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TREM2 Deficiency Disrupts Network Oscillations Leading to Epileptic Activity and Aggravates Aβ-related Hippocampal Pathophysiology in Mice

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Abstract

Background: Genetic mutations in triggering receptor expressed on myeloid cells-2 (TREM2) have been strongly associated with increased risk of developing Alzheimer's disease (AD) and other progressive dementias. In the brain, TREM2 protein is specifically expressed on microglia suggesting their active involvement in driving disease pathology. Using various transgenic AD models to interfere with microglial function through TREM2, several recent studies provided important data indicating a causal link between TREM2 and underlying amyloid- β (A β) and tau pathology. However, mechanisms by which TREM2 contributes to increased predisposition to clinical AD and influences its progression still remain largely unknown.

Methods: Spontaneous and brainstem nucleus pontis oralis stimulation-induced hippocampal oscillation paradigm was used to investigate the impact of TREM2 haploinsufficiency TREM2(Het) or total deficiency TREM2(Hom) on hippocampal network function in wild-type and amyloid-β overproducing Tg2576 mice under urethane anesthesia.

Results: Partial (TREM2(Het)) or total (TREM2(Hom)) deletion of TREM2, led to increased incidence of spontaneous epileptiform seizures in both wild-type and Tg2576 mice. Importantly, deficiency of TREM2 in Tg2576 mice significantly diminished power of theta oscillation in the hippocampus elicited by brainstem-stimulation compared to wild-type mice. However, it did not affect hippocampal theta-phase gamma-amplitude coupling significantly, since over a 60% reduction was found in coupling in Tg2576 mice regardless of TREM2 function.

Conclusion: Our findings indicate a role for TREM2-dependent microglial function in the hippocampal neuronal excitability in both wild type and $A\beta$ overproducing mice, whereas

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deficiency in TREM2 function exacerbates disruptive effects of $A\beta$ on hippocampal network oscillations.

Keywords

TREM2; microglia; hippocampus; seizure; theta oscillation; Alzheimer's disease

Introduction

The prevalence of Alzheimer's disease (AD), a devastating neurodegenerative disorder, is increasing worldwide and there is not yet any approved disease-modifying therapeutic that would affect disease progression [1]. Amyloid pathology has long been the major focus of drug development, since the spread and accumulation of amyloid- β (A β) in soluble misfolded oligomeric forms and in in extracellular deposits are thought to be the upstream initiating factor of the disease. However, clinical trials targeting A β have yielded disappointing results leading to increased efforts to identify new disease drivers and their possible interactions with the canonical AD hallmarks A β and hyperphosphorylated tau [2].

Evidence of the critical contribution of microglial activation in neuronal dysfunctions and cell death has fundamentally shifted our understanding of the pathogenesis of several neurodegenerative diseases, including AD [3, 4]. Microglia, the resident innate immune cells of the brain, have a pivotal role in providing homeostatic control by clearance of apoptotic debris and misfolded proteins, secretion of neurotrophic factors, inflammatory response to injury, and remodeling neuronal connections [5]. Recent experimental studies also demonstrated that microglia are involved in regulation of higher brain functions and behavior through modulation of neuronal activity [6, 7]. Unfortunately, the most of these important microglia functions can significantly deteriorate as a result of genetic mutations or ageing process [8–10]. Genome-wide association studies have linked coding variants in triggering receptor expressed on myeloid cells-2 (TREM2), which is exclusively found on microglia in the brain, with increased risk for AD, frontotemporal dementia and Nasu-Hakola disease characterized with progressive presenile dementia and bone cysts [11-13]. In particular, naturally occurring missense variants p.R47H and p.R62H that confer a partial loss of function of TREM2 has been implicated in developing late onset AD [14]. Interestingly, carriers of these TREM2 variants also showed faster rate of cognitive decline compared with non-carriers [15]. In addition, patients with a rare homozygous mutations p.Thr66Met in TREM2 had early onset frontotemporal-like dementia, with hippocampal deficits and presence of epileptic seizures [16].

To investigate genotype-phenotype correlation between TREM2 and AD, transgenic mouse lines modeling AD pathology with partial or total TREM2 deficiency were generated. In these mice, it has been observed that TREM2 haploinsufficiency caused a strong reduction in the number of microglia surrounding plaques [17], with decreased plaque compaction and severe axonal dystrophy, similarly to results from postmortem brain samples of AD patients with TREM2 mutation [18]. Interestingly, total deficiency of TREM2 signaling was shown to affect amyloid pathology in transgenic APP/PS1 mice differently depending on the stage of disease progression, as it reduced amyloid accumulation at early stage and intensified

amyloid accumulation at late stage of disease [19]. In the context of tau pathology, findings on TREM2 deficiency are also discordant. Decreased TREM2 function exacerbates tau hyperphosphorylation [20] and facilitates seeding and spreading of tau aggregates around A β plaques in the early phase of AD [21, 22], but during the later symptomatic phase appears to slow down tau-mediated neurodegeneration, as seen in TREM2 p.R47H variant carriers [23]. Although these studies have implicated distinctive TREM2-dependent microglial role in AD at the cellular level, data on its functional relevance are largely missing.

In the present neurophysiological study, we assessed hippocampal network oscillations in amyloid- β overproducing Tg2576 transgenic mice with normal and deficient TREM2 function with aim to elucidate potential contribution of this immunoreceptor on specific oscillatory dynamic changes associated with AD pathophysiology.

Methods

Mice

Transgenic Tg2576 (B6;SJL-Tg(APPSWE)2576Kha) hemizygous mice and their wild-type controls were obtained by Biogen through a Research Crossbreeding agreement with Taconic Biosciences Inc. (Germantown, NY). The Tg2576 mice model AD pathology by overexpressing a human amyloid precursor protein (isoform 695) with the Swedish mutation (KM670/671NL), which resulted in high levels of A β , extensive development of amyloid plaques, and cognitive deficits [24]. These mice were crossed with TREM2 deficient mice to yield Tg2576;TREM2 genotypes. Tg2576(Hemi);TREM2(WT/Het/Hom) experimental cohorts were generated by breeding Tg2576(Hemi);TREM2(Het) mice with Tg2576 (WT);TREM2(Het) mice and were maintained on a mixed B6/SJL background. TREM2 deficient mice were generated from Trem2tm1(KOMP)Vlcg founders obtained from UC Davis KOMP Repository and maintained on a C57BL/6 mice background. In these mice, a LacZ reporter cassette replaces the entire coding sequence of the endogenous Trem2 locus, resulting in a loss of TREM2 function [25]. Breeding was carried out at Biogen and The Jackson Laboratory (Bar Harbor, ME). All animals were littermates to ensure that genetic background and environment were homogeneous. For experiments, mice cohorts were transferred to Yale University where they were housed in Yale Animal Resources Center managed facility, in a temperature and humidity-controlled room with 12:12-h lightdark cycle and free access to food and water at all times.

A total of 65 male mice were used in the study (Tg2576(WT)TREM2(WT)

= 13; Tg2576(WT)TREM2(Het) = 9; Tg2576(WT)TREM2(Hom) =

12; Tg2576(Hemi)TREM2(WT) = 13; Tg2576(Hemi)TREM2(Het) = 9;

Tg2576(Hemi)TREM2(Hom) = 9). All animals were at the age between 9 and 12 months. Neurophysiological recordings were done after acclimation period of at least 30 days, with all measures taken to minimize pain or discomfort of animals. All procedures were performed in accordance with National Institutes of Health guidelines (NIH publications No. 86–23, revised 1996) and were approved by the Institutional Animal Care and Use Committees of each respective institution.

Neurophysiological recordings

Hippocampal neuronal network integrity was tested using brainstem stimulation-induced theta oscillation paradigm, which was employed in previous neurophysiological studies of transgenic rodent models of AD pathology [26–29].

Briefly, mice were anesthetized with urethane (1.5 g/kg) given intraperitoneally and placed in a Kopf stereotaxic frame (Tujunga, CA) on a temperature-regulated heating pad set to maintain body temperature at 37°C (Physitemp Instruments Inc., Clifton, NJ). After achieving surgical plane of anesthesia which was controlled by the absence of the tail pinch reaction, left parietal craniotomies were performed over the hippocampus and rostral brainstem for insertion of the two concentric stainless-steel bipolar electrodes (NE-100X, Rhodes Medical Instruments, Woodland Hills, CA). For recording of local field potentials (LFPs), one electrode was placed in the dorsal hippocampal CA1 region, while another was lowered in the nucleus pontis oralis (nPO) for electrical stimulation. Coordinates for both target areas were determined relative to bregma and brain surface and were as follows: CA1, anteroposterior -2.0 mm, lateral 1.5 mm, and dorsoventral 1.5 mm; nPO, anteroposterior -4.0 mm, lateral 1.2 mm, and dorsoventral 3.3 mm [30]. Following placements of the electrodes, animals remained in the stereotaxic frame for LFPs recording which began after stabilization period of at least 30 minutes. Each recording lasted for approximately 60 minutes, during which time initial dose of urethane was sufficient to provide stable plane of anesthesia [31]. At the end of each recording, mice were sacrificed, brain tissue collected and histologically processed for verification of electrodes positions. The nPO stimulation protocol consisted of a train of 0.3 ms square pulses delivered over a period of 6 seconds at a rate of 250 Hz, which was provided by Isoflex stimulus isolator unit (A.M.P.I. Instruments, Jerusalem, Israel) and repeated every 100 seconds. The stimulation-response relationship was determined in each animal using 3 cycles of increasing current in steps of 20 µA, with power and frequencies averaged for each stimulation intensity. For quantitative analyses, LFPs were amplified by an A-M System (Carlsborg, WA) with filters set between 1 Hz and 500 Hz, digitized at a rate of 2 kHz, and collected on a computer via a CED Micro1401-3 interface and Spike2 software (Cambridge Electronic Design, Cambridge, UK).

Stimulation-induced hippocampal oscillation episodes were analyzed for total theta power and peak theta frequency using a fast Fourier transform in Spike2 at a spectral resolution of 0.24 Hz. Total theta power was calculated by summing power values between 3 Hz and 9 Hz, and peak frequency was identified by determining where the peak power occurred. Estimating phase-amplitude coupling between theta and gamma oscillations, a neural signature of coordinated memory formation and storage, was performed using custom scripts written in Matlab (Mathworks, Natick, MA). All offline filtering was done using eegfilt.m from EEGLAB toolbox [32]. To avoid possible stimulation artefacts, the first second of each 6-second-long stimulation episode was omitted in all analyses. Seizures were identified visually by sudden large positive and/or negative deflections of hippocampal oscillations by neurophysiologist during EEG recordings, and confirmed during off-line analysis by a computational neuroscientist as instantaneous increase in spontaneous LFPs wideband power (4 z-score difference for 2 s or longer). The strength of theta-gamma coupling was assessed using modulation index (MI) as described previously [28, 33, 34].

Specifically, using the Hilbert transformation we extracted the instantaneous phase in LFP signal containing oscillations from 3 to 10 Hz (phase frequencies), and the amplitude envelope in signal containing oscillations from 20 to 100 Hz (amplitude frequencies). For each animal an MI value was estimated for each frequency pair composed by one slow and one fast oscillation of the LFPs recorded during stimulation with current intensity inducing 50% of the maximal theta oscillation power in the stimulation-response curves. An important characteristic of this metric is that it is not influenced by the absolute amplitude of theta or gamma oscillations, but depends on the relative changes of the gamma oscillation envelope as a function of the theta oscillation phase [33]. To generate the heat-maps, the MI values over 3 stimulations from each animal in a group were averaged and expressed using a color-coded scale.

Statistical analysis

All data were initially determined to be suitable for parametric analysis according to normality and homoscedasticity using the Shapiro-Wilk and Brown–Forsythe tests, respectively. Statistical analyses were performed by 1-way ANOVA with Tukey's post hoc test for assessing stimulus-response relationship in elicited theta power or peak frequency, or with Dunnett's post hoc test for comparisons of average MI values between the groups (GraphPad Prism Software, Inc, La Jolla, CA). Data are expressed as the mean \pm standard error of the mean (SEM). Differences were considered significant when P < 0.05.

Results

Behavioral observations

Inspecting the animals before neurophysiological experiments revealed a striking pathologic self-grooming in individually housed TREM2 homozygous and heterozygous mutant mice. Due to excessive grooming behavior these mice displayed large hairless patches on their lateral body surface (Fig. 1A). Although the extent of severe fur loss varied between these animals, close physical examination of the affected area by researcher and attending veterinarian from animal facility did not find any significant skin ulceration or inflammation that would have required particular treatment. The similar signs of compulsory over-grooming behavior were not detected in corresponding wild-type or transgenic Tg2576 mice with intact TREM2 function.

Epileptiform brain activity

TREM2 homozygous and heterozygous mutant mice had high susceptibility to spontaneous epileptic seizure activity. During hippocampal EEG recordings we observed in some of these mice suddenly occurring episodes of synchronous large amplitude spike discharges lasting several seconds (average duration 15.61 ± 1.56 s) and followed by a depression of neuronal field potentials (Fig. 1B). These epileptiform seizures were present in 44% of Tg2576(WT)TREM2(Het) (4 out of 9 mice), 33% of Tg2576(WT)TREM2(Hom) (4 out of 12 mice), and 22% of either Tg2576(Hemi)TREM2(Het) or Tg2576(Hemi)TREM2(Hom) (2 out of 9 mice, in each) groups (Fig. 1C). In contrast, no spontaneous paroxysmal spiking discharges were seen in the hippocampus of mice with intact TREM2 function, regardless of

the presence of Tg2576(Hemi) and Tg2576(WT) genotypes. These results indicate increased neuronal excitability only in mice with impaired TREM2 function.

Changes in hippocampal oscillations

Evaluation of elicited hippocampal theta oscillation and theta-phase gamma-amplitude coupling in rodents under anesthesia was shown to provide valuable insights into hippocampus network functioning related to memory and cognitive processes in behaviorindependent manner [35, 36]. Therefore, similarly to our previous studies, we applied high-frequency electrical stimulation of the brainstem nPO to induce regular theta oscillation (3–9 Hz) in the hippocampal CA1 region with a current-dependent increase of peak frequency in urethane anesthetized mice (Fig. 2A). In these experiments analyses were performed only in animals in which hippocampal epileptiform activity did not develop during baseline recording or at any stage of brainstem stimulation thus potentially resulting in instability of neurophysiological activity. Quantitative input-output analyses of elicited hippocampal oscillation showed a significant difference in absolute theta power ($F_{(5,48)}$ = 16.55; P < 0.001), but not peak theta frequency $(F_{(5,48)} = 0.51; P = 0.769)$ in response to varying stimulus intensities between the groups (Fig. 2B,D). Further AUC analysis with post hoc comparisons for theta power changes (stimulation intensities 0.02 -0.12 mA) showed a marked reduction in theta power in Tg2576(Hemi)TREM2(Het) and even more significant decrease in Tg2576(Hemi)TREM2(Hom) mice when compared with control mice (Tg2576(WT)TREM2(WT) vs. Tg2576(Hemi)TREM2(Het), P = 0.047; Tg2576(WT)TREM2(WT) vs. Tg2576(Hemi)TREM2(Hom), P = 0.003, Tukey's post hoc test), indicating that deficiency of TREM2, and particularly its elimination leads to disruption of theta oscillation in the hippocampus of transgenic AB overproducing mice (Fig. 2C).

In further assessment of oscillatory dynamics in the hippocampus during stimulation, thetaphase gamma amplitude coupling was measured in subsets of wild-type and transgenic Tg2576 mice with intact and total TREM2 deletion. In line with our previous studies, a strong coupling between phase of theta and amplitude of low gamma (30–60 Hz) was observed in Tg2576(WT)TREM2(WT) mice, whereas in other groups of mice this coupling was attenuated (Fig. 3A). Comparing the degree of theta-phase gamma-amplitude coupling calculated as modulation index (MI) significant difference was found among groups ($F_{(3,26)}$ = 3.65, P = 0.026). Further post hoc analysis using Dunnett's test showed significantly lower coupling in Tg2576(Hemi)TREM2(WT) mice (P = 0.021) and Tg2576(Hemi)TREM2(Hom) mice (P = 0.048) comparing to wild-type controls (Tg2576(WT)TREM2(WT), (Fig. 3B). TREM2 deficiency in Tg2576(WT)TREM2(Hom) mice indicated a trend to reduced coupling, but it did not reach a significant level.

Discussion

Microglial TREM2-mediated signaling impairment has been proposed as a risk factor for the development and progression of AD, yet its contribution to different aspects of disease-related neuronal dysfunctions remains ambiguous. Here we provide in vivo evidence that TREM2 deficiency increases susceptibility for spontaneous epileptiform seizures in

the hippocampus of both control and A β overproducing Tg2576 mice. TREM2 deficiency also significantly aggravates compromised hippocampal theta oscillation activity in these transgenic AD mice but without further influencing their already markedly diminished theta-phase gamma-amplitude coupling as compared to control counterparts.

The epileptiform activity has been recognized as an important clinical entity of AD, since several retrospective, prospective and cross-sectional studies identified a significantly higher incidence of unprovoked seizures in AD patients than in the age-matched control subjects [reviewed in 37, 38]. Furthermore, these seizures were found to be predominantly nonconvulsive with focal temporolimbic localization, occurring even at an early, prodromal stage of AD [39, 40], and more frequently in male patients [41]. Importantly, evidence suggests that increased subclinical epileptiform activity in AD patients, which is indicative of neuronal hyperexcitability, might contribute to their accelerated cognitive decline [38]. Similar network hyperexcitability and spontaneous non-convulsive seizures were also detected in several transgenic models of AD with overt A β expression in the brain [29, 42, 43] after chronic cortical and hippocampal EEG monitoring. Strikingly, in this study, we found a higher propensity to epileptiform activity in both partial and total TREM2 deficient mice regardless of AB pathology. Intact TREM2 function is required for microglia-mediated remodeling of synaptic connections by phagocytic elimination of supernumerary dendritic spines, which is an important process for the maturation of neural networks and maintenance of delicate balance between excitation and inhibition during development and in adulthood [5]. Accordingly, it has been shown that mice with ablated TREM2 on microglia have a higher density of dendritic spines due to suppressed phagocytic capacity and consequently defective synapse pruning [44]. Since spine synapses closely correspond to the excitatory synapses [45], an increase in their number can induce an excitatory gain offsetting the balance between excitation and inhibition in the complex neuronal networks, thus modulating neurotransmission and behavioral responses. Recent studies revealed a spectrum of abnormalities in TREM2 deficient mice including enhanced excitatory neurotransmission and reduced synaptic plasticity in the hippocampus, impaired long-range functional connectivity between cortical and hippocampal regions, and behavioral alterations such as excessive self-grooming, a trait typically associated with animals' increased anxiety and perseveration [44, 46, 47]. This TREM2-dependent amplification of excitatory input in the network due to microglial failure to prune synapses may also lower the threshold for seizure activity facilitating spontaneous epileptogenesis as observed here. Indeed, genetically determined defects in microglial elimination of excessive excitatory synapses in the neocortex and hippocampus during development [48, 49], or brain-wide pharmacological depletion of microglial phagocytosis in adult mice [6] both resulted in increased susceptibility to behavioral seizures. Our finding of a high rate of seizures among TREM2 haploinsufficient mice suggests that even a partial deficit in microglial TREM2 function is enough to induce significant perturbation in this mechanism of controlling synaptic balance.

Over the course of AD development, aberrant network excitability has been linked to $A\beta$ pathology [50]. Recent data suggest that this dysfunction is likely initiated by preexisting neuronal activity, stemming from impaired baseline excitatory transmission, which triggers a vicious cycle where the soluble $A\beta$ species induce hyperexcitability before the emergence

of plaques [51]. This hyperexcitability not only increases the potential for epileptic seizures but might also exacerbate AD pathology by increasing diffusion and deposition of AB [52], and impairing the function of microglia [53]. It has been demonstrated that neuronal hyperactivity-evoked widespread release of adenosine triphosphate (ATP) may damage microglial phagocytosis and apoptotic debris clearance, and enhance proinflammatory cytokines secretions [53]. Interestingly, new evidence showed that microglial sensing of elevated extracellular ATP can in turn activate negative feedback mechanism to suppress neuronal activity via adenosine [6], further remodeling circuit function. In this view, an increase in TREM2 that parallels with the rise of A β levels seen in transgenic mice [12], and at early AD stage in patients [54] might have the compensatory function to prevent or mitigate alterations due to this initial detrimental events. However, during disease progression microglia in response to oligometric and plaque A β challenge tend to undergo substantial ultrastructural changes leading to aberrant activation involved in Aβ-mediated neuronal dysfunctions and pathologic loss of synapses in cognitive-related brain regions [55, 56]. This aligns with the notion of the negative impact of impaired TREM2 signaling on Aβ pathology during the late stage of AD [14, 19].

Neurophysiological studies revealed a close association between profound alterations in EEG signals and cognitive dysfunctions in AD patients and transgenic animals carrying pathological genes for AB overproduction and tau hyperphosphorylation [57]. Specifically, generalized EEG slowing with spectral profile shift to the low frequencies has been linked to cognitive deterioration in patients [58, 59], whereas in transgenic models with demonstrated learning and memory deficits, multiple oscillatory alterations in cortical and hippocampal networks have been identified [29, 42, 43]. Previously, we reported several distinctive hippocampal oscillatory abnormalities that clearly reflected AD-related impairment of functional network integrity, such as the reduction in theta band power and weakening of theta-phase gamma-amplitude coupling in transgenic A β overproducing APP/PS1 and 5xFAD mice [26, 28] and TgF344-AD rats [29], as well as tau Tg4510 mice [27] recorded under urethane anesthesia. We also observed that these oscillatory dysfunctions are age-dependent and directly correlate with plaque load in the hippocampus [26, 29]. Hippocampal theta oscillation has been linked to cognitive processes, and the correlation between hippocampal theta activity and learning and memory has been indicated in both experimental animals and humans [60-63]. Hence, it has been proposed that theta oscillation could serve as a surrogate measure to quantify the cognitive abilities that depend on hippocampal function, but also to predict cognitive effects of drugs [35, 63]. Accordingly, pharmaco-EEG experimental studies showed that drugs currently in use for symptomatic AD treatment, donepezil and memantine, both increase the power of elicited hippocampal theta oscillation [64, 65] in anesthetized rodents. Further, the coupling between the phase of theta and amplitude of gamma oscillations in the hippocampus has been consistently shown to implicate effective memory encoding and consolidation in humans [66] and experimental animals [33, 67, 68]. In line with these findings, multiple studies found a reduction in hippocampal theta-gamma coupling in relation to AD pathology [28, 29, 69, 70]. Consistent with these observations, here we find that Tg2576 mice also showed compromised elicited theta oscillation in response to brainstem stimulation, and importantly that this deficit is significantly aggravated in animals with deficient TREM2 function. Moreover, we

observed that transgenic Tg2576 mice displayed weakened hippocampal theta-phase gamma amplitude coupling as compared to their littermate controls, but this impairment contrary to theta power was not dependent on TREM2 function. Therefore, current findings implicate microglial TREM2 important role in slowing down certain robust aspects of AD-related oscillatory impairments such as hippocampal theta power decline, but not in mitigating other more subtle oscillatory abnormalities such as a reduction in theta-gamma coupling that depends on fine-tuned interaction between multiple neurotransmitter systems and cell types, and thereby is more vulnerable to disease pathology.

It is important to note that all neurophysiological recordings in this study were performed under urethane anesthesia, which may have confounding influences on reported findings. Even though urethane does not markedly affect neurotransmission inducing spontaneous brain state alterations that resemble physiological sleep [31], it has been shown that subthreshold excitability in the hippocampus can be compromised given its suppressive effects on neuronal spike activity and spike synchrony therein [71, 72]. This might be the reason why we haven't observed EEG epileptiform activity in Tg2576 mice with intact TREM2 function unlike previous studies when this transgenic model was analyzed in nonanesthetized condition [42, 73]. Nevertheless, our results of high seizure susceptibility in hippocampal EEG of Tg2576 TREM2 deficient mice, even during urethane anesthesia, indicate that diminished TREM2 function leads to enhanced neuronal excitability and aggravated A β -induced hippocampal oscillatory deficit. Studies on mice with TREM2 deficiency could contribute to our understanding the pathological role of microglia function in heightened neuronal excitability in its contribution to AD disease progression, and potentially identify novel therapeutic targets.

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Fig.1.

Pathological behavior and increased seizure susceptibility in control and Tg2576 mice with partial or complete TREM2 loss of function mutations. A) Photographs showing large patches of missing hair caused by excessive self-grooming of individually housed TREM2 homozygous mutants crossed with transgenic AD mice (Tg2576(Hemi); upper panel) and their littermate wild-type controls (Tg2576(WT); lower panel). The same compulsive hair removal feature was found in TREM2 heterozygous mice of either genotype. B) Typical EEG recording of spontaneous epileptiform activity in the hippocampal CA1 region of TREM2 deficient mice under anesthesia. Trace shows aberrant large amplitude spiking discharges from Tg2576(WT)TREM2(Het) mouse lasting over 15 seconds. Enlarged segment of the trace (marked in red) is at the bottom. C) Distribution of susceptibility to epileptic seizures between different genotype groups of mice. In wild-type controls, seizures were detected in 44% of TREM2 heterozygous and 33% of TREM2 heterozygous and homozygous groups. Scale bar: 0.02 mV, 1 s (upper trace); 0.02 mV, 0.2 s (lower trace).

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Fig.2.

Disruption of elicited hippocampal oscillations in amyloid- β overproducing Tg2576 mice with partial or complete TREM2 loss of function mutations. A) High-frequency electrical stimulation of brainstem nucleus pontis oralis (nPO) induces highly regular theta oscillations in the hippocampal CA1 region of anesthetized mice, characterized by relatively constant frequency and spectral content as shown in the time-frequency decomposition map (horizontal black bar under the map marks the period of stimulation; the color-coded scale denotes EEG power). B-D) Stimulus-response relationship for absolute theta power and peak frequency over increasing stimulus intensities, and area under the curve (AUC) analyses showed significant decline in elicited theta power (B,C), but not peak theta frequency (D) in transgenic Tg2576 mice with both TREM2 heterozygous and homozygous mutations compared to respective wild-type controls. (*P < 0.05, **P < 0.01, 1-way ANOVA with Tukey's post hoc test). Scale bar: 0.2 mV, 1 s.



Fig.3.

Reduction of phase-amplitude coupling between hippocampal theta and gamma oscillations in amyloid- β overproducing Tg2576 mice with both intact and deficient TREM2 function. Theta phase-gamma amplitude coupling during nPO-stimulation induced oscillations in hippocampus of anesthetized mice is calculated as Modulation Index (MI). A) Heat-maps with averaged theta-gamma coupling over all animals in each group indicate the strength of theta phase-nested distribution of gamma power (30–60 Hz). B) Comparison of average MI between groups revealed significantly decreased coupling in transgenic Tg2576 mice, regardless whether TREM2 was present or knock-out, compared to controls. (*P < 0.05, 1-way ANOVA with Dunnett's post hoc test).