

# Protein Tyrosine Phosphatase Receptor Type Z Dephosphorylates TrkA Receptors and Attenuates NGF-dependent Neurite Outgrowth of PC12 Cells

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**Protein tyrosine phosphatase receptor type Z (Ptpz/Ptp $\zeta$ /RPTP $\beta$ ) is a receptor-like protein tyrosine phosphatase (RPTP) which is predominantly expressed in the central nervous system. Tropomyosin-related kinases (Trks) are single-pass transmembrane molecules that are highly expressed in the developing nervous system. Upon the ligand binding of neurotrophins, Trk receptors are activated through autophosphorylation of tyrosine residues; however, the PTPs responsible for the negative regulation of Trk receptors have not been fully elucidated. Here, we identified Ptpz as a specific PTP that efficiently dephosphorylates TrkA as a substrate. Co-expression of Ptpz with Trk receptors in 293T cells showed that Ptpz suppresses the ligand-independent tyrosine phosphorylation of TrkA, but not of TrkB or TrkC, and that Ptpz attenuates TrkA activation induced by nerve growth factor (NGF). Co-expression analyses with TrkA mutants revealed that Ptpz dephosphorylates phosphotyrosine residues in the activation loop of the kinase domain, which are requisite for activation of the TrkA receptor. Consistent with these findings, forced expression of Ptpz in PC12D cells markedly inhibited neurite extension induced by a low dose of NGF. In addition, an increment in the tyrosine phosphorylation of TrkA was observed in the brain of Ptpz-deficient mice. Ptpz thus appears to be one of the PTPs which regulate the activation and signalling of TrkA receptors.**

**Key words:** Ptpz, TrkA, NGF, dephosphorylation, neurite extension.

Abbreviations: CNS, central nervous system; FRS2, fibroblast growth factor receptor substrate 2; Grb2, growth factor receptor-bound protein 2; ICR, intra-cellular region; NGF, nerve growth factor; PLC- $\gamma$ , phospholipase C- $\gamma$ ; PSD-95, postsynaptic density-95; PTK, protein tyrosine kinase; PTP, protein tyrosine phosphatase; Ptpz, protein tyrosine phosphatase receptor type Z; RPTK, receptor PTK; RPTP, receptor-like PTP; Shc, Src-homology and collagen; Trk, tropomyosin-related kinase.

Activation of a receptor protein tyrosine kinase (RPTK) by a given extra-cellular stimulus, such as a growth factor, initiates a highly orchestrated signalling cascade. On binding a ligand, a RPTK becomes phosphorylated at multiple tyrosine residues, which serve as docking sites for distinct signalling proteins (1). The spectrum of signalling molecules that associate with the RPTK determine the nature of the response following stimulation by the ligand (1). On the other hand, protein tyrosine phosphatases (PTPs) are assumed to be key regulatory molecules in kinase cascades (2). In some studies including ours, receptor-like PTPs (RPTPs) have been shown to dephosphorylate a particular phosphorylated tyrosine residue in a RPTK and thereby determine the specificity of a signalling event rather than simply acting as an 'off-switch' (3, 4). Identifying PTPs that regulate (attenuate) RPTKs should reveal novel mechanisms by which extra-cellular stimuli and intrinsic programs regulate RPTK function.

Neurotrophin signalling is critical for normal nervous system development and function, including survival, differentiation, axonal and dendritic growth, neurotransmitter secretion and neuronal plasticity (5). In general, each neurotrophin binds to a specific tropomyosin-related kinase (Trk): nerve growth factor (NGF) to TrkA; brain-derived neurotrophic factor (BDNF) and neurotrophin-4/5 (NT-4/5) to TrkB; and neurotrophin-3 (NT-3) to TrkC (5). The intra-cellular pathways stimulated by Trk receptors share many common substrates with other receptor tyrosine kinases, such as Src-homology and collagen (Shc), growth factor receptor-bound protein 2 (Grb2), fibroblast growth factor receptor substrate 2 (FRS2) and phospholipase C $\gamma$  (PLC- $\gamma$ ) (6). While the signal transduction pathways used by the Trk receptors are well understood, the PTPs responsible for the regulation of Trk receptors have not been fully elucidated.

Protein tyrosine phosphatase receptor type Z and type G (Ptpz/Ptp $\zeta$ /RPTP $\beta$  and Ptpg/Ptp $\gamma$ , respectively) structurally resemble each other and form the R5 subfamily of RPTPs: Both molecules contain a carbonic anhydrase (CAH)-like domain and a fibronectin type III-like domain extra-cellularly, and two

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phosphatase domains intracellularly (7). Three and four splicing variants are known for Ptpz (Ptpz-A, -B and -S) and Ptpg (Ptpg-A, -B, -C and -S), respectively (8, 9). Ptpz and Ptpg distribute predominantly in the nervous system from the early embryonic stage through adulthood, suggesting their involvement in neuronal cell migration, differentiation, circuit formation and regulation of neuronal plasticity (10–15). Ptpz is known to bind several extra-cellular matrix proteins and cell-adhesion molecules such as contactin, tenascin, L1, NCAM and TAG1 (16). In addition, we found that heparin-binding growth factors, pleiotrophin/HB-GAM and midkine, bind to the extra-cellular region of Ptpz (17, 18) and inactivate its PTP activity through oligomerization (19).

For elucidation of the physiological functions of Ptpz, we attempted to identify its specific substrates by developing a novel screening system for PTP substrates, named the 'yeast substrate-trapping system'. We successfully identified several intra-cellular substrates for Ptpz, such as G protein-coupled receptor kinase-interactor 1 (Git1), p190 RhoGAP, golgi-associated PDZ and coiled-coil motif containing (GOPC/PIST) and MAGI-1 (20, 21). *Ptpz*-deficient mice exhibit abnormalities in hippocampus-dependent memory formation (22, 23), in that aberrant phosphorylation of p190 RhoGAP was observed (23). Very recently, we identified ErbB4 as a novel substrate for Ptpz by pull-down experiments using a carboxyl terminal region of Ptpz, and revealed that the tyrosine phosphorylation level of ErbB4 is increased in *Ptpz*-deficient mice (24). Both Ptpz and ErbB4 bind to post-synaptic density-95 (PSD-95) through the PDZ-binding motif at their C-terminal ends (20, 24, 25).

We previously reported that Ptpg but not Ptpz strongly inhibits NGF-induced neurite outgrowth in PC12D cells (26). Although we revealed modulation of the p13<sup>suc1</sup> complex by Ptpg, it remained to be explored whether activation of TrkA is directly affected by Ptpg or Ptpz. In the present study, we examined this possibility, and unexpectedly found that Ptpz but not Ptpg efficiently dephosphorylates the TrkA receptor. Ptpz preferentially dephosphorylated the receptor at Y674 and Y675, which are responsible for the regulation of Trk tyrosine kinase activity. Consistent with these findings, the tyrosine phosphorylation of TrkA was increased in *Ptpz*-deficient mice. In addition, we demonstrated that Ptpz indeed modulates the sensitivity of cells to NGF by neurite extension assays using PC12D cells.

#### MATERIALS AND METHODS

**Antibodies**—Anti-Ptpz-S was described previously (27). Anti-pan-Trk (C-14) and anti-TrkA (763) antibodies were purchased from SantaCruz. Anti-phosphotyrosine antibody (4G10) was from UBI. Anti-Myc and anti-HA antibodies were from Sigma. Anti-GFP antibody was from Nacalai. Alexa-conjugated secondary antibodies were from Invitrogen.

**Cell Culture**—293T cells were maintained in DMEM/F-12 medium (Sigma) supplemented with 10% fetal bovine serum (FCS). PC12D cells were maintained in DMEM supplemented with 5% fetal bovine serum and 10%

heat-treated horse serum (HS). Transfection was performed using LipofectAMINE PLUS (Invitrogen) according to the manufacturer's protocol.

**DNA Constructs**—Expression constructs encoding the full-length rat Ptpz-B (pZeo-PTP $\zeta$ ) and its PTP-inactive DA mutant (pZeo-PTP $\zeta$ D1902A) were described previously (19, 20). HA-Ptpg(WT), an expression construct encoding a fusion protein comprising the signal peptide sequence of murine Ig  $\kappa$ -chain (amino acid residues, 1–21), a HA-epitope tag (SYPYDVPDYAS) and the full-length form of mouse Ptpg (residues, 34 to the end), was prepared with the expression vector pDisplay (Invitrogen) using *Bgl*II and *Not*I sites. The full-length mouse TrkA, TrkB and TrkC cDNAs were cloned using the total RNA from the embryonic Day 18 mouse brain as a template by RT-PCR. These cDNA fragments were then cloned into the expression vector pcDNA3.1 (Invitrogen) using *Eco*RI and *Xho*I sites. Point mutants of TrkA were also generated using PCR-based site-directed mutagenesis.

**Immunoprecipitation and Immunoblotting**—Cells were lysed in a lysis buffer, which consists of 20 mM HEPES, pH 7.0, 120 mM NaCl, 1% NP-40, 50 mM NaF, 0.5 mM Na<sub>3</sub>VO<sub>4</sub> and a protease inhibitor mixture (10  $\mu$ g/ml leupeptin, 1  $\mu$ g/ml pepstatin A and 1 mM phenylmethylsulphonyl fluoride). Cell lysates were clarified by centrifugation, and subjected to immunoprecipitation with various antibodies bound to protein G-Sepharose CL-4B (Amersham Biosciences). Immunoprecipitates were solubilized with SDS sample buffer and subjected to SDS-PAGE. Proteins were transferred onto Immobilon-P membranes (Millipore), stained with specific primary antibodies and peroxidase-linked secondary antibodies (Amersham Biosciences), and visualized by chemiluminescence using ECL plus (Amersham Biosciences). For the detection, the lumino-image analyzer LAS-1000plus (Fujifilm) was used.

**In Vitro Dephosphorylation Assay**—GST-PtpzICR encoding the entire intra-cellular region of rat Ptpz-A was described previously (20, 21). For *in vitro* dephosphorylation, we first prepared the autophosphorylated TrkA proteins as the substrate. Myc-tagged ICR of TrkA was expressed in 293T cells grown on a 35-mm culture dish, and purified by immunoprecipitation with an anti-Myc antibody and protein G-Sepharose CL-4B. The TrkA-ICR was then autophosphorylated *in vitro* with 100  $\mu$ M of ATP in 100  $\mu$ l of the PTK buffer: 20 mM Tris-HCl, pH7.2, 10 mM MgCl<sub>2</sub>, 10 mM MnCl<sub>2</sub> and 1 mM DTT. Protein G beads were washed once and re-suspended in 100  $\mu$ l of 10 mM Tris-HCl, pH 7.0, containing 5 mM DTT, 5 mM EDTA and 100  $\mu$ g/ml bovine serum albumin (PTP buffer). For the dephosphorylation assay, 10 ng of GST-PtpzICR or GST alone was reacted with 10  $\mu$ l of Myc-TrkA-ICR at 30°C. At the indicated time, the reaction was stopped by adding SDS sample buffer. The samples were separated by SDS-PAGE followed by immunoblotting with anti-phosphotyrosine antibody.

**Morphological Analysis**—PC12D cells were seeded in 35-mm dishes precoated with 1 mg/ml of collagen-I (Nitta gelatin) at a density of  $5 \times 10^4$  cells/dish in DMEM containing 5% FCS and 10% HS. After 24 h, transfection was performed using LipofectAMINE PLUS (Invitrogen)

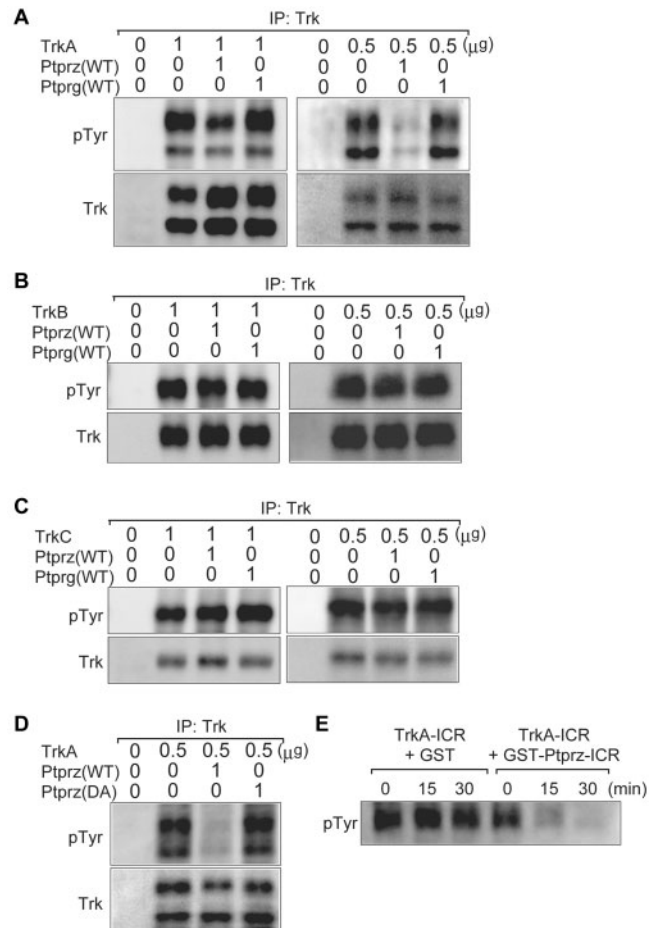
according to the manufacturer's protocol. Expression constructs used were pZeo-PTP $\zeta$ , pZeo-PTP $\zeta$ D1902A and HA-Ptpz(WT). To visualize Ptpz-transfected cells, quarter the amount of EGFP-N1 (Clontech) was cotransfected. At 24 h after transfection, the medium was changed to fresh medium containing 10 or 100 ng/ml of 7S NGF. After another 48 h, cells were fixed with 4% paraformaldehyde in phosphate-buffered saline, and then with anti-GFP antibody and Alexa488-conjugated secondary antibody. EGFP-positive neurites were analyzed with a BX51TRF microscope (Olympus) equipped with a DP-70 digital CCD camera (Olympus). Neurite lengths were measured from 200 cells selected at random in three wells. In colonies, only peripheral cells were examined. Processes of more than 150  $\mu$ m were scored as long neurites.

**Tyrosine Phosphorylation of TrkA in Mouse Brain**—The cerebral cortex of mice (5 weeks old) were homogenized with 20 vol of ice-cold 0.32 M sucrose in 4 mM HEPES, pH 7.4, containing 1 mM vanadate and 10 mM NaF. After removal of cell nucleus and debris by centrifugation at 800g for 10 min, crude synaptosomes were prepared by centrifugation at 9,000g for 15 min. Synaptosomes were then subjected to the immunoprecipitation and immunoblotting experiments as above.

## RESULTS

**Ptpz but not Ptpzg Dephosphorylates TrkA Receptors**—To test whether Ptpz or Ptpzg can dephosphorylate Trk receptors by using a cell culture system, expression constructs for full-length Trk receptors (TrkA, B and C) were co-transfected into 293T cells with the full-length wild-type construct of Ptpz or Ptpzg. Immunostaining with anti-Ptpz, anti-HA and anti-Trk antibodies revealed that about 70% of the cells expressed the exogenous genes equally (data not shown). Western blot analysis of immunoprecipitates with anti-pan-Trk antibody showed TrkA proteins (140 and 110 kDa), a TrkB protein (130 kDa) and a TrkC protein (130 kDa) (Fig. 1A–C). Highly tyrosyl-phosphorylated Trk receptor proteins were detected when each Trk receptor alone was expressed (Fig. 1A–C, lane 2). However, when the TrkA, TrkB or TrkC construct was independently co-transfected with wild-type Ptpz [Ptpz(WT)], the tyrosyl-phosphorylation of TrkA was significantly suppressed (Fig. 1A, lane 3; 1D, lane 3). In contrast, the phosphorylation of the TrkA receptor was not affected when the phosphatase-inactive DA mutant of Ptpz [Ptpz(DA)] was cotransfected (Fig. 1D, lane 4), indicating that the suppression of TrkA phosphorylation is attributable to the catalytic activity of Ptpz. On the other hand, Ptpzg exerted no effect on the phosphorylation of any Trk receptor when co-expressed (Fig. 1A–C, lanes 4).

We then tested whether Ptpz directly dephosphorylates the TrkA receptor *in vitro* using purified proteins. A Myc-tagged intra-cellular region of TrkA (TrkA-ICR) was expressed in 293T cells, and purified by immunoprecipitation with an anti-Myc antibody. The TrkA receptor was autophosphorylated *in vitro*, and then incubated with GST-Ptpz-ICR or GST. GST-Ptpz-ICR

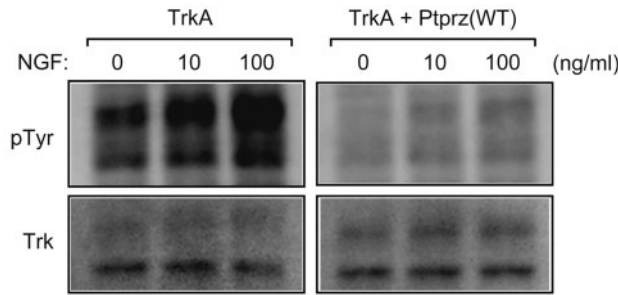


**Fig. 1. Dephosphorylation of TrkA receptors by Ptpz.** (A) Tyrosine dephosphorylation assay of TrkA. 293T cells were co-transfected with the indicated amounts ( $\mu$ g) of expression plasmids. Cell lysates were immunoprecipitated with anti-pan-Trk antibody, and the protein complexes obtained were analysed by SDS-PAGE and immunoblotting using anti-phosphotyrosine antibody (pTyr). Immunoblots were stripped and reprobed with anti-Trk antibody (Trk). The molecular mass of the full-length TrkA was estimated to be 140 and 110 kDa. These two proteins reportedly represent the fully matured form, gp140<sup>TrkA</sup>, and glycosylated precursor form, gp110<sup>TrkA</sup>, respectively (49). Lane 1 is a negative control with the empty vectors. (B) Tyrosine dephosphorylation assay of TrkB. The molecular mass of the full-length TrkB was estimated to be 130 kDa. (C) Tyrosine dephosphorylation assay of TrkC. The molecular mass of the full-length TrkC was estimated to be 130 kDa. (D) Dephosphorylation of TrkA by Ptpz(WT), but not Ptpz(DA). (E) *In vitro* dephosphorylation analysis of the TrkA receptor. Tyrosyl-phosphorylated TrkA-ICR was incubated with an equal amount of GST or GST-Ptpz-ICR. After the reaction, proteins were separated by SDS-PAGE and subjected to immunoblotting using anti-phosphotyrosine antibody (pTyr).

efficiently dephosphorylated TrkA, whereas GST alone did not (Fig. 1E), indicating that TrkA is a direct substrate for Ptpz.

Next, we examined the effects of the co-expression of Ptpz on the tyrosyl-phosphorylation of TrkA receptors when stimulated with NGF in living cells. When NGF was applied to serum-starved 293T cells, the tyrosyl-phosphorylation of TrkA was significantly enhanced in a



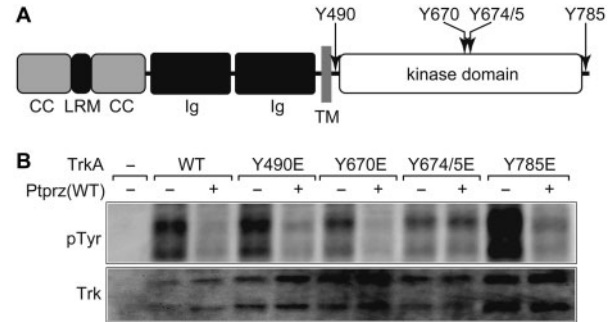


**Fig. 2. Inhibition of NGF-induced tyrosine phosphorylation of TrkA receptors by Pptrz.** 293T cells were transfected with 0.5  $\mu$ g of TrkA-expression construct and either 1  $\mu$ g of empty vector or 1  $\mu$ g of Pptrz(WT)-expression construct. Cells were serum-starved for 24 h. Tyrosine phosphorylation levels of TrkA were analysed after stimulation with NGF. Cells were lysed and immunoprecipitated with anti-Trk. Protein complexes were analysed by immunoblotting using anti-phosphotyrosine (pTyr). Immunoblots were re-probed with anti-Trk (Trk).

dose-dependent manner (Fig. 2, lanes 1–3). In contrast, when wild-type Pptrz was co-expressed, the enhancement of the phosphorylation of the TrkA receptor by NGF was significantly suppressed (Fig. 2, lanes 4–6), indicating that Pptrz can attenuate TrkA's activation by NGF.

**Pptrz Dephosphorylates TrkA at Specific Sites**—Upon binding NGF, TrkA is autophosphorylated at several distinct tyrosine residues (Y490, Y670, Y674, Y675 and Y785) (Fig. 3A) (6). We next attempted to identify the tyrosine residue on TrkA that is dephosphorylated by Pptrz. For this purpose, we generated TrkA mutants in which each candidate tyrosine was mutated to glutamate one by one: Glutamate mimics phosphorylated tyrosine but is not dephosphorylated by PTPs. 293T cells were transfected with these mutants, with or without the wild-type Pptrz construct. When wild-type Pptrz was co-expressed, the tyrosyl-phosphorylation of wild-type TrkA, TrkA(Y490E), TrkA(Y670E) and TrkA(Y785E) was markedly suppressed, but that of TrkA(Y674/675E) was not (Fig. 3B). These results indicate that Pptrz preferentially dephosphorylates Y674 and/or Y675 in the kinase domain of the TrkA receptor. This finding suggests that Pptrz functions in inhibiting the activation of TrkA receptors by NGF, because phosphorylation at these residues is implicated in the activation of TrkA receptors (6).

**Pptrz Attenuates NGF-induced Neurite Extension in PC12D Cells**—We next examined whether Pptrz expression affects the outgrowth of neurites induced by NGF in PC12D cells. PC12D cells, a subclone of PC12, extend neurites immediately in response to NGF (28, 29). PC12D cells were transiently transfected with various expression constructs; Pptrz(WT), Pptrz(DA) and Pptrz(WT). In the absence of NGF, none of the transfected cells nor the parental cells extended neurites longer than 150  $\mu$ m (data not shown). On the other hand, at 48 h after treatment with 10 ng/ml of NGF, the mock-transfected cells extended long neurites vigorously (Fig. 4A). In contrast, neurite outgrowth was considerably attenuated in Pptrz-expressing (Fig. 4B) and Pptrz-expressing (Fig. 4C) cells. Figure 4D summarizes the

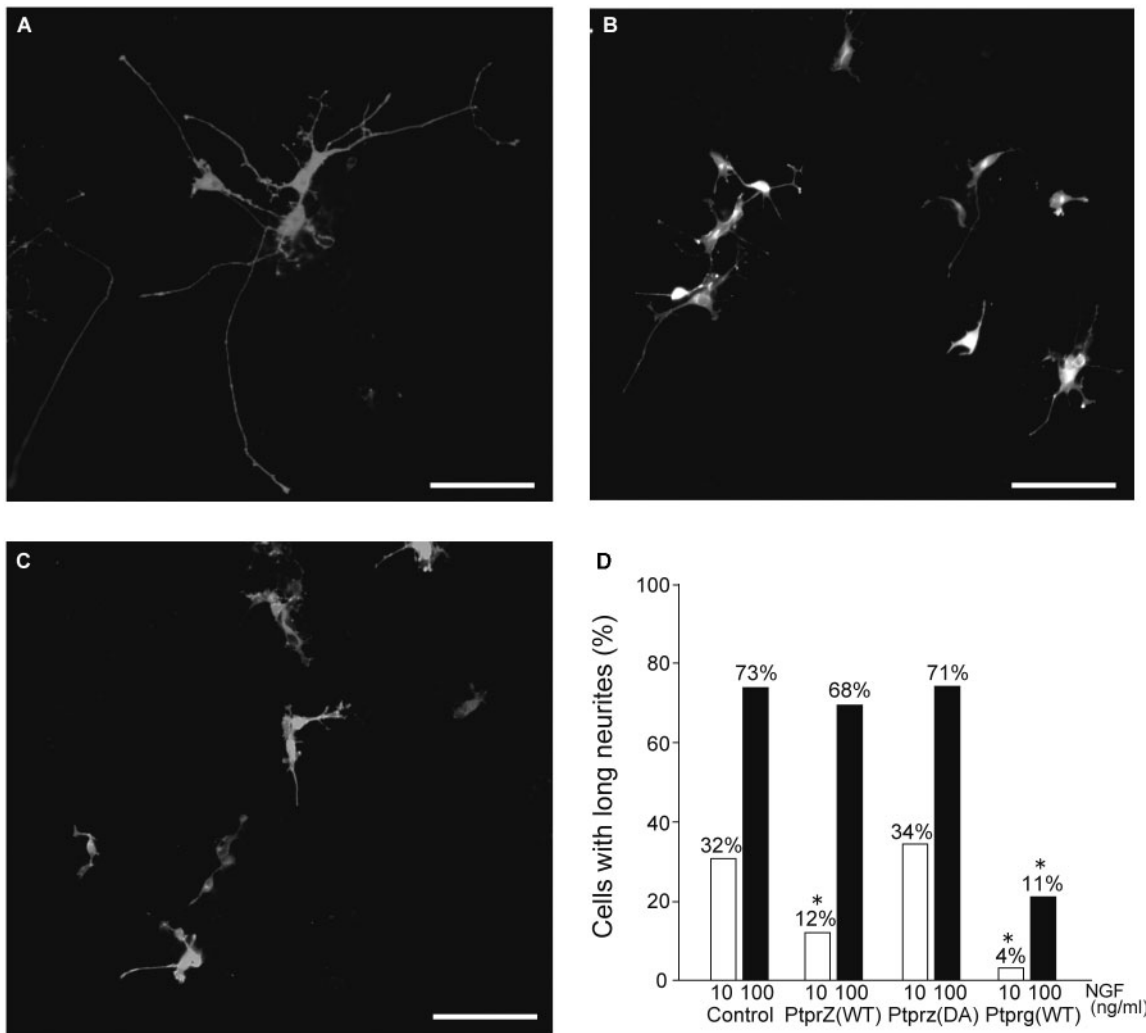


**Fig. 3. Preferential dephosphorylation of Y674/675 in the kinase domain of the TrkA receptor by Pptrz.** (A) Schematic representation of the TrkA receptor. Positions of major phosphorylated tyrosine residues are shown. CC, cysteine cluster; LRM, leucine-rich motif; Ig, immunoglobulin domain; TM, transmembrane region. (B) 293T cells were transfected with the indicated expression plasmids. Cell lysates were subjected to immunoprecipitation with anti-Trk antibody to isolate TrkA receptors. The tyrosine phosphorylation levels on TrkA receptors were analysed by SDS-PAGE and immunoblotting using anti-phosphotyrosine antibody (pTyr). Immunoblots were stripped and re-probed with anti-Trk antibody (Trk). Lane 1 is a negative control with the empty vector.

results on the extension of neurites in the transfected cells. Among mock-transfected cells, ~30% extended neurites longer than 150  $\mu$ m at 10 ng/ml of NGF. In contrast, only ~10% of Pptrz-expressing cells had long neurites ( $P < 0.01$ , by Mann–Whitney U-test). This inhibitory effect is dependent on the PTPase activity of Pptrz, because PC12D cells expressing Pptrz(DA) showed significant neurite extension like the mock-transfected cells (Fig. 4C).

As previously reported, a drastic suppression of neurite outgrowth was observed when PC12D cells over-expressed Pptrz(WT) even at a high dose of NGF (100 ng/ml) ( $P < 0.01$ , by Mann–Whitney U-test). In contrast, at 100 ng/ml of NGF, no significant difference was observed between mock-transfected cells and Pptrz(WT)-expressing cells, indicating that Pptrz under our conditions did not suppress the outgrowth of neurites in PC12D cells at a high dose of NGF. This is probably because the relatively low transfection efficiency and low level of Pptrz caused by the transient expression are not enough to completely suppress the activation of TrkA in PC12D cells as a whole: Immunostaining with an anti-Pptrz antibody revealed that about 3% of the transfected cells expressed the exogenous RPTP gene, and the expression of Pptrz in the individual PC12D cells was markedly small (~1/25) as compared with 293T cells (data not shown).

**Increased Tyrosine Phosphorylation of TrkA in Pptrz-Deficient Mice**—We examined the tyrosine phosphorylation of TrkA in the synaptosomal fraction of the cerebral cortex of wild-type and Pptrz-deficient mice (Fig. 5A). Pptrz-B (250 kDa), a major isoform of Pptrz, was detected in cerebral synaptosomes of the wild-type mice (Fig. 5A, left; see also ref. 24). Wild-type and Pptrz-deficient mice showed a similar expression level of TrkA in the synaptosomal fractions of the cerebral cortex (Fig. 5A, right). However, immunoblotting with anti-phosphotyrosine antibody revealed that the tyrosine



**Fig. 4. Attenuation of NGF-induced neurite extension in PC12D cells by Ptpz.** (A) Morphology of mock-transfected cells after treatment with NGF (10 ng/ml). (B) Morphology of Ptpz-expressing cells after treatment with NGF (10 ng/ml). (C) Morphology of Ptprg-expressing cells after treatment with

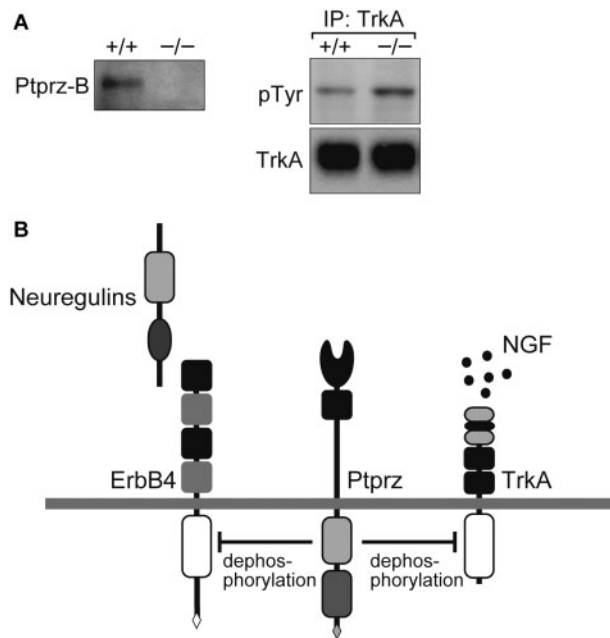
NGF (10 ng/ml). Scale bars in (A)–(C): 100  $\mu$ m. (D) Summary of neurite outgrowth in PC12D cells. Percentages of cells with neurites longer than 150  $\mu$ m are shown. \* $P < 0.01$ , when compared with mock-transfected cells (Mann–Whitney U-test).

phosphorylation of TrkA was significantly increased in *Ptpz*-deficient mice (Fig. 5A, right; 1.8-fold increase,  $P < 0.05$ ,  $n = 4$  by unpaired Student's *t*-test). Taken altogether, we concluded that the TrkA receptor is a physiological substrate for Ptpz.

#### DISCUSSION

It has been postulated that the activation of RTKs is negatively regulated by PTPs (2). In this study, we found that Ptpz specifically recognizes the TrkA receptor as a substrate. Our data indicated that Ptpz recognizes particular phosphorylation sites within the activation loop of the kinase domain, and modulates activation of the TrkA receptor. In line with this finding, the expression of Ptpz in PC12D cells attenuated the NGF-induced extension of neurites and the tyrosine phosphorylation of TrkA was significantly increased in *Ptpz*-deficient mice.

TrkA rapidly dimerizes upon binding NGF and becomes phosphorylated at several distinct tyrosine residues (Y490, Y670, Y674, Y675 and Y785) (6). This allows the recruitment and activation of a number of specific signalling proteins, such as Shc, FRS2 and phospholipase C- $\gamma$  (PLC- $\gamma$ ) (6). Tyrosine residues Y490 and Y785 are responsible for the direct association with and tyrosine phosphorylation of Shc and PLC- $\gamma$ , respectively (30–33). Activation of the MAP kinase pathway largely depends on the activation of Shc (6). Shc has been shown to be essential for mediating the activation of PI3-kinase (34), which is critical for the ability of Trk receptors to mediate survival through the activation of Akt (6). On the other hand, activation of PLC- $\gamma$  leads to the release of calcium ion from intra-cellular stores through production of inositol(1,4,5)triphosphate. Phosphorylation at residues Y670, Y674 and Y675 in the activation loop of the kinase domain is implicated in the activation of the tyrosine kinase of TrkA (35).



**Fig. 5. Comparison of the tyrosine phosphorylation of TrkA between wild-type and *Ptpzr*-deficient mice.** (A) The tyrosine phosphorylation level of TrkA in the synaptosomal fractions of wild-type and *Ptpzr*-deficient mice. Expression of Ptpzr was analysed by immunoblotting after treatment with chondroitinase ABC as described (8). TrkA was immunoprecipitated with anti-TrkA antibody, and the tyrosine phosphorylation level of TrkA was analysed by SDS-PAGE and immunoblotting using anti-phosphotyrosine (pTyr) antibody. Immunoblots were stripped and reprobed with anti-Trk antibody (TrkA). (B) Schematic drawing of regulation of TrkA and ErbB4 by Ptpzr. ErbB4 and TrkA are activated by their specific ligands, neuregulins and NGF, respectively. Ptpzr attenuates the activation of ErbB4 (24) and TrkA through dephosphorylation.

In this study, we found that Ptpzr preferentially dephosphorylates Y674 and/or Y675 of TrkA (Fig. 3). This suggests that Ptpzr functions in inhibiting the activation of TrkA receptors after they bind NGF. Several PTPs have been reported to dephosphorylate TrkA (36–38). Among them, SHP-1, a SH2 domain-containing cytosolic PTP, forms a complex with TrkA at Y490, and dephosphorylates it at Y674 and/or Y675 like Ptpzr (38). Therefore, Ptpzr may regulate the activation of TrkA in coordination with SHP-1 *in vivo*. Here, it is not clear why the substrate-trapping mutant (DA mutant) of Ptpzr showed no effect on NGF-induced neurite outgrowth in PC12D cells.

RPTPs have been proposed to be inactivated by ligand-induced dimerization. Since Ptpzr is inactivated through oligomerization including dimerization by pleiotrophin/HB-GAM (21), it is possible that the activation of TrkA is regulated by pleiotrophin/HB-GAM or Midkine. Thus, the identification of RPTPs as TrkA phosphatases would shed light on the novel mechanisms by which extracellular ligands for RPTPs influence neurotrophic function. From this point of view, the discovery of RPTPs that specifically dephosphorylate Y490 or Y785 is of importance.

Another subfamily member, Ptprg, does not dephosphorylate Trk receptors (Fig. 1), however, Ptprg strongly

inhibits NGF-induced neurite outgrowth in PC12D cells (Fig. 4C, see also ref. 26). This suppression is probably achieved through modulation of the p13<sup>suc1</sup> complex (26), which comprises FRS2 (SNT) and associated proteins (39). FRS2 is an important signalling molecule that binds to the NPXY motif on Trks, and considered to play an important role in neuronal differentiation and neurite outgrowth (39). Neurotrophin induces phosphorylation of FRS2, and the phosphorylated FRS2 then provides binding sites for adaptor proteins including Grb2, Crk and Src (6). We postulated that some of the constituents in this complex are substrates for Ptpzr (26). Thus, the substrate specificity would be the cause of the functional difference between Ptpzr and Ptprg.

In the mature brain, the expression of Ptpzr persists in specific regions including the hippocampus and cerebral cortex (8, 15), where the structure and function of synapses are continually modified throughout life. We recently reported that Ptpzr forms a complex with ErbB4 through PSD-95 and dephosphorylates it as a substrate (24). ErbB4 is a member of the ErbB family of RPTKs activated by neuregulins and other growth factors, and probably modulates synaptic plasticity (40–42). NGF is also expressed in the adult CNS, with the highest levels being present in the hippocampus and cerebral cortex (43–45). The TrkA receptor is also broadly distributed in the brain including the hippocampus (46). The application of anti-NGF antibody inhibits differentiation into the cholinergic phenotype (47) and blocks the sprouting of acetylcholinesterase-positive branches in the hippocampus (48). In addition, neurotrophic factors and their receptors are implicated in neuronal plasticity (5). Consistent with these findings, mutant mice deficient in *Ptpzr* exhibited enhanced long-term potentiation (LTP) in the CA1 region in hippocampal slices (22), and impaired hippocampus-dependent memory formation (22, 23). Taken altogether, these observations suggest an important role for Ptpzr in the development and regulation of synapses through the control of activities of TrkA and ErbB4 (Fig. 5B).

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