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著者: 松川 嘉

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Netrin-G/NGL Complexes Encode Functional Synaptic Diversification

Hiroshi Matsukawa, Sachiko Akiyoshi-Nishimura, Qi Zhang, Rafael Luján, Kazuhiko Yamaguchi, Hiromichi Goto, Kunio Yaguchi, Tsutomu Hashikawa, Chie Sano, Ryuichi Shigemoto, Toshiaki Nakashiba, and Shigeyoshi Itohara

1Laboratory for Behavioral Genetics, RIKEN Brain Science Institute (BSI), Wako, Saitama, 351-0198, Japan; 2DINE, Departamento de Ciencias Médicas, Facultad de Medicina, Universidad Castilla-La Mancha, 13071 Albacete, Spain; 3Research Resource Center, RIKEN BSI, Wako, Saitama, 351-0198, Japan; 4Division of Cerebral Structure, National Institute for Physiological Science, Okazaki 444-8787, Japan; and 5IST Austria, 3400 Klosterneuburg, Austria

Synaptic cell adhesion molecules are increasingly gaining attention for conferring specific properties to individual synapses. Netrin-G1 and netrin-G2 are trans-synaptic adhesion molecules that distribute on distinct axons, and their presence restricts the expression of their cognate receptors, NGL1 and NGL2, respectively, to specific subendodermal segments of target neurons. However, the neural circuits and functional roles of netrin-G isoform complexes remain unclear. Here, we use netrin-G-KO and NGL-KO mice to reveal that netrin-G1/NGL1 and netrin-G2/NGL2 interactions specify excitatory synapses in independent hippocampal pathways. In the hippocampal CA1 area, netrin-G1/NGL1 and netrin-G2/NGL2 were expressed in the temporoammonic and Schaffer collateral pathways, respectively. The lack of presynaptic netrin-Gs led to the dispersion of NGLs from postsynaptic membranes. In accord, netrin-G mutant synapses displayed opposing phenotypes in long-term and short-term plasticity through discrete biochemical pathways. The plasticity phenotypes in netrin-G-KOs were phenocopied in NGL-KOs, with a corresponding loss of netrin-Gs from presynaptic membranes. Our findings show that netrin-G/NGL interactions differentially control synaptic plasticity in distinct circuits via retrograde signaling mechanisms and explain how synaptic inputs are diversified to control neuronal activity.

Key words: excitatory synapse; mice; netrin-G1; netrin-G2; pathway specificity; trans-synaptic adhesion molecule

Introduction

Information processing in the brain depends on proper neuronal connectivity, which rests on a foundation of synaptic communication. For precise information processing between connected neurons, individual neurons express a variety of synaptic molecules that constitute functional assemblies necessary for the maturation of synaptic connectivity (Sheng and Kim, 2011; Südhof, 2012). Accumulating evidence suggests that synaptic cell adhesion proteins serve not only to maintain aligned synaptic membrane structures but also to control presynaptic and postsynaptic properties (Yamagata et al., 2003; Dalva et al., 2007; Han and Kim, 2008; Siddiqui and Craig, 2011). For example, neurexin-1 and neurexin-2 in vertebrates are preferentially distributed onto excitatory and inhibitory postsynaptic membranes, respectively, and thus differentially regulate excitatory and inhibitory responses (Chubykin et al., 2007; Blundell et al., 2010). Another class of cognate trans-synaptic ligand/receptor pairs, netrin-Gs/NGLs, is thought to have a unique role in neural circuit functions elaborated in vertebrates. Netrin-G1 and netrin-G2 (also known as laminet1 and laminet2) are vertebrate-specific and glycosylphosphatidylinositol (GPI)-anchored subfamily members of the UNC-6/netrin family (Nakashiba et al., 2000, 2002; Yin et al., 2002). NGL1 and NGL2 are also known as LRRRC4c and LRRRC6) are vertebrate-specific and are type I transmembrane proteins containing a C-terminal intracellular postsynaptic density-95/disks large/zona occludens-1 (PDZ) binding motif, which interacts with scaffolding proteins such as PSD-95 family members (Lin et al., 2003; Kim et al., 2006; Woo et al., 2009). Netrin-G1 and netrin-G2 selectively bind to NGL1 and NGL2, respectively. Crystallographic complex studies confirmed the direct contact of those proteins (Seiradake et al., 2011). In cultured neurons, exogenous NGLs induce presynaptic differentiation and netrin-G clustering (Kim et al., 2006; Song et al., 2013). During development, NGL2 deletion leads to a lower spine density in hippocampal pyramidal neurons (DeNardo et al., 2012). Netrin-G1 and netrin-G2 are expressed in complementary neuron groups (Nakashiba et al., 2002; Yin et al., 2002; Nishimura-Akiyoshi et al., 2007; Woo et al., 2009), and distribute on axons segregated differentially along cortical laminar structures (Nakashiba et al., 2002; Nishimura-Akiyoshi et al., 2007). Axonal netrin-G1 and
netrin-G2 constrain NGL1 and NGL2, respectively, into subden- dritic segments to establish circuit specificity in a single neuron. This suggests that netrin-G/NGL trans-synaptic interactions help to diversify information processing. However, the role of netrin- Gs/NGLs interactions in synaptic functions remains largely unknown.

Here we studied the roles of netrin-G1/NGL1 and netrin-G2/ NGL2 interactions in hippocampal excitatory circuits using a full set of KO mice for netrin-Gs and NGLs. Immunoelectron microscopy (iEM) supported the requirement of these ligand/receptor pairs to stably localize presynaptic netrin-Gs and postsynaptic NGLs in a pathway-selective manner. Moreover, netrin-G/NGL trans- synaptic interactions regulate synaptic plasticity by a retrograde sig- naling mechanism. These data reveal the significance of netrin-Gs/ NGLs systems in vertebrate brain.

Materials and Methods

Animals. All experimental procedures were performed in accordance with the guidelines of the RIKEN Institutional Animal Care and Experimentation Committee. Netrin-G1 (Ntng1–1)–KO and netrin-G2 (Ntng2–2)–KO mice were generated previously and maintained as C57BL/6J congenic heterozygotes (Nishimura-Akiyoshi et al., 2007). For Ntng1–1 disruption, short sequences including the translation initiation codon in the third exon were replaced with a loxP site. For Ntng2–2 disruption, the entire cording sequence and downstream exons were replaced with N-LacZ and a loxP site. Ntng1–1 (5'-AGGGTCTCCACAGGTAATATCC-3') and Ntng2–2 (5'-TGGAGCTCATTTGAGCAGC-3') were generated by homologous recombination using C57BL/6-derived embryonic stem cells, CMT1–2 (Mercilllipore), and MS12 (Kawase et al., 1994), respectively, and maintained as C57BL/6J congenic heterozygotes. Due to an inefficient recombination rate at the Ntng1 locus, a long targeting vector based on bacterial artificial chromosome was used. Technical details will be described elsewhere (K. Yaguchi and S.I.). In all cases, homozygous KO and WT control littersmates were obtained by crossing the heterozygotes. Adult mice (3–11 months old, otherwise described) were used in the present study. Mouse genotypes were deter- mined by PCR using primers as follows: Ntng1–1 and Ntng2–2 (described) were used in the present study. Mouse genotypes were deter- mined by PCR using primers as follows: Ntng1–1 and Ntng2–2 (described) were used in the present study. Mouse genotypes were deter- mined by PCR using primers as follows: Ntng1–1 and Ntng2–2 (described) were used in the present study. Mouse genotypes were deter- mined by PCR using primers as follows: Ntng1–1 and Ntng2–2 (described) were used in the present study. Mouse genotypes were deter- mined by PCR using primers as follows: Ntng1–1 and Ntng2–2 (described) were used in the present study. Mouse genotypes were deter- mined by PCR using primers as follows: Ntng1–1 and Ntng2–2 (described) were used in the present study. Mouse genotypes were deter- 

Electron microscopy. For post-embedding iEM, adult mice were deeply anesthetized by intraperitoneal injection of 2.5% Avertin (Sigma-Aldrich; 0.5 ml/kg body weight), and perfused with 0.1 M PBS, pH 7.4, containing 4% PFA (TAAB), 0.1% glutaraldehyde (GA; Nacalai), and 15% saturated picric acid (Nacalai). After 4–6 h post fix in 4% PFA-0.1 M PBS, the brains were coronally cut in 0.5-μm-thick slices, cryoprotected in 30% sucrose-0.1 M PBS, and freeze-thawed by dipping into liquid nitrogen for better anti- body penetration. After washing in TBS, the free-floating sections were incubated in 10% normal goat serum (NGS) diluted in TBS for 1 h at RT. Sections were then incubated with anti-netrin-G1 diluted 1:4000 (~0.1 μg/ml) in 1% NGS-TBS for >48 h at 4°C. After several washes in TBS, the sections were incubated with goat anti-rabbit IgG coupled to 1.4 nm gold (Probes) diluted 1:100 in 1% NGS-TBS overnight at 4°C. After several washes in PBS, the immunoreacted sections were post fixed in 1% GA-PBS for 10 min. They were then washed in double distilled water, followed by silver enhancement of the gold particles with an HQ Silver kit (Probes). The sections were then treated with 1% osmium tetroxide–0.1 M PB, stained with 1% uranyl acetate, dehydrated, and flat embedded in Durcupan resin (Fluka) on glass slides. Ultrastructural sections on the grids were counterstained with 1% uranyl acetate and Reynolds’ lead citrate. Ultrastructural analyses were performed in transmission EM (JEM-1010; JEOL).

For conventional transmission EM experiments, netrin-G1–KO, netrin- G2–KO, and their WT littersmates (males, 6 weeks old) were perfused with a physiologic saline (2.5 ml) followed by a mixture of 2% PFA and 2.5% GA in 0.1% cadoxylate buffer, pH 7.4, 2.5 ml, under deep anesthesia with 0.15–0.20 ml of Nembutal (sodium pentobarbital; 50 mg/ml). The brains were fixed further in 2.5% GA at 4°C overnight. Then brain slices containing the desired hippocampal areas in each animal were metal stained with lead citrate and uranyl acetate and examined under the electron microscope (1200EX; JEOL). Electron micrographs were obtained from the middle layers of the CA1 stratum radiatum (SR) and stratum lacunosum moleculare (SLM) of the dorsal hippocampus.

Light microscopic immunohistochemistry. Free-floating brain sections (50 μm thick), which had been incubated with anti-netrin-G1 (1:4000) and anti-netrin-G2 (1:400) antibodies through the same procedure as described above in the pre-embedding protocol, were incubated with HRP-conjugated anti-rabbit IgG (Vector Laboratories) diluted 1:100 in 1% NGS-TBS overnight at 4°C. After several washes in PBS, sections were incubated with avidin–biotin peroxidase complex (ABC–Elite; Vector Laboratories) diluted 1:100 in PBS for 1 h at RT. After washing three times in PBS and subsequently in 50 ml Tris-HCl, pH 7.4, the sections were then incubated in the same buffer including 0.02% DAB (Dojindo) for 5 min. DAB reaction was initiated by 0.003% hydrogen peroxide and terminated by addition of excess amounts of PBS. The sections were mounted on surface-coated slides, dehydrated, and pre- served in a quick-drying mounting medium (Eukitt) for later image acquisition. The bright-field images were taken with a digital slide scanner (NanoZoomer; Hamamatsu Photonics). For double staining of netrin-G1 and netrin-G2, cryosections (40 μm thick) were cut and incubated with rabbit polyclonal anti-netrin-G2 (1:400) and mouse mAb for netrin-G1 (1:200) overnight at 4°C. The sections were then incubated with a mixture of secondary antibodies—Alexa Fluor 546-conjugated anti-rabbit IgG (1:1000; Life Technologies) and Alexa Fluor 488-conjugated anti-mouse IgG (1:2000; Life Technologies)—for 2 h at RT, and counterstained with DAPI (1:10,000; Invitrogen).
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Figure 1. Presynaptic localization of netrin-G proteins and postsynaptic localization of NGL proteins in distinct hippocampal layers. A, Double-staining fluorescence image showing layer-selective distributions of netrin-G subtypes in the hippocampal CA1 and DG regions. SO, stratum oriens. B, Schematics of entorhino-hippocampal/intrahippocampal projections expressing netrin-Gs (red) and the local synaptic connections (left). In the hippocampus, CA1 pyramidal and DG granule cells (PC and GC) receive netrin-G1 (red)-expressing and netrin-G2 (green)-expressing inputs originating from entorhinal cortex (EC) layers II/III and CA3 areas. Scale bar, 1 mm. C, Confocal images showing double immunostaining for Bassoon and netrin-G1, netrin-G2, NGL1, and NGL2, as detected using a post-embedding immunogold technique. D, Immunogold particles (arrows) for netrin-G1 and netrin-G2 were detected mainly along the presynaptic axon terminals (at) making excitatory synapses on dendritic spines (s) of the pyramidal cells in SLM and SR, respectively. E, Immunoparticles for NGL1 and NGL2 were mostly detected postsynaptically in the vicinity of the postsynaptic density of dendritic spines in SLM and SR, respectively. Scale bars, 200 nm. F, Relative distance of the immune particles to presynaptic membrane structure was allocated to 10 nm wide bins and expressed as counts. Positive location was defined as extracellular face from presynaptic terminal (at 0 nm). Frequencies of particle distribution for netrin-G1 and netrin-G2 peaked within the synaptic cleft, while that of NGL1 nestled postsynaptically. The mean synaptic cleft size was 19.16 nm, as determined by 17 representative synapses. G, A pre-embedding immunogold technique showed multiple immunoparticles for netrin-G1 (arrows) exclusively along the longitudinally running axon structure in SLM. Scale bar, 200 nm.

Triple fluorescence images were captured and tiled to build a large-field image with a fluorescence microscope (BZ-X700; Keyence).

For double staining of netrin-G1 with Bas- soon (mouse mAb VAM-P5003; StressGen), thin sections from paraffin-embedded samples were deparaffinized, rehydrated, and processed for heat-induced epitope retrieval. Blocking was done for 1 h at RT. Then the slices were incubated with primary antibodies for 12 h and fluorescence-conjugated secondary antibodies for 4 h at RT. The confocal images were captured with 100 × oil lens by Leica TCS SP8 STEDE 3× microscope.

Dendrite morphology analysis. Mice were anesthetized with tribromoethanol (125 mg/kg, i.p.) and then perfused transcardially with 0.9% NaCl and 4% PFA in 0.1 m PB. Coronal brain slices at 150 μm thickness were prepared by a microslicer (Dosaka). To label fine structure of neurons (Gan et al., 2000; Mataga et al., 2004), tungsten particles coated with lipophilic dye (DiI; Life Technologies) were delivered to the brain slices using the Helios Gene Gun (Bio-Rad). A confocal microscope (FluoView; Olympus) was used to image secondary apical dendrites of typical pyramidal neurons at the hippocampal area CA1. Dendritic protrusions were counted in z-stacks with NIH Image.

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Electrophysiology. Mice were deeply anesthe- tized with halothane (Takeda) and killed by decapitation. The whole brain was quickly re- moved into an ice-cold cutting solution containing the following (in mM): 200 sucrose, 4 KCl, 1 NaH2PO4, 0.2 CaCl2, 10 MgCl2, 26.2 NaHCO3, 11 l-(-)glutamate, 0.1 l-(-)-aspartic acid, and 0.5 sodium pyruvate, saturated with 95% O2 and 5% CO2. Each brain hemisphere glued on a piece of agar block (4% in saline) was sliced along the longer axis of hippocampus at a thickness of 400 μm, using a microslicer (Lin- earSlicer Pro7; DOSAKA). Hippocampi were dissected out of whole-brain slices and recovered in a submerged holding chamber containing a physiologic medium (aCSF) containing the following (in mM): 119 NaCl, 2.5 KCl, 1 NaH2PO4, 2.5 CaCl2, 1.3 MgCl2, 26.2 NaHCO3, 11 l-(-)glutamate, 0.1 l-(-)-aspartic acid, and 0.5 sodium pyruvate, continuously bubbled with 95% O2 and 5% CO2. Each slice, which had been equilibrated for >2 h, was transferred to an immersion-type recording chamber and superfused with well bubbled aCSF at a rate of 2.5 μl/min. All physiologic experiments were performed at 25°C. Extracel- lular field potential recordings were performed using borosilicate glass electrodes filled with 3 m NaCl to obtain a low pipette resistance, resulting in high signal-to-noise ratio. Excit- atory synaptic potentials were evoked with tungsten bipolar stimulating electrodes (WPI), which were connected to an isolated box (ISO- Flex; A.M.P.I.) delivering constant voltage pulses of 200 μs duration to input fibers. External recording solutions contained the following (in μM): 100 picrotoxin and 1 CGBP58845 to block GABA_A and GABA_B receptors, respectively. CA3–CA1 fibers were surgically disconnected to prevent emergence of
Basil synaptic properties were examined in hippocampal CA1 and DG synapses. AMPAR-mediated fEPSPs were recorded by stimulating axon fibers. TA, SC, LPP, and MPP. Recording and bipolar stimulating electrodes were placed in the middle of each layer. Relative postsynaptic responsiveness to presynaptic activity was not significantly different between netrin-G1-KO and WT control mice. The average slopes of fEPSPs (10 consecutive recordings) were plotted as a function of the average amplitude of PSFV. The paired-pulse ratio (PPR; percentage) was not statistically different between netrin-G1-KO and WT mice. The slope ratios were obtained by average fEPSPs (10 consecutive recordings) and plotted as a function of the interstimulus interval (ISI).

Figure 2. Deficiency of netrin-G2 attenuated synaptic transmission in the specific pathway. A, Schematics of field recording. Basal synaptic properties were examined in hippocampal CA1 and DG synapses. AMPAR-mediated fEPSPs were recorded by stimulating axon fibers. TA, SC, LPP, and MPP. Recording and bipolar stimulating electrodes were placed in the middle of each layer. Relative postsynaptic responsiveness to presynaptic activity was not significantly different between netrin-G1-KO and WT control mice. The average slopes of fEPSPs (10 consecutive recordings) were plotted as a function of the average amplitude of PSFV. The paired-pulse ratio (PPR; percentage) was not statistically different between netrin-G1-KO and WT mice. The slope ratios were obtained by average fEPSPs (10 consecutive recordings) and plotted as a function of the interstimulus interval (ISI). B, Basal synaptic transmission was significantly reduced only at MPP-DG synapses, but normal at the SC-CA1 synapses of netrin-G2-KO mice. E, MPP-DG synapses of netrin-G2-KO mice showed significantly increased PPR at all interstimulus intervals, while the SC-CA1 synapses showed comparable PPR to WT mice. Absence of netrin-G1 or netrin-G2 did not affect the basal synaptic properties of the neighboring pathways. F, G, Representative sample traces of mEPSC events recorded from basal synaptic properties of the neighboring pathways. The incubation of the slices with CNQX (1 μM) resulted in inhibition of ~80% and helped to avoid the nonlinear increment of SC-CA1 fEPSPs (Shimuta et al., 2001). As an exception, the input–output test for SC-CA1 of NGL2-KO mice was performed without CNQX. The examinations for input–output relations and short-term synaptic plasticity including paired-pulse facilitation/depression (PPF/PPD), post-tetanic potentiation (PTP), and post-theta burst potentiation (PTBuP) were performed in the presence of r-AP5 (25 μM or 50 μM). To induce PTP, a single tetanic train (100 Hz, 1 s) was given to SC or TA. For LTP recordings of TA-CA1 and SC-CA1 synapses, different TBS protocols were used to obtain reliable potentiation. TA-CA1 synapses were conditioned with four TBS trains at 20 s intervals after making a 50 min baseline. Each TBS consisted of five bursts at 5 Hz, each of which has eight stimuli at 100 Hz. Note that stimulus duration was doubled into 400 μs when TA-CA1 synapses were conditioned. For the induction of LTP at SC-CA1 synapses, a single TBS train (three bursts at 5 Hz, each of which has four stimuli at 100 Hz) was given to SC. The same patterns of TBS were used to induce PTTBuP. Synaptic transmission was pharmacologically potentiated by bath application of PDBu or forskolin/IBMX. Slope of the fEPSP was analyzed for the comparison between genotypes.

Whole-cell voltage-clamp configuration was obtained by a blind access with a patch pipette (3.5–8 MΩ). The composition of the pipette solution contained the following (in mM):
Figure 3. Altered PTP in netrin-G1-KO and netrin-G2-KO caused by distinct signaling defects. A, B, PTP at the TA-CA1 synapses was impaired in netrin-G1-KO. In contrast, PTP at the SC-CA1 synapses was facilitated in netrin-G2-KO SC-CA1. C, D, EPSP potentiation by PDBu (0.4 μM) was reduced at TA-CA1 synapses in netrin-G1-KO, but normal at SC-CA1 synapses in netrin-G2-KO. The slope of EPSP was normalized to the average value before the onset of PDBu application. Insets show superimposed average traces (during 2 min) before and after the treatment at the times indicated by the numbers. The average of potentiation during the last 5 min of recording was statistically compared between genotypes (WT vs netrin-G1-KO, **p < 0.005; WT vs netrin-G2-KO, p = 0.86). E, F, EPSP potentiation induced by combination of forskolin (20 μM) and IBMX (50 μM) was not statistically different between netrin-G1 mutant and WT mice (the average of potentiation during the last 5 min of recording: WT vs netrin-G1-KO, p = 0.19; WT vs netrin-G2-KO, p = 0.16). G, Treatment of slices with PKC inhibitor, Bis (2 μM) attenuated PTP generation in both WT and netrin-G1-KO mice. H, Comparison of TA-CA1 PTP magnitudes at 20 and 40 s (left and right, respectively), following tetanic stimulation in the absence or presence of Bis. Values of bar graphs were extracted from A and G. I, EPSP potentiation by PDBu was indistinguishable in the presence of Bis between netrin-G1-KO and WT (the average of potentiation during the last 5 min of recording: p = 0.80). Insets show superimposed average traces (during 2 min) before and after the treatment at the times indicated by the numbers. J, Comparison of the PDBu-induced synaptic potentiation in the absence (left) and presence (right) of Bis. Values are averages of the last 5 min of recording in each condition. The numbers of slices (left) and mice (right) used in each experiment are shown in parentheses. Data are shown as mean ± SEM. Student’s t-test; *p < 0.05, **p < 0.01, ***p < 0.001, and absolute P-values.

122.5 Cs gluconate, 17.5 CsCl, 10 HEPES, 0.2 EGTA, 8 NaCl, 2 Mg-ATP, 0.3 Na3-GTP, and 5 QX-314, pH 7.3, 290–300 mOsM. Miniature EPSCs were recorded from DG granule cells in the presence of picrotoxin (100 μM), CGP55845 (1 μM), and tetrodotoxin (1 μM), at −80 mV. AMPA receptor and NMDA receptor currents (AMPA-EPSC and NMDAR-EPSC) were recorded at −70 mV and +50 mV/−20 mV, respectively. NMDAR-EPSCs were recorded after pharmacologic isolation with NBQX (20 μM). It should be noted that NMDAR-EPSC was evaluated by depleting membrane potential at −20 mV, not more positive potential, when a TA pathway was stimulated, because TA-evoked NMDAR-EPSC exhibits inward rectification, as previously reported (Otmakhova et al., 2002). Also, in our slice experimental conditions, the absolute amount of NMDAR-EPSC at −20 mV was equivalent to or usually greater than that recorded at +50 mV at TA-CA1 synapses (data not shown). Amplitude of the EPSC was analyzed for the comparison between genotypes. Series resistances were <30 MΩ and there was no significant difference in series resistance between experimental groups. The monitored synaptic responses were filtered at 1–2 kHz using Axon 200B and digitized at 10 kHz using Clampex software of pClamp 9 suite (Molecular Devices). Membrane potentials in whole-cell patch were represented without being compensated for liquid junction potential.

Drugs were purchased from the following sources: picrotoxin, forskolin, tetrodotoxin, and IBMX from Nacalai Tesque; CGP55845, t-AP5, CNQX, and NBQX from Tocris Bioscience; QX-314 and PDBu from Sigma; and bisindolylmaleimide I (Bis) from Calbiochem.

Immunoblot analysis. Fresh forebrains were homogenized in 10 volumes of ice-cold HEPEBS-buffered sucrose solution containing the following (in mM): 320 sucrose, 8 HEPEBS, pH 7.4, containing protease inhibitor cocktail (Roche). After centrifuging the homogenate at 1400 × g, crude membrane (P2) fraction was pelleted by further centrifuging the supernatant (S1) at 12,000 × g. The pellet was suspended and lysed with 6 mM Tris buffer, pH 8.4, on ice for 30 min. The lysate was again pelletted at 25,000 × g and resuspended with HEPEBS-buffered sucrose solution; synaptic plasma membrane (SPM) was fractionated with layered sucrose gradient (0.8 M/1.0 M/1.2 M sucrose plus 5 mM HEPEBS) at 82,700 × g. Interface between 1.0 M and 1.2 M sucrose was retrieved. Proteins of P2
(25 μg) and SPM (15 μg) fractions were separated by SDS-PAGE (10%), then transferred onto PVDF membrane (Immobil; Millipore). After blocking in TBST containing 5% skim milk, the blot membrane was incubated at 4°C with rabbit primary antibodies at appropriate dilution as follows (P2/SPM): anti-netrin-G1 (1:1000/1:2000) and anti-netrin-G2 (1:200/1:500). For normalizing protein amounts, anti-actin (Millipore) or anti-GAPDH (Ambion) antibodies were used.

Statistics. All values are expressed as mean ± SEM. The data were analyzed by the one-way ANOVA and Student’s two-tailed t test with IBM SPSS Statistics (version 21).

Results

Synaptic localization of netrin-Gs and NGLs

Immunohistochemical findings at the light microscopic level indicate that netrin-G1 and netrin-G2 distribute along distinct axons, and their receptors, NGL1 and NGL2, are concentrated on subdendritic segments in a layer-selective manner (Nishimura-Akiyoshi et al., 2007). However, subcellular localizations of netrin-Gs and NGLs in vivo remain obscure. We examined their localization using subtype-specific antibodies for these molecules. In the hippocampal CA1 and DG, the distal layer [SLM and outer molecular layer (OML)] and proximal/middle layer [SR and middle molecular layer (MML)] were clearly distinguished by immunoreactivity for netrin-G1 and netrin-G2, respectively, representing distinct inputs to CA1 pyramidal and DG granule cells (Fig. 1A, B). Confocal microscopic analysis showed close localization of netrin-G1 and a presynaptic marker Bassoon in the OML (Fig. 1C). The subcellular distributions of netrin-G and NGL proteins in the hippocampus were further examined by post-embedding iEM. Excitatory asymmetric synapses immunolabeled with netrin-G1 and netrin-G2 antibodies were detected in the SLM and SR, respectively (Fig. 1D), and immunosignals for NGL1 and NGL2 were detected along PSDs of the SLM and SR, respectively (Fig. 1E). The specificity of the antibodies used in the post-embedding preparation was assessed by counting synapses with one or more immunogold particles in a given observation area between WT and KO ultra-thin sections, and between the distinct layers of WT samples (Fig. 7C,D; data not shown). The location of immunoparticles at synapses was determined by measuring the radial distance of each gold particle in relation to the presynaptic membrane. Netrin-G1 and netrin-G2 immunoparticles had peak radial distances within the first external bin from the presynaptic membrane (i.e., within the synaptic cleft), while the frequency of NGL1 labeling was highest in the postsynaptic face (Fig. 1F). The data for NGL2 were excluded from this analysis, because of its modest signal-to-noise ratio in the post-embedding experiments. The pre-embedding iEM revealed distribution of netrin-G1 along the axon (Fig. 1G). These findings indicate that netrin-Gs are presynaptic proteins and NGLs are postsynaptic proteins.
Netrin-G supports synaptic transmission at complementary pathways

To determine whether the lack of netrin-G leads to structurally abnormal synapses, synapse density and size were analyzed in CA1 subregions, the SLM and SR, of netrin-G1-KO and netrin-G2-KO mice by conventional transmission EM. The differences in these parameters among adult WT, netrin-G1-KO, and netrin-G2-KO mice were not significant [SR synapse density (average ± SEM)/μm²: WT, 0.94 ± 0.04; G1-KO, 0.95 ± 0.04; G2-KO, 1.03 ± 0.04; F(2,216) = 1.345, p = 0.263, one-way ANOVA; SLM synapse density/μm²: WT, 0.60 ± 0.04; G1-KO, 0.56 ± 0.03; G2-KO, 0.63 ± 0.04; F(2,199) = 0.931, p = 0.492; SR synapse size (width, μm): WT, 192.0 ± 2.87; G1-KO, 198.9 ± 3.41; G2-KO, 200.1 ± 2.91; F(2,199) = 2.091, p = 0.124; SLM synapse size: WT, 229.3 ± 4.6; G1-KO, 238.8 ± 6.4; G2-KO, 235.6 ± 5.8; F(2,792) = 0.683, p = 0.506].

To investigate the functional consequence of disrupted netrin-G/NGL interactions at synapses, we first examined synaptic transmission of hippocampal circuits in netrin-G-KO mice. AMPAR-mediated fEPSPs were recorded in the hippocampal CA1 and DG upon stimulation of SC/TA and MPP/LPP pathways, respectively (Fig. 2A). In the netrin-G1-KO, the relation between the presynaptic fiber volley activity (PSFV) and fEPSP was not altered at all synapses tested (Fig. 2B). Furthermore, synaptic facilitation in response to paired-pulse stimuli, which is inversely correlated with release probability (Zucker and Regehr, 2002), was not significantly distinguishable between genotypes (Fig. 2C). Netrin-G2 deficiency affected basal synaptic transmission in a pathway-specific manner. Basal synaptic transmission was significantly reduced at MPP-DG synapses, whereas it was normal at other synapses (Fig. 2D). PPD was also attenuated specifically at MPP-DG synapses in netrin-G2-KO mice (Fig. 2E), suggesting impaired release probability. The frequency and amplitude of the mEPSCs, from DG granule cells, however, did not differ between the netrin-Gs mutants and control WT mice (Fig. 2F, G), suggesting a significant role of netrin-G2 on evoked vesicular release. Loss of the netrin-G1 or netrin-G2 subtype in one pathway did not affect the basal synaptic properties at the neighboring netrin-G1-negative or netrin-G2-negative hippocampal circuits.

Presynaptic plasticity is differentially regulated by netrin-G1 and netrin-G2 complexes

We wondered if presynaptic deficits in netrin-Gs-KO might manifest under physiologically challenging paradigms in the TA-CA1 and SC-CA1 circuits that showed no abnormalities in basal synaptic transmission. We induced synaptic plasticity and investigated PTP under the presence of NMDAR antagonist, which is a mostly presynapse-dependent short-term plasticity (Zucker and Regehr, 2002; Fioravante et al., 2011). PTP induction was attenuated at TA-CA1 synapses of netrin-G1-KO and facilitated at SC-CA1 synapses of netrin-G2-KO mice (Fig. 3A, B). Pharmacologic evidence suggests that PTP is largely mediated by the activation of presynaptic PKC induced by repetitive synaptic firing. Bath application of phorbol ester potentiated evoked transmitter

Figure 5. Impaired short-term plasticity induced by TBuS in netrin-Gs-KO mice. A, TBuS train used for TA-CA1 LTP induction. Each TBuS composed five bursts (1-5, 5 Hz) of eight pulses (100 Hz). Development of TA-CA1 fEPSP by the consecutive four trains of TBuS (1-4), in the presence of o-AP5 (50 μM), was impaired in netrin-G1-KO, compared with the WT mice. Superimposed are representative TA-CA1 fEPSPs traces during the burst (1) of each TBuS (blue arrow), scaling with the peak amplitude of fEPSP evoked by initial stimulus of TBuS (1). B, Relative slope of TA-CA1 fEPSP in multiple TBuS conditionings in A was plotted with the number of TBuS. C, Synaptic potentiation following consecutive four TBuS trains (PTBuS) was decreased in netrin-G1-KO, compared with WT mice. D, TBuS train used for SC-CA1 LTP induction, which comprises three bursts (1-3, 5 Hz) of four pulses (100 Hz). Superimposed are representative fEPSP traces elicited by three bursts (1-3) in the presence of o-AP5 (50 μM), scaling with the peak amplitude of fEPSP evoked by initial stimulus of burst (1). Intraburst synaptic facilitation was aberrantly enhanced in netrin-G2-KO, when compared with WT mice. E, Relative amplitude of SC-CA1 fEPSP during each burst in D was plotted with the number of stimulus pulse (1-4). F, PTBuS was slightly but significantly enhanced in netrin-G2-KO, compared with WT mice. The numbers of slices (left) and mice (right) used in each experiment are shown in parentheses. Data are represented as mean ± SEM. Student’s t test, *P < 0.05, **P < 0.01, ***P < 0.001.
release at many synapses (Malenka et al., 1986; Shapira et al., 1987; Hori et al., 1999; Lou et al., 2005; Korogod et al., 2007; Wierda et al., 2007; Lou et al., 2008) and occludes PTP generation (Korogod et al., 2007), and PKC antagonists diminish PTP (Brager et al., 2003; Beierlein et al., 2007; Korogod et al., 2007; Lee et al., 2007). Therefore, we investigated the enhancing effect of a phorbol ester derivative, PDBu, which activates PKC as well as presynaptic active-zone protein Munc13s (Hori et al., 1999; Lou et al., 2008), toward TACa1 and SC-Ca1 synaptic responses, to examine whether the differentially impaired PTP observed in netrin-G1-KO and netrin-G2-KO mice was due to the disruption of an intrinsic signaling pathway through PKC. Both TACa1 and SC-Ca1 EPSPs gradually increased upon PDBu (0.4 μm) application. Synaptic responsiveness to the drug was significantly impaired at netrin-G1-KO TACa1 synapses (Fig. 3C), but unaltered at netrin-G2-KO SC-Ca1 synapses (Fig. 3D). We next examined the effect of a combination of forskolin (20 μm) and IBMX (50 μm), which promotes PKA and guanine nucleotide exchange factor directly activated by cAMP (cAMP-GEF, RapGEF, or Epac) signaling processes by raising intracellular cAMP levels, leading to enhanced neurotransmission (Chavez-Noriega and Stevens, 1994; Sakaba and Neher, 2001; Kaneko and Takahashi, 2004; Huang and Hsu, 2006). Chemical potentiation by forskolin/IBMX was not statistically distinguishable between netrin-G-KO and WT mice (Fig. 3E,F). These findings indicate that netrin-G1 deficiency at TACa1 synapse selectively impairs the signaling pathway downstream of diacylglycerol, which bifurcates PKC-dependent (through phosphorylation by PKC) and PKC-independent (through diacylglycerol binding substrates such as Munc-13s) signaling components. Pre-incubation with a selective PKC inhibitor, bisindolylmaleimide I (Bis; 2 μm), diminished the difference in the magnitude of PTP at TACa1 synapses (Fig. 3G,H), and the PDBu-dependent potentiation between netrin-G1-KO and WT mice (Fig. 3I,J). These findings suggest that intrinsic phosphorylation processes mediated by PKC are hampered or that the functionality of the PKC-targeted substrate is impaired at netrin-G1-deficient TA presynaptic terminals, whereas netrin-G2 is coupled to distinct signaling from PKC, in hippocampal CA1 circuits. Interestingly, forskolin treatment became phenotypic in the PPF at the netrin-G2-KO SC-Ca1 synapses (Fig. 3K), while no differences were noted under naive conditions (Fig. 2E). Moreover, forskolin treatment abolished the PPD phenotype in netrin-G2-KO MPP-DG synapses (Fig. 3L). These findings suggest that netrin-G2 has a role in regulating presynaptic mechanisms involving CAMP. These findings suggest that both netrin-G subtypes are functionally involved in synaptic efficacy modulation induced by intense synaptic activities, such as high-frequency stimuli, although they are not necessarily required to maintain basal transmission in hippocampal CA1 circuits.

**Hippocampal LTP is differently supported by netrin-G1 and netrin-G2 complexes**

We further analyzed LTP induced by TBU5 in hippocampal CA1 circuits of netrin-G1-KO and netrin-G2-KO mice. LTP was induced in an NMDAR-dependent manner (Figs. 4A, B, 5C,F), and the magnitudes were progressively increased by multiple conditionings in a stepwise manner (Fig. 4A, B). Although initial conditioning of the TA-Ca1 synapse was insufficient to cause a significant difference in the magnitude of LTP, subsequently applied multiple conditionings revealed impaired LTP in netrin-G1-KO mice (Fig. 4A, top). In contrast, SC-Ca1 LTP was facilitated in netrin-G2-KO mice (Fig. 4B, bottom). The effects of KO on LTP were specific to the pathway where the protein was expressed. At the pathway where the protein was not expressed, LTP was intact (Fig. 4A, bottom; B, top).
basal synaptic transmission mediated by AMPAR activation was NMDAR in the brain (95, PSD-93, SAP-102, and SAP-97, via C-terminal PDZ binding to the ligand/receptor interaction).

To gain insights into the alternative possibilities, we generated NGL1-KO and NGL2-KO mice and examined synaptic plasticity in these mice. We observed that PTP was attenuated at TA-CA1 synapses of adult NGL1-KO mice, while basal transmission and PPF were normal (Fig. 6A–C), similar to the phenotypes of adult netrin-G1-KO mice. In adult NGL2-KO mice, we observed normal basal synaptic transmission at the SC-CA1 synapses and attenuated PPD at MMP-DG synapses (Fig. 6D,E), similar to the phenotypes of netrin-G2-KO mice. These findings suggest that netrin-G/NGL interactions are required for the presynaptic function of netrin-G proteins.

NGLs potentially interact with PSD proteins, including PSD-95, PSD-93, SAP-102, and SAP-97, via C-terminal PDZ binding sequences, and NGL2 forms macromolecules with PSD-95 and NMDAR in the brain (Kim et al., 2006). The lack of netrin-G1 and netrin-G2 causes diffusion of NGL1 and NGL2 from a circuit-selective subendritic segment to the entire dendrite in vivo (Nishimura-Akiyoshi et al., 2007). Thus, postsynaptic protein–protein network function, including NMDARs, might be involved in the LTP phenotypes. NMDAR activation is essential for LTP induction. We therefore examined whether postsynaptic NMDARs are functionally altered in mice lacking one of the netrin-Gs. Netrin-G deficiency, however, did not affect the amplitude ratio of NMDAR- and AMPAR-mediated EPSCs (Fig. 4C,D) or the charge transfer ratio (data not shown). Because the basal synaptic transmission mediated by AMPAR activation was indistinguishable in the hippocampal CA1 area between genotypes (Fig. 2B,D), it is unlikely that NMDAR functionality is impaired at synapses that are netrin-G deficient. Therefore, opposing LTP phenotypes, decreased or increased in netrin-G1–KO and netrin-G2–KO synapses, respectively, could not be attributed to NMDAR malfunction.

Figure 7. Disappearance of presynaptic netrin-Gs in NGL-KO mice. A, B, Subregional iEM analyses for netrin-Gs in NGL-KO mice. Ultrathin sections from NGL1-KO and NGL2-KO mice were immunoreacted with antibodies against netrin-G1 and netrin-G2, respectively. C, D, Similar iEM analyses for netrin-Gs in netrin-G-KO mice revealed specificity of immune signals. Counts of synapses labeled with one or more immunogold(s) are shown on top of each bar.?

PTBuP, short-term potentiation after TBuS, was also attenuated at TA-CA1 synapses of netrin-G1–KO mice and facilitated at the SC-CA1 synapses of netrin-G2–KO mice (Fig. 5C,F). These results support the notion that the LTP phenotypes are at least in part due to functional consequences at the presynaptic site.

Presynaptic short-term plasticity is regulated by NGLs similar to netrin-Gs

The lack of netrin-G1 or netrin-G2 caused circuit-selective abnormalities in synaptic plasticity, such as attenuated PTP in the netrin-G1–KO TA-CA1 pathway and attenuated PPD in the netrin-G2–KO MMP-DG pathway. The synaptic abnormalities could be caused by disruption of the netrin-G/NGL interactions or loss of the cell-autonomous functions of netrin-Gs unrelated to the ligand/receptor interaction.

To test these hypotheses, we performed iEM with specific antibodies against netrin-G1 and netrin-G2 in hippocampal subregions of NGL1-KO and NGL2-KO mice. Compared with the frequency of synapses immunolabeled on WT ultrathin sections, those labeled with anti-netrin-G1 and anti-netrin-G2 antibodies were negligible on NGL1-KO and NGL2-KO sections (Fig. 7). These findings suggest that postsynaptic NGLs are required to retain netrin-Gs on presynaptic membranes, which is a mechanistic explanation for the nearly identical presynaptic phenotypes between netrin-G-KO and NGL-KO mice.

Synaptic localization of netrin-Gs is regulated by NGLs

Our results thus far suggested that netrin-G/NGL interactions are required for the presynaptic function of netrin-G proteins. One possible mechanism is that localization of netrin-Gs within presynaptic membrane is regulated by postsynaptic NGLs at the synaptic cleft. To examine this possibility, we performed iEM with specific antibodies against netrin-G1 and netrin-G2 in hippocampal subregions of NGL1-KO and NGL2-KO mice. Compared with the frequency of synapses immunolabeled on WT ultrathin sections, those labeled with anti-netrin-G1 and anti-netrin-G2 antibodies were negligible on NGL1-KO and NGL2-KO sections (Fig. 7). These findings suggest that postsynaptic NGLs are required to retain netrin-Gs on presynaptic membranes, which is a mechanistic explanation for the nearly identical presynaptic phenotypes between netrin-G-KO and NGL-KO mice.

To gain further insight into the netrin-G/NGL interaction in vivo, we quantified the amount of netrin-G attached to the plasma membrane (P2-netrin-G) and synaptically localized netrin-G (SPM-netrin-G) in NGL-KO mice. We previously demonstrated that a lack of netrin-G1 and netrin-G2 affects the distribution but not the amount of NGL1 and NGL2 (Nishimura-Akiyoshi et al., 2007). Interestingly, total amounts of

NGLs potentially interact with PSD proteins, including PSD-95, PSD-93, SAP-102, and SAP-97, via C-terminal PDZ binding sequences, and NGL2 forms macromolecules with PSD-95 and NMDAR in the brain (Kim et al., 2006). The lack of netrin-G1 and netrin-G2 causes diffusion of NGL1 and NGL2 from a circuit-selective subendritic segment to the entire dendrite in vivo (Nishimura-Akiyoshi et al., 2007). Thus, postsynaptic protein–protein network function, including NMDARs, might be involved in the LTP phenotypes. NMDAR activation is essential for LTP induction. We therefore examined whether postsynaptic NMDARs are functionally altered in mice lacking one of the netrin-Gs. Netrin-G deficiency, however, did not affect the amplitude ratio of NMDAR- and AMPAR-mediated EPSCs (Fig. 4C,D) or the charge transfer ratio (data not shown). Because the basal synaptic transmission mediated by AMPAR activation was indistinguishable in the hippocampal CA1 area between genotypes (Fig. 2B,D), it is unlikely that NMDAR functionality is impaired at synapses that are netrin-G deficient. Therefore, opposing LTP phenotypes, decreased or increased in netrin-G1–KO and netrin-G2–KO synapses, respectively, could not be attributed to NMDAR malfunction.

The fEPSPs just after the first TBuS conditioning were proportionally attenuated at the TA-CA1 synapses of netrin-G1–KO mice, and significantly potentiated at the SC-CA1 synapses of netrin-G2–KO mice, implying a presynaptic mechanism(s) underlying the LTP phenotypes. To test this possibility, we applied the same stimulation protocol for inducing LTP under the presence of NMDAR antagonist D-AP5 (Fig. 5A,D). As expected, the enhancement during the TBuS(s) was significantly attenuated at TA-CA1 synapses of netrin-G1–KO mice and facilitated at the SC-CA1 synapses of netrin-G2–KO mice (Fig. 5B,E), The
P2-netrin-G1 were decreased in NGL1-KO mice (Fig. 8A). The reduction of SPM-netrin-G1 was more evident in NGL1-KO mice (Fig. 8B). Consistent with the biochemical data, immunohistochemical analysis revealed a markedly reduced intensity for netrin-G1 in NGL1-KO mice (Fig. 8C). Multiple netrin-G1 bands in the Western blot (WB) analysis represent isoforms generated by alternative splicing, and these isoforms are differentially expressed in distinct neuronal subsets (Nakashiba et al., 2000). The reduction rate of the netrin-G1 isoforms varied in P2 and SPM fractions. These results might suggest the presence of an isoform-selective mechanism for maintaining presynaptic netrin-G1. Similar to the case of NGL1-KO mice, WB analysis revealed clear reduction of netrin-G2 in P2 and SPM fractions of NGL2-KO mice (Fig. 8D,E). The reduction rate was greater in SPM fraction than P2 fraction. NGL2-KO mice, however, showed an almost comparable intensity and pattern of immunosignals for netrin-G2 in immunohistochemical analysis (Fig. 8F). The discrepancy between biochemical and immunohistochemical analyses for netrin-G2 may reflect the powerful signal amplification effects of the method used. Alternatively, relative amounts of membrane-bound and unbound (secreted) isoforms of netrin-G2 could be changed in NGL2-KO mice. In any case, these findings indicate that the trans-synaptic netrin-G/NGL interactions not only localize, but also stabilize, netrin-Gs at the presynaptic sites (Fig. 9A).

**Discussion**

Netrin-G1 and netrin-G2 are expressed throughout the brain in distinct but complementary regions (Yaguchi et al., 2014), and they form specific molecular complexes with NGL1 and NGL2, respectively. We provide evidence that netrin-Gs and NGLs localize at presynaptic and postsynaptic membranes, respectively, and that trans-synaptic netrin-G/NGL interactions play a crucial role in differentially controlling synaptic plasticity in distinct excitatory circuits in the hippocampus.

**Molecular interaction of netrin-Gs and NGLs guides synaptic transmission**

Loss of presynaptic netrin-Gs results in the mislocalization of NGLs throughout the entire dendritic tree (Nishimura-Akiyoshi et al., 2007). Our iEM and biochemical analyses of NGL-KO mice indicated that netrin-Gs required postsynaptic NGLs to be stably localized within presynaptic membranes for maintaining the synaptic integrity. Unlike NGLs in netrin-G-KO mice (Nishimura-Akiyoshi et al., 2007), netrin-G1 and netrin-G2 proteins were significantly reduced in NGL1-KO and NGL2-KO mice, respectively.

In vitro experiments suggest that the lack of netrin-G/NGL interactions causes deficits in synaptogenesis (Kim et al., 2006). An in vivo study revealed reduced synapse density in the CA1 of NGL2-KO juvenile mice, suggesting a role of netrin-G2/NGL2 interactions in synapse formation (DeNardo et al., 2012). No morphologic abnormalities, however, were evident in the CA1 of adult netrin-G KO mice. Moreover, we observed unaltered basal synaptic transmission in CA1 circuits in both netrin-G-KO mice. Unlike DeNardo et al. (2012), we observed normal synaptic transmission in the CA1 circuits of our adult NGL2-KO mice. We also observed no differences in CA1 pyramidal neuron spine den-
Imbalance and/or mislocalization of NGLs potentially form a macromolecular complex with NMDARs and other postsynaptic molecules via scaffolding proteins, PSD-95, PSD-93, SAP102, and SAP97, which are abundantly localized at the PSDs of excitatory synapses (Kim et al., 2006). The associations are mediated via the C-terminal PDZ-binding motif, which is common to both NGL1 and NGL2 (Lin et al., 2003). Mutant mouse studies demonstrated that these PSD proteins exhibit distinct roles in LTP induction, e.g., both PSD-95-KO and SAP-102-KO mice show enhanced LTP, while PSD-93-KO mice have impaired LTP (Migaud et al., 2006; Cuthbert et al., 2007; Carlisle et al., 2008). Additionally, whirlin, a PDZ domain-containing adaptor protein, binds the C-terminal PDZ domain of NGL1 (Delprat et al., 2005). Whirlin is expressed in various tissues, including the brain, and is involved in actin polymerization in hair cells (Mbure et al., 2003; van Wijk et al., 2006). The lack of netrin-G1 and netrin-G2 might disturb the protein–protein network at a circuit-specific PSD environment due to the resultant removal of NGLs from postsynaptic structures (Nishimura-Akiyoshi et al., 2007). Imbalance and/or mislocalization of NGLs-associating PSD proteins in a specific circuit (a subdendritic segment) may explain some of the mechanisms underlying differential LTP abnormalities between netrin-G1-KO and netrin-G2-KO mice (Fig. 4). Postsynaptic NGLs might be involved in membranous remodeling of the spines mediated by a dynamic transition in equilibrium between F-actin and G-actin, which is correlated with the expression of long-term plasticity (Okamoto et al., 2004). Precise roles of netrin-G/NGL/PSD interactions at the PSD environment remain to be determined.

**Netrin-G1/NGL1 and netrin-G2/NGL2 interactions diversify presynaptic plasticity**

Basal synaptic transmission and short-term plasticity evoked by paired stimuli in hippocampal CA1 circuits were not obviously altered in netrin-G1-KO or netrin-G2-KO mice (Fig. 2). Repetitive synaptic stimulations, such as TBuS and tetanus, applied to CA1 subregions produced opposing synaptic phenotypes between netrin-G1-KO and netrin-G2-KO mice (Figs. 3–5). How are these netrin-G/NGL interactions involved in the induction of such activity-dependent synaptic efficacy modifications? At TA-CA1 synapses of netrin-G1-KO mice, PTP, PTBuP, and LTP were all attenuated. On the other hand, at SC-CA1 synapses of netrin-G2-KO mice, PTP, PTBuP, and LTP were all enhanced. PTP is widely accepted as a short-term plasticity induced by presynaptic mechanisms (Zucker and Regehr, 2002; Fioravante et al., 2011). Consistency in the directions of the abnormalities may suggest that the same presynaptic mechanism underlies the short-term and long-term plasticity. This notion was supported by observations during TBuS to induce LTP, in which EPSPs were attenuated in netrin-G1-KO TA-CA1, but enhanced in netrin-G2-KO SC-CA1.

Pharmacologic experiments suggested that presynaptic netrin-G1 is coupled with PKC signaling. Netrin-G1 might be involved in the process of PKC translocation onto the juxtamembrane or regulation of the PKC target(s), such as SNAP-25, munc-18, and Ca2+ channels, which have crucial roles in release mechanisms (Nagy et al., 2002; Lou et al., 2005; Korogod et al., 2007; Wierda et al., 2007; Shu et al., 2008). A novel cis-binding partner for netrin-G1/NGL1, LAR, might have an important role in this mechanism as a signal transducer, as netrin-G1 has no...
cytoplasmic domain. LAR possesses two tyrosine phosphatase domains (D1 and D2). D1 is catalytically active, whereas D2 is inactive but can physically interact with liprin-α and β-catenin (Serra-Pagés et al., 1995; Kypta et al., 1996). As demonstrated in dissociated hippocampal neurons, cis-interactions between netrin-G1 and LAR are required for presynaptic differentiation, but are independent of LAR phosphatase activity (Song et al., 2013). Liprin-α is an especially important binding partner of LAR and implicated in active zone assembly (Zhen and Jin, 1999; Kaufmann et al., 2002; Dai et al., 2006). Thus, the netrin-G1/NG1/LAR complex might situate vesicle fusion machineries near vesicular release sites and control accessibility of these proteins to PKC depending on synaptic activity.

Netrin-G2 is also coupled to a neuronal activity-dependent biochemical pathway that differs from the netrin-G1-dependent pathway at the TA-CA1 synapses. Netrin-G2-KO SC-CA1 synapses exhibited facilitated PTP, PTBuP, and LTP, PDBu induced no difference between netrin-G2-KO and control mice. Interestingly, the netrin-G2-KO SC-CA1 synapses showed impaired PPF under the forskolin treatment, while no differences were noted under naive conditions. Forskolin treatment abolished the PPD phenotype in netrin-G2-KO MPP-DG synapses (Fig. 3). These findings suggest that netrin-G2 has a role in regulating presynaptic mechanisms involving cAMP. The cAMP-signaling cascade has a crucial role in release machineries (Lu et al., 2006). A GPI-linked isoform of the IgD receptor mediates B-cell activation through cAMP-dependent signaling pathways (Chaturvedi et al., 2002). Ephrin-A family members are GPI-anchored cell adhesion molecules that associate in cis with two different BDNF receptors, TrkB and p57NTR, which are type I transmembrane proteins that regulate axon elongation and retraction, respectively (Lim et al., 2008; Marler et al., 2008). Thus, we hypothesize that the opposite presynaptic phenotypes between netrin-G1-KO and netrin-G2-KO mice are due to the biochemical properties of the specific cis-partners. Future studies aimed at identifying the predicted cis-binding partners will help to clarify the mechanisms depending on presynaptic netrin-Gs in transmitter release and presynaptic plasticity.

The presynaptic plasticity is regulated by a retrograde mechanism of netrin-G/NGL interaction. A similar mechanism for regulating presynaptic release machineries has been suggested in the neurexin/neuroligin system (Futai et al., 2007). Among related systems, the netrin-G/NGL system is unique in its role to control differentially distinct excitatory inputs for integrating information into a single cell (Fig. 9B).

Netrin-G/NGL trans-synaptic interactions specify circuit properties

Netrin-Gs/NGLs trans-synaptic interactions have regulatory roles in hippocampal synaptic transmission and plasticity (Figs. 2–5). LTP is generally believed to be the basis for information storage, such as episodic memory (Rumpel et al., 2005; Whitlock et al., 2006). PTP is also thought to influence working memory and information processing (Silva et al., 1996; Abbott and Regehr, 2004). In hippocampal and cortical laminar structures, principal neurons widely extend their dendritic branches, traversing multiple laminar structures and allowing them to receive distinct information flows. These distinct synaptic inputs are precisely integrated within the arborized dendrite architectures for generating output spiking (Spruston, 2008). Input location is a key factor for interactions across parallel distinct layers. For example, distal inputs alone only slightly affect somatic spikes, but can modulate proximal input for neuronal output (Remondes and Schuman, 2002; Larkum et al., 2004; Dudman et al., 2007). Thus, the relative balance between parallel excitatory inputs is likely critical to associatively optimize the mode of somatic output. Altered remodeling function of the input strength contributes to aberrant modulation of the output. In mouse brain, netrin-G1 is strongly expressed in distal laminar layers, while netrin-G2 is preferentially expressed in the proximal layers (Nishimura-Akiyoshi et al., 2007). The anatomic characteristics suggest that netrin-G1/NGL1 interactions serve as an intermodule regulator, while netrin-G2/NGL2 interactions serve as an intramodule regulator.

The results of our study indicate that netrin-G1/NGL1 and netrin-G2/NGL2 interactions trans-synaptically and differentially regulate synaptic plasticity of selective excitatory inputs. This netrin-G code for synaptic plasticity offers insight into how diversity in neural circuits is generated (Fig. 9B).

By acquiring differential expression patterns of paralogs netrin-G1 and netrin-G2 and having one-to-one relationships with binding partners, these ligand/receptor pairs had unique roles in elaborating higher vertebrate brain function. Several human studies suggest the involvement of netrin-Gs in mental and neurologic disorders, such as schizophrenia (Aoki-Suzuki et al., 2005), bipolar disease (Eastwood and Harrison, 2008), and autism (O’Roak et al., 2012). Further studies are needed to determine how the interactions integrate distinct information.

References


