Cellular/Molecular

Auxiliary Subunit GSG1L Acts to Suppress Calcium-Permeable AMPA Receptor Function

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AMPA-type glutamate receptors are ligand-gated cation channels responsible for a majority of the fast excitatory synaptic transmission in the brain. Their behavior and calcium permeability depends critically on their subunit composition and the identity of associated auxiliary proteins. Calcium-permeable AMPA receptors (CP-AMPARs) contribute to various forms of synaptic plasticity, and their dysfunction underlies a number of serious neurological conditions. For CP-AMPARs, the prototypical transmembrane AMPAR regulatory protein stargazin, which acts as an auxiliary subunit, enhances receptor function by increasing single-channel conductance, slowing channel gating, increasing calcium permeability, and relieving the voltage-dependent block by endogenous intracellular polyamines. We find that, in contrast, GSG1L, a transmembrane auxiliary protein identified recently as being part of the AMPAR proteome, acts to reduce the weighted mean single-channel conductance and calcium permeability of recombinant CP-AMPARs, while increasing polyaminedependent rectification. To examine the effects of GSG1L on native AMPARs, we manipulated its expression in cerebellar and hippocampal neurons. Transfection of GSG1L into mouse cultured cerebellar stellate cells that lack this protein increased the inward rectification of mEPSCs. Conversely, shRNA-mediated knockdown of endogenous GSG1L in rat cultured hippocampal pyramidal neurons led to an increase in mEPSC amplitude and in the underlying weighted mean single-channel conductance, revealing that GSG1L acts to suppress current flow through native CP-AMPARs. Thus, our data suggest that GSG1L extends the functional repertoire of AMPAR auxiliary subunits, which can act not only to enhance but also diminish current flow through their associated AMPARs.

Key words: auxiliary AMPAR subunits; calcium-permeable AMPARs; glutamatergic synapses; GSG1L; polyamine block; single channels

Significance Statement

Calcium-permeable AMPA receptors (CP-AMPARs) are an important group of receptors for the neurotransmitter glutamate. These receptors contribute to various forms of synaptic plasticity, and alterations in their expression or regulation are also seen in a number of serious neurological conditions, including stroke, motor neuron disease, and cocaine addiction. Several groups of auxiliary transmembrane proteins have been described that enhance the function and cell-surface expression of AMPARs. We now report that the recently identified auxiliary protein GSG1L decreases weighted mean channel conductance and calcium permeability of CP-AMPARs while increasing polyamine-dependent rectification by diminishing outward current. Our experiments reveal that GSG1L is an auxiliary subunit that can markedly suppress CP-AMPAR function, in both recombinant systems and central neurons.

Introduction

Fast excitatory signaling in the mammalian brain relies mainly on AMPA receptors (AMPARs) that mediate rapid point-to-point transmission. These receptors also promote synapse formation and development and are themselves subject to various activitydependent modifications. Thus, they represent a central element in the processes of synaptic plasticity, learning, and memory (Newpher and Ehlers, 2008; Kessels and Malinow, 2009). AM-PARs are homo- and hetero-tetramers assembled from combina-

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tions of the pore-forming subunits GluA1-GluA4 (Traynelis et al., 2010). Although a majority of central AMPARs contain the GluA2 subunit, those lacking GluA2 constitute an important subtype and endow central synapses with distinct functional characteristics. Notably, the absence of GluA2 renders AMPARs permeable to calcium; these receptors also exhibit a high singlechannel conductance and sensitivity to block by endogenous intracellular polyamines, resulting in inwardly rectifying EPSCs. The activity-dependent regulation of calcium-permeable AMPARs (CP-AMPARs) is known to underlie various forms of synaptic plasticity (Liu and Cull-Candy, 2000; Isaac et al., 2007; Clem and Huganir, 2010; Traynelis et al., 2010; Szabo et al., 2012). Furthermore, the dysfunction of CP-AMPARs contributes to deleterious neurological processes, including cell damage after stroke and postsynaptic changes associated with drug abuse (Cull-Candy et al., 2006; Liu and Zukin, 2007; Bellone and Lüscher, 2012).

Auxiliary subunits of the stargazin (transmembrane AMPAR regulatory protein; TARP) and cornichon families enhance synaptic transmission by promoting AMPAR function and forward trafficking (Jackson and Nicoll, 2011; Haering et al., 2014; Constals et al., 2015). For CP-AMPARs, these auxiliary subunits approximately double the average single-channel conductance, slow desensitization and deactivation, and enhance calcium permeability (Tomita et al., 2005; Kott et al., 2009; Soto et al., 2009, 2014; Coombs et al., 2012). They also attenuate intracellular polyamine block across all voltages, most obviously at depolarized potentials (Soto et al., 2007, 2014). The overall effect of these various modifications is to increase the size of the synaptic current and prolong its time course. Given that the conductance of CP-AMPARs is at least twice that of their GluA2-containing counterparts (Swanson et al., 1997; Soto et al., 2009; Coombs et al., 2012), the modulation of CP-AMPARs by auxiliary subunits results in substantial additional charge transfer during synaptic transmission.

GSG1L has been identified recently as an AMPAR auxiliary subunit with some structural similarity to TARPs (Schwenk et al., 2012, 2014; Shanks et al., 2012). Postembedding immunogold electron microscopy has shown that this protein is localized together with GluA2- or GluA4-containing AMPARs in dendritic spines of hippocampal pyramidal neurons, suggesting a possible role in synaptic transmission (Schwenk et al., 2012). Although there was previously no direct evidence that it influences EPSC properties, GSG1L has been shown to resemble TARPs in moderately slowing both deactivation and desensitization of recombinant AMPARs, while increasing the surface expression of GluA2 (Schwenk et al., 2012, 2014; Shanks et al., 2012). However, unlike the TARPs, GSG1L is known to slow AMPAR recovery from desensitization (Schwenk et al., 2012; Shanks et al., 2012), raising the possibility that other differences may exist between GSG1L and previously investigated auxiliary AMPAR subunits.

Here, we find that GSG1L contrasts markedly with other auxiliary AMPAR subunits so far described in decreasing the mean single-channel conductance and calcium permeability of GluA2lacking AMPARs, while greatly suppressing outward current flow in the presence of intracellular polyamines. Consistent with this, transfection of cerebellar stellate cells with GSG1L increased inward rectification of synaptic CP-AMPARs. Moreover, shorthairpin RNA (shRNA)-mediated knockdown of GSG1L in hippocampal neurons accelerated EPSC decay and increased synaptic single-channel conductance, demonstrating that GSG1L can contribute to the functional regulation of synaptic CP-AMPARs in neurons.

Materials and Methods

tsA201 cell culture. Cells were maintained as described previously (Soto et al., 2007) and transfected using Lipofectamine 2000 (Invitrogen). Rat AMPAR subunits (flip isoforms) were transfected either alone or with rat GSG1L. After 4-6 h transfection, cells were washed with PBS, and the medium was replaced. Some transfected cells were maintained in the presence of 50 µM NBQX to limit cytotoxic effects. Electrophysiological recordings were made 24-48 h later. AMPAR subunit cDNAs were a gift from S. Heinemann (Salk Institute, La Jolla, CA) and P. Seeburg (Max Planck Institute, Heidelberg, Germany). GSG1L (pcDNA3.1 vector) was courtesy of B. Fakler (University Freiburg, Freiburg, Germany). AMPARs were subcloned into the pIRES2-GFP vector and GFP used to identify transfected cells. GSG1L was transfected in excess of AMPAR subunits (2:1) to ensure coassembly. Tandem constructs of GSG1L and GluA1 or GluA2(Q) were made by directly linking the N terminus of GSG1L to the C-terminal tail of the GluA subunit, incorporating a 9 aa linker (GluA-GGGGGEFAT-GSG1L; Soto et al., 2014).

Fast agonist application to excised patches. We recorded macroscopic currents at 23-26°C from outside-out patches using an Axopatch 200A amplifier, as described previously (Soto et al., 2007). Records were lowpass filtered at 5 kHz and digitized at 20 kHz using WINWCP software (John Dempster, University of Strathclyde, Glasgow, UK) and analyzed using Igor Pro 6.35 (WaveMetrics) with NeuroMatic 2.8v (http://www. neuromatic.thinkrandom.com/). Pipettes were pulled from thick-walled borosilicate glass (1.5 mm outer diameter, 0.86 mm inner diameter; Harvard Apparatus), coated with Sylgard resin (Dow Corning 184), and fire polished to a final resistance of \sim 4–8 M Ω . The external solution contained the following (in mM): 145 NaCl, 2.5 KCl, 1 CaCl₂, 1 MgCl₂, 10 glucose, and 10 HEPES buffer, pH 7.3 with NaOH. Rapid application of glutamate (10 mM) and, when indicated, cyclothiazide (CTZ; 50 µM; Tocris Bioscience), was achieved by piezoelectric translation of a thetabarrel application tool. The internal (pipette) solution contained the following (in mM): 145 CsCl, 2.5 NaCl, 1 Cs-EGTA, 4 Mg-ATP, 10 HEPES, pH 7.3 with CsOH, and 0.1 spermine tetrahydrochloride (Tocris Bioscience). In some recordings, spermine was omitted and endogenous polyamines were chelated by adding 20 mM Na2ATP to the intracellular solution (Watanabe et al., 1991). To extract channel properties from macroscopic responses, we used nonstationary fluctuation analysis (NSFA) routines written in Igor Pro, as described previously (Soto et al., 2007). The weighted mean single-channel current (i) and the total number of channels (N) were determined by plotting the ensemble variance (σ^2) against mean current (\overline{I}) and fitting with the following equation:

$$\sigma^2 = i\bar{I} - \bar{I}^2/N + \sigma_B^2.$$

Channel openings in the tail of macroscopic patch currents (filtered at 2 kHz) were also analyzed using QuB (version 2.0.0.7; http://www.qub. buffalo.edu/). Clear single-channel events (lasting longer than 2 ms) were selected by eye and preprocessed with piecewise linear baseline correction. For each selected record, an all-point amplitude histogram was generated and fitted with two Gaussians to determine the amplitude of the single-channel current. In each case, the mean amplitude was at least 2.5 times the SD of the background noise (0.1–1 pA). On average, 53 channel events (range, 28–145) were measured from each patch. Current–voltage (*I–V*) relationships were examined from -80 to +100 mV, and rectification index (RI_{+80/-80}) values were calculated as the ratio of the peak current at 80 and -80 mV. For philanthotoxin-433 (PhTx-433; Sigma) block experiments, patches were exposed to glutamate (1 mM) and CTZ (50 μ M) before being switched rapidly to a solution with glutamate, CTZ, and PhTX-433.

Measurement of calcium permeability. This was performed as described previously (Soto et al., 2007). Ramps from -80 to +80 mV were applied in "low" and "high" Ca²⁺ solutions (in mM): low Ca²⁺, 145 NaCl, 2.5 KCl, 1 CaCl₂, 1 MgCl₂, and 10 HEPES, pH 7.3 with NaOH; high Ca²⁺, 30 CaCl₂, 110 *N*-methyl-D-glucamine, 5 HEPES, and 50 sucrose, pH 7.3 with HCl. Control ramps were recorded in each solution and then in the same solutions containing 10 mM glutamate and 50 μ M CTZ. The control *I*–*V* relationships were subtracted from those obtained in agonist. In these experiments, the pipette solution contained either 100 μ M or no

added spermine. As no differences in the reversal potentials were observed and the data were pooled. The relative Ca^{2+} permeability, P_{Ca}/P_{Na} , was determined from the reversal potentials in low and high Ca^{2+} extracellular solutions using the following equation:

$$P_{\rm Ca}/P_{\rm Na} = \frac{a_{\rm Na}}{4a_{\rm Ca}} \left(\exp \frac{\left(2V_{\rm rev\,Ca} - V_{\rm rev\,Na}\right)F}{RT} + \exp \frac{\left(V_{\rm rev\,Ca} - V_{\rm rev\,Na}\right)F}{RT} \right),$$

where a_{Na} and a_{Ca} represent the activities of Na⁺ and Ca²⁺ in the extracellular solutions (108.7 and 16.5, calculated from activity coefficients of 0.75 and 0.55, respectively; Geiger et al., 1995), and *R*, *T*, and *F* have their conventional meanings.

Cerebellar cultures. Primary dissociated cultures were prepared from male and female postnatal day 5 (P5) to P6 C57BL/6 mice (Bats et al., 2012). All procedures for the care and treatment of mice were in accordance with the Animals (Scientific Procedures) Act 1986. After dissociation, the cells in suspension were transfected with GFP or GSG1L:IRES: GFP plasmids by electroporation using the Amaxa nucleofection system and Amaxa mouse neuron nucleofector kit (Lonza). Neurons were then plated on poly-L-lysine-coated glass coverslips and grown in a humidified 5% CO2-containing atmosphere at 37°C in Basal Medium Eagle (Gibco) supplemented with KCl (25 mM final concentration), 10% fetal bovine serum (v/v; Gibco), 20 µg/ml gentamicin, and 2 mM L-glutamine for 3 d. Unlike our previous study (Bats et al., 2012), after 3 d in vitro we replaced the growing medium with a serum-free low-potassium medium to prevent glial cell proliferation and promote synaptic activity [Eagle's Minimum Essential Medium (Gibco) supplemented with 5 mg/ml glucose, 0.1 mg/ml transferrin, 0.025 mg/ml insulin, 2 mM glutamine, and 20 μ g/ml gentamicin]. Recordings were made from stellate cells after 8–11 d in vitro.

Hippocampal cultures. Dissociated hippocampal neurons were prepared from P0-P1 female Sprague Dawley rat pups and isolated as described previously (Beaudoin et al., 2012). Briefly, hippocampi were dissected and then digested with warm papain solution for 1 h at 37°C [20 U/ml papain, 0.5 mM EDTA, and 1 mM L-cysteine (Worthington Biochemical) in Earle's balanced salt solution (EBSS; Sigma)]. Tissue digest was halted with albumin-ovomucoid inhibitor solution (2.7 ml of EBSS plus 300 µl reconstituted inhibitor; Worthington Biochemical). Hippocampal tissue was then triturated with a fire-polished Pasteur pipette and resuspended in basal medium Eagle (Gibco) supplemented with 10% fetal bovine serum (v/v; Gibco), 0.45% glucose (w/v), 1 mM sodium pyruvate, 2 mM GlutaMax (Gibco), and 50 µg/ml penicillin/streptomycin (Gibco). Neurons were plated onto 24-well tissue culture dishes coated with poly-L-lysine at a cell density of $8-10 \times 10^5$ cells per well in neuronal medium. After 4 h, the medium was replaced with serum-free Neurobasal A medium supplemented with B27 (Life Technologies). Cells were grown at 37°C in a humidified environment of 5% CO₂, and approximately half of the medium was exchanged every 3 d. Chronic activity blockade was achieved using a medium supplemented with 1 μ M TTX and 150 µM D-AP5 (Ascent Scientific) from day 6 in vitro. Neurons were transfected using Lipofectamine 2000 (Invitrogen), with shRNA against GSG1L or a scrambled control. Coexpression of GFP was used to identify transfected cells. After 45 min transfection, cells were washed with PBS, and the medium was replaced. Electrophysiological recordings were made 48-72 h later from pyramidal-shaped neurons. shRNA used to knockdown GSG1L and a nonsilencing scrambled control were selected from the Thermo Fisher Scientific Open Biosystems GIPZmir library (V3LHS_348573).

Quantification of GSG1L shRNA efficiency. HEK293 cells were transfected (as described above) with GSG1L and either GSG1L–shRNA or the scrambled control. After 2 d, they were fixed for 10 min in a solution of 4% paraformaldehyde in PBS. Cells were washed three times in PBS and then blocked with 10% goat serum, 0.5% bovine serum albumin, and 0.2% Triton X-100 (all in PBS) for 10 min. Coverslips were incubated in blocking solution with GSG1L-specific antibody (rabbit, 0.5 μ g/ml; catalog #121432; Abcam) for 60 min, washed five times, and then incubated with an Alexa Fluor 647-conjugated goat rabbit-specific secondary antibody (4 μ g/ml; catalog #A21244; Life Technologies) for an additional 45 min. After five additional washes, coverslips were mounted on slides

using Prolong Gold antifade mounting medium (Invitrogen) and imaged using a confocal microscope (Leica) with a 40× oil-immersion objective. We identified isolated cells successfully transfected with shRNA or the scrambled control (expressing GFP) along with GSG1L (Alexa Fluor 647 fluorescence), and, for each cell, we recorded the corrected mean fluorescence intensity (minus background) of the somatic Alexa Fluor 647 signal. In cells that expressed shRNA, GSG1L protein was reduced by 72% compared with cells expressing the scrambled shRNA (25 and 57 cells, respectively, from two independent experiments). Note that this is likely an underestimation of the extent of knockdown, because the selection of HEK293 cells positive for both GSG1L and GFP excluded any cells in which GSG1L expression was eliminated completely by the shRNA.

Neuronal electrophysiology. Cells were viewed using upright or inverted microscopes (BX50WI or IX71; Olympus) and perfused at a rate of 2-3 ml/min (2 ml bath volume). The extracellular solution contained the following (in mM): 145 NaCl, 2.5 KCl, 2 CaCl₂, 1 MgCl₂, 10 glucose, and 10 HEPES, adjusted to pH 7.3 with NaOH. To block voltage-gated sodium channels, NMDA and GABA_{A} receptors, 0.5 $\mu\textsc{m}$ TTX, 20 $\mu\textsc{m}$ D-AP5, and 20 µM SR-95531 [2-(3-carboxypropyl)-3-amino-6-(4-methoxyphenyl)pyridazinium bromide] were included. For recordings from cultured hippocampal neurons, glycine receptors were blocked using 1 µM strychnine (Ascent Scientific). Pipettes were filled with a solution containing the following (in mM): 128 CsCl, 10 HEPES, 10 EGTA, 10 tetraethylammonium (TEA)-Cl, 2 MgATP, 1 CaCl₂, 2 NaCl, 1 QX-314 [N-(2,6-dimethylphenylcarbamoylmethyl) triethylammonium bromide], and 0.1 spermine tetrahydrochloride (both from Tocris Bioscience), pH 7.4 with CsOH (final resistance of \sim 4–8 M Ω). Recordings were made using an Axopatch 200A amplifier, low-pass filtered at 5 kHz, and digitized at 20 kHz using WINEDR software (John Dempster, University of Strathclyde). For cerebellar recordings, the internal solution contained the following (in mM): 140 CsCl, 10 HEPES, 5 EGTA, 2 MgATP, 0.5 CaCl₂, 4 NaCl, and 0.1 mM spermine tetrahydrochloride, pH 7.4 with CsOH. Recordings were made using an Axopatch 200B amplifier, low-pass filtered at 5 kHz, and digitized at 25-50 kHz using Igor Pro (with NeuroMatic NClamp).

mEPSC analysis. We detected mEPSCs using amplitude threshold crossing, with the thresholds typically \sim 10 pA for cultured hippocampal neurons and \sim 6 pA for cultured cerebellar neurons. To calculate the RI of stellate cell mEPSCs, we used a method similar to that described previously (Bats et al., 2012). Because any mEPSCs carried predominantly by CP-AMPARs might fall below the detection threshold at positive potentials as a result of complete (or near-complete) block by intracellular spermine, simply comparing the amplitudes of detected events would lead to an underestimate of rectification. Accordingly, we calculated RI as follows:

$$RI = \frac{g_{mean+60} \times f_{+60}}{g_{mean-80} \times f_{-80}},$$

where g_{mean} is the peak conductance of the mean mEPSC at the given voltage, and f is the frequency of the detected events. This is equivalent to the ratio of summed conductances for recordings of equal length, and, in the absence of rectification, RI would equal 1. When analyzing mEPSC amplitudes, all events with a monotonic rise were included regardless of overlapping decays. For channel conductance (below) and kinetic analyses, only events that showed a monotonic rise and an uncontaminated decay were included. Such events were aligned on their rising phase before averaging. The decay of averaged mEPSCs was described by the sum of two exponentials, from which the weighted time constant of decay ($\tau_{w,decay}$) was calculated as the sum of the fast and slow time constants weighted by their fractional amplitudes. Peak-scaled NSFA (psNSFA) was used to estimate the weighted mean single-channel conductance of synaptic receptors (Traynelis et al., 1993). Each mEPSC was divided into 80 bins of equal amplitude, and the variance of the mEPSC about the scaled average in each bin was computed. The variance was plotted against the mean current, and the weighted mean single-channel current was estimated by fitting the full parabolic relationship with the following equation:



Figure 1. GSG1L attenuates single-channel conductance and calcium permeability of homomeric AMPARs. *a*, Individual currents (gray) and averages (black) evoked by glutamate (10 mM, 100 ms) applied to patches (-60 mV) from cells expressing GluA2(Q), with and without GSG1L. Current–variance relationships show estimated weighted mean single-channel conductances for illustrated patches. Symbols show mean and error bars SEM. Dashed lines denote baseline variance. *b*, Pooled data from NSFA. Box-and-whisker plots indicate median (line), 25th to 75th percentiles (box), range of data within 1.5 × interquartile range of box (whiskers) and mean (open circles). *p < 0.05, ***p < 0.001 (unpaired Welch two-sample *t* tests, n = 7-14). *c*, Resolved single-channel openings in the tail of macroscopic currents (truncated; 2 kHz filter; left). All-point amplitude histograms from indicated individual channel events (middle). Cumulative probability distributions for conductances from four GluA2(Q) patches and three GluA2(Q)/GSG1L patches (right). Averaged distributions (bold lines) and SEM (fills). *d*, Representative *I*–*V* plots for GluA2(Q)–GSG1L tandem constructs (voltage ramp in presence of 1 mm glutamate plus 50 μ m CTZ; no internal spermine) in low (1 mm) and high (30 mm) Ca²⁺. Enlargement shows reversal potential shift. Box-and-whisker plots (as in *b*) show the effect of GSG1L on P_{Ga}/P_{Na} (***p < 0.001, Welch *t* test, n = 7 and 9).

$$\sigma_{\rm PS}^2 = i\bar{I} - \bar{I}^2/N_{\rm p} + \sigma_{\rm B}^2$$

where $\sigma_{\rm PS}^2$ is the peak-scaled variance, \bar{I} is the mean current, *i* is the weighted mean single-channel current, $N_{\rm p}$ is the number of channels open at the peak of the EPSC, and $\sigma_{\rm B}^2$ is the background variance.

Statistical analysis. Summary data are presented in the text as mean \pm SEM from n cells. Comparisons involving two datasets only were performed using a two-sided Welch two-sample t test that does not assume equal variance (normality was not tested statistically but judged from Q-Q plots and/or density histograms). Analyses involving data from three or more groups were performed using two-way ANOVA (Welch heteroscedastic F test), followed by pairwise comparisons using twosided Welch two-sample t tests. For PhTx-433 concentration-response curves fit using a four-parameter logistic equation (sigmoidal dose response with variable slope), best-fit values for log EC_{50} and maximum were compared using extra sum-of-squares F tests. Exact p values are presented to two decimal places, except when p < 0.0001. Differences were considered significant at p < 0.05. Statistical tests were performed using Prism version 6 (GraphPad Software) or R (version 3.1.2; the R Foundation for Statistical Computing; http://www.r-project.org/) and R Studio (version 0.98.1103; RStudio). No statistical test was used to predetermine sample sizes; these were based on standards of the field. No blinding or randomization was used.

Results

GSG1L reduces mean single-channel conductance and calcium permeability of CP-AMPARs

We initially examined the ability of GSG1L to regulate CP-AMPARs by recording currents activated by fast application of glutamate to outside-out membrane patches from tsA201 cells expressing GluA2(Q), GluA1, or GluA4, together with GSG1L (Fig. 1*a*; see Materials and Methods). NSFA revealed that, in all cases, GSG1L reduced the weighted mean single-channel conductance of CP-AMPARs by ~35% (Fig. 1*b*). Two-way ANOVA showed a main effect of AMPAR subtype ($F_{(2,51)} = 5.45$, p = 0.0072), a main effect of GSG1L ($F_{(1,51)} = 34.21$, p < 0.0001), and no interaction ($F_{(2,51)} = 0.20$, p = 0.82; n = 7-14). Mean singlechannel conductance was reduced from 19.9 ± 1.1 to 13.1 ± 1.3 pS for GluA2(Q) (n = 14 and 10; $t_{(20.48)} = 4.34$, p = 0.00030), from 16.9 ± 1.7 to 11.4 ± 0.4 pS for GluA1 (n = 7 and 8; $t_{(6.74)} =$ 3.22, p = 0.015), and from 21.2 ± 2.0 to 14.3 ± 0.9 pS for GluA4 (both n = 9; $t_{(11,15)} = 3.09$, p = 0.010).

This reduction in the mean single-channel conductance estimated by NSFA was mirrored by a decrease in mean amplitude of directly resolved single-channel events in the tail of glutamate-evoked macroscopic currents (Fig. 1*c*). Mean channel conductance was reduced from 24.0 ± 2.0 to 13.7 ± 0.7 pS (n = 4 and 3, respectively; $t_{(3.63)} = 4.87$, p = 0.010). Because recombinant AMPARs typically open to three or four closely spaced subconductance levels in the absence of auxiliary subunits (Swanson et al. 1997; Jin et al., 2003; Zhang et al., 2008; Prieto and Wollmuth, 2010), the effect of GSG1L could reflect an alteration in the relative probability of openings to these preexisting levels (Tomita et al. 2005). Alternatively, it may reflect a change in the absolute conductance levels (Shelley et al., 2012). We have not attempted to distinguish between these possibilities.

To determine the effect of GSG1L on relative Ca²⁺ permeability (P_{Ca}/P_{Na}), we compared the reversal potentials of glutamateactivated currents in solutions containing low or high external Ca²⁺ (1 or 30 mM; Fig. 1*d*). Although a change from low to high Ca²⁺ had little effect on GluA2(Q) homomers, it shifted the reversal potential of GSG1L-containing receptors to more negative values. From the



Figure 2. GSG1L enhances CP-AMPAR block by intracellular spermine. *a*, Representative glutamate-evoked currents (10 m/m, 100 ms) and normalized peak /–V relationships (global average; n = 6 and 8), showing increased rectification at positive potentials in the presence of GSG1L (100 μ M spermine). Error bars denote SEM, and curves are weighted fits of seventh-order polynomials. *b*, Box-and-whisker plots (as in Fig. 1) showing pooled RI data. ***p < 0.001 (unpaired Welch two-sample *t* tests, n = 6-12). *c*, Normalized *I*–*V* relationships for homomeric GluA2(Q) receptors with and without GSG1L in the absence of spermine (n = 5 and 4, respectively; symbols as in *a*). Vertical error bars denote SEM, and fitted lines are fourth-order polynomials. Inset shows a family of GluA2(Q)/GSG1L currents (-80 to +80 mV) in response to application of 10 m/m glutamate. Right, Normalized conductance against voltage for the data shown in the *I*–*V* relationships. Fitted lines are drawn according to $G = G_{min} + (G_0 - G_{min})\exp(V_m/V_c)$, where G_0 is the conductance at 0 mV, G_{min} is the minimal conductance (normalized value of 1), and V_c is a constant. Without GSG1L, G_0 was 1.11, and V_c was 36.9 mV. With GSG1L, the corresponding values were 1.20 and 65.7 mV. *d*, As in *a*, but for heteromeric GluA1/GluA2(R) in absence and presence of GSG1L (as tandem). Note the unchanged near-linear peak /–V relationship (third-order polynomial fits; n = 5 and 3).

reversal potential shift, we estimated that GSG1L reduced P_{Ca}/P_{Na} by ~40% (from 3.9 ± 0.2 to 2.2 ± 0.2; n = 7 and 9, respectively; $t_{(12.83)} = 7.35$, p < 0.0001). These actions of GSG1L on mean channel conductance and relative calcium permeability contrast markedly with those of both TARPs and cornichons.

GSG1L enhances spermine-dependent rectification of CP-AMPARs

The voltage-dependent block by intracellular polyamines such as spermine, giving rise to inward rectification, has been used widely to identify the presence of synaptic CP-AMPARs (Liu and Cull-Candy, 2000; Noh et al., 2005; Bats et al., 2012). To determine whether, like TARPs (Soto et al., 2007), GSG1L reduces CP-AMPAR rectification, we constructed I-V plots for currents obtained from outside-out patches in response to rapid application of glutamate. As expected, homomeric CP-AMPARs displayed inwardly rectifying I-V plots in the presence of added intracellular spermine (100 μm; Fig. 2a). Strikingly, in the presence of GSG1L, the current at positive potentials was reduced, thus the rectification was enhanced rather than diminished (Fig. 2a,b). Two-way ANOVA showed no main effect of AMPAR subtype on the rectification index (RI_{+80/-80}; $F_{(3,61)} = 0.23$, p =0.87), a main effect of GSG1L ($F_{(1,61)} = 169.40, p < 0.0001$), and no interaction ($F_{(3,61)} = 1.37$, p = 0.26; n = 6-12). RI_{+80/-80} was reduced from 0.21 ± 0.01 to 0.03 ± 0.01 for GluA2(Q) (n = 7 and 9; $t_{(11.25)} = 10.37, p < 0.0001$), from 0.14 ± 0.01 to 0.04 ± 0.01 for GluA1 (n = 12 and 11; $t_{(11.72)} = 6.17$, p < 0.0001), from 0.18 \pm 0.03 to 0.05 ± 0.01 for GluA3 (n = 6 and 7; $t_{(5.75)} = 5.23$, p = 0.0022), and from 0.18 \pm 0.03 to 0.04 \pm 0.01 for GluA4 (n = 6 and 10; $t_{(5,19)} =$ 5.05, p = 0.0036).



Figure 3. GSG1L does not shift voltage for half-maximal block of CP-AMPARs by intracellular spermine. *a*, Plots of normalized conductance against voltage for the data shown in Figure 2*a* (corrected for the outward rectification observed in the absence of spermine; Fig. 2*c*). Lines are fits at negative voltages to a Boltzmann function, $G = G_{max} \left(\frac{1}{1 + \exp(V_m - V_{1/2}/k)} \right)$, where G_{max} is the maximal glutamate-activated conductance at hyperpolarized voltages, $V_{1/2}$ is the voltage at which spermine block is half-maximal, and *k* is a slope factor describing the membrane potential shift necessary to cause an *e*-fold change in conductance. For the fits of the pooled data, the $V_{1/2}$ and slope values were similar with and without GSG1L: -58.0 mV and 10.3 for GluA2(Q); -56.0 mV and 12.2 for GluA2(Q)/GSG1L *b*, Box-and-whisker plots (as in Fig. 1) showing pooled $V_{1/2}$ and *k* values from Boltzmann fits to *G*-*V* plots for individual patches: n = 6 GluA2(Q) and 9 GluA2(Q)/GSG1L patches.

To decide whether the effect of GSG1L on rectification was mediated by a modulation of polyamine action rather than a change in the intrinsic properties of AMPAR gating, we examined the I-Vrelationship of GluA2(Q), with and without GSG1L, in the absence of intracellular polyamines. Recordings were obtained a minimum of 15 min after patch excision to allow complete chelation of endogenous polyamines by the added Na_2ATP (see Materials and Methods). The control *I–V* relationship and that obtained with GSG1L both showed clear outward rectification and were overlapping (Fig. 2c). This result suggests that the effect of GSG1L on inward rectification was entirely polyamine dependent.

Heteromeric GluA2(R)-containing AMPARs are unaffected by intracellular polyamines. Accordingly, GSG1L might be expected to have no effect on their normally linear I-V relationship. To test this, we examined the effect of GSG1L on Ca²⁺impermeable GluA1/GluA2(R) (Fig. 2d). In this case, we compared coexpressed GluA1 and GluA2(R) with coexpressed tandem GluA1-GSG1L and GluA2(R) (see Materials and Methods). Consistent with our results for CP-AMPARs, we found that I-Vplots from GluA2(R)-containing calciumimpermeable (CI)-AMPARs remained unchanged in the presence of GSG1L. Similar results were obtained with GluA4/ GluA2(R) receptors (data not shown).

Although GSG1L increased rectification as assessed using the measure of RI, we noted that the influence of GSG1L was apparent only at positive voltages. To examine this more closely, we next plotted conductance (G) against voltage. As predicted from the overlap of the I-V plots at negative voltages, Boltzmann fits to G-V plots revealed that GSG1L produced no significant shift in the voltage for half-maximal block by spermine $[V_{1/2}; -64.8 \pm 6.3 \text{ mV vs}]$ -67.9 ± 5.3 mV; n = 6 for GluA2(Q) and 9 for GluA2(Q)/GSG1L; $t_{(11,01)} =$ -0.38, p = 0.71] or in the slope factor $(k; 13.8 \pm 1.1 \text{ vs } 16.6 \pm 0.6; t_{(7.89)} =$ -2.24, p = 0.056; Fig. 3). This suggests that GSG1L primarily hinders the voltage-dependent relief of spermine block seen normally at positive potentials, which is thought to reflect permeation of the polyamine (Bowie and Mayer, 1995).

PhTx-433 block of CP-AMPARs is attenuated by GSG1L

Like polyamine-dependent inward rectification, the use-dependent block produced by extracellular application of the

polyamine toxin PhTx-433 (Jackson et al., 2011) is generally considered a defining feature of CP-AMPARs (Liu and Cull-Candy, 2000; Noh et al., 2005; Bats et al., 2012; but see Bowie, 2012). Although the effects of this toxin on glutamate-activated CP-AMPAR-mediated currents are unaffected by TARPs (Jackson et al., 2011), we found that GSG1L attenuated PhTx-433



Figure 4. GSG1L reduces CP-AMPAR block by PhTx-433. *a*, Representative glutamate-evoked patch currents (-60 mV; 50 μ M (TZ) showing block after rapid jump into solution containing 100 nM PhTx-433 (0.2 kHz filter, for display). Steady-state block (denoted by vertical colored lines) produced by extracellular PhTx-433 (gray shading) is attenuated by GSG1L. *b*, Concentration-response curves (mean \pm SEM; n = 3-7 cells) show that GSG1L reduces PhTx-433 potency (log EC₅₀) and maximum block. Solid lines are fits to the Hill equation, giving the following values for GluA4 and GluA4 plus GSG1L: maximum block, 94.8 and 84.1%; EC₅₀, 5.9 nM and 19.9 nM; Hill slope, 1.2 and 1.2.



Figure 5. GSG1L expression alters mEPSC rectification in cultured cerebellar neurons. *a*, Representative recordings of mEPSCs from a cultured stellate cell (control cell; GFP transfected) at -80 and +60 mV. Traces are filtered at 2 kHz for display. Middle, Averaged mEPSCs. Right, "Composite" mEPSCs showing the average waveforms scaled by the frequency of events at each voltage together with the value of $RI_{+60/-80}$ calculated as described in Materials and Methods. *b*, As in *a*, but for a representative GSG1L-transfected cell. *c*, Pooled data showing the peak mEPSC amplitudes at -80 and +60 mV. *d*, Pooled data showing the peak conductance ratios and frequency ratios at -80 mV and +60 mV. *e*, Pooled data showing the increased rectification with GSG1L. Box-and-whisker plots as in Figure 1. *p < 0.05, **p < 0.01, unpaired Welch *t* test, n = 7 and 8.

block of GluA4 CP-AMPARs (Fig. 4). Analysis of PhTX-433 concentration-response curves showed that GSG1L reduced both PhTx-433 potency (EC₅₀ reduced from 5.9 to 19.9 nM; $F_{(1,31)} = 27.47$, p < 0.0001) and the maximum block (from 94.8 to 84.1%; $F_{(1,31)} = 5.92$, p = 0.029; extra sum-of-squares *F* test).



Figure 6. GSG1L knockdown alters properties of mEPSCs in cultured hippocampal neurons. *a*, Representative recordings of mEPSCs from cultured hippocampal neurons (-60 mV) treated with TTX and p-AP5 to promote the expression of synaptic CP-AMPARs (see Materials and Methods). Traces are from cells expressing scrambled shRNA or shRNA against GSG1L. *b*, Averaged mEPSCs from cells shown in *a. c*, Averaged probability density histograms of mEPSC peak amplitude from six control cells (scrambled shRNA; 235–2011 mEPSCs per cell) and seven GSG1L–shRNA cells (66-1496 mEPSCs per cell). Solid lines are lognormal fits, with parameters μ and σ being 0.36 and 3.31 for control and 0.46 and 3.67 for shRNA–GSG1L. *d*, Pooled data showing the effect of GSG1L knockdown on mEPSC amplitude, mEPSC decay (weighted time constant $\tau_{w,decay}$), and synaptic channel conductance (from psNSFA). Box-and-whisker plots as in Figure 1. *p < 0.05, ***p < 0.001, unpaired Welch *t* test, n = 5-8.

GSG1L modifies the properties of neuronal CP-AMPARs

We next examined whether GSG1L could similarly modulate synaptic CP-AMPARs. Cerebellar stellate cells are known to express a mixture of CI- and CP-AMPARs (Liu and Cull-Candy, 2000). These cells express TARPs (Fukaya et al., 2005) but conveniently lack GSG1L (Lein et al., 2007), allowing us to examine the effect of GSG1L on native AMPARs. We recorded mEPSCs at -80 and +60 mV from cultured stellate cells transfected with GFP (control) or GSG1L (Fig. 5a,b). In our culture conditions, mEPSCs from control cells showed moderate inward rectification, consistent with the presence of CP-AMPARs. Transfection with GSG1L had no effect on mEPSC amplitude at -80 mV $(-43.6 \pm 7.7 \text{ vs} - 35.5 \pm 4.3 \text{ pA}, n = 7 \text{ and } 8 \text{ for GFP and GSG1L}$ transfected cells, respectively; $t_{(9.52)} = 0.94$, p = 0.37) but reduced the average amplitude at +60 mV (from 32.7 \pm 3.3 to 22.1 \pm 3.2 pA; $t_{(9.52)} = 0.94$, p = 0.037; Fig. 5c). However, because the AMPAR number and subtype may vary from one synapse to another within an individual cell, those mEPSCs carried primarily by CP-AMPARs might be undetectable at positive potentials because of complete (or near-complete) block by intracellular spermine. Thus, both the amplitude and frequency of mEPSCs at +60 mV could be affected by spermine such that a simple comparison of the amplitudes of the detected events would lead to an underestimate of rectification. In GSG1L-transfected cells, although the ratio of mEPSC peak conductances (+60/-80 mV)was not reduced significantly ($t_{(11.14)} = 1.60, p = 0.14$), the ratio of frequencies was changed ($t_{(12.60)} = 3.65, p = 0.0031$; Fig. 5*d*). Accordingly, $\mathrm{RI}_{+60/-80}$ (see Materials and Methods) was reduced from 0.62 \pm 0.08 to 0.24 \pm 0.07 (n = 7 and 8; $t_{(12.13)}$ = 3.69, P = 0.0031; Fig. 5e). Of note, when we transfected GSG1L into cerebellar granule cells, in which synaptic currents are mediated solely by CI-AMPARs, mEPSC frequency, amplitude, and RI was unchanged

(data not shown). The failure of GSG1L to promote the expression of synaptic CP-AMPARs in granule cells suggests that the increased rectification we observed in stellate cells, rather than reflecting an increased proportion of synaptic CP-AMPARs (or a decreased proportion of CI-AMPARs), likely reflects the reduced relief of spermine block at existing CP-AMPARs.

To establish the effects in neurons of endogenous GSG1L (Schwenk et al., 2012; Shanks et al., 2012), we next examined cultured hippocampal neurons treated with shRNA against GSG1L (see Materials and Methods; Fig. 6a,b). Although a majority of the AMPARs in hippocampal CA1 pyramidal cells are thought to be calcium impermeable, the expression of CP-AMPARs in these cells has been identified clearly (Rozov et al., 2012; Mattison et al., 2014). Furthermore, CP-AMPAR expression is known to increase in cultured hippocampal pyramidal neurons after blockade of synaptic activity (Ju et al., 2004). Therefore, to increase the prevalence of CP-AMPARs in our cells, we maintained them in a condition of reduced activity (1 µM TTX, 150 µM D-AP5) and confirmed the presence of inwardly rectifying mEPSCs (data not shown). In pyramidal-shaped cells, knockdown of

GSG1L led to an acceleration of mEPSC decay, from 2.9 ± 0.3 to 2.1 ± 0.1 ms (Fig. 6*b*,*d*). Because the deactivation of recombinant GluA1/GluA2(R) receptors is known to be prolonged in the presence of GSG1L (Schwenk et al., 2012), this could reflect an effect on CI-AMPARs, or CP-AMPARs, or both.

GSG1L knockdown also resulted in an increase in mEPSC amplitude (Fig. 6b), which was accompanied by a broadening of the amplitude distribution, suggesting a heterogeneity across synapses (Fig. 6c). On average, mEPSC amplitude was increased by ~80%, from -28.3 ± 3.2 to -52.0 ± 7.6 pA (n = 6 and 7; $t_{(7.98)} = -2.89, p = 0.020$; Fig. 6d). This was accompanied by a proportional increase in the mean synaptic single-channel conductance estimated from psNSFA (see Materials and Methods) from 29.6 \pm 5.1 to 51.4 \pm 3.5 pS (n = 5 and 8; $t_{(10.80)} = -5.24$, p = 0.0012; Fig. 6*d*). The estimated number of channels activated during a quantal event was not changed after GSG1L knockdown $(34 \pm 4 \text{ vs } 30 \pm 5)$, consistent with the view that the increased mEPSC amplitude reflected an increase in mean single-channel conductance associated with loss of GSG1L. The combined effects of GSG1L knockdown could reflect changes in both CI- and CP-AMPAR properties. However, in studies of recombinant receptors, we found GSG1L to have no effect of the mean channel conductance of heteromeric, GluA1-GSG1L/GluA2(R) CI-AMPARs (n = 6 and 7; $t_{(9.95)} = -0.26$, p = 0.79). This contrasts with the effect of GSG1L on the mean channel conductance of homomeric CP-AMPARs (Fig. 1b) and suggests that synaptic CP-AMPARs are altered after GSG1L knockdown.

Discussion

In this study, we report novel features of AMPAR modulation by GSG1L. Our focus was on CP-AMPARs. We found that GSG1L

suppressed recombinant CP-AMPAR function by decreasing average single-channel conductance, decreasing calcium permeability, and inhibiting the normal voltage-dependent relief of intracellular polyamine block that occurs at positive potentials. Outward current flow was thus greatly diminished. Although GSG1L has been shown previously to slow deactivation and desensitization of CP-AMPARs (Shanks et al., 2012) in a manner similar to that of both type 1 TARPs (γ -2, γ -3, γ -4, and γ -8) and cornichons (CNIH2/3), its combined effects differ fundamentally from those seen with these canonical AMPAR auxiliary subunits. Importantly, the influence of GSG1L on CP-AMPAR channel conductance and calcium permeability are opposite to those of both TARPs and cornichons. Furthermore, its effects on polyamine block appear unconventional. With TARPed AMPARs, in the presence of intracellular polyamines, outward current is detected at approximately +30 mV (Soto et al., 2007), likely reflecting polyamine permeation and relief of block (Bowie et al., 1998). In contrast, in the presence of GSG1L, outward current is greatly decreased even at very depolarized potentials, implying that polyamine permeation is inhibited. Conversely, block at negative voltages appears unaffected. Overall, our study suggests that auxiliary subunits can act to suppress and enhance current flow through AMPARs.

Consistent with our data on recombinant AMPARs, we find that overexpression of GSG1L in cerebellar interneurons that express synaptic CP-AMPARs results in an increase in mEPSC rectification. Moreover, we find that knockdown of GSG1L in cultured hippocampal neurons increases the peak amplitude of mEPSCs and enhances the conductance of the underlying channels, suggesting that, in these cells, GSG1L regulates the properties of synaptic AMPARs. Although it is possible that the influence of GSG1L on CP-AMPARs may vary in other brain areas, because of its presence together with different subsets of auxiliary subunits (Schwenk et al., 2014), this finding emphasizes the contrast with canonical auxiliary subunits, the genetic disruption of which typically results in a reduction of AMPAR expression and activity (Hashimoto et al., 1999; Rouach et al., 2005; Fukaya et al., 2006; Bats et al., 2012). It will be interesting to determine whether suppressive effects of GSG1L on native CP-AMPARs involve direct modulation of channel function or whether its effects are indirect and result from negation of the actions of other auxiliary subunits. The fact that we observed clear suppressive effects of GSG1L with heterologously expressed CP-AMPARs in the absence of TARPs or CNIHs indicates that a direct effect can occur.

An additional difference between GSG1L and other auxiliary subunits is seen in its effects on extracellular block by the polyamine toxin PhTx-433 (Jackson et al., 2011). This toxin has been used widely to identify the presence of neuronal CP-AMPARs (Isaac et al., 2007). Whereas the blocking action of PhTx-433 on glutamate-gated currents is unaffected by TARPs γ -2 or γ -8 (Jackson et al., 2011), its ability to block, and hence identify, CP-AMPARs is diminished by GSG1L. This observation raises the intriguing question as to whether the presence of an auxiliary subunit with GSG1L-like properties could account for reports of CP-AMPARs in cortical oligodendrocyte lineage cells and retinal amacrine cells that exhibit a relatively low sensitivity to block by polyamine toxins (Meucci et al., 1996; Osswald et al., 2007; Diamond, 2011).

Calcium entry via CP-AMPARs has been implicated in the induction of synaptic plasticity (Gu et al., 1996; Mahanty and Sah, 1998; Laezza et al., 1999; Liu and Cull-Candy, 2000) and

neuronal damage (Carriedo et al., 1998; Noh et al., 2005). Moreover, the interaction of intracellular polyamines with CP-AMPARs contributes to short-term plasticity (Rozov and Burnashev, 1999; Shin et al., 2005). Here we show that GSG1L attenuates the calcium permeability and increases intracellular polyamine block of CP-AMPARs. These effects, together with the marked slowing of recovery from desensitization of homomeric CP-AMPARs (Shanks et al., 2012), highlight the potential importance of regulated GSG1L expression. Specifically, the regulation of its expression could provide a mechanism for the dynamic control of CP-AMPAR function at a cellular level. In this context, GSG1L could mitigate damaging CP-AMPAR-mediated calcium entry in certain pathological conditions.

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