

Brain-Derived Neurotrophic Factor Modulates Cerebellar Plasticity and Synaptic Ultrastructure

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Neurotrophins are key regulators of neuronal survival and function. Here we show that TrkB, the receptor for brain-derived neurotrophic factor (BDNF), is located at parallel fiber to Purkinje cell (PF/PC) synapses of the cerebellum. To determine the effects of TrkB receptor activation on synapse formation and function, we examined the parallel fiber to Purkinje cell synapses of mice with a targeted deletion of the BDNF gene. Although Purkinje cell dendrites are abnormal in BDNF $-/-$ mice, PF/PC synapses are still able to form. Immunohistochemical analysis of mutant animals revealed the formation of numerous PF/PC synapses with the appropriate apposition of presynaptic and postsynaptic proteins. These synapses are functional, and no differences were detected in the waveform of

evoked EPSCs, the amplitude of spontaneous mini-EPSCs, or the response to prolonged 10 Hz stimulus trains. However, paired-pulse facilitation, a form of short-term plasticity, is significantly decreased in BDNF $-/-$ mice. Detailed ultrastructural analysis of the presynaptic terminals demonstrated that this change in synaptic function is accompanied by an increase in the total number of synaptic vesicles in mutant mice and a decrease in the proportion of vesicles that are docked. These data suggest that BDNF regulates both the mechanisms that underlie short-term synaptic plasticity and the steady-state relationship between different vesicle pools within the terminal.

Key words: neurotrophin; BDNF; Trk receptor; cerebellum; facilitation; ultrastructure; presynaptic plasticity

Information processing depends on the ability of neurons to modify the strength of their connections. Considerable evidence now implicates the neurotrophins as an important class of endogenous modulators of synaptic function. Neurotrophin expression and release are upregulated by neuronal activity (Canossa et al., 1997), and neurotrophins potentiate neurotransmission and plasticity (Lu and Chow, 1999; McAllister et al., 1999; Schuman, 1999). Brain-derived neurotrophic factor (BDNF) acting through the receptor tyrosine kinase TrkB has been implicated in the plasticity of central synapses (Lu and Figurov, 1997). Adding BDNF to dissociated rat hippocampal cultures (Lessmann et al., 1994; Levine et al., 1995) potentiates synaptic activity. A long-lasting potentiation is seen with the addition of BDNF to rat hippocampal slices (Kang and Schuman, 1995a,b). In addition, BDNF plays a role in short-term plasticity, including paired-pulse facilitation (PPF) (Patterson et al., 1996; Stoop and Poo, 1996; Gottschalk et al., 1998), and the response to repetitive stimulation (Figurov et al., 1996; Gottschalk et al., 1998; Pozzo-Miller et al., 1999). Both long- and short-term plasticity are impaired in the hippocampus of BDNF $-/-$ mice (Korte et al., 1995), and the

addition of BDNF reverses these defects (Korte et al., 1996a,b; Patterson et al., 1996; Pozzo-Miller et al., 1999). Because presynaptic mechanisms are thought to underlie paired-pulse facilitation (Katz and Miledi, 1968; Zucker, 1989; Schulz et al., 1994; Atluri and Regehr, 1996) and contribute to the response to repetitive stimulation (Kreitzer and Regehr, 2000), BDNF may modulate presynaptic function (Li et al., 1998; Martinez et al., 1998; Pozzo-Miller et al., 1999).

Synapses with low probability of release can undergo a pronounced modulation of synaptic activity (Bolshakov and Siegelbaum, 1995) and may be more likely to reveal the functional effects of neurotrophins (Berninger et al., 1999). In the cerebellum, granule cell axons, called parallel fibers, form numerous synapses onto Purkinje cell dendrites in the molecular layer. These synapses are excitatory, exhibit short-term plasticity in the form of facilitation, and characteristically exhibit a low probability of release (Atluri and Regehr, 1996; Sabatini and Regehr, 1997). Because granule cells synthesize BDNF (Hofer et al., 1990) and because the receptor TrkB is expressed by both granule cells (Klein et al., 1990; Segal et al., 1995) and Purkinje cells (Klein et al., 1990), the parallel fiber to Purkinje cell (PF/PC) synapse is well suited for studying the role of BDNF on short-term synaptic plasticity.

We examined the effect of BDNF deficiency at the PF/PC synapse in the mouse cerebellum using animals with a targeted deletion of BDNF (Ernfors et al., 1994). Numerous functional PF/PC synapses form in these animals. The finding that TrkB is located at PF/PC synapses supports the idea that BDNF can directly modulate short-term synaptic plasticity. Electrophysiological analysis of the PF/PC synapse of BDNF $-/-$ mice revealed preserved neurotransmission but a significant decrease in PPF. This specific impairment of PPF is associated with an

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accumulation of synaptic vesicles and a lengthening of the active zone. These observations implicate BDNF as an important endogenous synaptic factor that regulates short-term synaptic plasticity as well as synaptic ultrastructure. Identification of the structural changes that accompany changes in plasticity contributes to our understanding of the molecular mechanisms underlying activity-dependent synaptic plasticity.

MATERIALS AND METHODS

BDNF mutant mice. Breeding pairs of BDNF +/- mice (Ernfors et al., 1994) were purchased from Jackson Laboratories (Bar Harbor, ME). Genotypes of animals were determined via PCR using three oligonucleotide primers: oIMR132: 5'-GGG AAC TTC CTG ACT AGG GG-3'; oIMR133: 5'-ATG AAA GAA GTA AAC GTC CAC-3'; and oIMR134: 5'-CCA GCA GAA AGA GTA GAG GAG-3'.

The PCR was run under standard conditions. Primers 132 and 133 amplify a ~340 bp product from the targeted BDNF allele. Primers 133 and 134 amplify a ~275 bp band from the wild-type (WT) BDNF allele. PCR of heterozygous animals yielded both amplification products. BDNF -/- pups weighed one-third to two-thirds as much as littermates, depending on age, and were severely ataxic. BDNF -/- pups generally survive until the third postnatal week.

Immunohistochemistry. Midsagittal cerebellar sections (8 μ m) were obtained from postnatal day (P) 8, P15, and P24 wild-type and BDNF -/- mice as described (Schwartz et al., 1997). Anesthetized mice were perfused with 4% paraformaldehyde in 0.2 M phosphate buffer, and cryoprotected brains were cut on a Leica CM 3050 cryostat. For immunohistochemistry, cerebellar sections were incubated in a PBS blocking solution containing 5% normal goat serum and 0.1% Triton X-100. Incubations in primary antibody were performed in the same blocking solution. The antibodies used were as follows: anti-calbindin-D, 1:500 (CL-300, Sigma, St. Louis, MO); mouse anti-SV2, 1:200 (kind gift of K. M. Buckley, Harvard Medical School); anti-TrkB, 1:2000 (Chemicon, Temecula, CA); anti-synapsin, 1:100 (Chemicon); and anti-GluR δ 1/2, 1:25 (Chemicon). Incubations with primary antibodies were performed overnight at 4°C. Immunostaining was visualized using cy3-conjugated goat anti-mouse IgG, 1:200 (Jackson Immunochemicals, West Grove, PA); Alexa Fluor 488 goat anti-mouse IgG green, 1:500 (Molecular Probes, Eugene, OR); or Alexa Fluor 568 goat anti-rabbit IgG red, 1:500 (Molecular Probes) for 90–120 min at room temperature. Sections were washed and then mounted in GelTol aqueous mounting medium (Immunotech, Marseille, France). DeltaVision restoration fluorescence microscopy was performed on an Olympus fluorescence microscope configured with a DeltaVision stage and software (Applied Precision Inc., Issaquah, WA). Immunofluorescent sections were viewed with a 100 \times objective (numerical aperture 1.4). Z-series (20 0.2 μ m serial optical sections) were acquired, and softWoRx imaging software (Applied Precision) was used to perform a blind deconvolution on a Silicon Graphics workstation. Final images were representative z-series rendered in softWoRx Volume Viewer unless indicated otherwise. Statistical significance for the correlation between staining patterns was established by calculating for each wavelength the number of pixels with an intensity 1 SD above the mean within a known area. In double-labeled samples, synaptic structures were selected in only one wavelength without knowledge of the second wavelength, which was then restored after the selection of structures. Staining in both wavelengths was considered associated only if no measurable distance existed between the two structures within the single optical section where they were defined or within an adjacent optical section. The probability of two pixels for each wavelength being randomly located in the same position or in one of the eight adjacent positions is given by: (pixels positive for wavelength 1/total pixels) \times (pixels positive for wavelength 2/total pixels) \times 9. Statistical significance was determined by a two-tailed unpaired *t* test assuming unequal variances.

Stensaas-modified Golgi stain. Age P15 BDNF +/+ and BDNF -/- mice were perfused transcardially with 4% paraformaldehyde in 0.1 M sodium phosphate. Brains were immediately removed and immersed 48 hr in Stensaas solution containing 85 mM potassium dichromate, 151 mM chloral hydrate, 3.7% formaldehyde, and 0.5% glutaraldehyde. Tissue was rinsed in 0.5% silver nitrate and transferred to 1.25% silver nitrate for 4 d. After progressive dehydrations in ethanol, tissue was transferred to low viscosity nitrocellulose, and chloroform was added to harden the nitrocellulose. Sections (100 μ m) were cut on a sledge microtome and

collected in 80% ethanol, rinsed in 95% ethanol, and transferred to terpineol. Sections were rinsed in xylene, covered in mounting media, and coverslipped. Images were taken with a Nikon Eclipse E800 microscope at 100 \times magnification. Multiple images in different focal planes were taken of individual Purkinje cells. The focused portions of each image were combined to bring the entire dendritic arbor into focus.

Electrophysiology. Synaptic currents were recorded from freshly cut transverse slices of cerebellar vermis of P14–P17 BDNF -/- mice and their wild-type littermates as described previously (Atluri and Regehr, 1996; Chen and Regehr, 1997). Briefly, slices were maintained in an oxygenated (95% O₂/5% CO₂) external solution containing (in mM): 125 NaCl, 2.5 KCl, 2.6 NaHCO₃, 1.25 NaH₂PO₄, 25 glucose, 2 CaCl₂, and 1 MgCl₂. The external recording solution also contained 20 μ M bicuculline to block IPSCs (Sigma). Whole-cell patch-clamp recordings of Purkinje cells were obtained using 1.0–1.5 M Ω electrodes containing (in mM): 35 CsF, 100 CsCl, 10 EGTA, 10 HEPES, and 0.1 D600, pH 7.4. Holding potential was maintained at -40 mV to inactivate voltage-dependent calcium conductances. Access resistance (<5 M Ω after series resistance compensation), leak current (<100 pA), and the time course of decay of the EPSC (<6 msec) were monitored continuously, and cells that did not meet these criteria were rejected from analysis. Parallel fibers were stimulated with a glass electrode filled with extracellular solution and positioned several hundred micrometers from the recording electrode.

Paired-pulse facilitation protocol involved measurement of the peak amplitude of EPSCs elicited by pairs of stimuli, separated by a randomized interstimulus interval (ISI) ranging from 10 to 500 msec. Pairs of EPSCs were interleaved with EPSCs elicited by a single stimulus. The average of three to five trials was used to calculate the percentage facilitation [$100 \times (A_2 - A_1)/A_1$]. A₁ and A₂ are the average peak amplitude of the first and second EPSC, respectively, in response to a single stimulation. At intervals shorter than 100 msec, A₂ is measured from the waveform obtained by subtracting the average single EPSC response from the average EPSC response to a pair of stimuli.

Mini-EPSCs (mEPSCs) were recorded as described previously (Chen and Regehr, 1997). Extracellular solution contained 0.25 μ M TTX and 20 μ M bicuculline. Data were recorded in 20 sec epochs, sampled at 2.5 kHz, and filtered digitally at 500 Hz. Inclusion criteria were an 8 pA threshold, a minimum rate of rise of 0.4 pA/msec, and a decay time constant between 3 and 20 msec. Experiments used to calculate the average cumulative histogram distributions of mEPSC amplitude contained >4000 mEPSC events. Changes in mEPSC frequencies have been used as an indicator of a presynaptic process when the same population of synapses is studied before and after a pharmacological manipulation. However, the frequency of mEPSCs cannot be used as a presynaptic indicator when comparing population of synapses from different slices. The angle of the slice and the number of intact presynaptic fibers vary significantly from slice to slice and would thus make mEPSC frequency comparison between BDNF -/- and WT mice difficult to interpret.

Because trains of stimuli can result in recurrent excitation and activation of endogenous G-protein-mediated modulation of synaptic currents, experiments using repetitive stimulation were recorded in an external solution containing the GABA_B receptor antagonist CGP55845a (2 μ M), the adenosine A₁ receptor antagonist 8-cyclopentyl-1,3-dipropylxanthine (5 μ M), and the mGlu-RIII antagonist (RS)- α -cyclopropyl-4-phosphonophenylglycine (30 μ M). Frequencies >10 Hz resulted in desynchronization of EPSCs during a train of stimuli, attributable to accumulation of presynaptic residual calcium (Atluri and Regehr, 1998; Kreitzer and Regehr, 2000). Therefore we did not analyze the synaptic response to higher frequency stimulation. All recordings were digitized with a 16 bit digital-to-analog converter (Instrutech, Great Neck, NY), with pulse control software, and analyzed using Igor Pro software (Wavemetrics, Lake Oswego, OR) and custom macros. All experiments were performed at 25°C.

Electron microscopy. Brains were obtained from P14 wild-type and BDNF -/- mice anesthetized and perfused with 4% paraformaldehyde and 2% glutaraldehyde in 0.2 M phosphate buffer. Blocks from midsagittal cerebellar vermis in folia V were stained with 1% osmium tetroxide/1.5% potassium ferrocyanide, bathed in 1% uranyl acetate, and ethanol dehydrated and placed in propylene oxide for 1 hr. Infiltration with Epon/Araldite mixed 1:1 with propylene oxide was performed overnight. Samples were embedded in Epon/Araldite. Thin sections from sample were photographed under a JEOL 1200EX electron microscope. Negatives were scanned into Adobe Photoshop. Images of individual synapses were analyzed using NIH Image 3.1 (National Institutes of Health, Bethesda, MD). Scale bars were taken from scans of original electron

micrograph negatives. All synapses were analyzed without knowledge of genotype. For each synapse the number of synaptic vesicles and the length of the postsynaptic density (PSD) were quantitated. Vesicle distribution was quantitated by measuring the distance from the center of each synaptic vesicle to the closest point at the active zone. Docked vesicles were defined as those located within 50 nm (one vesicle diameter) of the active zone.

RESULTS

The TrkB receptor is located at PF/PC synapses in wild-type mice

The BDNF receptor TrkB is expressed by both granule cells and Purkinje cells. To determine whether TrkB is appropriately located to mediate synapse-specific effects of BDNF, the expression pattern of TrkB protein in the cerebellum was analyzed. In both wild-type (Fig. 1*A*) and mutant mice (data not shown), significant TrkB immunostaining was seen throughout the molecular layer and in Purkinje cell bodies and main dendrites. To clarify the subcellular localization of TrkB, cerebellar sections of wild-type mice were double labeled with the TrkB antibody and an antibody against calbindin, which is expressed specifically in Purkinje cells. Staining was visualized by high-resolution wide-field restoration microscopy. Nanomotor-guided focusing allows for image acquisition at precise distance intervals throughout the volume of the sample. On the basis of the information in the stack of optical sections, a deconvolution algorithm identifies out-of-focus light and returns it to its source using a point-spread function. Although the original optical sections reveal the general dendritic morphology of a Purkinje cell (Fig. 1*B*), the increased contrast, decreased background, and sharper borders of the deconvolved optical sections (Fig. 1*C*) allow for the identification of details on the order of 175 nm.

Double labeling for calbindin and TrkB in wild-type cerebellum (Fig. 1*D–F*) revealed the presence of numerous TrkB-positive puncta (*red*) adjacent to and partially overlapping with the tips of calbindin-positive dendritic spines (*green*). There is a high degree of association between calbindin-positive spines and TrkB puncta. We found that 75% ($\pm 6\%$ SEM) of calbindin-positive spines ($n = 116$) were associated with TrkB puncta and that 79% ($\pm 1\%$ SEM) of TrkB puncta ($n = 121$) were associated with calbindin-positive spines. This degree of association is significantly higher than that predicted by chance alone ($20 \pm 0.5\%$ SEM; $p < 0.01$; t test).

The localization of TrkB immunostaining at spine tips is consistent with a synaptic localization of TrkB at PF/PC synapses. Double labeling was performed using an antibody against the glutamate receptor delta 2 subunit (GluR $\delta 2$) and an antibody against calbindin. GluR $\delta 2$ is a useful marker for PF/PC synapses because after an initial period of expression throughout Purkinje cells, this receptor subunit becomes targeted specifically to the spines receiving parallel fiber afferents (Landsend et al., 1997). Double labeling with an antibody against calbindin (*green*) and an antibody against GluR $\delta 2$ (*red*) (Fig. 1*G–I*) yielded numerous GluR $\delta 2$ puncta that overlapped significantly with calbindin-positive dendrites and spines. We found that 74% ($\pm 5\%$ SEM) of calbindin-positive spines ($n = 91$) are associated with GluR $\delta 2$, whereas 65% ($\pm 4\%$ SEM) of GluR $\delta 2$ puncta ($n = 100$) are associated with calbindin-positive spines. This degree of association is significantly higher than that predicted by chance alone ($14 \pm 1\%$ SEM; $p < 0.01$; t test). The finding that the degree of association between calbindin-positive spines and TrkB puncta, and between spines and GluR $\delta 2$, is comparable and that TrkB and GluR $\delta 2$ have similar staining patterns relative to calbindin

strongly suggests that TrkB, like GluR $\delta 2$, is localized to PF/PC synapses. In fact, TrkB staining often appeared to be even more specifically associated with the very tips of dendritic spines than did GluR $\delta 2$ (stronger *yellow* signal in Fig. 1*I* than in Fig. 1*F*). The absence of antibodies of the appropriate species and sensitivity currently make it difficult to determine conclusively the relative positions of TrkB and GluR $\delta 2$.

To further confirm the synaptic localization of TrkB, double labeling with an antibody against the presynaptic protein SV2, a component of synaptic vesicle membranes, and the antibody against TrkB was performed (Fig. 1*J–L*). We found that 75% ($\pm 6.5\%$ SEM) of SV2 puncta ($n = 158$) are associated with TrkB-positive spines. This degree of association is significantly higher than that predicted by chance alone ($15 \pm 1.1\%$ SEM; $p < 0.01$; t test). Nonetheless, the synaptic localization of the BDNF receptor is appropriate for BDNF to directly regulate PF/PC synapse development and function. As shown here, there is some overlap of TrkB and SV2 puncta, and there is also some overlap of TrkB and calbindin staining. Because of the narrow width of the synaptic cleft (~ 30 nm) and the fact that the TrkB antibody recognizes the extracellular domain of the receptor, we cannot determine conclusively on which side of the synaptic cleft TrkB resides.

BDNF is not required for PF/PC synaptogenesis

The synaptic localization of the BDNF receptor suggests that BDNF may play a crucial role in synapse formation and development. Therefore, the absence of BDNF could preclude PF/PC synapse formation. Previously, the addition of BDNF or the removal of endogenous BDNF with TrkB IgG has been shown to alter Purkinje cell dendritic spine number and morphology (Morrison and Mason, 1998; Shimada et al., 1998). Furthermore, analysis of cerebella in BDNF $-/-$ mice revealed that the dendrites of Purkinje cells are severely atrophied and disorganized in P8 animals (Schwartz et al., 1997). To determine whether the absence of BDNF precludes the formation of synapses onto Purkinje cell dendrites, we first examined Purkinje cell morphology in P8, P15, and P24 wild-type and BDNF mutant mice using an antibody against the Purkinje cell-specific marker calbindin (Fig. 2). Compared with Purkinje cell dendrites of BDNF $+/+$ animals, dendrites of BDNF $-/-$ animals remain less extensive at all ages. Accordingly, this defect is associated with a persistently thinner molecular layer. Nonetheless, there is continued development of Purkinje cell dendrites even in the absence of BDNF because primary, secondary, and tertiary branches are visible in P15 and P24 BDNF $-/-$ mice.

Because the overlap between adjacent dendrites makes it difficult to appreciate the morphology of individual dendritic arbors, cerebella from P15 BDNF $+/+$ and BDNF $-/-$ mice were stained using a rapid Golgi method (Fig. 3) to visualize individual Purkinje cell arbors. As illustrated, the extent of dendritic arbors in individual BDNF $-/-$ Purkinje cells is reduced when compared with wild-type Purkinje cells (Fig. 3). In BDNF $-/-$ cells, the primary dendrite is shorter and divides into fewer secondary and tertiary branches. Despite their abnormal morphology, BDNF $-/-$ dendrites are studded with numerous spines (Fig. 3*B*, *inset*, *C*, *inset*) as in wild-type dendrites (Fig. 3*A*, *inset*). Because these dendritic spines are the postsynaptic targets of granule cell axons, Purkinje cell dendrites may develop sufficiently to receive synapses from parallel fibers despite the significant effects of BDNF deficit on dendritic morphology.

To determine whether synapses form on Purkinje cell dendritic

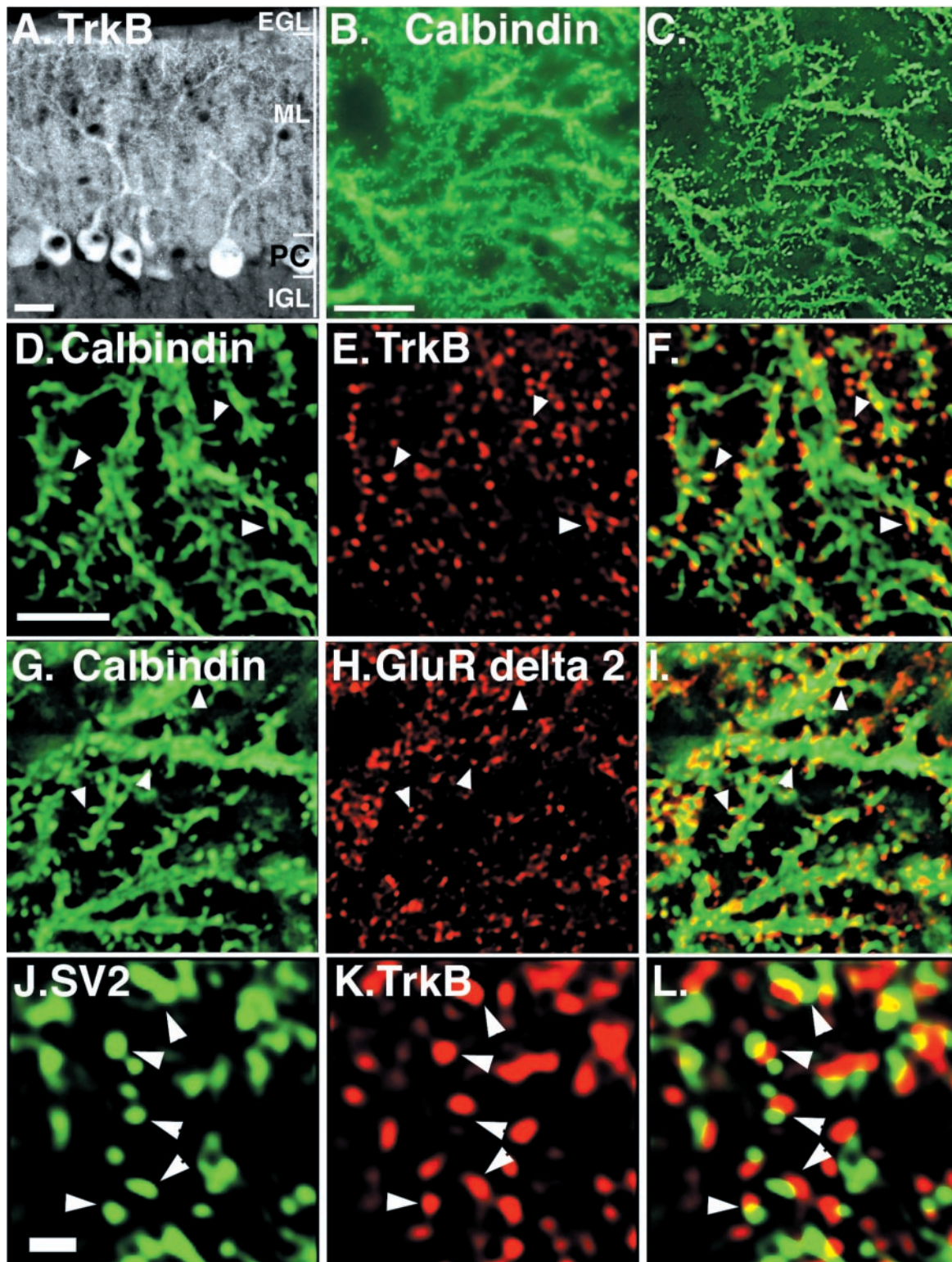


Figure 1. The BDNF receptor, TrkB, is located at PF/PC synapses. *A*, P15 cerebellum immunolabeled with anti-TrkB. TrkB staining is seen throughout the molecular layer (ML) and in Purkinje cell bodies (PC) and dendrites in BDNF $+/+$ mice. EGL, External granule cell layer; IGL, internal granule cell layer. *B*, *C*, Single optical section through the molecular layer of P15 wild-type cerebellum labeled with anti-calbindin before deconvolution (*B*) and after deconvolution (*C*). *D–F*, Cerebellar molecular layer of P15 wild-type mouse double labeled with anti-calbindin (*D*, green) and anti-TrkB (*E*, red) antibodies. The merged image (*F*) shows numerous TrkB puncta juxtaposed to and sometimes overlapping with calbindin-positive Purkinje cell dendritic spines (arrowheads). *G–I*, Cerebellar molecular layer of P15 wild-type mouse double labeled with anti-calbindin (*G*, green) and anti-GluR δ 2 (*GluR delta 2*) (*H*, red). Merged image (*I*) shows GluR δ 2-positive overlapping Purkinje cell dendrites and capping Purkinje cell dendritic spines (arrowheads). *J–L*, Cerebellar molecular layer of P15 wild-type mouse double labeled with anti-SV2 (*J*, green) and anti-TrkB (*K*, red). Merged image (*L*) shows SV2-positive puncta juxtaposed to and partially overlapping with TrkB-positive puncta (arrowheads). *D–L* were reconstructed from 5–10 adjacent optical sections. Scale bars: *A*, 20 μ m; *B*, *C*, 10 μ m; *D–I*, 5 μ m; *J–L*, 5 μ m.

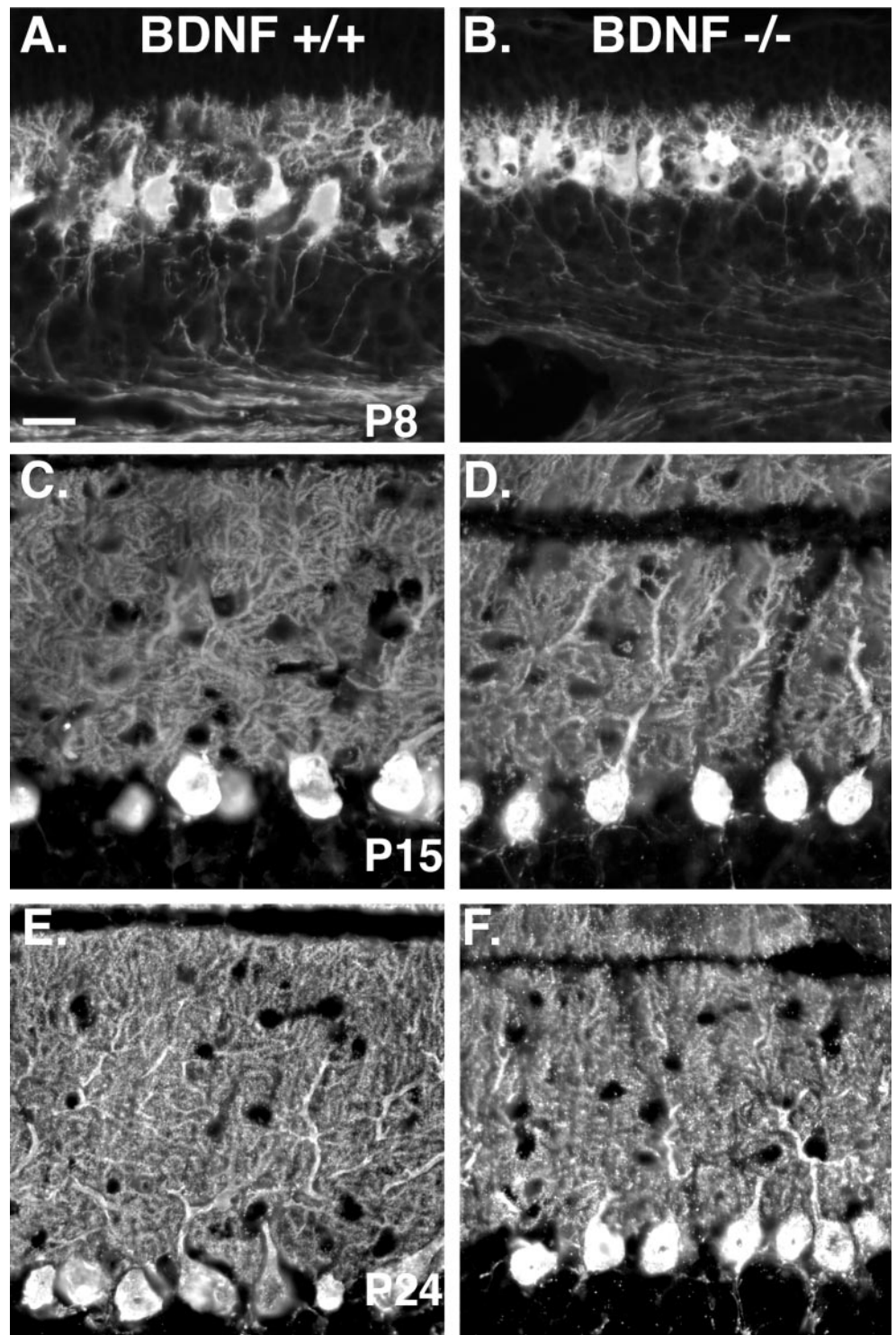


Figure 2. The development of Purkinje cell dendritic arbors in wild-type (*A, C, D*) and BDNF $-/-$ mice (*B, D, F*). Parasagittal cerebellar sections from P8 (*A, B*), P15 (*C, D*), and P24 (*E, F*) mice were immunolabeled with anti-calbindin antibody. In the absence of BDNF, Purkinje cell dendrites are stunted at P8 and undergo significant development over the following 2 weeks, but remain less extensive than in WT mice. Scale bar, 20 μ m.

spines from BDNF $-/-$ mice, immunohistochemistry was performed using an antibody against calbindin and a second antibody against glutamate receptor subunit GluR δ 2. Double labeling shows puncta of GluR δ 2 staining that cap calbindin-positive spines (Fig. 4*A*), suggesting that many of these spines in fact do receive synapses as in the wild-type mouse (Fig. 1*I*). Quantification of the staining patterns confirmed that 70% of spines were capped with GluR δ 2 puncta ($n = 94$) and 72% of GluR δ 2 puncta were found at the tips of calbindin-positive spines ($n = 90$). These values are very close to those determined for BDNF $+/+$ cere-

bella. To visualize presynaptic boutons, antibodies against the presynaptic terminal proteins SV2 and synapsin were used (Fig. 4*B*). In the molecular layer of the BDNF $-/-$ cerebellum, numerous puncta immunopositive for both SV2 and synapsin were seen, resulting in a strong yellow signal. These correspond to individual presynaptic terminals. Quantitation revealed that 90% of SV2 puncta were synapsin positive ($n = 106$) and that 92% of synapsin puncta were SV2 positive ($n = 125$). This degree of association is significantly higher than that predicted by chance alone ($1 \pm 0.1\%$ SEM; $p < 0.01$; t test). Double-positive puncta

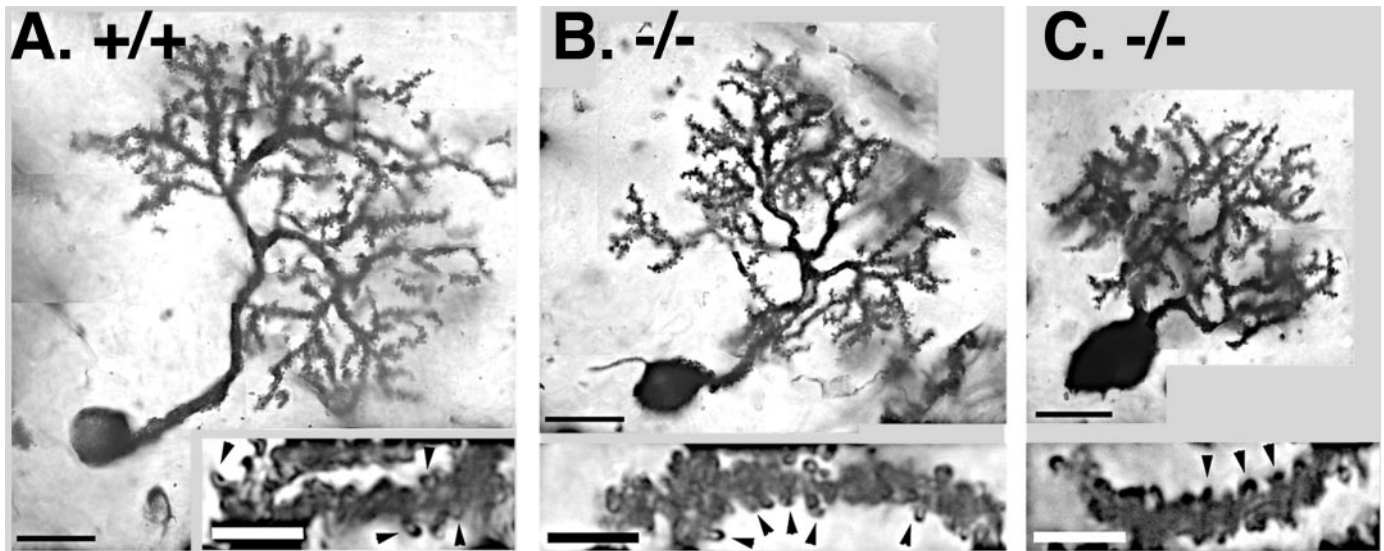


Figure 3. Individual Purkinje cell dendrite morphology in the absence of BDNF. *A*, Modified Golgi stain resolves the morphology of an individual Purkinje cell dendritic arbor in a P15 wild-type mouse. *Inset* shows a higher magnification view of a dendritic segment studded with numerous spines (*arrowheads*). *B*, *C*, Two individual Purkinje cells from a P15 BDNF $-/-$ mouse exhibit a significant decrease in the extent of the dendritic arbor. *Insets* show that dendrites nonetheless carry numerous spines (*arrowheads*). Scale bars: *A–C*, 20 μm ; *insets*, 5 μm .

were defined as those that overlapped by at least 50% in the same optical section. To confirm that these accumulations of presynaptic protein correspond to bona fide synapses, we used the antibody against the postsynaptic marker, GluR δ 2. Double labeling revealed that numerous GluR δ 2 puncta were present and juxtaposed to SV2 puncta in the molecular layer of BDNF $-/-$ mice (Fig. 4C). As predicted, SV2 staining and GluR δ 2 staining overlap noticeably less than staining for the two presynaptic markers SV2 and synapsin, resulting in much less yellow signal. The staining patterns for SV2, synapsin, calbindin, and GluR δ 2 seen in BDNF $-/-$ mice were very similar to staining patterns observed in BDNF $+/+$ cerebella (data not shown). This demonstrates that even in the absence of BDNF there is development and juxtaposition of the presynaptic and postsynaptic components of the PF/PC synapse.

BDNF may regulate synapse number in several systems. Therefore, a quantitative analysis of synapse density in the molecular layer of BDNF $+/+$ and $-/-$ mice was performed. In cerebellar sections double labeled with an antibody against SV2 and an antibody against GluR δ 2, synapses were defined by the juxtaposition or overlap of puncta of SV2 and GluR δ 2 staining. Synaptic density per 300 μ^3 was determined from four deconvolved optical z-series in each of three cerebellar subregions (caudal, dorsal, and rostral) in wild-type ($n = 3$) and mutant mice ($n = 3$). A small difference was seen in the overall density of PF/PC synapses between wild-type mice (48.4 synapses/300 $\mu^3 \pm 4.1$ SEM) and mutant mice (40.7 synapses/300 $\mu^3 \pm 1.9$ SEM; $p = 0.07$). Thus, in spite of the marked effect of BDNF on Purkinje cell dendritic morphology and molecular layer thickness, PF/PC synapses can form, although their numbers are slightly reduced.

BDNF deficit leads to impaired short-term plasticity

BDNF enhances neurotransmission and potentiates activity-dependent plasticity at hippocampal synapses (Kang and Schuman, 1995a,b; Korte et al., 1995; Patterson et al., 1996). We analyzed neurotransmission across PF/PC synapses in thin cerebellar slices from BDNF $-/-$ and wild-type mice aged P14–P17.

We recorded the spontaneous mEPSC as well as the AMPA receptor-evoked EPSCs from Purkinje cells (Chen and Regehr, 1997). Analysis of mEPSC amplitudes showed no difference between BDNF $+/+$ and BDNF $-/-$ synapses (Fig. 5). In addition, evoked EPSCs are similar at BDNF $+/+$ and $-/-$ synapses (Fig. 6A). These observations suggest that synaptic function is preserved despite the BDNF deficit.

Short-term synaptic plasticity at the PF/PC synapse in BDNF $-/-$ mice was examined using pairs of stimuli, separated by varying interstimulus intervals. At the PF/PC synapse, pairs of closely spaced stimuli result in the facilitation of synaptic strength (Atluri and Regehr, 1996). This paired-pulse facilitation can be quantified as the change in the peak amplitude of the second EPSC relative to the first (percentage facilitation; see Materials and Methods). For brief interstimulus intervals, the degree of facilitation in BDNF $-/-$ mice is approximately half that of control levels ($p < 0.01$; ANOVA) (Fig. 6B). At longer interstimulus intervals (>400 msec), there is no facilitation in either wild-type or mutant animals. This result suggests that there is an alteration of short-term plasticity at the PF/PC synapse in BDNF $-/-$ mice.

Response to repetitive stimulation is another index of presynaptic function. Parallel fibers were stimulated with a 10 Hz train for 10 or 50 trials. The ratios of the peak amplitude of the 10th/1st EPSC and the 50th/1st EPSC were determined (Fig. 6C). The similarity in the 50th/1st impulse between wild-type (0.70 ± 0.21 SEM; $n = 6$) and BDNF $-/-$ mice (0.53 ± 0.03 SEM; $n = 5$) indicates that the synaptic apparatus in the mutant mouse is capable of maintaining neurotransmission over long periods of repetitive stimulation. In contrast, after a short train of stimulation, the 10th/1st ratio is 1.40 (± 0.08 SEM; $n = 7$) in wild-type mice and 1.11 (± 0.06 SEM; $n = 6$) in BDNF $-/-$ mice. This difference is statistically significant ($p < 0.05$; t test). Because PPF is decreased in BDNF $-/-$ mice at an ISI of 100 msec, which corresponds to a frequency of 10 Hz, altered short-term plasticity must contribute to this difference seen in the 10th/1st ratio. Thus

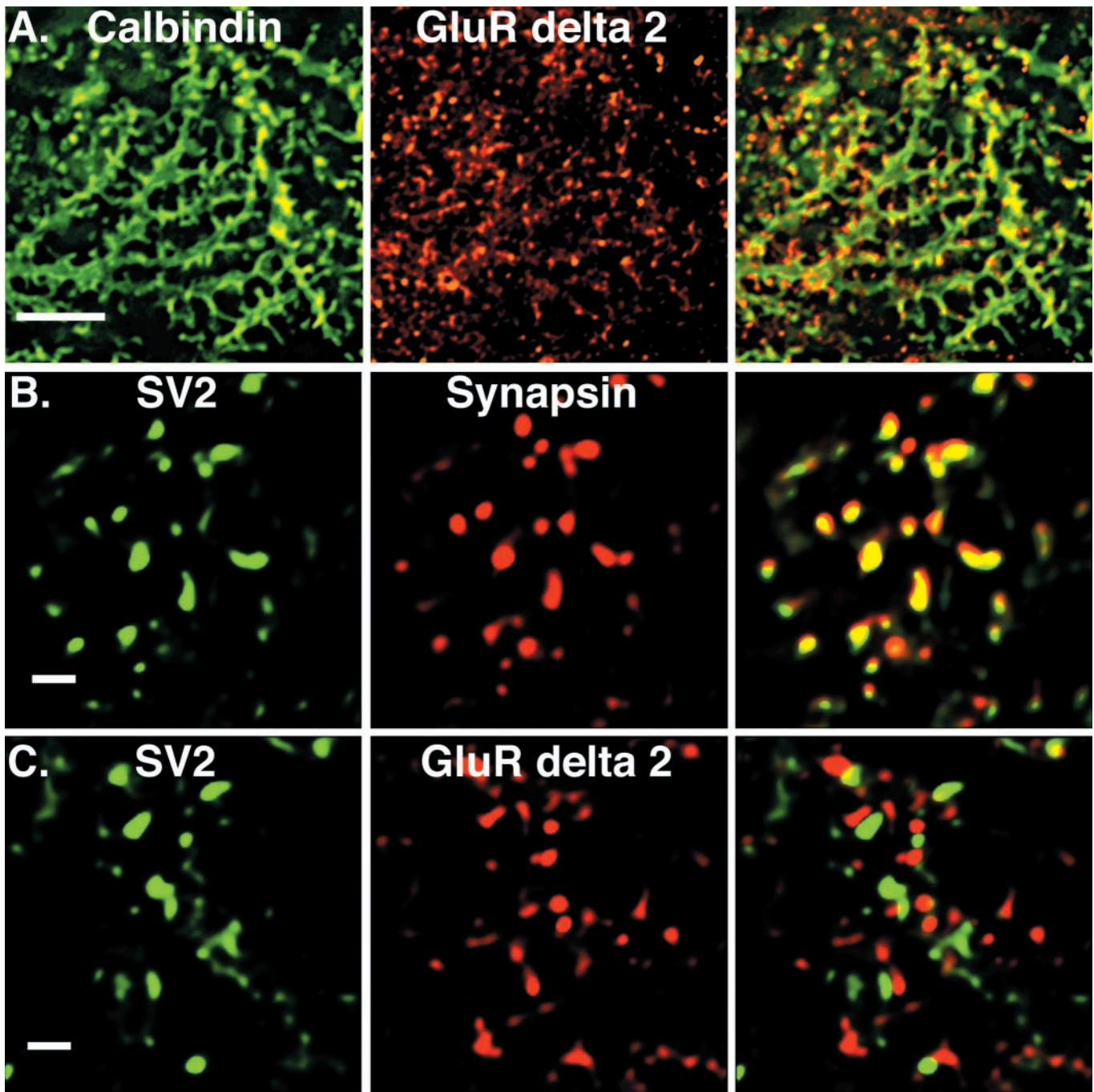


Figure 4. Synaptogenesis in the molecular layer of BDNF $-/-$ cerebellum. *A*, P15 cerebellum immunolabeled with antibodies against calbindin (*green*) and the glutamate receptor subunit GluR δ 2 (*GluR delta 2*) (*red*). Numerous calbindin-positive spines are capped by puncta of GluR δ 2 staining characteristic of functional synapses. *B*, Individual presynaptic terminals in the molecular layer immunolabeled with antibodies against the presynaptic terminal proteins SV2 (*green*) and synapsin (*red*) resulting in identical staining patterns. *C*, Immunolabeling with antibodies against SV2 (*green*) and GluR δ 2 (*red*) results in staining patterns where the puncta are juxtaposed with much less overlap, consistent with the association of presynaptic terminals with their postsynaptic targets. Images were reconstructed from 5–10 adjacent optical sections. Scale bars: *A*, 5 μ m; *B*, *C*, 1 μ m.

the evidence for impaired paired-pulse facilitation in the context of normal mEPSCs, normal evoked EPSC, and equivalent response to sustained trains of stimulation argues for a selective deficit in short-term plasticity in BDNF $-/-$ mice.

To determine whether the effects of BDNF deficit on presynaptic function are reversible, exogenous BDNF was applied to cerebellar slices from BDNF $-/-$ mice. Under the experimental

conditions used (see Materials and Methods), only presynaptic effects of BDNF addition would be detected. The functional deficit in short-term synaptic plasticity evident in the decreased PPF was not reversed by bath application of BDNF after either 30 min or 3 hr of BDNF exposure (data not shown). To be certain that BDNF is capable of stimulating responsive cells in the slice under these conditions, Trk phosphorylation was analyzed. Treat-

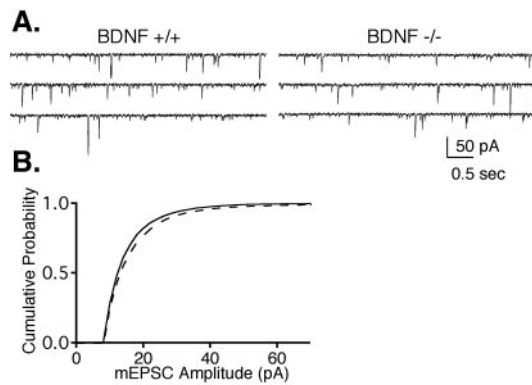


Figure 5. *A*, Representative consecutive mEPSC traces are shown from a wild-type (*left*) and a BDNF $-/-$ (*right*) mouse. *B*, The normalized cumulative amplitude distribution for BDNF $+/+$ ($n = 4$; *solid trace*) and BDNF $-/-$ ($n = 4$; *dashed trace*) are similar ($p = 1.0$ by Kolmogorov–Smirnov test).

ment of wild-type or BDNF $-/-$ cerebellar slices with 200 ng/ml BDNF for 1 hr was able to induce Trk receptor phosphorylation (data not shown).

Synaptic vesicle distribution is altered in the absence of BDNF

Under the experimental conditions used in this study, changes in the evoked response to pairs of stimuli are consistent with a presynaptic mechanism that regulates synaptic vesicle exocytosis. Recent studies have revealed the existence of distinct vesicle pools and have begun to shed light on their relationship to the dynamics of vesicle exocytosis and endocytosis (Sudhof, 1995; Betz and Angleson, 1998; Fernandez-Chacon and Sudhof, 1999; Jahn and Sudhof, 1999; Li and Schwarz, 1999; Doussau and Augustine, 2000; Sudhof, 2000). Because presynaptic function is impaired in BDNF $-/-$ mice, we hypothesized that this functional impairment may also be reflected in the synaptic ultrastructure.

Ultrastructural changes have previously been associated with changes in synaptic function. We performed a detailed analysis of the PF/PC synapse in wild-type and BDNF $-/-$ mice via electron microscopy. Electron micrographs of the molecular layer reveal the main ultrastructural components of the PF/PC synapse in both genotypes (Fig. 7). These include a presynaptic swelling of the parallel fiber terminal, which contains numerous round, loosely packed synaptic vesicles. The postsynaptic dendritic spine from a Purkinje cell dendrite is also clearly visible in cross section. The asymmetric electron-dense band between the terminal and the dendritic spine is characteristic of excitatory synapses and is often used to demarcate the extent of the active zone. Although immunohistochemistry revealed the appropriate localization of important synaptic proteins, electron micrographs demonstrated conclusively that PF/PC synapses display the basic structural features of synapses even in the absence of BDNF.

A quantitative analysis revealed a significant increase in the number of synaptic vesicles in the terminals of BDNF $-/-$ mice compared with those of BDNF $+/+$ mice (Table 1). Synaptic terminals were analyzed in longitudinal and cross-sectional orientations. The increase in vesicle number in BDNF $-/-$ terminals was seen in both of these orientations. Investigations of the structure and dynamics of the vesicle pool have yielded evidence that it can be subdivided into different functional pools (Rosenmund and Stevens, 1996) that may exist in a steady-state relation-

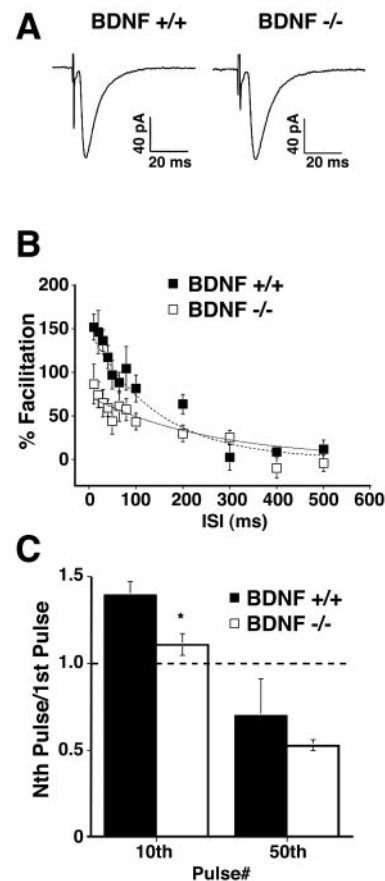


Figure 6. Specific impairment of short-term synaptic plasticity in BDNF $-/-$ mice. *A*, Evoked AMPA receptor-mediated EPSCs were elicited by stimulating parallel fibers and recording from an individual Purkinje cell in voltage clamp. The decay times of the evoked EPSCs are indistinguishable between WT and BDNF $-/-$ mice. Traces are the averages of 5–10 trials. *B*, The mean percentage facilitation is plotted as a function of interstimulus intervals for BDNF $-/-$ (■; $n = 6$) and wild-type littermates (□; $n = 6$). BDNF $-/-$ mice exhibit decreased paired-pulse facilitation at interstimulus intervals <200 msec ($p < 0.05$). *C*, The response of Purkinje cells to repetitive 10 Hz stimulation is illustrated by plotting the ratio of the amplitude of the response to the 10th and 50th pulses to the amplitude of the response to the first pulse. Although 10th/1st ratio is significantly decreased in BDNF $-/-$ mice because of impaired PPF, there is no significant difference in the 50th/1st. This indicates that there is no difference in the response to prolonged 10 Hz stimulation in the absence of BDNF.

ship. Therefore, to determine whether the increase in the number of vesicles in BDNF $-/-$ mice was restricted to a specific region of the vesicle pool, we plotted the number of vesicles per 25 nm bin as a function of distance from the synaptic cleft (Fig. 8*A,B*). This revealed that within the first 100 nm of the synaptic cleft there is no difference in vesicle number. However, beyond 100 nm there is a significant increase in the number of vesicles in BDNF $-/-$ terminals ($D \ll D_{0.01}$, by Kolmogorov–Smirnov). When we calculated the fraction of all the vesicles that was located within 50 nm of the active zone, commonly referred to as the docked vesicle pool, we found a significant decrease in the absence of BDNF (Table 1). These findings show that BDNF deficit leads to a selective increase in the vesicle pool that lies distal to the active zone.

Vesicle number may change as a function of synapse size. We found that PSD length was significantly increased in longitudinal

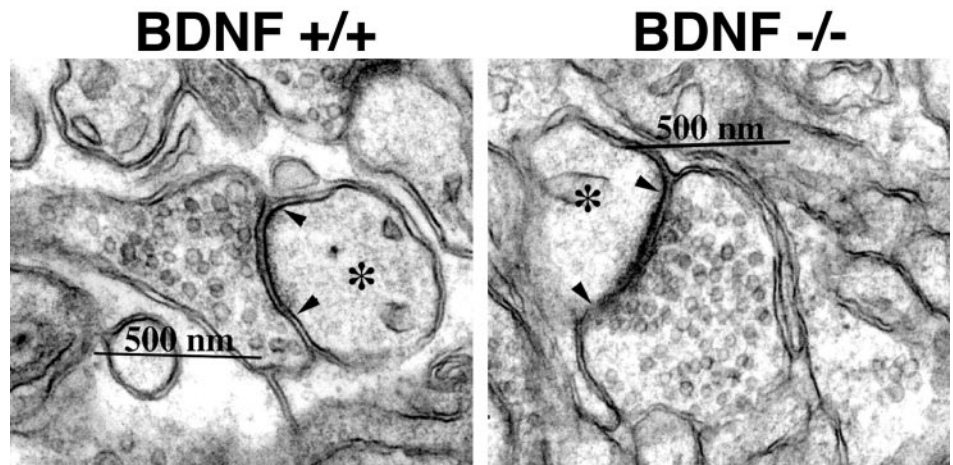


Figure 7. Electron micrographs of the PF/PC synapse in P15 WT and BDNF $-/-$ mice. Individual synapses exhibit a dendritic spine (*), postsynaptic density demarcated by arrowheads, and presynaptic terminal filled with numerous round, loosely packed vesicles in both wild-type and BDNF $-/-$ animals.

Table 1. Ultrastructural analysis of PF/PC synapses in wild-type and BDNF $-/-$ mice

	Vesicles/synapse (\pm SEM)	PSD (nm)	DV/ μ m PSD	Ratio docked/ total	Vesicles/ μ m ²
Longitudinal sections					
+/+	49.8 \pm 2.9	455.8 \pm 12.1	20.9 \pm 0.7	0.247 \pm 0.01	346.1 \pm 8.4
-/-	64.4 \pm 3.2*	517.1 \pm 13.4*	20.5 \pm 0.7	0.193 \pm 0.01*	307.8 \pm 5.8*
* p < 0.01; BDNF +/+ (172 synapses); BDNF -/- (177 synapses)					
Cross sections					
+/+	18.9 \pm 1.8	355.6 \pm 14.2	17.4 \pm 0.9	0.373 \pm 0.02	347.8 \pm 11.9
-/-	33.1 \pm 2.2*	357.0 \pm 9.4	15.9 \pm 0.7	0.214 \pm 0.01*	250.1 \pm 5.9*
* p < 0.01; BDNF +/+ (51 synapses); BDNF -/- (129 synapses)					

BDNF $-/-$ mice have a significant increase in the number of synaptic vesicles per terminal (Vesicles/synapse) and in longitudinal PSD length compared with wild type. However, no difference in the density of docked vesicles was detected (DV/ μ m PSD). Despite an elongated PSD, BDNF $-/-$ terminals had a smaller fraction of all of their vesicles that was docked (Ratio docked/total). The overall packing density of vesicles in BDNF $-/-$ terminals was also significantly decreased (Vesicles/ μ m²). Vesicles/synapse, Total vesicles per synapse; PSD (nm), postsynaptic density length; DV/ μ m PSD, docked vesicles per micrometer of active zone; Ves/ μ m², overall vesicle packing density. Numbers are mean \pm SEM; * p < 0.01 by t test.

sections of BDNF $-/-$ terminals, suggesting an overall elongation of presynaptic terminals in the absence of BDNF. To correct for the effect of synapse size, the number of docked vesicles was normalized to the length of the active zone. Similarly, the total number of vesicles was normalized to the total area occupied by the vesicle pool (Table 1). We found that the density of docked vesicles was not significantly different between BDNF +/+ and BDNF -/- terminals (Table 1). In contrast, the overall packing density of vesicles was significantly decreased in the terminals of BDNF $-/-$ mice compared with terminals in wild-type mice (Table 1). Therefore, parallel fibers terminals are not simply larger in BDNF $-/-$ mice, but exhibit an altered ultrastructure and abnormal distribution of vesicles that may reflect the change in synaptic function demonstrated here.

DISCUSSION

In the absence of BDNF, numerous PF/PC synapses form despite severe and persistent disturbances of Purkinje cell dendritic morphology. Although neurotransmission is remarkably well preserved, these synapses exhibit a specific defect in short-term synaptic plasticity. Although the amplitude of spontaneous mEPSCs, the waveforms of evoked EPSCs, and the response of the synapse to repetitive stimulation all appear normal, paired-pulse facilitation is significantly decreased in BDNF $-/-$ mice. Changes in paired-pulse facilitation are often used as a measure of changes in presynaptic function. We found that this change is associated with a significant increase in synaptic vesicle number

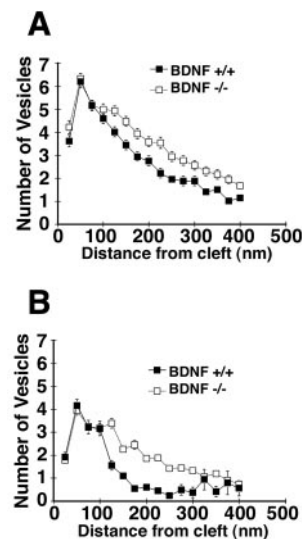


Figure 8. Increased vesicle number in BDNF $-/-$ synapses is restricted to vesicles located farthest from the active zone. The number of vesicles within consecutive 25 nm bins was plotted as a function of distance from the synaptic cleft. Within the first 100 nm from the synaptic cleft there is no difference in the number of vesicles between the wild-type mice (■) and BDNF $-/-$ mice (□) in either longitudinal sections (A) or cross sections (B). However, beyond 100 nm, BDNF $-/-$ mice exhibit a significant increase in the number of vesicles compared with WT mice. * $D \ll D_{0.01}$ by Kolmogorov–Smirnov.

and a significant decrease in the overall synaptic vesicle packing density in BDNF $-/-$ mice. The data presented here support a model in which BDNF *in vivo* acts on the PF/PC synapse to modify short-term synaptic plasticity and regulate synaptic vesicle number and distribution.

Correlating vesicle pools with synaptic function

As is the case in the hippocampus in BDNF $-/-$ mice (Pozzo-Miller et al., 1999) or TrkB $-/-$ mice (Martinez et al., 1998), BDNF is not absolutely required for synaptogenesis between granule cells and Purkinje cells to take place. In fact, in the absence of BDNF, we detected only a slight decrease in the overall PF/PC synapse density. Accordingly, BDNF $-/-$ Purkinje cell dendrites were studded with numerous spines as in wild-type cells, and most of these spines were the sites of PF:PC synapses.

Although synaptogenesis proceeds, specific structural and functional abnormalities remain in BDNF $-/-$ mice. PF:PC synapses in BDNF $-/-$ mice have more vesicles and longer PSDs than do wild-type synapses, whereas immature synapses are characterized by fewer vesicles and shorter PSDs. Thus the structural and physiologic differences seen in the PF/PC synapses of BDNF $-/-$ mice are not characteristics of developmentally delayed synapses but represent an aberrant phenotype. As demonstrated here for the cerebellum, decreased paired-pulse facilitation has been shown to be associated with structural changes in presynaptic terminals in the hippocampus of BDNF $-/-$ (Pozzo-Miller et al., 1999) and TrkB $-/-$ mice (Martinez et al., 1998). However, the nature of the ultrastructural changes is different in the two systems. Pozzo-Miller et al. (1999) observed a decrease in the number of docked vesicles in BDNF $-/-$ mice. More recently, Tyler and Pozzo-Miller (2001) have demonstrated that BDNF applied to organotypic hippocampal slices increases docked vesicle number and neurotransmitter release. In contrast, in the cerebellum, we observed no change in docked vesicle density but rather an increase in vesicles distant from the active zone. How can we reconcile these apparent differences between the hippocampus and the cerebellum? In both systems the absence of BDNF results in a significant decrease in the proportion of synaptic vesicles that are docked relative to the total number of synaptic vesicles. Different vesicle pools are likely to exist in a dynamic steady state within the terminal (Li and Schwarz, 1999). Thus, one interpretation that reconciles the ultrastructural findings of the BDNF $-/-$ hippocampus and cerebellum is that neurotrophins shift the balance between the different vesicle pools in favor of the docked pool. For example, BDNF could regulate the movement of synaptic vesicles within the terminal by modulating the interactions between motor proteins like myosin V, cytoskeletal actin, and vesicle-associated proteins such as synapsin (Hilfiker et al., 1999; Doussau and Augustine, 2000).

This study of BDNF $-/-$ cerebellar synapses provides a new example in which changes in presynaptic function correlate with alterations in vesicle number or vesicle distribution. In various systems, the degree of paired-pulse facilitation is inversely related to the number of docked vesicles (Dobrunz and Stevens, 1997; Murthy et al., 1997; Jiang and Abrams, 1998; Schikorski and Stevens, 1999). There are several examples in which mutations that alter the distribution of synaptic vesicles also lead to changes in presynaptic function. For example, mutations in synaptotagmin, the presynaptic calcium sensor, impair evoked neurotransmitter release (DiAntonio and Schwarz, 1994) and decrease the number of morphologically docked vesicles in *Drosophila* (Reist

et al., 1998). Hippocampal synapses deficient in synapsin I, an abundant presynaptic phosphoprotein, exhibit increased paired-pulse facilitation (Rosahl et al., 1993) and a significant decrease in the number of vesicles distal to the active zone (Takei et al., 1995). The unconventional myosin, myosin Va, is a motor protein that may contribute to actin-based vesicle movement in synaptic terminals (Prekeris and Terrian, 1997). In mice, mutations in myosin Va lead to progressive ataxia, loss of balance, seizures, and death (Evans et al., 1997), and in humans they lead to a rare immunodeficiency and pigmentation disorder, Griscelli syndrome (Kumar et al., 2001). Cerebellar granule cells in myosin Va mutant animals have enlarged synaptic terminals with increased numbers of vesicles suggesting the abnormal accumulation of presynaptic components (Bridgman, 1999). Acute changes in synaptic proteins can likewise alter both vesicle distribution and presynaptic function. The injection of anti-synapsin antibodies into lamprey axons depletes the distal pool of vesicles and results in a pronounced decrease in EPSP amplitude in response to high-frequency but not low-frequency stimulation (Pieribone et al., 1995). These studies are essential in helping to elucidate the relationship between synaptic vesicle distribution and synaptic function at various types of synapses. Here we have shown that facilitation is impaired in enlarged terminals where vesicles accumulate in the absence of BDNF. The increase in parallel fiber terminal size and decrease in overall vesicle density in the absence of BDNF are similar to the changes observed in the terminals of hippocampal afferents in TrkB $-/-$ mice (Martinez et al., 1998) as well as in parallel fiber terminals of myosin Va mutant mice (Bridgman, 1999) and suggest a common underlying abnormality.

Chronic and acute effects of BDNF

The changes in BDNF $-/-$ mice reported here could reflect either the long-term effects of BDNF deficit on synapse development or the acute effects of BDNF deficit on synaptic function. Neurotrophins are powerful differentiation factors and may guide the recruitment and assembly of the molecular components required for synaptic function and/or plasticity. Thus alterations in short-term plasticity and synaptic ultrastructure may represent abnormal development. This idea is consistent with the finding that exogenous BDNF fails to reverse the defect in PPF. If the changes in BDNF $-/-$ mice are caused by long-term BDNF deprivation, then BDNF rescue may require days.

Alternatively, the effects of BDNF on synaptic plasticity could be very acute and depend on the timing of endogenous BDNF release at active synapses. Bath application of BDNF may fail to reproduce the *in vivo* time course and hence fail to reverse the defect in PPF. Recent studies have demonstrated that BDNF is located within vesicles at the presynaptic terminal (Haubensak et al., 1998) and can be rapidly released in response to activity (Canossa et al., 1997; Marini et al., 1998). BDNF can also potentiate neurotransmitter release from cerebellar granule neurons (Numakawa et al., 1999; Yamagishi et al., 2000) and can cause a rapid depolarization of Purkinje cells (Kafitz et al., 1999). These rapid and local effects of BDNF may not be mimicked by prolonged bath application of neurotrophin.

Finally, the inability of exogenous BDNF to alter facilitation could reflect a postsynaptic role of BDNF. The experimental conditions used here detect changes in presynaptic function exclusively (see Materials and Methods). Thus, if endogenous BDNF acts on postsynaptic TrkB receptors and induces the release of a *trans*-synaptic retrograde signal such as a cannabinoid

(Kreitzer and Regehr, 2001; Maejima et al., 2001) or adenosine to alter paired-pulse facilitation, we would not detect any change in PPF after application of exogenous BDNF.

Taken together, the data presented here suggest that BDNF can regulate the degree of short-term synaptic plasticity that can be elicited at a synapse. Given that neurotrophin expression and release are activity dependent, neurotrophins may be important mediators of hebbian plasticity by providing neuronal circuits with a mechanism by which plasticity is favored at more active synapses. Such second-order plasticity, or metaplasticity, constitutes a powerful additional dimension in which neurons can adjust the strength of their connections to achieve their information-processing goals.

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