

TrkB signaling in parvalbumin-positive interneurons is critical for gamma-band network synchronization in hippocampus

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Although brain-derived neurotrophic factor (BDNF) is known to regulate circuit development and synaptic plasticity, its exact role in neuronal network activity remains elusive. Using mutant mice (TrkB-PV^{-/-}) in which the gene for the BDNF receptor, tyrosine kinase B receptor (*trkB*), has been specifically deleted in parvalbumin-expressing, fast-spiking GABAergic (PV+) interneurons, we show that TrkB is structurally and functionally important for the integrity of the hippocampal network. The amplitude of glutamatergic inputs to PV+ interneurons and the frequency of GABAergic inputs to excitatory pyramidal cells were reduced in the TrkB-PV^{-/-} mice. Functionally, rhythmic network activity in the gamma-frequency band (30–80 Hz) was significantly decreased in hippocampal area CA1. This decrease was caused by a desynchronization and overall reduction in frequency of action potentials generated in PV+ interneurons of TrkB-PV^{-/-} mice. Our results show that the integration of PV+ interneurons into the hippocampal microcircuit is impaired in TrkB-PV^{-/-} mice, resulting in decreased rhythmic network activity in the gamma-frequency band.

gamma oscillations | synaptic transmission | Cre recombinase | dendrite | slice

Tyrosine kinase B receptor (TrkB), the cognate receptor for brain-derived neurotrophic factor (BDNF) and neurotrophin-4, mediates key signaling events that control many aspects of neuronal development and function (1–4), including the maturation of parvalbumin-positive (PV+) interneurons in the hippocampal microcircuit. BDNF is preferentially synthesized in, and secreted from glutamatergic neurons, whereas *trkB* is expressed in both glutamatergic and γ -aminobutyric acid (GABA)-ergic neurons in hippocampus (5). Among cortical interneurons, PV+ interneurons express *trkB* abundantly (6). This anatomical organization of the BDNF signaling components and the known importance of feedback and feedforward communication between principal cells and interneurons (7–11) suggest a potential role for TrkB signaling in modulating neuronal network function.

Rhythmic activity in cortical networks is important for the formation of neuronal assemblies (12–14). Of particular interest is rhythmic network activity in the gamma-frequency band (30–80 Hz, gamma oscillations) (15–17). Gamma oscillations are a result of the synchronized electrical activity of the neurons within a network and are thought to be important for temporal encoding, binding of sensory features, and memory storage and retrieval (18–22). Moreover, gamma oscillations are altered in several brain disorders, such as Alzheimer's disease (23–25), schizophrenia (24, 26–31), and epilepsy (24, 32, 33). Gamma oscillations are exquisitely susceptible to modulation of the cellular and synaptic mechanisms underlying the rhythmic activity. Fast-spiking PV+ interneurons are the main recipient of recurrent glutamatergic innervations in the hippocampal circuitry, and their role in gamma-frequency synchronization in cortical

and hippocampal networks is well-established (34, 35). Despite extensive knowledge of the role of BDNF-TrkB signaling in neuronal development and synaptic function, the question of how BDNF might modulate and control rhythmic activity has so far not been answered.

To determine whether BDNF regulates synchronized network activity through PV+ interneurons (36, 37), we used a mutant mouse line, in which the *trkB* gene has been selectively ablated in PV+ interneurons. By inducing gamma oscillations in the hippocampal slice preparation with the muscarinic acetylcholine receptor agonist carbamoylcholine (CCh), we found that the power of gamma oscillations was dramatically reduced in the hippocampal area CA1 of mutant mice compared with control (Ctr) littermates. Morphological analysis and concomitant whole-cell patch-clamp and extracellular field recordings were used to determine, whether the decrease in gamma oscillation power was caused by morphological and/or functional alterations in PV+ interneurons. We also showed that the removal of TrkB signaling leads to a desynchronization and overall reduction in frequency of action potentials generated in PV+ interneurons, which is consistent with a reduction in gamma oscillation power. Taken together, our results demonstrate a critical role of BDNF-TrkB signaling in PV+ interneurons for hippocampal network synchrony in the gamma-frequency band.

Results

Ablation of the *trkB* Gene in PV+ Interneurons Using the PV-Cre Transgene. The calcium-binding protein PV serves as a marker of a subpopulation of GABAergic neurons with fast-spiking properties (38). To determine whether, and at what time point, PV+ interneurons express *trkB* during hippocampal development, we performed PV immunohistochemical staining on sections from *TrkB^{LacZ/+}* knockin mice, in which β -galactosidase (β -gal) recapitulates the expression pattern of *trkB* (39). We found that the CA1 region starts to express *trkB* before postnatal day (P) 7 and PV between P7 and P14 (Fig. 1*A* and *B*). In this brain region, *trkB* was expressed in nearly all PV+ interneurons (54 of 56) at P14 (Fig. 1*B*) and in all PV+ interneurons at P21 (Fig. 1*C*; 37 of 37) and at P28 (Fig. 1*D*; 24 of 24). Thus, CA1 PV+ interneurons express *trkB* during postnatal development and in adulthood.

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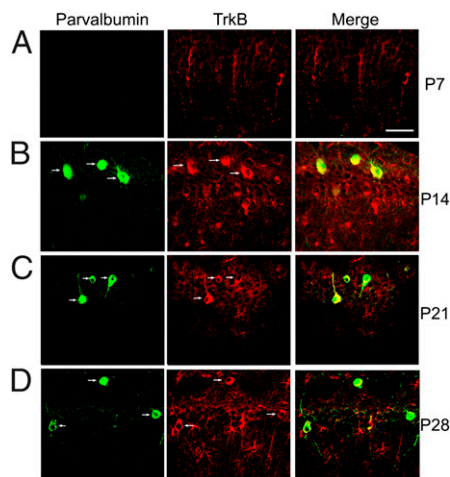


Fig. 1. Colocalization of TrkB and parvalbumin in the hippocampal CA1 region. Expression patterns of TrkB and parvalbumin in the hippocampal CA1 region were revealed by immunohistochemistry against parvalbumin and β -galactosidase in *TrkB^{LacZ/+}* mice at P7 (A), P14 (B), P21 (C), and P28 (D). Arrows denote neurons expressing both parvalbumin and TrkB. (Scale bar: 50 μ m.)

We used a *Cre* transgene driven by the PV promoter (*PV-Cre*) to delete the *trkB* gene in PV-expressing cells. To determine the efficiency of gene deletion mediated by *PV-Cre* in PV+ interneurons, we introduced this *PV-Cre* transgene into *Rosa26* reporter (*R26R*) mice, in which the *Rosa26* locus expresses β -galactosidase once Cre-mediated recombination has occurred (40). In *R26R/+;PV-Cre* mice, β -gal was expressed in the majority (22 of 31) of PV+ interneurons in hippocampal area CA1. Importantly, the Cre recombination activity was strictly confined to PV+ interneurons, because we did not detect any neurons that expressed β -gal but not PV (Fig. 2A).

We then crossed *PV-Cre* with floxed *trkB* (*fB*) mice to generate Ctr mice (*fB/fB*) and conditional *trkB* knockout mice (*fB/fB;PV-Cre*,

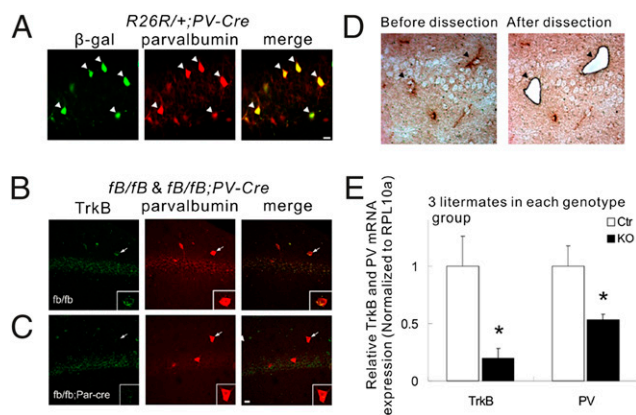


Fig. 2. Specific ablation of TrkB receptor in parvalbumin-positive interneurons in the hippocampal CA1 area. (A) Cell type-specific recombination of the *Rosa26* reporter (*R26R*). The floxed *R26R* allele recombination was detected in the vast majority of PV+ neurons (white arrowheads) at P28, as revealed by immunohistochemistry against PV and β -galactosidase in *R26R/+;PV-Cre* mice. (B and C) Dual fluorescent in situ hybridization with probe for TrkB mRNA and immunohistochemistry with antibody against PV were carried out on brain sections from control mice (*fB/fB*) and *trkB* KO mice (*fB/fB;PV-Cre*). White arrows indicate the same neurons. Individual neurons are magnified in *Insets*. (D) Images of the CA1 region before and after LCM. The section was briefly stained with antibodies against PV (dark arrowheads). (E) qPCR using the mRNA isolated from the captured cells. Data are presented as means \pm SEM. * $P < 0.05$. (Scale bars: 20 μ m.)

also called *TrkB-PV^{-/-}*). Two approaches were used to determine whether the *trkB* gene was eliminated in PV+ interneurons of *TrkB-PV^{-/-}* mice. First, a double-staining experiment was performed to detect TrkB mRNA by in situ hybridization (ISH) using fluorescence-labeled antisense oligonucleotides as a probe, and PV immunoreactivity by immunohistochemistry using an anti-PV antibody. We found that although many PV+ interneurons expressed TrkB mRNA in Ctr mice (Fig. 2B), there was hardly any TrkB signal in PV+ interneurons in *TrkB-PV^{-/-}* mice (Fig. 2C). Importantly, *trkB* ISH signals were detectable in PV-negative cells, suggesting that the deletion of *trkB* gene was selective for PV+ interneurons. For the second approach, laser capture microdissection (LCM) was used to capture PV+ interneurons in hippocampal sections, followed by quantitative reverse-transcriptase PCR (qPCR) to measure TrkB mRNA levels in the captured tissue samples (Fig. 2D). We show that TrkB mRNA in CA1 PV+ interneurons isolated from *TrkB-PV^{-/-}* mice was reduced to $19.7 \pm 8.4\%$ compared with those isolated from Ctr mice ($P < 0.05$; Fig. 2E). Together, these two approaches demonstrate a selective ablation of *trkB* gene in CA1 PV+ interneurons of *TrkB-PV^{-/-}* mice. Interestingly, there was a $47.1 \pm 4.9\%$ reduction in the level of PV mRNA in CA1 PV+ interneurons isolated from *TrkB-PV^{-/-}* mice compared with those from Ctr mice ($P < 0.05$; Fig. 2E), suggesting a reduction in PV expression even in PV+ cells.

Reduction in the Density of PV+ Interneurons and Their Dendrites. In agreement with a previous study (41), the vast majority of PV+ interneurons were found in the strata pyramidale (SP) and oriens (SO) of the hippocampal CA1–3 regions. We counted the number of PV+ interneurons within SO and SP of the CA1 area at different ages. The density of PV+ interneurons in the CA1 region was not different in *TrkB-PV^{-/-}* mice versus Ctr mice at P14. However, the density was reduced by 16.3% at P28 (from 0.0295 to 0.0247 cell per μ m³; $P < 0.05$) and by 19.2% at 6 wk of age (from 0.0285 to 0.023 cell per μ m³; $P < 0.01$) (Fig. 3A and B).

The dendritic morphology of PV+ interneurons, as measured by the density of PV+ neuropil, was examined in the CA1 region in P28 Ctr and *TrkB-PV^{-/-}* mice after staining with the PV antibody. The dendrites of PV+ interneurons exhibited a $14.6 \pm 5.9\%$ reduction in length in strata radiatum (SR) ($P < 0.05$) and a $19.8 \pm 4.9\%$ reduction in SP ($P < 0.05$) in *TrkB-PV^{-/-}* mice, respectively, compared with Ctr mice (Fig. 3C and D; 4 littermates in each group). Taken together, deletion of the *trkB* gene elicited a modest effect on the dendrites and the number of PV+ interneurons in the hippocampal area CA1.

Decreased Excitatory Inputs to PV+ Interneurons and Inhibitory Inputs to Principal Neurons. The consequences of *trkB* ablation on the glutamatergic synaptic input onto PV+ interneurons were determined by recordings of miniature excitatory postsynaptic currents (mEPSCs). First, the PV+ interneurons were characterized based on their fast-spiking discharge pattern exhibited upon depolarization induced by current injection (Fig. 4A and B) (38). The resting membrane potential and the number of depolarization-induced repetitive action potentials (AP) did not show a significant difference between Ctr and *TrkB-PV^{-/-}* mice (Fig. 4B). Spontaneous mEPSCs were recorded in the fast-spiking interneurons in the presence of 2 μ M tetrodotoxin (TTX) (Fig. 4C). The mEPSC amplitude in *TrkB-PV^{-/-}* mice (17.61 ± 2.62 pA; $n = 13$ cells from 13 slices in six mice) was significantly reduced ($P < 0.01$) compared with Ctr mice (24.31 ± 4.66 pA; $n = 12$ cells from 12 slices in six mice) (Fig. 4D). However, the ablation of *trkB* did not seem to alter mEPSC frequency. Thus, the frequency of mEPSCs was 13.11 ± 4.22 Hz for Ctr and 16.4 ± 4.96 Hz for *TrkB-PV^{-/-}* mice ($P = 0.078$; Fig. 4E). These results suggest that the ablation of *trkB* in PV+ interneurons pre-

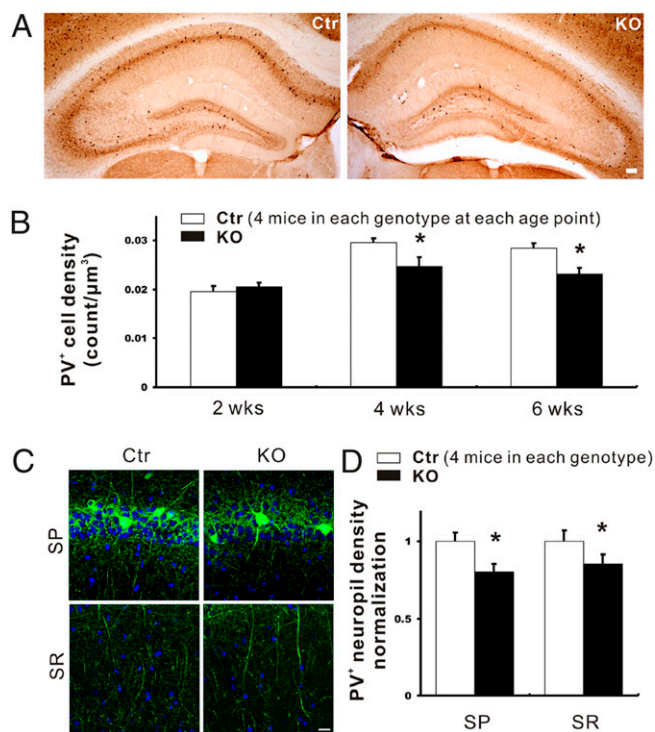


Fig. 3. PV+ cell number and neuropil density in the hippocampal CA1 area of conditional *TrkB* knockout (*TrkB-PV^{-/-}*, or KO) and control (Ctr) mice. (A) Representative images showing PV immunoreactivity in hippocampi of 4-wk-old Ctr and KO mice. (Scale bar: 100 μm.) (B) Density of PV+ cells in the CA1 regions of Ctr and KO mice at 2, 4, and 6 wk of age. (C) Representative fluorescence images showing immunoreactive neuropils in SP and SR of the CA1 area. (Scale bar: 50 μm.) (D) Density of PV-immunoreactive neuropil in the CA1 area. Data are presented as means ± SEM. **P* < 0.05.

dominantly altered the postsynaptic AMPA receptor function rather than their presynaptic glutamatergic inputs (42).

We next asked whether the outputs of the PV+ interneurons in the *TrkB-PV^{-/-}* mice were impaired. Spontaneous miniature inhibitory postsynaptic currents (mIPSCs), which reflect inhibitory inputs from GABAergic interneurons, including PV+ interneurons, were recorded in pyramidal neurons identified by their shape under an infrared-differential interference contrast microscope (43). All recordings were performed within the SP layer in the presence of 2 μM TTX, 25 μM 2R-amino-5-phosphonovaleric acid (Apv), and 25 μM 6-cyano-7-nitroquinoxaline-2,3-dione (CNQX) (Fig. 4F). The mIPSC amplitude showed no significant difference between Ctr mice (42.31 ± 11.32 pA; *n* = 22 cells from 22 slices in seven mice) and *TrkB-PV^{-/-}* mice (36.59 ± 8.46 pA; *n* = 20 cells from 20 slices in seven mice; *P* = 0.069) (Fig. 4G). However, the ablation of *trkB* significantly reduced the mIPSC frequency from 19.2 ± 2.84 Hz in Ctr mice to 14.42 ± 1.91 Hz in *TrkB-PV^{-/-}* mice (*P* < 0.01) (Fig. 4H). A large fraction of the total GABAergic cell population in the hippocampus is composed of the PV-expressing chandelier and basket cells, which contribute more substantially to the mIPSCs recorded in pyramidal neurons because of their largely somatic synaptic target profile. Moreover, *trkB* was selectively deleted in these PV+ interneurons but not in other GABAergic neurons. Thus, although we cannot exclude the possible contributions of other GABAergic interneurons, it is likely that the decrease in mIPSC frequency recorded in pyramidal neurons is due mostly to a decrease in GABA release from PV+ interneurons.

Decreased Power of Gamma Oscillations and Impaired Neuronal Synchrony. Changes in overall synaptic input and output of PV+ interneurons may alter hippocampal network activity. To test this possibility, CCh-induced extracellular gamma oscillations were recorded in the CA1 SR layer from acute hippocampal slices. In either Ctr or *TrkB-PV^{-/-}* hippocampal slices, bath application of CCh (25 μM) induced a persistent network rhythm in the gamma-frequency band (peak frequency: 36.31 ± 3.06 Hz). As shown in Fig. 5A, the average peak power of the extracellular gamma oscillations in area CA1 was $5.80 \pm 4.29e^{-5}$ in Ctr but $0.78 \pm 0.29e^{-5}$ in *TrkB-PV^{-/-}* slices. We next calculated power spectra and integral power for 20–80 Hz frequency range for 60-s-long recording segments as described in *Materials and Methods*. Both gamma oscillation power and power area were significantly reduced in area CA1 of *TrkB-PV^{-/-}* slices compared with Ctr slices (Fig. 5B and C). Specifically, the gamma band power recorded in *TrkB-PV^{-/-}* slices was $3.986 \pm 0.998e^{-5}$ mV² (five slices from five mice), compared with $17.81 \pm 7.883e^{-5}$ mV² in Ctr slices (six slices from six mice; *P* < 0.01; Fig. 5C). There was no significant change in the oscillation peak frequency (*TrkB-PV^{-/-}*: 36.316 ± 3.058 Hz; Ctr: 36.031 ± 2.9 Hz; Fig. 5D).

To investigate the mechanism underlying the observed reduction of gamma oscillation amplitude and power, whole-cell patch clamp recordings of PV+ interneurons were performed concomitantly with extracellular field recordings of ongoing gamma oscillations (Fig. 6A and B). Both in Ctr and *TrkB-PV^{-/-}* recordings, the majority of action potentials occurred tightly phase-locked (phase angle $\pi/4$) shortly after the positive peak of the field oscillation, which was defined as phase angle 0° (Fig. 6C and F). Although the peak phase angle in recordings from Ctr and *TrkB-PV^{-/-}* cells was the same, the overall number of action potentials per gamma oscillation cycle generated by PV+ interneurons was significantly lower in *TrkB-PV^{-/-}* mice (0.541 ± 0.131 ; *n* = 6 slices from six mice) compared with Ctr mice (0.992 ± 0.14 ; *n* = 6 slices from six mice; *P* < 0.01) (Fig. 6E), resulting in a relative desynchronization.

Discussion

Previous in vitro and in vivo studies have shown that the synaptic connection between pyramidal cells and PV+ interneurons, both locally and interregionally, is crucial for the emergence of gamma oscillations in CA1 (10, 44). Although BDNF-TrkB signaling is known to regulate synaptic formation and transmission tone between glutamatergic principal neurons and interneurons (1–4, 11), its specific role in network function remains unclear. Using the *TrkB-PV^{-/-}* mice, in which the *trkB* gene has been specifically deleted in PV+ interneurons, we show that BDNF-TrkB signaling in PV+ interneurons is critical for the gamma band rhythmic activity in the hippocampal network.

The *TrkB-PV^{-/-}* genotype littermates have a relatively normal PV+ interneuron population during early developmental stages, and the reduction of PV+ interneuron density was only found after P14. Given that *trkB* gene was not deleted in PV+ interneurons until after P14, the PV+ interneuron circuit development should be relatively normal. In young adults (4 or 6 wk old), there was a mild reduction in the density of PV+ neuropil, paralleling a small decrease in the number of PV+ cells. Laser capture of PV+ interneurons from hippocampal tissues combined with qPCR demonstrated that the *TrkB* mRNA expression is reduced, but not totally absent. Explanations include (i) the laser captured tissues may contain some nearby pyramidal cells, which also express *trkB*; (ii) *TrkB* mRNA may still be present in a small portion of PV cells that lacked Cre recombinase as shown by negative β-gal staining (Fig. 2A); and (iii) even in PV+ interneurons expressing Cre recombinase, the homologous recombination efficiency may not have been 100%, leaving some neurons with intact *trkB*. The moderate effect on PV+ in-

with the Arcturus LCM system from Molecular Devices (33, 34). Total RNA was isolated from the captured PV interneurons by using Ambion RNAqueous Microkit (Ambion and Applied Biosystems) (35). First-strand cDNA was reverse-transcribed by using total RNA and Super-Script II (Invitrogen) according to the manufacturer's protocol. qPCR was performed by using Step-One thermal cycler (Applied Biosystems) with SYBR Green PCR master mix (Roche). Levels of TrkB and PV mRNAs were normalized over the RPL10a mRNA level in the same sample. Detailed methods are provided in *SI Materials and Methods*.

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