frontiers in IMMUNOLOGY



Dickkopf-3, a tissue-derived modulator of local T-cell responses

Michael Meister¹, Maria Papatriantafyllou¹, Viola Nordström², Varun Kumar³, Julia Ludwig¹, Kathy O. Lui⁴, Ashleigh S. Boyd⁴, Zoran V. Popovic^{2,5}, Thomas Henry Fleming³, Gerhard Moldenhauer¹, Peter P. Nawroth³, Hermann-Josef Gröne², Herman Waldmann⁴, Thilo Oelert¹ and Bernd Arnold¹*

¹ Department of Molecular Immunology, German Cancer Research Center, Heidelberg, Germany

² Department of Molecular Pathology, German Cancer Research Center, Heidelberg, Germany

³ Department of Medicine I and Clinical Chemistry, University of Heidelberg, Heidelberg, Germany

⁴ Therapeutic Immunology Group, Sir William Dunn School of Pathology, University of Oxford, Oxford, UK

⁵ Department of Pathology, University Medical Center Mannheim, University of Heidelberg, Mannheim, Germany

Edited by:

Shohei Hori, RIKEN, Japan

Reviewed by:

Alain Le Moine, Université Libre de Bruxelles, Belgium Dennis O. Adeegbe, Dana Farber Cancer Institute and Harvard Medical School, USA

*Correspondence:

Bernd Arnold, Department of Molecular Immunology, German Cancer Research Center (DKFZ), Im Neuenheimer Feld 280, 69120 Heidelberg, Germany e-mail: b.arnold@dkfz.de The adaptive immune system protects organisms from harmful environmental insults. In parallel, regulatory mechanisms control immune responses in order to assure preservation of organ integrity. Yet, molecules involved in the control of T-cell responses in peripheral tissues are poorly characterized. Here, we investigated the function of Dickkopf-3 in the modulation of local T-cell reactivity. Dkk3 is a secreted, mainly tissue-derived protein with highest expression in organs considered as immune-privileged such as the eye, embryo, placenta, and brain. While T-cell development and activation status in naïve Dkk3-deficient mice was comparable to littermate controls, we found that Dkk3 contributes to the immunosuppressive microenvironment that protects transplanted, class-I mismatched embryoid bodies from T-cell-mediated rejection. Moreover, genetic deletion or antibody-mediated neutralization of Dkk3 led to an exacerbated experimental autoimmune encephalomyelitis (EAE). This phenotype was accompanied by a change of T-cell polarization displayed by an increase of IFNy-producing T cells within the central nervous system. In the wildtype situation, Dkk3 expression in the brain was up-regulated during the course of EAE in an IFNy-dependent manner. In turn, Dkk3 decreased IFNy activity and served as part of a negative feedback mechanism. Thus, our findings suggest that Dkk3 functions as a tissue-derived modulator of local CD4⁺ and CD8⁺ T-cell responses.

Keywords: Dickkopf-3, T cells, immune privilege, transplantation, autoimmunity

INTRODUCTION

Adaptive immune responses in vital organs have to be precisely coordinated in order to assure proper pathogen clearance and at the same time, to prevent extended immune-mediated inflammation that might impair their function. In particular, this is essential for tissues with limited regenerative capacity such as the central nervous system (CNS), the eye, and the pregnant uterus. In these organs, extensive inflammation is prevented locally by mechanisms such as immune deviation away from destructive immune responses or by active suppression of inflammation.

Besides the self-regulating capacity of the immune system, tissue-resident cells also play an important role in the surveillance of T-cell responses by production of immunosuppressive molecules. For example, programed death-ligand 1 (PD-L1), FasL, and indoleamine 2,3-dioxygenase (IDO) have already been reported to be expressed in peripheral tissues associated with an immuno-suppressive micromilieu. PD-L1, a mediator of peripheral T-cell tolerance, is expressed by endothelial cells in the heart, muscle cells, β -cells in the pancreas, and glial cells in the inflamed brain (1–5). Via interaction with its receptor, PD-L1 induces co-inhibitory signals in activated T cells and promotes T-cell apoptosis, anergy, and functional exhaustion (6, 7). Furthermore, FasL, which induces

apoptosis of activated T cells, is expressed by neurons, astrocytes, oligodendrocytes, microglia (8), the stroma cells of the retina (9), and on fetal cytotrophoblasts and maternal decidual cells of the placenta (10). Moreover, trophoblasts were shown to produce the enzyme IDO, which catabolizes the amino acid tryptophan, necessary for T-cell survival. The presence of IDO in the trophoblast results in inactivation and apoptosis of T lymphocytes through tryptophan deprivation (11). IDO is also produced by several cell types in the eye (12, 13) and in the brain (14).

Dickkopf-3 (*Dkk3*) belongs to an evolutionary conserved small gene family, which encodes five secreted glycoproteins, Dkk1-4, and Dkk11 that share two conserved cysteine-rich domains (15). While Dkk1, 2, and 4 have been shown to induce head formation during embryogenesis via modulation of the Wnt-pathway, Dkk3 is not involved in head development (15, 16). There is though, increasing evidence that it may also interfere with Wnt signaling (17–19). Expression studies revealed highest Dkk3 levels in tissues such as brain, spinal cord, eye, embryo, ovary, and uterus (15, 17).

In a T-cell receptor (TCR)–MHC class-I double transgenic mouse system, we previously identified Dkk3 as a modulator of CD8 T-cell responses. Under these conditions, the induction of

antigen-specific, peripheral CD8 T-cell tolerance was dependent on the presence of Dkk3 (20). However, the immunological function of Dickkopf-3 in the physiological situation of a polyclonal T-cell repertoire still remains unclear. Here, we show that Dkk3 functions as a tissue-derived modulator of T-cell responses that locally controls activation and differentiation of T lymphocytes. Our findings provide strong evidence that Dkk3 contributes to the immunoregulatory properties of the tissue microenvironment and therefore may provide a valuable target for immune intervention in autoimmunity, transplantation, and inflammatory disease.

MATERIALS AND METHODS

MICE

C57BL/6N and IFN γ R1^{-/-} were obtained from Jackson Laboratories. *Dkk3*^{-/-} mice were provided by Niehrs (21) and backcrossed on C57BL/6N background for more than 10 times. All mice were kept under specific pathogen-free conditions (SPF) at the animal facility of the German Cancer Research Center and used at the age of 8–14 weeks. Animal care was in concordance with the instructions of the regulating authorities.

FLOW CYTOMETRY

Surface-antigen staining was performed according to standard protocols using the following antibodies: CD3-PerCpCy5.5 (BioLegend), CD4-PB (BioLegend), CD8-APC-Cy7 (eBioscience), CD25-PE (BD Bioscience), CD44-PeCy7 (BioLegend), CD62L-APC (eBioscience), anti-TCR-Vβ antibodies (BD Bioscience). For intracellular cytokine staining, 1×10^6 lymphocytes were restimulated with 50 µg/ml myelin oligodendrocyte-derived glycoprotein (MOG)₃₃₋₅₅ peptide in the presence of Golgi Plug (BD Bioscience) for 6 h. The staining was performed with Cytofix/Cytoperm reagents (BD Bioscience) according to the manufacturer's protocol. The following antibodies were used: IL-17-APC (BD Biosciences), GM-CSF-APC (BD Biosciences), IFNy-APC (BD Biosciences). Intracellular transcription factor staining was performed using a FoxP3 staining buffer set (eBioscience) and a FoxP3-APC antibody (eBioscience) according to the manufacturer's protocol. Flow cytometry was performed on a FACSCanto (BD Bioscience) with FACS Diva software (BD Bioscience) and analyzed with FlowJo 9.6 (Treestar).

EMBRYOID BODY GENERATION AND TRANSPLANTATION

Embryoid bodies (EBs) were generated for transplantation as described previously (22). In brief, ES cell lines were cultured on mitotically inactivated embryonic fibroblasts. Before differentiation, ES cells were passaged on gelatinized flasks in the presence of leukemia-inhibitory factor to maintain their pluripotency. After two passages in the absence of fibroblasts, single-cell suspensions were plated onto dishes of bacteriological plastic to generate EB ready for transplantation at day 14 of culture. Transplantation of EB was carried out as described previously (22). Briefly, two to four EB were grafted under the kidney capsule, and groups of mice were killed for analysis at specified time points. Acceptance of the grafted tissues was defined as an increase in diameter to >5 mm, vascularization, and the lack of leukocyte infiltrate and tissue damage, revealed by histology.

EXPERIMENTAL AUTOIMMUNE ENCEPHALOMYELITIS

Experimental autoimmune encephalomyelitis (EAE) was induced by injecting 200 μ g MOG_{33–55} peptide emulsified in 100 μ l complete Freund's adjuvant containing 200 μ g *Mycobacterium tuberculosis* (Thomas Geyer) s.c. At days 0 and 2 after immunization, each mouse received 200 ng pertussis toxin i.p. (Merck Biosciences). Clinical symptoms were scored as follows: 0, normal; 1, limp tail or hind limb weakness; 2, limp tail and hind limb weakness; 3, partial hind limp paralysis; 4, complete hind limb paralysis; 5, dead or moribund, killed by investigator.

IMMUNOHISTOLOGY

Mice were heart-perfused and CNS tissues were fixed with phosphate-buffered 4% formaldehyde. Three-micrometer paraffin sections were de-paraffinized and rehydrated, before staining. For immunohistochemistry, the TSA-Indirect Kit was used (NEN Life Science Products). For fluorescence microscopy, sections were stained with anti-Dkk3, anti-NeuN (Millipore), or anti-GFAP (Millipore) antibodies. As a standard negative control, anti-Dkk3 was substituted by equal amounts of normal mouse IgG (Santa Cruz Biotechnology). Pictures were generated on a cell observer microscope (Zeiss).

ISOLATION OF LYMPHOCYTES FROM CNS

Experimental autoimmune encephalomyelitis-diseased Dkk3^{-/-} and wild-type (WT) mice were heart-perfused with PBS. Brain and spinal cord was removed and minced in ice-cold PBS with 7% FCS. Remaining pieces were digested in 2.5 mg/ml collagenase D (Roche) and 1 mg/ml DNAse I (Roche) in PBS for 30 min at 37°C and three times mashed through a 40- μ m nylon sieve (Falcon). Finally, lymphocytes were isolated by a Percoll gradient (GE Healthcare).

qRT-PCR

Total RNA was extracted from skin using a FastPrep tissue homogenizer (ThermoScan) and the RNeasy kit (Qiagen). Purified RNA was reverse transcribed using the SuperScript II kit (Invitrogen). Quantitative real-time PCR was performed on a 7500 RT-PCR System (Applied Biosystems) using Absolute qPCR SYBR Green ROX Mix (Thermo Scientific) with a final primer concentration of 200 nM. Primer sequences: Actb primer: 5'-TGACAGGATGC AGAAGGAGATTA-3'/5'-AGCCACCGATCCACAGA-3'; Dkk3 primer: 5'-TCCCATTGCCACCTTTGG-3'/5'-CCAGTTCTCCA GCTTCAAGTACAC-3'; Cxcl9 primer: 5'-CTTCGAGGAACCC TAGTGATAAGG-3'/5'-CCTCGGCTGGTGCTGATG-3'; Cxcl10 primer: 5'-GACGGTCCGCTGCAACTG-3'/5'-CCCTATGGCCC TCATTCTCA-3'; Ccl2 primer: 5'-AGCAGGTGTCCCAAAGAAG CT-3'/5'-GGGTCAGCACAGACCTCTCTCT-3'; Ccl5 primer: 5'-CTGCTTGCTCTAGTCCA-3'/5'-ATGCTGATTTCTTGGGTTT-3'; Gbp2 primer: 5'-GATGAACAAGCTAGCTGGGAAGAG-3'/5'-CCTTGGTGTGAGACTGCACAGT-3'. Data were calculated relative to the housekeeping gene Actb by using the $2^{-\Delta\Delta Ct}$ method.

ENZYME-LINKED IMMUNOSORBENT ASSAY

A flexible assay plate (BD Biosciences) was coated with anti-Dkk3-4.22 antibody ($1 \mu g/ml$). Tissues were homogenized in ProteoJET mammalian cell lysis reagent (Fermentas) supplemented with Complete, EDTA-free, protease inhibitor cocktail (Roche). Biotinylated goat anti-mouse Dkk3 antibody (R&D Systems, Minneapolis, MN, USA) at a final concentration of 1μ g/ml was used as detection antibody. Streptavidin-peroxidase (Jackson ImmunoResearch) diluted in PBS-Tween was added for 30 min at room temperature. After washing, orthophenylene diamine substrate at a concentration of 1 mg/ml in 100 mM sodium hydrogen phosphate buffer containing 0.03% peroxide was applied. The reaction was stopped with 2 M sulfuric acid. Optical densities were analyzed with a Viktor 2 photometer (Perkin Elmer) at 490 nm.

DORSAL ROOT GANGLIA

L2-L6 dorsal route ganglia (DRG) neurons isolation and cultured was performed as described (23). DRGs were excised and roots and membranes were removed and placed in sterile calcium- and magnesium-free PBS. Ganglia were collected and resuspended in DMEM/F12 (1:1) containing 10% horse serum, 2% penicillinstreptomycin solution, 1% L-glutamine, and 0.8% D-glucose (w/v). 0.1% collagenase was added and the tissue incubated at 37°C for 45 min. The tissue was incubated for a further 45 min in growth medium containing 0.05% trypsin at 37°C. Cells were dissociated by trituration through a fire-polished Pasteur pipette. After centrifugation at $250 \times g$ for 5 min, the resultant pellet was washed twice in growth medium. Finally, cells were plated immediately, either onto a six-well plate or glass cover-slips which had previously been coated with poly L-ornithine $(1 \mu g/ml)$ and laminin (25 µg/ml) supplemented with murine nerve growth factor (100 ng/ml). After 24 h incubation, the culture medium was supplemented with cytosine arabinoside $(10 \,\mu M)$ and incubated for 12 h, after which time, culture medium was changed every 2 days until 70-80% confluence was reached.

IFN γ INJECTION INTO THE HYPOTHALAMIC ARC

Bilateral stereotaxic injections were performed as described (25). Hundred nanograms of recombinant murine IFN γ (Peprotech) was injected into the hippocampus of each hemisphere (two injections per hemisphere: (1) caudal to bregma: -2.2 mm, lateral: -2.5 mm, ventral: 2.4 mm; (2) caudal to bregma: -2.0 mm, lateral: -2.0 mm, ventral: 1.6 mm). Twenty-four hours after injection, hippocampi of both hemispheres were isolated and used for analysis.

STATISTICAL ANALYSIS

All data are represented as mean \pm SEM. Statistical analysis was performed using the GraphPad Prism 6 software. *p* values were calculated using an unpaired two-tailed Student's *t*-test. *p* < 0.05 was considered significant.

RESULTS

Dkk3 DEFICIENCY DOES NOT INFLUENCE T-CELL DEVELOPMENT, ACTIVATION, AND MEMORY FORMATION IN THE STEADY STATE

In order to identify the impact of Dkk3 on T cells of a polyclonal repertoire, we initially analyzed T-cell development in Dkk3-deficient mice. Flow cytometric analysis revealed similar percentages (**Figure 1A**) and numbers (**Figure 1B**) of thymocyte subsets in Dkk3^{-/-} and WT control mice. Additionally, cell surface expression of CD69, which is a marker for the engagement

of the TCR-mediated positive selection process in the thymus, was identical on the different thymocyte subsets in both strains (Figure 1C). Frequencies of CD4⁺ CD25⁺ Foxp3⁺ regulatory T cells in the thymus were not altered in $Dkk3^{-/-}$ mice (Figure 1D). The ratio and numbers of CD4⁺ and CD8⁺ T cells in spleen (Figures 1E,F) and peripheral lymph nodes (Figures 1G,H) were also comparable. Moreover, analysis of TCR-VB expression showed that genetic Dkk3 deletion did not strongly affect the development of the TCR repertoire (Table 1). In addition to their developmental properties, we analyzed the activation and memory status of T cells in naïve Dkk3^{-/-} mice. Surface expression of the activation markers CD25 and CD69 on CD4⁺ as well as CD8⁺ T cells from spleen (Figure 1I) and lymph nodes (Figure 1K) were comparable in Dkk3^{-/-} and WT control mice. Similarly, Dkk3 deficiency did not influence the expression of the memory markers CD44 and CD62L on CD4⁺ and CD8⁺ T cells in spleen (Figure 1J) and lymph nodes (Figure 1L). Thus, these results illustrate that Dkk3 does not interfere with T-cell development, activation, and memory differentiation in the steady state.

Dkk3 CONTRIBUTES TO THE ACCEPTANCE OF EMBRYOID BODY GRAFTS

Next, we were interested in the influence of Dkk3 on T-cell function under stimulatory conditions. As we previously identified Dkk3 to be crucial for the establishment of peripheral CD8 Tcell tolerance in a TCR transgenic system (20), we wondered whether it may also influence CD8 T-cell responses in a polyclonal TCR repertoire. To address this question, we employed an EBs transplantation model. EBs are embryonic stem cell-derived composite tissues that have been previously reported to establish an immunosuppressive micromilieu (22). In previous work, it has been demonstrated that transplantation of male MHC class-I mismatched EBs under the kidney capsule of sex-matched recipient mice results in 50% graft acceptance (22). Here, male CBK-derived EBs were transplanted under the kidney capsule of male CBA recipient mice. The accepted and rejected transplants were analyzed at day 28 for Dkk3 expression. Strikingly, Dkk3 protein was detected in the accepted but not in the remains of the rejected grafts (Figure 2A). To dissect the contribution of Dkk3 to the establishment of the local immunosuppressive microenvironment of the EB grafts, we administered anti-Dkk3 neutralizing or isotype control antibody following transplantation and assessed its effect on graft survival. Interestingly, blocking of Dkk3 by the use of the monoclonal antibody resulted in an increased incidence graft rejection (Figure 2B). Taken together, our results demonstrate that Dkk3 expression correlates with the development of a local immunosuppressive micromilieu and the immunosuppressive function of Dkk3 contributes to the acceptance of MHC class-I mismatched EB grafts.

Dkk3^{-/-} MICE SUFFER FROM EXACERBATED EXPERIMENTAL AUTOIMMUNE ENCEPHALOMYELITIS

Dkk3 was reported to be strongly expressed in the brain (15). Detailed immune-histological analysis revealed a highly specific expression pattern within the organ. Prominent Dkk3 production was restricted to the hippocampus, medulla, and the visual area of the cerebral cortex (**Figure 3A**). Moreover, double-fluorescence



FIGURE 1 | Continued

(A) Thymocytes from naïve Dkk3^{-/-} mice and littermate controls were stained for CD4 and CD8 and analyzed by flow cytometry. Distribution of thymocyte subsets (DN: CD4⁻ CD8⁻ double negative; DP: CD4⁺ CD8⁺ double positive) is shown in a representative dot plot (left) and cumulative data of percentages of whole thymocytes (right; n = 4-5). (B) Absolute numbers of respective thymocyte subsets from (A), determined by flow cytometry (n = 4-5). (C) Flow cytometric analysis of CD69 cell surface expression on CD4⁺, CD8⁺, and CD4⁺ CD8⁺ double positive (DP) subsets from (A). One representative histogram is shown (n = 4-5). (D) Representative dot plot of CD25⁺ Foxp3⁺ cells among CD4 single

positive cells in the thymus of Dkk3^{-/-} and WT control mice. **(E)** Analysis of CD4⁺ and CD8⁺ T-cell subsets, among CD3 positive cells, in the spleen of Dkk3^{-/-} mice compared to littermate controls. One representative dot plot

(left) as well as respective cumulative data is shown. **(F)** Absolute numbers of lymphocyte subsets from **(D)**, determined by flow cytometry (n = 4-5). **(G)** Analysis of CD4⁺ and CD8⁺ Tcell subsets among CD3 positive cells in peripheral lymph nodes (LN) of Dkk3^{-/-} mice and WT controls. One representative dot plot (left) as well as respective cumulative data is shown. **(H)** Absolute numbers of lymphocyte subsets from **(F)**, determined by flow cytometry (n = 4-5). **(I+K)** Flow cytometric analysis of surface expression of the activation markers CD25 and CD69 on CD4⁺ (upper row) and CD8⁺ (lower row) T cells in **(I)** spleen and **(K)** peripheral lymph nodes of naïve Dkk3^{-/-} (red) and WT control (black) mice. Shown is one representative histogram (n = 10). **(J+L)** Flow cytometric analysis of the surface expression of CD44 and CD62L on CD4⁺ (upper row) and CD8⁺ (lower row) T cells in **(I)** peripheral lymph nodes of naïve Dkk3^{-/-} and WT control mice. Shown is one representative dot plot (n = 8).

| THYMUS (| CD4 | | | | | | | | | | | | | |
|---------------------|-----|-----|-----|---------|------|-----|---------|-----|-----|-----|-----|-----|-----|-----|
| TCR-Vβ | 2 | 3 | 4 | 5.1/5.2 | 6 | 7 | 8.1/8.2 | 9 | 10b | 11 | 12 | 13 | 14 | 17a |
| wт | 0.5 | 3.9 | 2.4 | 2.8 | 6.2 | 1.3 | 8.6 | 8.0 | 3.7 | 8.0 | 1.2 | 1.3 | 3.8 | 3.0 |
| Dkk3 ^{-/-} | 0.5 | 4.5 | 4.3 | 2.8 | 4.6 | 1.1 | 10.6 | 6.8 | 5.1 | 6.8 | 0.7 | 1.0 | 3.3 | 3.3 |
| THYMUS O | CD8 | | | | | | | | | | | | | |
| TCR-Vβ | 2 | 3 | 4 | 5.1/5.2 | 6 | 7 | 8.1/8.2 | 9 | 10b | 11 | 12 | 13 | 14 | 17a |
| WT | 0.1 | 4.3 | 1.4 | 5.8 | 11.5 | 1.7 | 8.4 | 5.8 | 1.0 | 0.9 | 0.5 | 0.5 | 2.0 | 2.1 |
| Dkk3 ^{-/-} | 0.1 | 5.5 | 3.3 | 4.6 | 9.5 | 1.6 | 9.5 | 6.1 | 1.6 | 1.2 | 0.2 | 0.2 | 2.5 | 1.6 |
| SPLEEN C | D4 | | | | | | | | | | | | | |
| TCR-Vβ | 2 | 3 | 4 | 5.1/5.2 | 6 | 7 | 8.1/8.2 | 9 | 10b | 11 | 12 | 13 | 14 | 17a |
| WT | 0.6 | 4.9 | 2.6 | 2.3 | 3.1 | 1.0 | 7.5 | 8.7 | 0.6 | 3.5 | 0.7 | 1.0 | 3.6 | 2.5 |
| Dkk3 ^{-/-} | 0.7 | 3.6 | 1.8 | 2.2 | 3.4 | 1.7 | 6.9 | 6.3 | 0.5 | 3.3 | 2.9 | 1.0 | 2.2 | 2.3 |
| SPLEEN C | D8 | | | | | | | | | | | | | |
| TCR-Vβ | 2 | 3 | 4 | 5.1/5.2 | 6 | 7 | 8.1/8.2 | 9 | 10b | 11 | 12 | 13 | 14 | 17a |
| WT | 0.2 | 5.7 | 1.6 | 4.7 | 11.7 | 0.9 | 6.2 | 6.7 | 0.2 | 0.6 | 0.5 | 0.3 | 1.9 | 0.9 |
| Dkk3 ^{-/-} | 0.3 | 4.2 | 0.7 | 5.2 | 8.0 | 0.6 | 5.5 | 4.3 | 1.1 | 0.9 | 1.5 | 0.6 | 1.1 | 0.5 |

Table 1 | Normal T-cell repertoire in Dkk3-deficient mice.

Flow cytometric analysis of TCR-Vβ chain expression on T cells. Percentages of CD4⁺ or CD8⁺ T cells in thymus and spleen carrying the respective TCR-Vβ.



embryoid bodies from rejection. (A) Male CBK mice-derived embryoid bodies were transplanted under the kidney capsule of male CBA mice. Twenty-eight days after transplantation, remaining embryoid body tissue was stained with anti-Dkk3 antibody. Shown is one representative staining of accepted and remaining elements of the rejected tissue

(n=3). **(B)** Male CBK mice-derived embryoid bodies were transplanted under the kidney capsule of male CBA mice. Seven days later, 1 mg anti-Dkk3 or isotype control antibody was administered, followed by 0.5 mg every fourth day. Twenty-eight days after transplantation, the acceptance rate of the transplants was determined (Log-rank test, n=24, *p < 0.05).



staining indicated that neurons (NeuN⁺) but not astrocytes (GFAP⁺) are the main source of Dkk3 in the brain (**Figure 3B**).

Based on our findings that Dkk3 in the microenvironment might influence CD8 T-cell responses (Figure 2), we wondered whether it acts on local CD4 T-cell reactivity as well. Due to the locally high expression of Dkk3 in the brain, we chose the EAE as a model system to answer this question. EAE is regarded as a mouse-model of human multiple sclerosis, in which auto-reactive CD4 T-cell-mediated demyelization is triggered in the CNS. Thus, EAE was induced by immunization with a MOG₃₃₋₅₅ peptide and CFA in Dkk3^{-/-} and WT mice and the disease course was followed. The onset of EAE and the disease incidence were comparable in the two mouse groups (Figure 4A; Table 2). Moreover, the severity of EAE in the acute phase was not significantly higher after genetic deletion of Dkk3 (Figure 4A; Table 2). However, Dkk3^{-/-} mice failed to recover and exhibited a more chronic disease when compared to WT controls (Figure 4A). Most of the EAE-affected WT mice recovered after initial paralysis and only 27.9% had clinical symptoms for longer than 20 days. In contrast,

the majority of diseased Dkk3^{-/-} mice exhibited increased disease duration, developing chronic disease or suffering from relapses, as 57.9% had clinical symptoms for longer than 20 days (**Figure 4B**).

We next sought to assess whether an established autoimmune process is also susceptible to the effects of Dkk3. For this, we treated mice 1 week after EAE induction with the neutralizing anti-Dkk3 antibody. Due to changes in the blood–brain barrier during EAE, antibodies can access inflammatory lesions in the CNS (24). We found that Dkk3 blockade starting 1 week after disease induction resulted in a significant increase and prolongation of EAE symptoms, as compared to isotype-treated controls (**Figure 4C**) suggesting that Dkk3 controls the potency and duration of the T-cell effector phase, locally in the CNS.

To explore the consequences of Dkk3 deficiency on T-cell function in EAE more detailed, we analyzed the T-cell phenotype in the CNS of diseased Dkk3^{-/-} and WT control mice. While IL-17- and GM-CSF-producing CD4⁺ T cells, have been shown to play an important role in EAE induction (25, 26), IFN γ was reported to have a pleiotropic effect in EAE. Depending on the



exacerbated EAE. (A) Dkk3^{-/-} and WT mice were immunized s.c. with MOG_{33-65} peptide in CFA. The mean EAE scores of one representative out of three experiments are shown (n = 9). (B) The percentage of living mice with disease duration longer than 20 days was determined. Results from

Table 2 | Onset and severity in the acute phase of experimental autoimmune encephalomyelitis in Dkk3-deficient mice was comparable to WT controls.

| | Incidence | % | Day of disease onset | Mean maximal disease severity | Dead mice |
|---------------------|-----------|------|-------------------------|----------------------------------|--------------|
| WT | 24/26 | 92.3 | 11.1 ± 2.6 | 2.4 ± 1.1 | 1 |
| Dkk3 ^{-/-} | 25/25 | 100 | 10.8 ± 2.3 | 3.1 ± 1.1 | 4 |

cell type of origin and time point of expression, it may mediate disease-promoting or disease-limiting effects (27). Therefore, we re-stimulated lymphocytes isolated from the CNS with MOG peptide and assessed the numbers of IL-17, GM-CSF, and IFNγproducing CD4 T cells. Since the MOG_{33–55} peptide also contains a CD8-epitope able to induce IFNγ production and EAE (28), we also analyzed CD8⁺ IFNγ⁺ T cells. The presence of CD4⁺ CD25⁺ Foxp3⁺ regulatory T cells, which are shown to be potent down-modulators of the T-cell response in EAE (29), was also determined.

At the peak of the disease, when EAE symptoms in Dkk $3^{-/-}$ and WT mice were still comparable (**Figure 5A**), the numbers

two independent experiments were pooled (n = 16, *p < 0.05). (C) C57BL6 mice received either anti-Dkk3 or isotype control antibody at days 7 (1 mg/mouse), 10, 13, 16, 21 (0.5 mg/mouse) after MOG₃₃₋₆₅/CFA immunization. Shown is one representative experiment out of three (n = 10, *p < 0.05).

of infiltrating CD4⁺ and CD8⁺ T cells in the CNS were similar (**Figure 5B**). However, while GM-CSF and IFN γ producing cells were not affected (**Figures 5E–G**), the proportion of IL-17-producing CD4⁺ T cells was slightly increased in the CNS of Dkk3-deficient mice (**Figure 5D**). Interestingly, at the same time, the proportion of CD4⁺ CD25⁺ Foxp3⁺ regulatory T cells was significantly elevated (**Figure 5C**). These differences were absent in spleen and draining lymph nodes (**Figures 5H–J**) indicating a local effect of Dkk3.

In contrast, during the recovery phase of the disease, when more severe EAE symptoms in Dkk3^{-/-} mice were manifest (**Figure 6A**), we found higher numbers of CD4⁺ and CD8⁺ T cells in the CNS but not in the draining lymph nodes (**Figure 6H**) of Dkk3-deficient mice compared to WT controls (**Figure 6B**). Furthermore, the ratio of IFN γ -producing CD4⁺ and CD8⁺ was significantly increased (**Figures 6F,G**) while the proportion of IL-17 and GM-CSF-producing CD4⁺ T cells (**Figures 6D,E**) as well as CD4⁺ CD25⁺ Foxp3⁺ cells (**Figure 6C**) was similar in the CNS of both strains. The impact on the T-cell phenotype was again a brain-specific effect, since Dkk3 deficiency did not interfere with the levels of IFN γ -producing CD4⁺ and CD8⁺ T cells in spleen and draining lymph nodes (**Figure 6I**). Thus, our data indicate



draining lymph nodes (Figure 6H) of Dkk3^{-/-} at the peak of EAE. (A) EAE was induced in Dkk3^{-/-} and WT mice by immunization with MOG_{33-65} in CFA. Mean clinical EAE score over time is displayed. At the indicated time point (black arrow, 14 days after EAE induction), half of the mice were sacrificed and lymphocytes were isolated from brain and spinal cord for analysis in (**B**-**G**) (n = 8). (**B**) Percentages of CD4⁺ and CD8⁺ T cells among leukocytes in the CNS of Dkk3^{-/-} and WT mice. (**C**) Proportion of CD25⁺ Foxp3⁺ of CD4⁺ cells (*p < 0.05). (**D**-**G**) Isolated lymphocytes from (**A**) were re-stimulated *in vitro* with 50 µg/ml MOG₃₃₋₆₅

that Dkk3 modulates T-cell activation and polarization during the advanced phase of EAE locally in the CNS and consequently promotes symptom recovery.

Dkk3 SERVES AS PART OF AN IFN γ NEGATIVE FEEDBACK LOOP IN THE BRAIN DURING EAE

IFN γ plays a multifaceted role in EAE. Its disease-limiting effect was suggested to be partially mediated by the induction of antiinflammatory mediators like IDO (14) or nitric oxide (NO) (30). Despite elevated IFN γ levels in the CNS, we observed a prolonged disease process in the Dkk3-deficient situation. Therefore, peptide for 6 h in the presence of Golgi Plug and intracellularly stained for the respective cytokines. Shown is one representative dot plot (left panel) and cumulative data of three individual experiments (n = 8). (D) Percentages of IL-17⁺ of CD4⁺ T cells (*p < 0.05). (E) Percentages of GM-CSF⁺ of CD4⁺ T cells. (F) Percentages of IFN γ^+ of CD4⁺ T cells. (G) Percentages of IFN γ^+ of CD8⁺ T cells. (H) Percentages of CD4⁺ and CD8⁺ T cells in spleen and peripheral lymph nodes (LN) of respective mice from (A). (J) Percentages of IL-17⁺ of CD4⁺ cells in spleen and draining lymph nodes (LN) of respective mice from (A). (J) Percentages of IL-17⁺ of CD4⁺ cells in spleen and draining lymph nodes (LN) of respective mice from (A).

we hypothesized that Dkk3 might be part of an IFN γ -mediated regulatory mechanism, that is lost in Dkk3^{-/-} mice. In order to test this, we initially analyzed the capacity of IFN γ to induce Dkk3 expression during EAE in the brain. Comparison of Dkk3 expression in the brain of healthy controls and EAE-bearing mice, at the peak of diseases, revealed a significant increase of Dkk3 levels in the inflamed brains of WT mice. However, such an elevation was not detectable in brains of IFN γ R1^{-/-} mice (**Figure 7A**), indicating an IFN γ -dependent induction of Dkk3 expression in EAE brains. In order to find out whether IFN γ is able to up-regulate Dkk3 expression in neuronal cells, we made use of *ex vivo* cultured



EAE diseased Dkk3^{-/-} mice in the recovery phase. (A) EAE was induced in Dkk3^{-/-} and WT mice by immunization with MOG_{33-55} in CFA. Mean clinical EAE score over time is displayed. At the indicated time point (black arrow, 20 days after EAE induction), mice were sacrificed and lymphocytes were isolated from brain and spinal cord for analysis in (B–G). (B) Percentages of CD4⁺ and CD8⁺T cells among leukocytes in the CNS of Dkk3^{-/-} and WT mice (*p < 0.05). (C) Proportion of CD25⁺ Foxp3⁺ of CD4⁺ cells. (D–G) Isolated lymphocytes from (A) were re-stimulated *in vitro* with 50 µg/ml MOG₃₃₋₅₅

peptide for 6 h in the presence of Golgi Plug and intracellularly stained for the respective cytokines. Shown is one representative dot plot (left panel) and cumulative data of three individual experiments (n = 8). (**D**) Percentages of IL₁7⁺ of CD4⁺ T cells. (**E**) Percentages of GM-CSF⁺ of CD4⁺ T cells. (**F**) Percentages of IFN γ^+ of CD4⁺ T cells (*p < 0.05). (**G**) Percentages of IFN γ^+ of CD8⁺ T cells (*p < 0.05). (**G**) Percentages of IFN γ^+ of CD8⁺ T cells (*p < 0.05). (**H**) Percentages of CD4⁺ and CD8⁺ T cells in spleen and draining lymph nodes (LN) of respective mice from (**A**). (**I**) Percentages of IFN γ^+ of CD4⁺ and CD8⁺ cells in spleen and draining lymph nodes (LN) of respective mice from (**A**).

primary DRG neurons. Histological analysis revealed, just as for neurons *in vivo*, that these cells express Dkk3 (**Figure 7B**). Moreover, IFN γ stimulation of DRGs increased the basal expression level (**Figure 7C**) and secretion of Dkk3 protein (**Figure 7D**) by nearly sixfold. As expected, this up-regulation was not observed in IFN γ R1^{-/-} control DRGs (**Figure 7E**). Thus, IFN γ up-regulates Dkk3 expression in neuronal cells.

In order to find out whether Dkk3 is, in turn, capable of influencing IFN γ activity during EAE, we analyzed IFN γ -induced gene expression in the brain. To enhance the pathological situation, in which the first IFN γ -producing T cells enter the brain 7 days after EAE induction (31), we performed the following experiment: 7 days after EAE induction in Dkk3^{-/-} and WT mice, IFN γ was stereotactically injected into the hippocampal arc of the mice, and expression of IFN γ target genes was analyzed 24 h after administration. This area was targeted because of the high local concentration of Dkk3 in the hippocampus (**Figure 3A**). As a control, hippocampi of non-injected animals were analyzed. QRT-PCR analysis revealed that IFN γ injection strongly induced IFN γ target gene expression in hippocampi of injected mice compared to non-injected controls. Furthermore, this induction was more prominent in Dkk3-deficient mice. In Dkk3^{-/-} hippocampi, the expression of Gbp2 and Cxcl10 was significantly enhanced upon IFN γ administration. However, Cxcl9, Ccl2, and Ccl5 expression



by ELISA. EAE was induced by immunization with MOG/CFA and minimal clinical EAE score of used mice was 4. Protein amount was related to organ weight (n = 5, **p < 0.01). (B) Histological analysis of *ex vivo* cultured, primary dorsal route ganglia (DRG) stained with anti-Dkk3 (upper row) or isotype control antibody (lower row). (C) QRT-PCR data showing relative *Dkk3* expression in cells from (C) (n = 3, **p < 0.01). (D) Dkk3 ELISA of supernatant from control and IFN γ stimulated primary DRG isolated from WT mice. DRG

was not much altered (**Figure 7F**). In control hippocampi, dkk3 deficiency did not cause an increase in IFN γ target gene expression (**Figure 7F**). In summary, these data suggest that Dkk3 negatively modulates IFN γ activity in the EAE brain.

DISCUSSION

Regulation of T-cell responses within tissues is essential for the preservation of organ integrity and function. In this context, the tissue microenvironment plays a pivotal role in the establishment of an immunological balance. In this study, we provide evidence that the secreted glycoprotein Dkk3 functions as a tissue-derived modulator of local T-cell responses. Our data show that Dkk3 (i) is part of the local microenvironment that protects transplanted MHC class-I mismatched embryo bodies from CD8⁺ T-cell-dependent rejection and (ii) contributes to the control of CD4⁺ T-cell-mediated autoimmune encephalomyelitis by limitation of T-cell activation and polarization as well as IFN γ activity in the CNS.

Our group previously demonstrated that the presence of Dkk3 is essential for the establishment of peripheral CD8⁺ T-cell tolerance in a TCR/MHC class-I double transgenic mouse system. Genetic deletion or antibody blockade of Dkk3 in this system led to an increased local CD8⁺ T-cell reactivity resulting in were stimulated for 24 h with 100 ng/ml IFN_Y (n = 3, **p < 0.01). (**E**) QRT-PCR data showing relative Dkk3 expression in control and IFN_Y stimulated primary DRG from IFN_YR1^{-/-} mice. DRG were stimulated for 24 h with 100 ng/ml IFN_Y (n = 3). (**F**) WT and Dkk3^{-/-} were immunized with MOG₃₃₋₆₅ in CFA. Seven days later, 100 ng recombinant IFN_Y in PBS was injected stereotacically into hippocampi of the respective mice. One day later, hippocampi were harvested and RNA was isolated. As controls, hippocampi of non-IFN_Y-injected mice, 8 days after MOG/CFA immunization were used. Shown are qRT-PCR data of the relative expression of *Gbp2*, *Cxcl10*, *Cxcl9*, *Ccl2* (MCP-1), *Ccl5* (RANTES) in respective hippocampi (n = 8, *p < 0.05, **p < 0.01).

an enhanced, antigen-specific eradication of tumors and skin grafts (20). The investigations performed here display a similar, immune-modulatory role of Dkk3 for the polyclonal T-cell repertoire *in vivo*. Transplanted EBs were shown to create an immune suppressive microenvironment by, for example, the release of transforming growth factor- β (TGF- β) (22). EB transplants were shown to express Dkk3 and antibody-mediated blockade of Dkk3 increased the frequency of their rejection, albeit some EB grafts remaining intact. We therefore, suggest that Dkk3 represents one constituent of the local, suppressive EB micromilieu suppressing cytotoxic CD8⁺ T-cell responses. This view is supported by the observation that the global absence of Dkk3 did not lead to a systemic hyperresponsiveness of cytotoxic CD8⁺ T cells *in vivo* in the polyclonal repertoire (20).

In contrast to the double transgenic system in which Dkk3 was expressed by a unique, tolerant CD8⁺ T-cell population, (20), Dkk3 was reported to be mainly produced by tissue cells but very rarely in the hematopoietic compartment in the physiological situation. Dkk3 expression in the hematopoietic system was only detected at a transcriptional level in a CD27 negative, IL-17-producing subset of $\gamma\delta$ T cells (32) and long-term memory CD8 T cells (33) but not in naïve or activated conventional $\alpha\beta$ T cells. Highest Dkk3 levels were found in organs like uterus,

placenta, eye, and brain (15, 17). These organs are classically considered to be immune-privileged, a situation in which the tissue is protected from otherwise damaging immune responses. This is of particular importance as their regenerative capacity is very low and cellular damage would strongly impair organ integrity and function (34, 35). Based on the immune suppressive capacity of Dkk3 shown here, one could speculate that Dkk3 is one contributor to the creation of such immunosuppressive microenvironments.

Indeed, our studies on EAE demonstrate an important role of Dkk3 in the regulation of CD4⁺ T-cell responses. Dkk3 deficiency resulted in a delayed EAE remission with exacerbated symptoms during recovery phase but did not lead to higher EAE susceptibility, premature disease onset nor increased severity at the peak of disease. These data indicate that Dkk3 mediates its function rather locally during the T-cell effector phase, rather than systemically in spleen and lymph nodes during the T-cell priming phase. This assumption is supported by the finding that blockade of Dkk3 by a monoclonal antibody, starting 7 days after immunization, is sufficient to induce a similar phenotype. The locally restricted function might also explain, at least in part, why Dkk3 deficiency does not lead to T-cell activation in secondary lymphoid organs in the steady state and thus, why Dkk3^{-/-} mice do not suffer from spontaneous autoimmunity (21).

The observed shift of T-cell polarization within the CNS of EAE-bearing Dkk3-deficient mice varied at different stages of the disease. At the peak of disease, a higher proportion of Th17 and Foxp3⁺ regulatory T cells was detected. While Th17 cells are crucial for the induction of EAE (25, 26), Foxp3⁺ Tregs were shown to suppress encephalitogenic T-cell reactivity (29). As both subsets were enriched in the Dkk3^{-/-} CNS, it could be that they compensate each others function, resulting in a similar EAE score compared to WT mice. However, in the recovery phase, the height-ened EAE severity in Dkk3-deficient mice was accompanied by elevated T-cell numbers and an increase in IFN γ producing CD4⁺ and CD8⁺ T cells in the CNS. Again, the altered T-cell polarization in the CNS of Dkk3-deficient mice upon EAE induction was restricted to the brain and not observed in the secondary lymphoid organs endorsing the hypothesis of local Dkk3 action.

IFNy has a multifaceted role in EAE. On the one hand, it has been positively correlated with disease severity (36), while on the other hand genetic or antibody-mediated neutralization exhibited disease-ameliorating effects (37-39). It was suggested that, after being pro-inflammatory and disease-promoting in the early phase, IFNy exerts an anti-inflammatory, beneficial function in the later phases (27). Its disease-providing properties are mediated by induction of antigen presentation, promotion of Th1 differentiation as well as up-regulation of adhesion molecules and the induction of a wide range of pro-inflammatory mediators (40). The anti-inflammatory function is attributed to the inhibition of T-cell expansion, induction of Foxp3 positive regulatory cells, and the up-regulation of anti-inflammatory mediators like IDO or the inducible nitric oxide synthase (iNOS) (27, 30, 41). Our studies reveal that Dkk3 expression is up-regulated in the brain of EAEbearing mice in an IFNy-dependent manner. Moreover, we found that Dkk3 is able to regulate the expression of certain IFNy target

genes in the brain. Together these results suggest that neuronderived Dkk3 might serve as a mediator of the anti-inflammatory arm of IFN γ during EAE, conducting its function by regulation of IFN γ activity in a negative feedback mechanism.

The immune modulator IDO also locally suppresses Tcell responses by the catabolic deprivation of tryptophan and production of kynurenine metabolites, and is thus, thought to play an important role in the induction of immune tolerance and the prevention of autoimmunity (42, 43). It is expressed in the brain and is further up-regulated by IFNy. Chemical inhibition or genetic deletion of IDO causes exacerbated EAE symptoms (14, 44). Moreover, IDO deficiency leads to increased numbers of IFNy-producing T cells in EAE-bearing mice and treatment of transferred EAE with kynurenines shifts the cellular immune response from Th1 to Th2 (45). These results led to the assumption that IDO expression in the CNS initiates a negative feedback loop to self-limit autoimmunity (44). In addition, via the interaction with its ligand PD-1, tissue-derived PD-L1 is important for the establishment of peripheral tolerance. In a fully MHC mismatched cardiac allograft model, blockade of PD-L1 abrogated CTLA4Ig-induced tolerance and accelerated organ rejection. This was associated with an increase in the frequency of IFNyproducing allo-reactive T cells and expansion of CD8 effector cells (46). High PD-L1expression was shown to be key for maintaining immune privilege in the eye (47) and feto-maternal tolerance in the placenta (48). PD-L1-deficient mice do not display T-cell activation *per se* or spontaneous autoimmunity in the steady state. However, upon MOG immunization, these mice suffer from exacerbated EAE (49). The functional characteristics of Dkk3 identified here seem quite similar to those of these well-characterized immune-modulatory molecules.

Mechanistically, Dkk3 does not conduct its function via interference with the development of the TCR repertoire. This is indicated by the findings that the TCR-VB frequencies are not dissimilar, that antibody blockade reflects the phenotype of genetic deletion, and that immune responses in the lymphoid compartments are not influenced by Dkk3 (20). Our finding that Dkk3 interferes with the expression of IFNy target genes like Gbp2 or Cxcl10 suggests that Dkk3 may not necessarily act directly on T cells but may also affect the function of ambient cells in the microenvironment which in turn modulate T-cell responses. For example, oligodendrocytes and astrocytes were shown to upregulate the expression of Gbp2 and Cxcl10 in response to IFNy during EAE (50, 51). Since Dkk3 may act as Wnt pathway modulator (17-19), it is possible that its immune-modulatory function also relies on this property, especially as Wnt signaling is known to influence T-cell effector function (52) and linage commitment (53-55). However, the distinct cellular and molecular mechanisms through which Dkk3 mediates its function warrant further investigation.

In summary, our results strongly suggest that Dkk3 contributes to the establishment of peripheral T-cell tolerance and the protection of tissues from self-destructive immune responses. These findings suggest that Dkk3 should be considered as a potential target for therapeutic approaches in transplant rejection and autoimmunity.

AUTHOR CONTRIBUTIONS

MM and MP contributed equally to this work. MM, MP, VN, VK, JL, KL, and AB performed experiments and analyzed data. MM, MP, TO, and BA designed experiments with contributions of TF, GM, PN, HG, and HW. MM, MP, and BA wrote the manuscript.

ACKNOWLEDGMENTS

We thank G. Küblebeck, A. Klevenz, S. Prokosch, S. Schmitt, and G. Pougialis for expert technical assistance.

REFERENCES

- Iwai Y, Terawaki S, Ikegawa M, Okazaki T, Honjo T. PD-1 inhibits antiviral immunity at the effector phase in the liver. J Exp Med (2003) 198:39–50. doi:10.1084/jem.20022235
- Ansari MJ, Salama AD, Chitnis T, Smith RN, Yagita H, Akiba H, et al. The programmed death-1 (PD-1) pathway regulates autoimmune diabetes in nonobese diabetic (NOD) mice. J Exp Med (2003) 198:63–9. doi:10.1084/jem. 20022125
- Salama AD, Chitnis T, Imitola J, Ansari MJ, Akiba H, Tushima F, et al. Critical role of the programmed death-1 (PD-1) pathway in regulation of experimental autoimmune encephalomyelitis. *J Exp Med* (2003) 198:71–8. doi:10.1084/jem. 20022119
- Liang SC, Latchman YE, Buhlmann JE, Tomczak MF, Horwitz BH, Freeman GJ, et al. Regulation of PD-1, PD-L1, and PD-L2 expression during normal and autoimmune responses. *Eur J Immunol* (2003) 33:2706–16. doi:10.1002/eji. 200324228
- Wiendl H, Mitsdoerffer M, Schneider D, Chen L, Lochmuller H, Melms A, et al. Human muscle cells express a B7-related molecule, B7-H1, with strong negative immune regulatory potential: a novel mechanism of counterbalancing the immune attack in idiopathic inflammatory myopathies. *FASEB J* (2003) 17:1892–4. doi:10.1096/fj.03-0039fje
- Butte MJ, Keir ME, Phamduy TB, Sharpe AH, Freeman GJ. Programmed death-1 ligand 1 interacts specifically with the B7-1 costimulatory molecule to inhibit T cell responses. *Immunity* (2007) 27:111–22. doi:10.1016/j.immuni.2007.05.016
- Francisco LM, Salinas VH, Brown KE, Vanguri VK, Freeman GJ, Kuchroo VK, et al. PD-L1 regulates the development, maintenance, and function of induced regulatory T cells. J Exp Med (2009) 206:3015–29. doi:10.1084/jem.20090847
- Choi C, Benveniste EN. Fas ligand/Fas system in the brain: regulator of immune and apoptotic responses. *Brain Res Brain Res Rev* (2004) 44:65–81. doi:10.1016/j.brainresrev.2003.08.007
- Griffith TS, Brunner T, Fletcher SM, Green DR, Ferguson TA. Fas ligand-induced apoptosis as a mechanism of immune privilege. *Science* (1995) 270:1189–92. doi:10.1126/science.270.5239.1189
- Jerzak M, Bischof P. Apoptosis in the first trimester human placenta: the role in maintaining immune privilege at the maternal-foetal interface and in the trophoblast remodelling. *Eur J Obstet Gynecol Reprod Biol* (2002) 100:138–42. doi:10.1016/S0301-2115(01)00431-6
- Munn DH, Zhou M, Attwood JT, Bondarev I, Conway SJ, Marshall B, et al. Prevention of allogeneic fetal rejection by tryptophan catabolism. *Science* (1998) 281:1191–3. doi:10.1126/science.281.5380.1191
- Malina HZ, Martin XD. Indoleamine 2,3-dioxygenase activity in the aqueous humor, iris/ciliary body, and retina of the bovine eye. *Graefes Arch Clin Exp Ophthalmol* (1993) 231:482–6. doi:10.1007/BF02044236
- Malina HZ, Martin XD. Indoleamine 2,3-dioxygenase: antioxidant enzyme in the human eye. *Graefes Arch Clin Exp Ophthalmol* (1996) 234:457–62. doi:10.1007/BF02539413
- Kwidzinski E, Bunse J, Aktas O, Richter D, Mutlu L, Zipp F, et al. Indoleamine 2,3dioxygenase is expressed in the CNS and down-regulates autoimmune inflammation. *FASEB J* (2005) 19:1347–9. doi:10.1096/fj.04-3228fje
- Niehrs C. Function and biological roles of the Dickkopf family of Wnt modulators. Oncogene (2006) 25:7469–81. doi:10.1038/sj.onc.1210054
- Glinka A, Wu W, Delius H, Monaghan AP, Blumenstock C, Niehrs C. Dickkopf-1 is a member of a new family of secreted proteins and functions in head induction. *Nature* (1998) **391**:357–62. doi:10.1038/34848
- Nakamura RE, Hunter DD, Yi H, Brunken WJ, Hackam AS. Identification of two novel activities of the Wnt signaling regulator Dickkopf 3 and characterization of its expression in the mouse retina. *BMC Cell Biol* (2007) 8:52. doi:10.1186/1471-2121-8-52

- Lee EJ, Jo M, Rho SB, Park K, Yoo YN, Park J, et al. Dkk3, downregulated in cervical cancer, functions as a negative regulator of beta-catenin. *Int J Cancer* (2009) 124:287–97. doi:10.1002/ijc.23913
- Das DS, Wadhwa N, Kunj N, Sarda K, Pradhan BS, Majumdar SS. Dickkopf homolog 3 (DKK3) plays a crucial role upstream of WNT/beta-CATENIN signaling for Sertoli cell mediated regulation of spermatogenesis. *PLoS One* (2013) 8:e63603. doi:10.1371/journal.pone.0063603
- Papatriantafyllou M, Moldenhauer G, Ludwig J, Tafuri A, Garbi N, Hollmann G, et al. Dickkopf-3, an immune modulator in peripheral CD8 T-cell tolerance. *Proc Natl Acad Sci U S A* (2012) 109:1631–6. doi:10.1073/pnas.1115980109
- Barrantes Idel B, Montero-Pedrazuela A, Guadano-Ferraz A, Obregon MJ, Martinez R, de Mena V, et al. Generation and characterization of dickkopf3 mutant mice. *Mol Cell Biol* (2006) 26:2317–26. doi:10.1128/MCB.26.6.2317-2326.2006
- 22. Robertson NJ, Brook FA, Gardner RL, Cobbold SP, Waldmann H, Fairchild PJ. Embryonic stem cell-derived tissues are immunogenic but their inherent immune privilege promotes the induction of tolerance. *Proc Natl Acad Sci U S A* (2007) **104**:20920–5. doi:10.1073/pnas.0710265105
- Bierhaus A, Fleming T, Stoyanov S, Leffler A, Babes A, Neacsu C, et al. Methylglyoxal modification of Nav1.8 facilitates nociceptive neuron firing and causes hyperalgesia in diabetic neuropathy. *Nat Med* (2012) 18:926–33. doi:10.1038/ nm.2750
- 24. Jorgensen SH, Storm N, Jensen PE, Laursen H, Sorensen PS. IVIG enters the central nervous system during treatment of experimental autoimmune encephalomyelitis and is localised to inflammatory lesions. *Exp Brain Res* (2007) 178:462–9. doi:10.1007/s00221-006-0752-8
- Langrish CL, Chen Y, Blumenschein WM, Mattson J, Basham B, Sedgwick JD, et al. IL-23 drives a pathogenic T cell population that induces autoimmune inflammation. J Exp Med (2005) 201:233–40. doi:10.1084/jem.20041257
- Komiyama Y, Nakae S, Matsuki T, Nambu A, Ishigame H, Kakuta S, et al. IL-17 plays an important role in the development of experimental autoimmune encephalomyelitis. *J Immunol* (2006) 177:566–73. doi:10.4049/jimmunol.177. 1.566
- Zhang J. Yin and yang interplay of IFN-gamma in inflammation and autoimmune disease. J Clin Invest (2007) 117:871–3. doi:10.1172/JCI31860
- Ford ML, Evavold BD. Specificity, magnitude, and kinetics of MOG-specific CD8+ T cell responses during experimental autoimmune encephalomyelitis. *Eur J Immunol* (2005) 35:76–85. doi:10.1002/eji.200425660
- 29. Kohm AP, Carpentier PA, Anger HA, Miller SD. Cutting edge: CD4+CD25+ regulatory T cells suppress antigen-specific autoreactive immune responses and central nervous system inflammation during active experimental autoimmune encephalomyelitis. *J Immunol* (2002) **169**:4712–6. doi:10.4049/jimmunol.169.9. 4712
- Willenborg DO, Staykova M, Fordham S, O'Brien N, Linares D. The contribution of nitric oxide and interferon gamma to the regulation of the neuroinflammation in experimental autoimmune encephalomyelitis. *J Neuroimmunol* (2007) 191:16–25. doi:10.1016/j.jneuroim.2007.09.007
- 31. Juedes AE, Hjelmstrom P, Bergman CM, Neild AL, Ruddle NH. Kinetics and cellular origin of cytokines in the central nervous system: insight into mechanisms of myelin oligodendrocyte glycoprotein-induced experimental autoimmune encephalomyelitis. *J Immunol* (2000) 164:419–26. doi:10.4049/jimmunol. 164.1.419
- Schmolka N, Serre K, Grosso AR, Rei M, Pennington DJ, Gomes AQ, et al. Epigenetic and transcriptional signatures of stable versus plastic differentiation of proinflammatory gammadelta T cell subsets. *Nat Immunol* (2013) 14:1093–100. doi:10.1038/ni.2702
- 33. Kaech SM, Hemby S, Kersh E, Ahmed R. Molecular and functional profiling of memory CD8 T cell differentiation. *Cell* (2002) 111:837–51. doi:10.1016/S0092-8674(02)01139-X
- Barker CF, Billingham RE. Immunologically privileged sites. Adv Immunol (1977) 25:1–54. doi:10.1016/S0065-2776(08)60930-X
- Simpson E. A historical perspective on immunological privilege. *Immunol Rev* (2006) 213:12–22. doi:10.1111/j.1600-065X.2006.00434.x
- Ando DG, Clayton J, Kono D, Urban JL, Sercarz EE. Encephalitogenic T cells in the B10.PL model of experimental allergic encephalomyelitis (EAE) are of the Th-1 lymphokine subtype. *Cell Immunol* (1989) 124:132–43. doi:10.1016/ 0008-8749(89)90117-2
- Duong TT, Finkelman FD, Singh B, Strejan GH. Effect of anti-interferon-gamma monoclonal antibody treatment on the development of experimental allergic encephalomyelitis in resistant mouse strains. *J Neuroimmunol* (1994) 53:101–7. doi:10.1016/0165-5728(94)90069-8

- Ferber IA, Brocke S, Taylor-Edwards C, Ridgway W, Dinisco C, Steinman L, et al. Mice with a disrupted IFN-gamma gene are susceptible to the induction of experimental autoimmune encephalomyelitis (EAE). J Immunol (1996) 156:5–7.
- Krakowski M, Owens T. Interferon-gamma confers resistance to experimental allergic encephalomyelitis. *Eur J Immunol* (1996) 26:1641–6. doi:10.1002/eji. 1830260735
- Boehm U, Klamp T, Groot M, Howard JC. Cellular responses to interferongamma. Annu Rev Immunol (1997) 15:749–95. doi:10.1146/annurev.immunol. 15.1.749
- Refaeli Y, Van Parijs L, Alexander SI, Abbas AK. Interferon gamma is required for activation-induced death of T lymphocytes. J Exp Med (2002) 196:999–1005. doi:10.1084/jem.20020666
- Mellor AL, Keskin DB, Johnson T, Chandler P, Munn DH. Cells expressing indoleamine 2,3-dioxygenase inhibit T cell responses. J Immunol (2002) 168:3771–6. doi:10.4049/jimmunol.168.8.3771
- Mellor AL, Munn DH. Tryptophan catabolism and T-cell tolerance: immunosuppression by starvation? *Immunol Today* (1999) 20:469–73. doi:10.1016/ S0167-5699(99)01520-0
- 44. Yan Y, Zhang GX, Gran B, Fallarino F, Yu S, Li H, et al. IDO upregulates regulatory T cells via tryptophan catabolite and suppresses encephalitogenic T cell responses in experimental autoimmune encephalomyelitis. *J Immunol* (2010) 185:5953–61. doi:10.4049/jimmunol.1001628
- Platten M, Ho PP, Youssef S, Fontoura P, Garren H, Hur EM, et al. Treatment of autoimmune neuroinflammation with a synthetic tryptophan metabolite. *Science* (2005) 310:850–5. doi:10.1126/science.1117634
- Tanaka K, Albin MJ, Yuan X, Yamaura K, Habicht A, Murayama T, et al. PDL1 is required for peripheral transplantation tolerance and protection from chronic allograft rejection. *J Immunol* (2007) **179**:5204–10. doi:10.4049/jimmunol.179. 8.5204
- Guleria I, Khosroshahi A, Ansari MJ, Habicht A, Azuma M, Yagita H, et al. A critical role for the programmed death ligand 1 in fetomaternal tolerance. *J Exp Med* (2005) 202:231–7. doi:10.1084/jem.20050019
- Hori J, Wang M, Miyashita M, Tanemoto K, Takahashi H, Takemori T, et al. B7-H1-induced apoptosis as a mechanism of immune privilege of corneal allografts. *J Immunol* (2006) 177:5928–35. doi:10.4049/jimmunol.177.9.5928
- Latchman YE, Liang SC, Wu Y, Chernova T, Sobel RA, Klemm M, et al. PD-L1deficient mice show that PD-L1 on T cells, antigen-presenting cells, and host tissues negatively regulates T cells. *Proc Natl Acad Sci USA* (2004) 101:10691–6. doi:10.1073/pnas.0307252101
- 50. Balabanov R, Strand K, Goswami R, McMahon E, Begolka W, Miller SD, et al. Interferon-gamma-oligodendrocyte interactions in the regulation of

experimental autoimmune encephalomyelitis. J Neurosci (2007) 27:2013–24. doi:10.1523/JNEUROSCI.4689-06.2007

- Hindinger C, Bergmann CC, Hinton DR, Phares TW, Parra GI, Hussain S, et al. IFN-gamma signaling to astrocytes protects from autoimmune mediated neurological disability. *PLoS One* (2012) 7:e42088. doi:10.1371/journal.pone.0042088
- Gattinoni L, Zhong XS, Palmer DC, Ji Y, Hinrichs CS, Yu Z, et al. Wnt signaling arrests effector T cell differentiation and generates CD8+ memory stem cells. *Nat Med* (2009) 15:808–13. doi:10.1038/nm.1982
- 53. Yu Q, Sharma A, Oh SY, Moon HG, Hossain MZ, Salay TM, et al. T cell factor 1 initiates the T helper type 2 fate by inducing the transcription factor GATA-3 and repressing interferon-gamma. *Nat Immunol* (2009) 10:992–9. doi:10.1038/ni.1762
- Ma J, Wang R, Fang X, Sun Z. Beta-catenin/TCF-1 pathway in T cell development and differentiation. *J Neuroimmune Pharmacol* (2012) 7:750–62. doi:10. 1007/s11481-012-9367-y
- van Loosdregt J, Fleskens V, Tiemessen MM, Mokry M, van Boxtel R, Meerding J, et al. Canonical Wnt signaling negatively modulates regulatory T cell function. *Immunity* (2013) **39**:298–310. doi:10.1016/j.immuni.2013.07.019

Conflict of Interest Statement: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

Received: 02 December 2014; accepted: 10 February 2015; published online: 24 February 2015.

Citation: Meister M, Papatriantafyllou M, Nordström V, Kumar V, Ludwig J, Lui KO, Boyd AS, Popovic ZV, Fleming TH, Moldenhauer G, Nawroth PP, Gröne H-J, Waldmann H, Oelert T and Arnold B (2015) Dickkopf-3, a tissue-derived modulator of local T-cell responses. Front. Immunol. **6**:78. doi: 10.3389/fimmu.2015.00078

This article was submitted to Immunological Tolerance, a section of the journal Frontiers in Immunology.

Copyright © 2015 Meister, Papatriantafyllou, Nordström, Kumar, Ludwig, Lui, Boyd, Popovic, Fleming, Moldenhauer, Nawroth, Gröne, Waldmann, Oelert and Arnold. This is an open-access article distributed under the terms of the Creative Commons Attribution License (CC BY). The use, distribution or reproduction in other forums is permitted, provided the original author(s) or licensor are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or reproduction is permitted which does not comply with these terms.