA_A receptors to the synapse and dynamic modulation of receptor number has been shown to be an important mechanism in synaptic plasticity (Nusser *et al.*, 1998; Wang *et al.*, 2003**b**; Kittler and Moss, 2003). When interpreting data relating to phosphorylation dependent changes in GABA_A receptor function it is important to note that different experimental approaches can reveal different functional effects (See Chapter 1). For instance, the mechanism of activation of a particular kinase or the time-scale of activation can have a significant influence on the effect observed (Angelotti *et al.*, 1993; Krishek *et al.*, 1994; Lin *et al.*, 1996; McDonald *et al.*, 1998). When activating any kinase signalling pathway it is very difficult to discount downstream activation of other kinases, as has been demonstrated in the current study (See Chapter 7). In addition, phosphorylation must be considered in the context of a dynamically changing balance of activity between different kinases and phosphatases. The background basal phosphorylation state of the GABA_A receptor in a given cell type may also alter the response observed in any given experiment (Brandon *et al.*, 2000; Brandon *et al.*, 2001).

The method of analysis also has significant implications in that measurement of whole-cell current amplitudes *per se*, may not identify subtle changes or differences in channel behaviour which could have a significant effect on synaptic currents (Hinkle and Macdonald, 2003). When analysing the effects of phosphorylation on a heterogeneous population of GABA_A receptor mediated IPSCs within a neuronal population, it is possible a significant effect on different subpopulations of events may be missed if only mean values are measured (See Chapters 5 and 8).

9.2.2 CaMK-II

When looking at CaMK-II dependent modification of receptor function these problems are pronounced by the complexity of the processes that modulate the kinase activity itself (Griffith *et al.*, 2003; Colbran, 2004). The current study indicates that pre-activated recombinant α -CaMK-II is capable of modulating GABA_A receptor function. However, in cerebellar granule cells maintained in depolarising conditions, endogenous β -CaMK-II may also be capable of basal phosphorylation of GABA_A receptors (See Chapter 4) as this is the predominant isoform in this cell type (Miller and Kennedy, 1985; Burgin *et al.*, 1990). It is currently not known what the role of different ratios of α to β CaMK-II subunits and/or translocation of CaMK-II to inhibitory as opposed to excitatory synapses might be in relation to GABA_A receptor modification.

The recombinant α -CaMK-II used in the current study can phosphorylate the same substrates as the full-length form of this subunit (Suzuki-Takeuchi *et al.*, 1992) but cannot form a holoenzyme (Shen and Meyer, 1998). Pre-activation of this recombinant form of α -CaMK-II and inclusion within the patch pipette has been used successfully to reveal CaMK-II dependent effects on the function of AMPA receptors (Lledo *et al.*, 1995; Barria *et al.*, 1997). Application of a pre-activated recombinant form of β -CaMK-II would be expected to have a similar effect using this protocol because of the similarity in substrate specificity between the two isoforms (Miller and Kennedy, 1985). The α and β subunits differ in their sensitivity to activation by Ca²⁺/CaM (Miller and Kennedy, 1985; Brocke *et al.*, 1999) which in this case is not relevant as CaMK-II was pre-activated *in vitro*.

It should be considered though, that localisation and targeting of recombinant constructs of CaMK-II may not be the same as that under physiological conditions after activation of an endogenous CaMK-II holoenzyme by Ca²⁺ signalling which, in itself, is a complex and highly regulated process (Meldolesi, 2001; Franks and Sejnowski, 2002; Augustine *et al.*, 2003). Pre-activated recombinant α -CaMK-II can

be expected to phosphorylate CaMK-II substrates within the cell but the role of $\alpha:\beta$ subunits within an endogenous heteromer cannot be determined using this protocol.

9.3 The roles of specific GABA_A receptor subtypes.

9.3.1 Specific synapses express discrete GABA_A receptor subtypes.

The current study has revealed a difference in the response to CaMK-II dependent phosphorylation of different GABA_A receptor subtypes (See Chapter 6 and Chapter 8). This finding is similar to the observed differences in response of different GABA_A receptor subtypes to PKA dependent phosphorylation (McDonald *et al.*, 1998). Such differences have been proposed to partly explain some of the cell type dependent differences in functional response to phosphorylation by PKA and PKC (Nusser *et al.*, 1999; Poisbeau *et al.*, 1999).

Recent evidence also suggests that different GABA_A receptor subtypes could be targeted to different locations within the same cell (Connolly *et al.*, 1996; Koulen *et al.*, 1996; Brünig *et al.*, 2002). Specific synapses within the same neurone may express specific GABA_A receptor subtypes (See Fig 9.2). Within rat or rabbit retinae, $\alpha 1$, $\alpha 2$ and $\alpha 3$ subunits were shown to not be co-localised within the same clusters. Individual neurones could express several isoforms of GABA_A receptor and these were targeted to specific postsynaptic sites (Koulen *et al.*, 1996; Brünig *et al.*, 2002). In dissociated cultures of hippocampal neurones, specific α subunits were targeted to discrete cellular regions in a different way depending on the cell type (Brünig *et al.*, 2002). Although these studies focused on α subunit expression it is known that certain α and β subunits preferentially assemble suggesting that the subcellular location of β subunits may also be tightly controlled (Benke *et al.*, 1994; Whiting, 2003). In support of this, in a recombinant system β subunits have been shown to target to different subcellular locations (Connolly *et al.*, 1996). Furthermore, specific targeting of GABA_A receptor subtypes to particular synapses appears to occur in the current study, in that β 2 subunit-containing receptors are targeted to form large IPSC amplitude synapses whereas the other β subunits, presumed to be predominantly β 3 in cerebellar granule cells, are targeted to smaller amplitude IPSC synapses (See Chapter 8).

Interneurones are able to form synapses at precise locations on the target cell (McBain and Fisahn, 2001; Di Cristo *et al.*, 2004; Farrant and Nusser, 2005; Somogyi, 2005). Different interneurones can form discrete synapses onto hippocampal pyramidal cells with different GABA_A receptor subtypes located postsynaptically (See Fig 9.2), even within a similar domain of the pyramidal cell. Pyramidal cells clearly have mechanisms to target GABA_A receptor subtypes to distinct synapses depending on the presynaptic cell type (Nyíri *et al.*, 2001).

The targeting of specific GABA_A receptor subtypes to specific sites is important because the channel properties of GABA_A receptors can differ according to the subunits within the receptor (See Fig 9.2). For example, the presence of different α subunits has been shown to alter the kinetics of a given receptor (Gingrich *et al.*, 1995). Different receptor subtypes may be targeted to specific locations so that the functional properties of a GABA_A receptor may be tailored to suit the specific synapse and functional role(s) that they play. In addition, these functional properties can be differentially modulated by phosphorylation depending on the GABA_A receptor subtype present (See Fig 9.2).

In the current study, $\beta 2$ subunit-containing receptors could not be modulated by CaMK-II in a recombinant system, whereas it was possible to observe modulation by CaMK-II of $\beta 3$ subunit-containing GABA_A receptors (Chapter 6). This suggests a

possible difference in the way GABA_A receptor subtypes are modulated by CaMK-II. CaMK-II dependent effects on β 3 subunit-containing receptors were thought to involve a change in decay time of IPSCs consistent with a change in the single channel properties of the receptor, whereas effects on β 2 subunit-containing receptors appeared to involve an increase in amplitude with no change in decay consistent with alterations in receptor number (See Chapter 8). The heterogeneity of response of different β subunits to phosphorylation may provide a mechanism to alter the function of specific synapses in precise ways according to their location and physiological role.

9.3.2 The physiological roles of $\beta 2$ and $\beta 3$ subunit-containing receptors.

Evidence suggests that certain α and β subunits preferentially assemble together (Benke *et al.*, 1994). It is thought that $\alpha 1\beta 2\gamma 2$ and $\alpha 2/3\beta 3\gamma 2$ receptors comprise the majority of receptors in the brain (Benke *et al.*, 1994; Whiting, 2003; Vicini and Ortinski, 2004). Although the $\alpha 1\beta 2\gamma 2$ receptor is the most widely expressed recent evidence suggests both $\beta 2$ and $\beta 3$ subunit-containing receptors have important physiological roles. The use of knock-in mice with mutations that render specific subtypes of receptor insensitive to selected drugs has enabled their contribution to the effects of these drugs to be ascertained (Vicini and Ortinski, 2004; Rudolph and Möhler, 2004; Wafford *et al.*, 2004; Wafford, 2005). Knock-in mutation of the $\beta 2$ and $\beta 3$ subunits, with mutations known to be critical for the action of general anaesthetics *in vitro*, has revealed that $\beta 2$ and $\beta 3$ subunit-containing receptors mediate different physiological effects of anaesthetics (Rudolph and Möhler, 2004). The $\beta 2$ subunit appears to be involved in mediating the sedative properties of anaesthetics whereas the β 3 subunit appears to mediate a loss of consciousness (Jurd *et al.*, 2003; Reynolds *et al.*, 2003).

The $\beta 2$ subunit also appears to mediate a significant proportion of the hypothermic effect of the general anaesthetic etomidate (Cirone *et al.*, 2004). The drug oleamide has sleep promoting effects in rodents but administration of this drug to $\beta 3$ knockout mice fails to have any affect suggesting that the action of this drug is solely mediated through actions on $\beta 3$ subunit-containing receptors (Laposky *et al.*, 2001). The use of $\alpha 1$ subunit and $\beta 2$ subunit knockout mice has also revealed that the behavioural effects of ethanol may be mediated by distinct GABA_A receptor subtypes (Blednov *et al.*, 2003).

9.3.3 Network oscillations.

Inhibitory synaptic transmission plays a critical role in the generation of network oscillations which are associated with behaviour (Gaiarsa *et al.*, 2002). During active exploration rat hippocampal neurones exhibit rhythmic theta and gamma frequency activity (McBain and Fisahn, 2001). Simulations involving interneuronal networks with fast and slow GABA_A receptor kinetics (fast and slow decaying IPSCs) have been shown to produce theta-gamma rhythmic activity (White *et al.*, 2000; McBain and Fisahn, 2001). The kinetics of GABA_A receptors have been shown to be important in shaping network oscillations in the cortex (Whittington *et al.*, 1995). Alteration of the decay times of IPSCs through phosphorylation may therefore be important in generating or modulating this rhythmic activity.

Subunit specific effects of phosphorylation on different subtypes may modulate receptor properties changing the decay times of IPSCs and therefore altering oscillations in the network. In addition mutations that affect the decay times

of IPSCs have been linked to pathological conditions such as epilepsy (Bianchi *et al.*, 2002; Fisher, 2004; Macdonald *et al.*, 2004) and have been shown to alter behaviour (Homanics *et al.*, 2005).

Specific GABA_A receptor subunits have been shown to play precise roles in the generation of oscillatory activity. The role of the β 3 subunit in network oscillations of the olfactory bulb has been investigated using the β 3 subunit knockout mouse. Alteration of normal synaptic inhibition resulted in a significant increase in the amplitude of theta and gamma oscillations recorded from this region in freely moving mice resulting in an alteration of normal olfactory discrimination (Nusser *et al.*, 2001). This implied an important role for β 3 subunit-containing receptors in modulating theta and gamma oscillations. In addition use of a knock-in mouse with a β 3 mutation rendering this subunit insensitive to the anaesthetic etomidate revealed that β 2 and β 3 subunit-containing receptors could exert opposing actions on theta activity in neocortical slice cultures (Drexler *et al.*, 2005).

It appears specific synapses formed at particular locations by specific interneurones may play precise roles in shaping network activity and oscillations (McBain and Fisahn, 2001; Klausberger *et al.*, 2004; Farrant and Nusser, 2005; Somogyi, 2005). It is becoming clear that the function of GABA_A receptors is not simply to provide "blanket" inhibition. If different inhibitory synapses play precise roles then modification of these synapses by phosphorylation may also differ in mechanism and functional effect. Phosphorylation dependent effects may occur through several different mechanisms with the functional effect on GABA_A receptor subtypes varying according to the subtype present in order to modulate and control the function of specific synapses and therefore patterns of activity related to behaviour.

Fig 9.2:



Fig 9.2: Specific GABA_A receptor subtypes are targeted to discrete subcellular locations.

Different classes of interneurone have been shown to form synapses at particular locations on the target cell. In addition there is evidence that specific GABA_A receptor subtypes are expressed at different synapses.

Different GABA_A receptor subtypes have different kinetic properties which shape the profile of an IPSC. Phosphorylation can have a differential effect on different GABA_A receptor subtypes.

Network oscillations relevant to behaviour are differentially affected by changes in activity at these different synapses.

IN = Interneurones. PC = Principal cell

9.4 Physiological role of GABA_A receptor modulation by CaMK-II.

In summary, evidence now suggests that GABA_A receptors can be composed of discrete sets of subunits which may be targeted to distinct subcellular locations (Koulen *et al.*, 1996; Brünig *et al.*, 2002). The different subtypes can be shown to play different physiological roles in that the different effects of certain drugs can be ascribed to specific GABA_A receptor subtypes (Rudolph and Möhler, 2004; Wafford *et al.*, 2004). In the current study phosphorylation appears able to affect the function of different GABA_A receptor subtypes in different ways (See Chapters 6 and 8). There are different mechanisms of phosphorylation-dependent modulation of function involving changes in receptor targeting and trafficking and also direct changes to channel properties. These phosphorylation-dependent changes are thought to be crucial for the functional role of GABA_A receptors at synapses. As a result CaMK-II dependent modulation of GABA_A receptors may play important roles in neurodevelopment, inhibitory plasticity and certain disease states.

9.4.1 Development.

The role of the β 3 subunit in neurodevelopment was highlighted by the severe behavioural deficits of the β 3 subunit knockout mouse (Homanics *et al.*, 1997; DeLorey *et al.*, 1998). In contrast, the β 2 subunit knockout mouse phenotype was relatively normal given the comparable loss of receptor number found in these mice (Sur *et al.*, 2001) suggesting each β subunit plays very different physiological roles in neurodevelopment. In support of this the β 3 subunit is the predominant subunit expressed early on in development (Laurie *et al.*, 1992). The current study indicated that β 3 subunit-containing receptors can be phosphorylated by CaMK-II resulting in a potentiation of function (See Chapters 6 and 7). In support of this electrophysiological evidence, biochemical studies have revealed a CaMK-II dependent phosphorylation of the β 3 subunit in embryonic cortical neurones. Application of muscimol in these cultures activates GABA_A receptors which, at this point in development, are excitatory. As a result of GABA_A receptor activation, there is a depolarisation and Ca²⁺ influx which results in phosphorylation of the β 3 subunit of the GABA_A receptor that can be blocked by KN-93. Further biochemical evidence suggests CaMK-II can bind to the β subunits of GABA_A receptors *in vitro* at a region within the first 30 amino acids of the TM3-4 loop (McAinsh K, Jovanovic JN and Moss SJ, personal communication). Application of pre-activated CaMK-II to the same embryonic cortical neuronal cultures resulted in a significant potentiation of the whole-cell current (See Chapter 4).

It appears from the biochemical evidence that CaMK-II dependent modulation of β 3 subunit-containing receptors may play a role in strengthening GABA_A receptor synaptic transmission early on in development possibly playing a role in activity dependent strengthening of inhibitory synaptic transmission. Certain Ca²⁺-dependent forms of inhibitory plasticity have been suggested to be dependent on the stage of development of a particular cell-type and may play a role in the functional maturation of GABAergic synaptic transmission (Komatsu and Iwakiri, 1993; Komatsu, 1994; Gubellini *et al.*, 2001). It could be speculated that CaMK-II dependent potentiation of GABA_A receptor function could provide a mechanism for these Ca²⁺-dependent forms of inhibitory plasticity.

9.4.2 Homeostatic plasticity and metaplasticity.

The role of excitatory synaptic plasticity, in particular LTP, in learning and memory has been extensively studied in recent years. However, it is now becoming apparent that inhibitory synapses are also capable of long-lasting synaptic plasticity and that changes in inhibition can modulate excitatory plasticity. Activity-dependent changes in synaptic plasticity altering the threshold of excitatory LTP induction has been termed metaplasticity (Abraham and Tate, 1997; Elgersma *et al.*, 2004). For example in CA1 pyramidal neurones induction of excitatory LTP activated a calcineurin-dependent LTD of GABA_A receptor-mediated synaptic transmission which was found to enhance the LTP of EPSPs (Lu *et al.*, 2000).

Inhibitory synapses have been demonstrated to change in strength in such a way as to enhance changes in excitatory transmission, but they have also been shown to adjust in opposition to changes in excitatory transmission in homeostatic forms of plasticity (Turrigiano and Nelson, 2004). Certain forms of inhibitory plasticity have been shown to be Ca^{2+} -dependent and bidirectional (Aizenman *et al.*, 1998) in a manner similar to excitatory LTP. CaMK-II has been shown to be critical in the potentiation of AMPA receptor-mediated synaptic transmission. It has also been shown that CaMK-II may play a role in inhibitory synaptic plasticity. CaMK-II modulation of GABA_A receptors may represent a possible mechanism of postsynaptic inhibitory plasticity that could be involved in homeostatic or metaplasticity.

Rebound potentiation is a form of CaMK-II dependent long-term synaptic plasticity that occurs in Purkinje cells in response to a strong stimulation of excitatory inputs onto the same cell. In this way it could be thought of as a homeostatic form of plasticity acting to increase GABAergic synaptic transmission

after strong excitation. Purkinje cells are thought to express pre-dominantly $\beta 2$ subunits over $\beta 3$ and $\beta 1$ (Persohn *et al.*, 1992). Depolarisation and Ca²⁺ entry after strong excitation activates CaMK-II which goes on to modify IPSCs by increasing their amplitude with no change in decay times (Kano *et al.*, 1992; Kano *et al.*, 1996; Kawaguchi and Hirano, 2002). This is consistent with the current studies observation of $\beta 2$ subunit-containing receptor mediated IPSCs increasing in amplitude with no change in decay times (See Chapter 8). It has been shown that rebound potentiation does not occur in the $\beta 2$ knockout mouse supporting the role of this subunit in this kind of plasticity (Duguid and Smart, personal communication).

9.5 The role of CaMK-II dependent modulation of GABA_A receptors in disease states.

9.5.1 Angelman syndrome and Autism.

The β 3 knockout mouse displays epilepsy and behavioural deficits that are similar to the phenotype of Angelman syndrome (DeLorey *et al.*, 1998) a neurodevelopmental disorder affecting humans that results from a maternal deletion/mutation of chromosome 15q11-13. In this region, there are two candidate genes thought to be potentially responsible for the neurodevelopmental deficits seen. The UBE3A gene which encodes a ubiquitin ligase, a protein involved in protein degradation (Weeber *et al.*, 2003) and the GABRB3 gene encoding the β 3 subunit of the GABA_A receptor. Both of these genes have been demonstrated to have reduced expression in Angelman syndrome patients (Samaco *et al.*, 2005) and disruption of both genes in mice has generated models of Angelman syndrome with many of the same behavioural deficits as that seen in humans (DeLorey *et al.*, 1998; Weeber *et al.*, 2003; Sinkkonen *et al.*, 2003).

The role of the β 3 subunit early on in development suggested that Angelman syndrome was caused by dysfunction in the normal development of inhibitory synaptic transmission. However, in another mouse model of Angelman syndrome involving maternal deletion of UBE3A, a reduction in CaMK-II activity and the total amount of CaMK-II associated with the postsynaptic density has been proposed to underlie the deficits in normal function associated with Angelman syndrome (Weeber *et al.*, 2003). It is interesting to speculate that disruption of an interaction between CaMK-II and the β 3 subunit of the GABA_A receptor may be involved in the neurodevelopmental deficits seen in this syndrome. This implies an interaction between CaMK-II and the β 3 subunit may be important in the development of inhibitory synaptic transmission.

Other conditions such as Rett syndrome, an X-linked dominant disorder caused by MECP2 (a transcriptional repressor protein) gene mutation and autism have also been demonstrated to involve a reduction in UPE3A and GABRB3 gene expression (Samaco *et al.*, 2005). If CaMK-II modulation of β 3 subunit-containing GABA_A receptors in the early stages of development plays a role in neurodevelopement this signalling pathway may be a potentially important target for future drug therapies.

9.5.2 Epilepsy

A number of mutations in $GABA_A$ receptor subunit genes have been identified in humans that have been associated with epilepsy (Jones-Davis and Macdonald, 2003; Macdonald *et al.*, 2004). Some of these mutations alter the properties of the channel itself and are proposed to alter the decay times of IPSCs (Bianchi *et al.*, 2002; Fisher, 2004). In addition, epilepsy has been associated with up and/or down-regulation of specific GABA_A receptor subunit mRNAs (Brooks-Kayal *et al.*, 1998; Meguro *et al.*, 2004) and increases in the number of GABA_A receptors at specific synapses (Nusser *et al.*, 1998).

During periods of enhanced neural activity the intracellular Ca^{2+} load would be expected to increase and subsequently CaMK-II activity may also be increased. Observed compensatory increases in GABA_A receptor function (Brooks-Kayal *et al.*, 1998; Nusser *et al.*, 1998) could be related to CaMK-II dependent modification of receptor function. The decay times of IPSCs and the number of GABA_A receptors at the synapse could both potentially be modulated by CaMK-II dependent phosphorylation.

In support of this epilepsy has also been linked to a decrease in the expression of α -CaMK-II. In human tissue of patients undergoing surgery who suffer from severe seizures immunocytochemical staining revealed a decrease in the expression of α -CaMK-II and α -CaMK-II autophosphorylated at Thr²⁸⁶ (Battaglia *et al.*, 2002). In hippocampal neuronal cultures, complimentary oligonucleotides for α -CaMK-II were used to decrease the expression of this subunit. Reduction in α -CaMK-II expression resulted in increased neuronal excitability and epileptiform discharges. The magnitude of the reduced α -CaMK-II expression could be correlated with increased neuronal excitability (Churn *et al.*, 2000). In addition, in a pilocarpineinduced rat model of epilepsy it has been demonstrated that there is a decrease in the activity of cortical and hippocampal CaMK-II (Singleton *et al.*, 2005). These results are in contrast to the fact that increased α -CaMK-II activity and expression is associated with increases in excitatory synaptic transmission (Ouyang *et al.*, 1997; Colbran and Brown, 2004; Collingridge *et al.*, 2004). It could be speculated that CaMK-II dependent effects on $GABA_A$ receptor mediated inhibitory synaptic transmission might be involved in epilepsy in a causative or compensatory manner.

9.6 CONCLUSIONS

The evidence presented in this study suggests that CaMK-II can have a different effect on the function of specific GABA_A receptor subtypes and that these effects may be mediated by different mechanisms. Although CaMK-II dependent effects on β 1/3 subunit-containing receptors appear to be related to direct phosphorylation of the receptor it is not yet known if CaMK-II modulation of β 2 subunit-containing receptors occurs through direct phosphorylation or another indirect process (See Fig 9.1). The different effects on IPSCs may be tailored to form specific roles. As β 2 and β 3 subunit-containing receptors appear to cluster at specific sets of synapses with specific physiological roles, phosphorylation may alter GABA_A receptor function of different subtypes according to the function they perform.

Modulation of GABA_A receptors by CaMK-II may represent a mechanism of synaptic plasticity that is involved in activity-dependent formation of inhibitory synapses during development. It may also represent a way of altering the decay of IPSCs in order to alter network oscillations crucial to behaviour. Alteration in IPSC decay times appears to form an important part of development and alteration of IPSC decay times has important pathological and physiological consequences (Macdonald *et al.*, 2004; Homanics *et al.*, 2005).

Long-term changes in the amplitude of IPSCs, as seen in rebound potentiation, may have important implications in the processes of learning and memory. Understanding the role of different $GABA_A$ receptor subtypes and their modulation by phosphorylation is crucial to understanding many physiological and pathological forms of inhibitory plasticity. In addition, understanding these processes may highlight important potential therapeutic targets for a number of diseases linked to alterations in the normal behaviour of GABA_A receptors and inhibitory synapses.

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Appendix 1:

A1.1 Oligonucleotide design for site-directed mutagenesis and knockout mouse genotyping.

Carried out in collaboration with Dr. A.M Hosie and Ms. H.M.A da Silva as follows:

Two complimentary oligonucleotides were designed containing the desired mutation for use as mutagenic primers in a PCR reaction. The design of the 5'-3' primer is shown in Table A1. Primers were synthesised by Invitrogen Life Technologies. The $\gamma 2S^{T357F, T359F}$ cDNA was a gift from Prof. S.J. Moss. As required some subunits were tagged with the 9E10 epitope (EQKLISEEDL) from c-myc, inserted between the fourth and fifth amino acid of the mature protein by site-directed mutagenesis (Connolly *et al.*, 1996).

Subunit	Sequence of primer #1:	Primer #2:
mutation	5' – 3'	
β3 ^{S383A}	gtataggaaacaggccatgcccaaggaag	cttccttgggcatggcctgtttcctatac
β3 ^{S408A,S409A}	ctacggaggagggctgcacagctcaaaatc	gattttgagctgtgcagccctcctccgtag
$\gamma 2S^{T348A,T350A}$	gatattcgtcccagagcagcagccattcaaatgaac	gttcatttgaatggctgctgctctgggacgaatatc
γ2S ^{Y357F,Y359F}	gagggatgaagaatttggctttgagtgtttggatg	catccaaacactcaaagccaaattcttcatccctc

Table A1: Primers designed for the construction of $GABA_A$ receptor subunitcDNAs containing the desired single point mutations.

Primers were designed by by Dr A Hosie. Letters indicate: a = adenine, g = guanine, c = cytosine, t = thymine.

A1.2 Transformation of competent *E-Coli* cells using TOP10 one shot Kit (Invitrogen).

Carried out by Ms. H.M.A da Silva as follows:

- An aliquot of cDNA PCR product was centrifuged and placed on ice.
- An aliquot of frozen competent cells was thawed and kept on ice.
- 0.5-1 μl of cDNA PCR product was mixed with the competent cells (25 μl) in a sterile microcentrifuge tube and this was left on ice for 30 min.
- The tube was then transferred to a 42°C water bath for 30 s and then placed back on ice.
- 80 μl of pre-warmed SOC media (Sigma, 0.5 % w/v yeast extract, 2.0 % w/v tryptone, 8.6 mM NaCl, 2.5 mM KCl, 20 mM MgSO₄, 20 mM glucose) was added to the cells and they were incubated for 60-90 min at 37°C shaking at ~900 rpm.
- 100 μl/transformation was then plated out on LB-agar plates containing
 50 μg/ml ampicillin (Sigma) and left overnight for colonies to develop.
- At least 2 colonies per mutant were selected from the agar plates and grown overnight in 2ml LB Broth base media (Sigma) with 10 % w/v ampicillin, the plasmid from these *E. Coli* cultures was purified before sequencing.

A1.3 Purification of plasmid DNA using the QIAPrep Spin Miniprep kit (Qiagen).

Carried out by Ms H.M.A da Silva as follows:

• 2 ml of bacterial suspension were transferred to a microcentrifuge tube and centrifuged for 1-2 min at 13,200 rpm and the supernatant discarded.

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- The pellet of bacterial cells was then re-suspended in 250 µl of Buffer P1 (as supplied).
- 250 µl of buffer P2 (as supplied) was added and again inverted 4-6 times to mix.
- $350 \ \mu l$ of buffer N3 (as supplied) was added and inverted 4-6 times.
- The mixture was then centrifuged at 16,000 rpm for 10 min, and the supernatant decanted into a spin column.
- The column was centrifuged at 13,200 rpm for 30-60 s.
- The flow through the column was discarded and 750 µl buffer PE was added and centrifuged for 30-60 s for an additional 2 min, to remove any residual washing buffer.
- The QIAprep column was placed into a clean microcentrifuge tube. To elute the DNA, 50 µl buffer EB (10 mM Tris-Cl @ pH 8.5) or water was added to the centre of the QIAprep column, allowed to stand for 1 min, centrifuged for 1 min.
- The concentration of purified cNDA was measured using a BioPhotometer (Eppendorf).

A1.4 Preparation of mouse tail DNA.

Carried out by Ms H.M.A da Silva as follows:

- Tails were placed into a 1.5 ml microcentrifuge tube and 700 μl of NTES (50 mM Tris HCl @ pH 8, 50 mM EDTA, 100 mM NaCl₂ and 1 % w/v SDS) was added along with 7.5 μl of 20 mg/ml Proteinase K (Sigma) per tube.
- This was incubated overnight at 55°C on a thermomixer at 300 rpm.

- The following day 600 µl of phenol/chloroform/isoamyl alcohol (25:24:1) was added and the tubes were left to shake for 10-15 min. They were then spun at 11,400 rpm in a microcentrifuge. The top phase (DNA) was transferred into a fresh tube avoiding the interface.
- Then 2.5 μ l of 20 mg/ml RNase was added and incubated for 1 hr at 37°C.
- Phenol/chloroform/iosamyl alcohol (25:24:1) was added for 5-10 min and then the tubes were microfuged again for 5 min.
- The top phase was transferred to a fresh tube and this was filled with 1 ml isopropanol. This was inverted a few times until a precipitate had formed and centrifuged for a further 10 min.
- The supernatant was aspirated and 100 µl 70 % v/v ethanol was added, followed by centrifugation for 5 min (12,000 rpm). The supernatant was aspirated again and the pellet left to air dry.
- 300 µl of TE buffer was added and left at 37°C at 300 rpm on a thermomixer overnight. The next day the solution could be transferred to the fridge until required.