

# The Secretory Granule-Associated Protein CAPS2 Regulates Neurotrophin Release and Cell Survival

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Neurotrophins are key modulators of various neuronal functions, including differentiation, survival, and synaptic plasticity, but the molecules that regulate their secretion are poorly understood. We isolated a clone that is predominantly expressed in granule cells of postnatally developing mouse cerebellum, which turned out to be a paralog of CAPS ( $\text{Ca}^{2+}$ -dependent activator protein for secretion), and named CAPS2. CAPS2 is enriched on vesicular structures of presynaptic parallel fiber terminals of granule cells connecting postsynaptic spines of Purkinje cell dendrites. Vesicle fractions affinity-purified by the CAPS2 antibody from mouse cerebella contained significant amounts of neurotrophin-3 (NT-3), brain-derived neurotrophic factor (BDNF), and chromogranin B but not marker proteins for synaptic vesicle synaptophysin and synaptotagmin. In cerebellar primary cultures, punctate CAPS2 immunoreactivities are primarily colocalized with those of NT-3 and BDNF and near those of a postsynaptic marker, postsynaptic density-95, around dendritic arborization of Purkinje cells. Exogenously expressed CAPS2 enhanced release of exogenous NT-3 and BDNF from PC12 cells and endogenous NT-3 from cultured granule cells in a depolarization-dependent manner. Moreover, the overexpression of CAPS2 in granule cells promotes the survival of Purkinje cells in cerebellar cultures. Thus, we suggest that CAPS2 mediates the depolarization-dependent release of NT-3 and BDNF from granule cells, leading to regulation in cell differentiation and survival during cerebellar development.

**Key words:** neurotrophin; BDNF; NT-3; granule cell; Purkinje cell; parallel fiber; cerebellum; CAPS; exocytosis; secretory granule

## Introduction

Neurotrophins play indispensable roles in regulating the differentiation and survival of neurons (Murphy et al., 1998; Miller and Kaplan, 2001; Ginty and Segal, 2002; Horch and Katz, 2002; Huang and Reichardt, 2002). Recent evidence indicates their modulatory roles in synaptic plasticity such as long-term potentiation relevant to learning and memory (Lu and Figurov, 1997; Kovalchuk et al., 2002; Egan et al., 2003) and in neuronal diseases, including Alzheimer's disease and Parkinson's disease (Murer et al., 2001). It was reported that increasing intracellular  $\text{Ca}^{2+}$  regulates activity dependency on the neurotrophin release from neuronal processes (Canossa et al., 2001; Balkowiec and Katz, 2002; Wang et al., 2002). However, the underlying molecular mechanism is poorly understood.

Many reports describe how neurotrophins play important roles in the postnatal development of the cerebellum (Lindholm et al., 1993, 1997; Segal et al., 1995; Schwartz et al., 1997; Doughty

et al., 1998; Bates et al., 1999). Neurotrophin-3 (NT-3) and brain-derived neurotrophic factor (BDNF) show a reversed expression pattern in cerebellar granule cells; between postnatal day 10 (P10) and P20 of rat cerebellum, the former is downregulated, whereas the latter is upregulated (Rocamora et al., 1993; Katoh-Semba et al., 1997, 2000; Das et al., 2001). Both the receptors TrkB (for BDNF) (Gao et al., 1995) and TrkC (for NT-3) (Velier et al., 1997) are expressed in postsynaptic Purkinje cells as well as presynaptic granule cells, and BDNF and NT-3 released from granule cells are thought to function in autocrine–paracrine and anterograde manners (Lindholm et al., 1993; Mount et al., 1994; Schwartz et al., 1997; Bates et al., 1999). Analyses of mice genetically lacking the gene for NT-3 (Bates et al., 1999) and BDNF (Schwartz et al., 1997) show that these neurotrophins play pivotal roles in the survival and differentiation of these neurons. Few studies, however, have addressed how they are released from neurons.

In this study, we isolated a molecule designated a1803 by differential screening of gene expression responsible for the postnatal development of mouse cerebellum. a1803 is a paralog of CAPS ( $\text{Ca}^{2+}$ -dependent activator protein for secretion), which has been characterized as controlling  $\text{Ca}^{2+}$ -dependent secretion from endocrine and neuroendocrine cells (Walent et al., 1992; Ann et al., 1997; Berwin et al., 1998; Tandon et al., 1998; Renden et al., 2001), and named CAPS2. There has been no direct evi-

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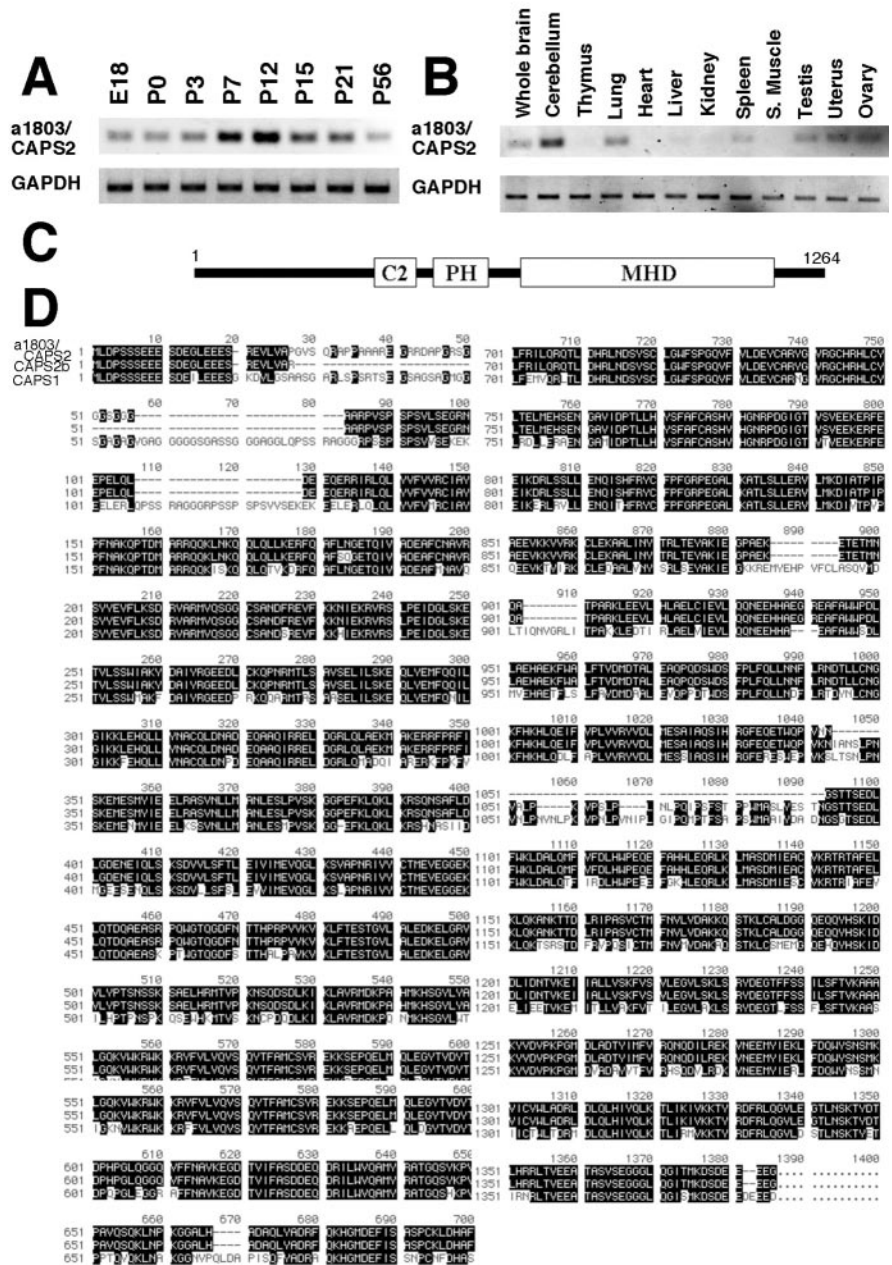
dence reported on involvement of CAPS1 in the neurotrophin release activity from neurons and neuronal cell function. In the present study, we indicate that CAPS2 is enriched on vesicular structures in the parallel fiber (PF) terminals of granule cells, and the CAPS2-associated vesicles are distinctly different from synaptic vesicles (SVs) and contain NT-3 and BDNF as well as a secretory granule (SG) marker, chromogranin B (CGB). Furthermore, we demonstrate that CAPS2 has neuronal activity that enhances neurotrophin release and promotes cell survival in cerebellar cultures.

**Materials and Methods**

**Differential display and molecular cloning.** Fluorescent differential display (FDD) and full-length cDNA cloning were performed essentially as described by Shiraishi et al. (1999). The FDD was performed using total RNAs obtained from mouse (ICR, Nihon SLC, Hamamatsu, Japan) cerebella at eight different developmental ages: embryonic day 18 (E18), P0, P3, P7, P12, P15, P21, and P56. The cDNA sequence of a1803/CAPS2a (GenBank accession number AB098623) was determined using a DNA sequencer (Prism 3700; Applied Biosystems, Foster City, CA).

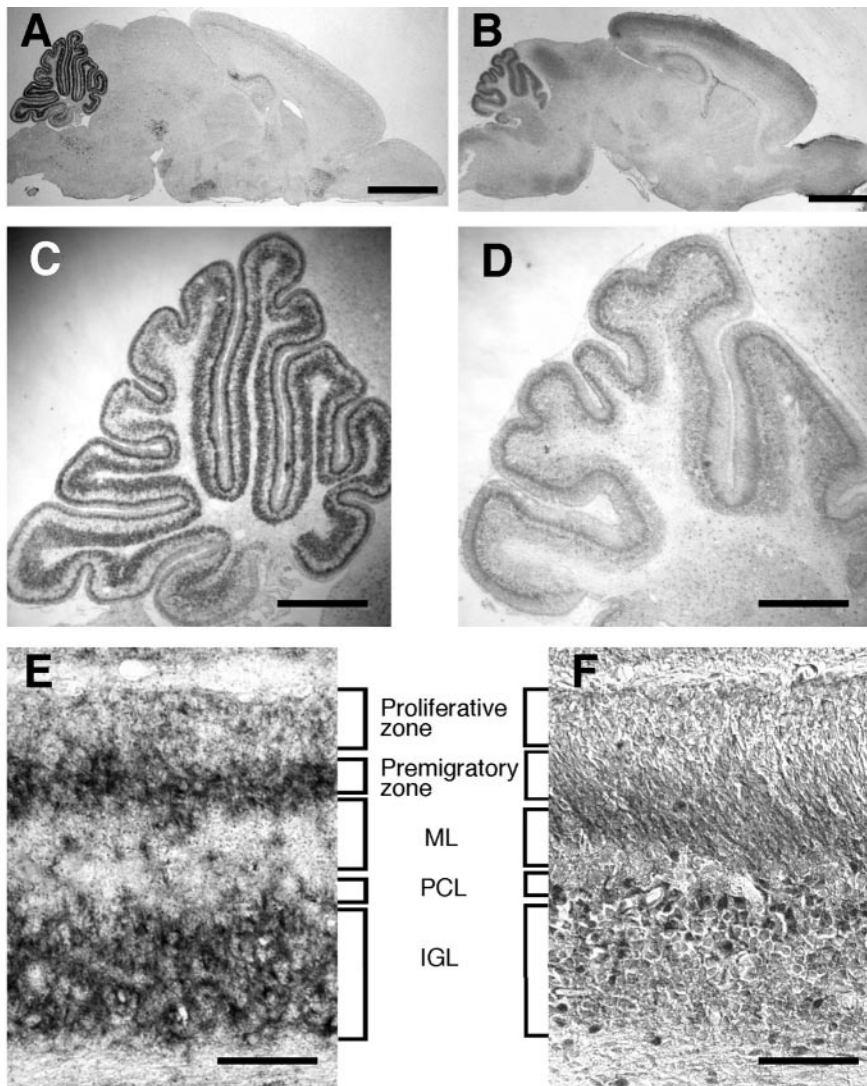
**In situ hybridization.** The cDNA sequence corresponding to amino acids 18–89 of a1803/CAPS2 was used as the template to prepare the antisense riboprobe. Cryosections of mouse brain (14 μm thick) were fixed in 4% paraformaldehyde for 5 min, washed twice in PBS, and treated with freshly prepared 10 μg/ml proteinase K (Invitrogen, Carlsbad, CA) at room temperature. After acetylation, sections were incubated in hybridization buffer containing 0.2 μg/ml digoxigenin-labeled riboprobes at 43°C overnight in a humid chamber. Hybridized sections were washed by successively immersing in 4× SSC (in mM: 150 NaCl and 15 sodium citrate, pH 7.0, room temperature), 2× SSC containing 50% formamide (50°C, 30 min), 2× SSC (37°C, 10 min), 2× SSC containing 20 μg/ml RNase A (37°C, 30 min), 2× SSC (37°C, 20 min), and 0.1× SSC (room temperature, 30 and 5 min). The hybridization signals were detected with the digoxigenin detection (Roche Diagnostics, Indianapolis, IN).

**Antibodies.** Rabbit and guinea pig polyclonal anti-a1803/CAPS2 and anti-CAPS1 antibodies were raised against the glutathione S-transferase-tagged a1803/CAPS2 (amino acids 18–89 and 235–336) and CAPS1 (amino acids 18–107) that were bacterially expressed and were affinity-purified against the maltose binding protein-tagged antigenic proteins that were covalently coupled to cyanogen bromide-activated Sepharose 4B. Rabbit anti-a1803/CAPS2 (amino acids 18–89) and anti-CAPS1 (amino acids 18–107) antibodies were used for Western blotting, immunocytochemistry, and immunohistochemistry. Guinea pig anti-a1803/CAPS2 (amino acids 18–89) antibody was used for immunocytochemistry. The rabbit anti-a1803/CAPS2 (amino acids 235–336) polyclonal antibody was used for immunoprecipitation. The mouse anti-CAPS1 monoclonal antibody (BD Biosciences, Franklin Lakes, NJ) was also used for immunoaf-



**Figure 1.** Expression and structure of a1803/CAPS2 (CAPS2a). *A*, The expression profile of a1803/CAPS2 in developing mouse cerebella at E18, P0, P3, P7, P12, P15, P21, and P56 was analyzed by RT-PCR. Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) was used as a control. *B*, The tissue distribution of a1803/CAPS2 expression in P56 mice was analyzed by RT-PCR. *C*, Schematic depiction of the predicted a1803/CAPS2 protein (amino acids 1–1264). The C2- and PH-like domain and the MHD are represented: amino acids 369–452, 494–596, and 677–1166, respectively. *D*, Alignment of mouse a1803/CAPS2 (CAPS2a; top), CAPS2b (middle), and CAPS1 (bottom) protein. Amino acids identical among the three sequences are shown against a black background.

finity purification. The cross-reactivity of both antibodies was examined by exogenous cDNA expression experiments in PC12 cells; the anti-a1803/CAPS2 (amino acids 235–336) antibody weakly cross-reacted with the endogenous CAPS1 protein of PC12 cells, whereas the anti-CAPS1 monoclonal antibody reacted with the endogenous CAPS1 of PC12 cells as indicated by the manufacturer but did not cross-react with the exogenously expressed CAPS2 protein. Anti-mouse polyclonal anti-vesicle-associated membrane protein (VAMP) antibody (StressGen Biotechnologies, San Diego, CA), mouse monoclonal anti-synaptophysin (Sigma, St. Louis, MO) and anti-chromogranin B (BD Biosciences)



**Figure 2.** Expression of CAPS2 mRNA and protein in P7 mouse brains. *A, C, E*, *In situ* hybridization analysis of CAPS2 mRNA distribution in parasagittal brain sections: whole brain (*A*), cerebellum (*C*), and cerebellar cortex (*E*) views. *B, D, F*, Immunohistochemical analysis of CAPS2 protein distribution in parasagittal brain sections: whole brain (*B*), cerebellum (*D*), and cerebellar cortex (*F*) views. PCL, Purkinje cell layer. Scale bars: *A, B*, 2 mm; *C, D*, 500  $\mu$ m; *E, F*, 50  $\mu$ m.

antibodies, and rabbit anti-synaptotagmin I and II (Fukuda et al., 2002a) and anti-BDNF (Santa Cruz Biotechnologies, Santa Cruz, CA) antibodies were used for Western blotting. Mouse monoclonal anti-calbindin (Sigma), anti-MAP-2 (Roche Diagnostics), anti-postsynaptic density-95 (PSD-95; Sigma), and anti- $\alpha$ -internexin (Chemicon, Temecula, CA) antibodies, rabbit anti-BDNF antibody (Kato-Semba et al., 1997), and sheep anti-NT-3 antibody (Chemicon) were used for immunocytochemistry.

**Immunoelectron microscopy.** For immunoelectron microscopy of paraformaldehyde-fixed sections, mice (postnatal day 15 in Fig. 5*A, B* and postnatal week 6 in Fig. 5*C*) were deeply anesthetized by ether and perfused with freshly prepared 2% paraformaldehyde in PBS for 15 min. Brains were removed by dissection and cut with a razor into several coronal sections (~2 mm thickness), which were soaked with the same fixative at 4°C for 2 hr. For cryoprotection, sections were placed into 20% sucrose solution for 2 hr and 25% sucrose solution overnight. The sections were frozen using liquid nitrogen. Serial 10- $\mu$ m-thick sections were cut in a cryostat. The samples were incubated with the rabbit polyclonal anti-CAPS2 antibody, followed by incubation with the anti-rabbit IgG antibody coupled with 1.4 nm gold particles (Nanoprobes Inc., Yaphank, NY). The sample-bound gold particles were silver-enhanced using an HQ-silver kit (Nanoprobes) at 18°C for 12 min. The samples were again washed and postfixed with 0.5% osmium oxide in a buffer containing

100 mM cacodylate buffer, pH 7.3. They were dehydrated by passage through a graded series of ethanol (50, 70, 90, and 100%) and propylene oxide and embedded in epoxy resin. From this sample, ultrathin sections were cut, stained with uranyl acetate and lead citrate, and then observed with an electron microscope (JEM-1200EX; JEOL, Tokyo, Japan).

**Subcellular fractionation.** Homogenates of P21 mice cerebella were centrifuged at 100,000  $\times$  *g* (Shiraishi et al., 1999), and the pellet fraction was further fractionated by centrifugation in continuous sucrose gradient from 0.3 to 1.8 M. After centrifugation, the subfractions were taken from the top (fraction 1) of the gradient to the bottom (fraction 16).

**Immunoaffinity purification of cerebellar vesicle fractions.** For immunoaffinity purification, 250  $\mu$ g of superparamagnetic polystyrene beads coated covalently with sheep anti-rabbit or anti-mouse IgG (Dynabeads M-280; Dynal, Lake Success, NY) were incubated overnight at 4°C in phosphate buffer (PBS, pH 7.4, and 0.1% BSA) containing the primary antibody (4  $\mu$ g), rabbit anti-a1803/CAPS2 (amino acids 235–556) antibody, mouse anti-synaptophysin antibody, mouse anti-CAPS1 monoclonal antibody, and control rabbit IgG and mouse IgG (Jackson ImmunoResearch, West Grove, PA). P15 mouse cerebella were dissected and homogenized in low-sucrose homogenization buffer (in mM: 5 HEPES, pH 7.4, 5 EDTA, and 30 sucrose and protease inhibitor mixture). Homogenates were centrifuged at 800  $\times$  *g* for 10 min, and the crude membrane in the supernatant was incubated overnight with 500  $\mu$ g of the magnetic beads for preabsorption. The resultant supernatant-containing vesicle fractions were incubated with the primary antibody-bound beads in incubation buffer (PBS, pH 7.4, 5% fetal bovine serum, and 2 mM EDTA) for 1 hr at 4°C. Vesicle-bound beads were collected and washed four times with incubation buffer and twice with PBS containing 2 mM EDTA for 15 min each. The collected vesicles were analyzed by ELISA or Western blotting.

**CAPS2, NT-3, and BDNF expression plasmids.** The mouse NT-3 and BDNF cDNA were subcloned into the pEF4/Myc-His plasmid vector containing the EF-1 $\alpha$  promoter (Invitrogen). The a1803/CAPS2 cDNA was subcloned into the pcDNA3 plasmid vector containing the cytomegalovirus (CMV) promoter (Invitrogen) to create the pcDNA3-CAPS2 [wild type (wt)]. The pcDNA3-CAPS2 ( $\Delta$ C2 + PH) had an internal deletion of amino acids 347–593, which corresponds to the C2 and pleckstrin homology (PH) domain. The cDNA encoding the C2 and PH domain (amino acids 344–598) was subcloned into the pCMV-hemagglutinin (HA) plasmid vector (Clontech, Palo Alto, CA) to create pCMV-HA-CAPS2 (C2 + PH), which expresses the HA-tagged C2 and PH domain of a1803/CAPS2.

**Preparation and infection of recombinant adenoviruses.** A replication-deficient adenovirus (Ad-CAPS2) was generated by the cosmid-adenovirus terminal protein complex method (Miyake et al., 1996). Briefly, the full-length a1803/CAPS2 cDNA was inserted into the CAG promoter expression unit of pAxCawt cosmid cassette (Takara Bio, Otsu, Shiga, Japan). Recombinant viruses were generated by homologous recombination between *Eco*T22I-digested Ad5-dlx DNA-terminal protein complex and recombinant cosmid vectors in human embryonic kidney 293 (HEK293) cells. The generated recombinant adenoviruses were propagated in HEK293 cells and then concentrated and purified by

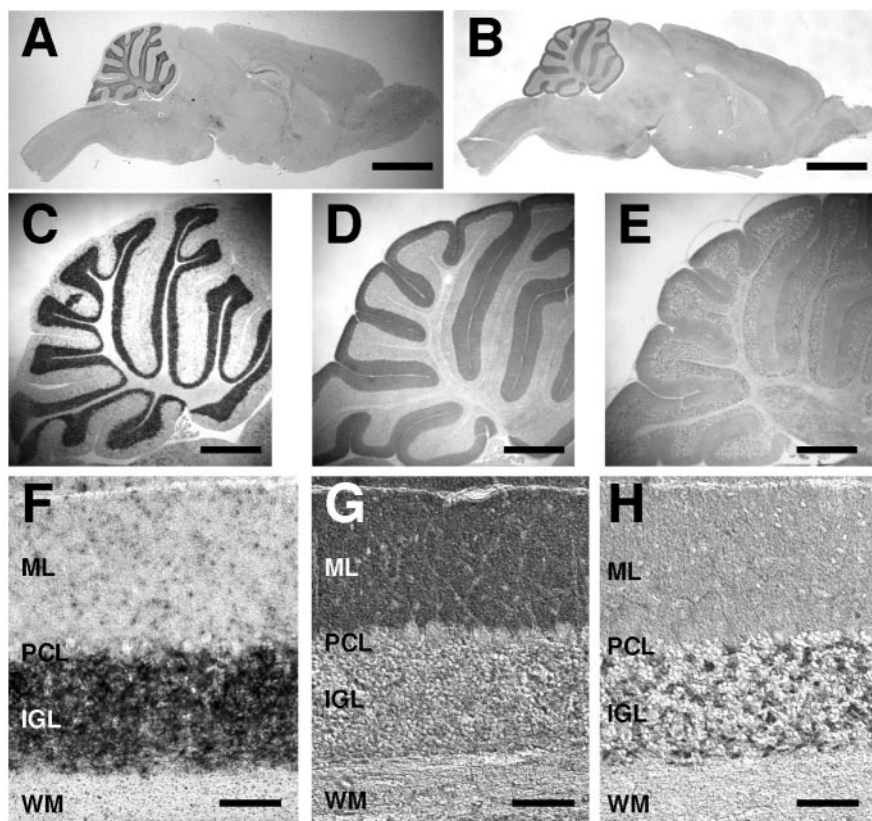
double-CsCl step gradient centrifugation. The virus titers were measured on HEK293 cells. The resulting Ad-CAPS2 was used for infection of primary cerebellar cultures (Shiraishi et al., 1999) at a multiplicity of infection of 30 for 1 hr at 37°C.

**Secretion assay.** Twenty-four hours after transfection with the expression plasmids as described above using Lipofectamine 2000 reagent (Invitrogen, Rockville, MD), PC12 cells were incubated in a fresh assay medium (DMEM containing 0.2% BSA) for 30 min and then incubated in a fresh assay medium with or without (in mM): 50 KCl, 10 EGTA, or both for a further 5 min. The control and high-KCl stimulation assay media were collected and the neurotrophin contents secreted were measured with an NT-3/BDNF Emax immunoassay system (Promega, Madison, WI) according to the manufacturer's instructions. When using modified HBSS (125 mM NaCl, 5 mM KCl, 1.2 mM  $\text{NaH}_2\text{PO}_4$ , 1.2 mM  $\text{MgCl}_2$ , 1 mM  $\text{CaCl}_2$ , 1  $\mu\text{M}$   $\text{ZnCl}_2$ , 10 mM glucose, 25 mM HEPES, and 0.2% BSA, pH 7.4) instead of DMEM, similar results were obtained by stimulating with the modified HBSS containing (in mM): 50 KCl and 80 NaCl. A secretion assay for cerebellar cultures was performed with medium incubated for 15 min.

## Results

### Differential developmental and cellular expression of CAPS2 in mouse cerebellum

To examine the genetic basis of mouse cerebellar development, we analyzed the developmental stage-specific gene expression using a differential display technique (Shiraishi et al., 1999) and isolated a developmentally regulated clone, a1803, with the expression peak at ~P12 (Fig. 1A), when various cellular developmental events are actively occurring, e.g., immense cell proliferation and migration and vigorous genesis of axons, dendrites, and synapses. Tissue distribution analysis of adult (P56) mice by reverse transcription (RT)-PCR showed that a1803 expression predominates in the cerebellum within the CNS and is detected at low levels in non-neuronal tissues, including lung, spleen, testis, uterus, and ovary (Fig. 1B). A predicted a1803 protein has structural features similar to the C2 and PH domains and the Munc13-1-homologous domain (MHD) containing a syntaxin-interacting domain (Fig. 1C) and shares 70.4% amino acid identity with CAPS, which is essential for  $\text{Ca}^{2+}$ -triggered SG exocytosis in permeabilized PC12 cells (Walent et al., 1992; Berwin et al., 1998; Tandon et al., 1998), as shown in Figure 1D. An open reading frame of mouse CAPS2 cDNA that has recently been registered on the public DNA database differs from that of a1803; there are two different stretches in either sequence, which probably results from alternative splicing events (Fig. 1D). Hereafter, if necessary, the a1803 splicing form is depicted as a1803/CAPS2 or CAPS2a, whereas the other CAPS2 variant is called CAPS2b. The original CAPS is called CAPS1. The RT-PCR analysis using the primers specific to the 5'-splicing segment showed that the mRNA for the a1803/CAPS2a splicing form containing the 5'-splicing segment was highly expressed in the postnatal and adult stages of the cerebellum, whereas that for CAPS2b lacking the 5'-splicing segment was undetectable (data not shown), indicating that



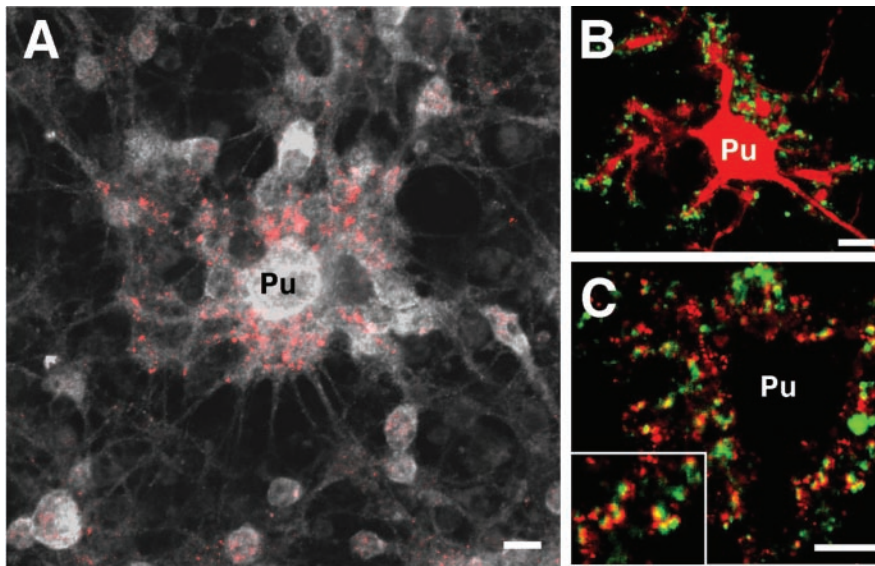
**Figure 3.** Expression of CAPS2 mRNA and protein in P21 mouse brains. *A, C, F*, *In situ* hybridization of the CAPS2 mRNA distribution in parasagittal brain sections: whole brain (*A*), cerebellum (*C*), and cerebellar (*F*) cortex views. *B, D, G*, Immunohistochemical analysis of CAPS2 protein distribution in parasagittal brain sections: whole brain (*B*), cerebellum (*D*), and cerebellar cortex (*G*) views. *E, H*, Immunohistochemical analysis of CAPS1 protein distribution in parasagittal brain sections: cerebellum (*E*) and cerebellar cortex (*H*) views. PCL, Purkinje cell layer; WM, white matter. Scale bars: *A, B*, 2 mm; *C–E*, 500  $\mu\text{m}$ ; *F–H*, 50  $\mu\text{m}$ .

the a1803/CAPS2a is the predominant form in the mouse cerebellum.

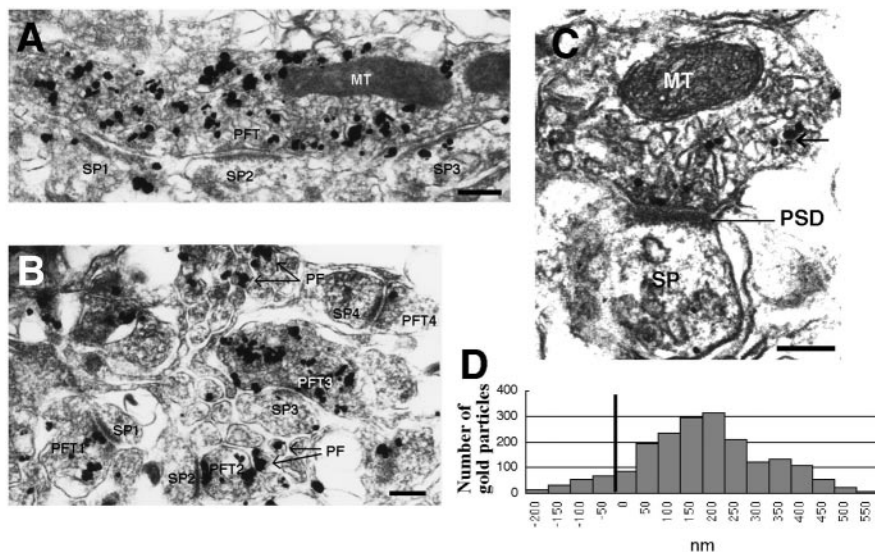
### CAPS2 is located in presynaptic parallel fiber terminals of granule cells

*In situ* hybridization analyses showed that in P7 mouse brains, the CAPS2 mRNA was predominantly expressed in the cerebellum (Fig. 2A) in which the postmitotic premigratory zone of the external granule layer (EGL) and the internal granule layer (IGL) were the predominant sites (Fig. 2C,E), indicating that CAPS2 is specific to differentiating and differentiated granule cells. In P21 brains, mRNA expression was detected at the highest level in cerebellar granule cells (Fig. 3A,C,F) and at lower levels in the hippocampal region and interpeduncular nucleus (Fig. 3A). Immunohistochemical analyses showed the predominant distribution of CAPS2 protein in the cerebellum at P7 (Fig. 2B,D,F) and P21 (Fig. 3B,D,G). In P21 cerebella, it was a marked contrast that the mRNA and protein were respectively localized to the IGL (Fig. 3F) and the molecular layer (ML) (Fig. 3G), indicating that the protein subcellularly distributes in granule cell axons (PFs). On the contrary, CAPS1 distributed over the whole brain area (data not shown). In the cerebellum, the CAPS1 mRNA was expressed in the IGL (data not shown), and the protein was located in glomerular rosettes of the IGL at a high level and Purkinje cells at a low level (Fig. 3E,H). These results indicate that CAPS2 and CAPS1 occupy distinct cellular distributions in the cerebellum.

We analyzed the subcellular localization of CAPS2 protein using cerebellar primary cultures at 14 d *in vitro* (DIV) (Fig. 4).



**Figure 4.** Subcellular localization of CAPS2 protein in cerebellar granule cells. *A*, Fluorescent confocal images of cerebellar cultures (14 DIV) immunostained with the anti-CAPS2 (red) and anti- $\alpha$ -internexin (white) antibodies. *B*, Fluorescent confocal images of cerebellar cultures (14 DIV) immunostained with the anti-CAPS2 (green) and anti-calbindin (red) antibodies. *C*, Fluorescent confocal images of cerebellar cultures (14 DIV) immunostained with the anti-CAPS2 (green) and anti-PSD-95 (red) antibodies. Inset, Higher magnification. Pu, Purkinje cell. Scale bars, 10  $\mu$ m.



**Figure 5.** Immunoelectron microscopic study of the CAPS2 protein in the mouse cerebellum. *A*, *B*, Electron micrographs of the molecular layer in the P15 mouse cerebellar coronal section (*A*) and sagittal section (*B*) reacted with the immunogold of anti-CAPS2 antibody. CAPS2 immunogold particles were predominantly detected in vesicular structures and were also present near the presynaptic and perisynaptic membranes and in spines. MT, Mitochondrion. *C*, Electron micrographs of the molecular layer of a 6-week-old mouse cerebellum reacted with the immunogold of anti-CAPS2 antibody. Extrasynaptic localization of immunogold particles apart from the active zone is indicated by an arrow. *D*, Quantification of CAPS2 distribution on the electron micrographs. The distance between CAPS2 immunogold particles and the active zone (position 0) was examined by calculating of 160 synapses in P15 cerebellar sagittal sections. The distance between immunogold particles and the active zone is indicated on the x-axis; numbers of immunogold particles are represented on the y-axis. +, Presynaptic side; -, postsynaptic side. Distribution of CAPS2 immunogold particles peaked at a presynaptic site 250 nm distant from the active zone. Scale bars, 200 nm.

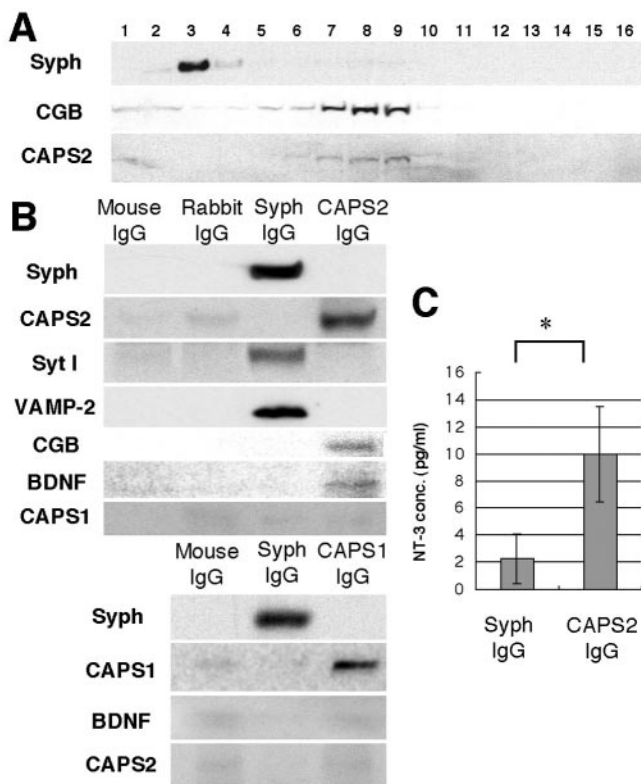
Punctate immunoreactivity of CAPS2 was primarily concentrated near Purkinje cells innervated by many granule cell axons immunostained for  $\alpha$ -internexin (Chien et al., 1996) (Fig. 4*A*). Coimmunostaining for calbindin, a marker for Purkinje cells, showed that the CAPS2 puncta clustered around the dendrites of Purkinje cells (Fig. 4*B*). In addition, coimmunostaining for PSD-95, a marker for postsynapses, showed that the puncta of CAPS2

were located near those of PSD-95, which immunostained the postsynaptic spines of Purkinje cells (Fig. 4*C*), suggesting the pre-synaptic localization of CAPS2 in PFs. To confirm this, we performed an immunogold electron microscopic observation (Fig. 5). In coronal sections of the P15 mouse cerebellum (longitudinal views of PFs) (Fig. 5*A*), immunogold particles for CAPS2 were predominantly distributed on vesicular structures in parallel fiber terminals (PFTs) connecting multiple spines (SP1–3, called en passant synapses), which are characteristic to the cerebellar PFTs, and were also present near the presynaptic (active zone) and perisynaptic membrane and spines. In sagittal sections of the P15 mouse cerebellum (transverse views of PFs; (Fig. 5*B*), CAPS2 immunogold particles were predominantly localized on vesicular structures and also detectable near the presynaptic (PFT1 and 3) and perisynaptic (PFT2 and 3) membrane and on postsynapses (SP3). In addition, CAPS2 immunogold particles were observed in transverse PFs. In 6-week-old mouse cerebella (Fig. 5*C*), CAPS2 immunogold particles were concentrated at the PFTs connecting dendritic spines of Purkinje cells and were detected on vesicular structures and in regions around the active zone and extrasynaptic sites. Distribution of CAPS2 was quantified by calculating the numbers of CAPS2 immunogold particles in 160 PF  $\rightarrow$  Purkinje cell synapses as a function of the distance from the active zone (Fig. 5*D*). As a result, CAPS2 was primarily distributed at presynaptic sites 250 nm distant from the active zone, which corresponded to the interval of five standard synaptic vesicles. It is noteworthy that there was significant CAPS2 distribution near the active zone (Fig. 5*D*, 0 to +50 nm).

#### CAPS2 is associated with vesicles distinct from synaptic vesicles

It was previously shown that CAPS1 associates with purified plasma membranes and SGs of rat brains (Berwin et al., 1998) and is essential for  $Ca^{2+}$ -triggered SG exocytosis in permeabilized PC12 cells (Tandon et al., 1998). A *Drosophila* mutant lacking a CAPS ortholog (dCAPS) was used to show that dCAPS is required for SG exocytosis in neuromuscular junctions (Renden et al., 2001). To examine the

property of subcellular fraction containing CAPS2, a crude synaptosomal P2 fraction was prepared from P21 mouse cerebella and further fractionated by floatation in continuous gradients of 0.3–1.8 M sucrose. In Figure 6*A*, Western blotting analyses of the resultant subfractions showed that immunoreactivity to an SV marker, synaptophysin (Synp), was recovered in lower-density fractions (fractions 2–4, from top to bottom), whereas that to a



**Figure 6.** Characterization of CAPS2-associated vesicular fractions. *A*, Western blotting analysis of subcellular vesicular fractions prepared from P21 mouse cerebella. The subfractions separated by 0.3–1.8 M sucrose gradient centrifugation were taken from the top (fraction 1) to the bottom (fraction 16). Each fraction was analyzed for immunoreactivity Syph, CGB, and CAPS2. *B*, Western blotting analysis of vesicle fractions that were immunoaffinity-purified using magnetic beads coated with the control mouse IgG, control rabbit IgG, anti-Syph antibody, and anti-CAPS2 antibody from P15 mouse cerebella. Each immunopurified fraction was analyzed for immunoreactivity to Syph, CAPS2, Syt I, VAMP-2, CGB, and BDNF. A vesicle fractions that was immunoaffinity-purified with anti-CAPS1 antibody was also analyzed for the content of synaptophysin, CAPS1, BDNF, and CAPS2. *C*, Sensitive two-site enzyme immunoassay for NT-3 in the Syph- and CAPS2-immunopurified vesicles. Each value represents mean  $\pm$  SD from four independent experiments. \* $p < 0.02$ .

SG marker CGB was in higher-density fractions (fractions 7–9) in which immunoreactivity of CAPS2 was cofractionated.

To further characterize each fraction, we purified vesicular fractions from P15 mouse cerebella by affinity for the antibody-coated magnetic beads (rabbit anti-CAPS2 IgG and control rabbit IgG and mouse anti-Syph IgG and control mouse IgG) and analyzed their protein components by Western blotting (Fig. 6*B*). The vesicular fraction immunopurified with the anti-Syph beads (Syph vesicle) contained SV proteins synaptotagmin (Syt) I (Fig. 6*B*) and II (data not shown) other than Syph, whereas that with the anti-CAPS2 beads (CAPS2-vesicle) was completely negative to such SV markers (Fig. 6*B*). A vesicle-associated membrane protein, VAMP-2, was present in the Syph vesicle (Fig. 6*B*). The presence of VAMP-2 in SGs is not definitive yet: present at a low level in SGs from chromaffin cells (Winkler, 1997) versus absent in SGs from rat brains (Zhai et al., 2001). In the CAPS2 vesicle, no trace amount of VAMP-2 was observed (Fig. 6*B*). Moreover, CGB was included in the CAPS2 vesicle but not in the Syph vesicle. Another SG protein, chromogranin A (CGA), was not detected in the CAPS2 vesicle (data not shown). Interestingly, there was no trace amount of CAPS1 detectable by this blot (Fig. 6*B*), although the anti-CAPS2 antibody used for this immunobead weakly cross-reacted with the CAPS1 protein (see Materials

and Methods), again indicating the predominant expression of CAPS2 in the cerebellum, as shown in Figure 3, *G* and *H*. Together, these results indicate that the Syph vesicle and the CAPS2 vesicle are molecularly separable.

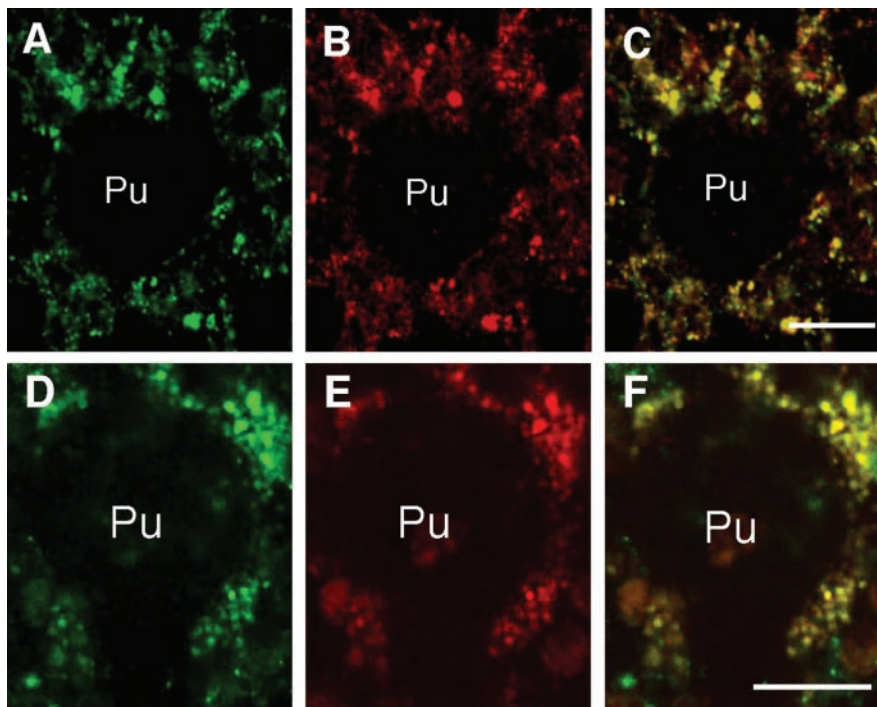
### CAPS2-associated vesicles contain neurotrophins

The Syph vesicle could be for neurotransmitters, e.g., glutamate and GABA, in the cerebellum. This raises the question of what the CAPS2 vesicle is. We examined various candidates that are known secretory substances produced by granule cells and found that the CAPS2 vesicle contains BDNF and NT-3, which are important neurotrophins for cerebellar development and maintenance. Western blotting for BDNF showed that only the CAPS2 vesicle had apparent immunoreactivity to a 14 kDa band corresponding to BDNF (Fig. 6*B*). We could not detect NT-3 by this Western blotting, which was probably attributable to its amount, the detection limit of the anti-NT-3 antibody we used, or both. However, using the highly sensitive two-site enzyme immunoassay (Katoh-Semba et al., 2000), we were able to detect significant amounts of NT-3 in the CAPS2 vesicle but not in the Syph vesicle (Fig. 6*C*). These results indicate that CAPS2 is associated with vesicles containing BDNF and NT-3, which are separable from typical SVs. On the contrary, the immunisolated vesicle with anti-CAPS1 monoclonal antibody, which did not cross-react with the CAPS2 protein, and did not contain BDNF, CAPS2 (Fig. 6*B*), or NT-3 (data not shown).

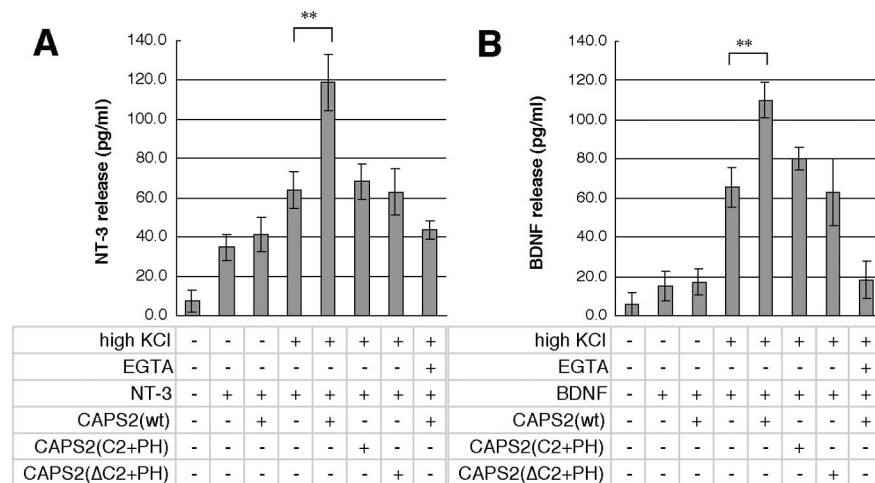
To ascertain the coincidence in subcellular localization between CAPS2 and neurotrophins, we conducted immunocytochemical analyses of cultured cerebellar neurons (14 DIV) (Fig. 7). Most immunoreactive puncta of CAPS2 were primarily colocalized with those of NT-3 (Fig. 7*A–C*) and of BDNF (Fig. 7*D–F*) in regions surrounding dendritic arborizations of Purkinje cells.

### CAPS2 has activity to enhance depolarization-induced release of neurotrophins

To investigate the functional role of CAPS2, we heterologously expressed the CAPS2 constructs together with NT-3 or BDNF in PC12 cells and analyzed NT-3 or BDNF secretion by ELISA. PC12 cells express CAPS1 but not CAPS2, NT-3, or BDNF. As shown in Figure 8, *A* and *B*, by expressing NT-3 or BDNF in PC12 cells, the immunoreactivity for NT-3 or BDNF in the culture media was slightly increased without stimulation, which may be via a constitutive secretion pathway or may be stimulated by unknown factors under the present culture conditions. On high-KCl stimulation, immunoreactivity was increased approximately twofold (for NT-3) (Fig. 8*A*) and fourfold (for BDNF) (Fig. 8*B*) from the basal levels, which is consistent with previous reports describing the depolarization-dependent secretion of exogenously expressed BDNF (Goodman et al., 1996) and NT-3 (Heymach et al., 1996) in PC12 cells probably via CAPS1. By coexpressing the full-length CAPS2 (wt), the high-KCl-induced immunoreactivity to NT-3 and BDNF was enhanced  $\sim$ 2- and 1.7-fold, respectively, compared with expressing NT-3 or BDNF alone. This CAPS2-mediated enhancement of NT-3 and BDNF immunoreactivity in culture media depended on high-KCl stimulation, and extracellular EGTA in the presence of high KCl blocked this enhancement, suggesting the importance of membrane depolarization-induced  $\text{Ca}^{2+}$  influx, which is generally elicited by high-KCl stimulation. On the other hand, coexpression of the CAPS2 mutant carrying the C2 and PH domain alone (C2 + PH) or lacking the C2 and PH domain ( $\Delta$ C2 + PH) did not confer this enhancement, indicating the importance of these structural domains for this activity.



**Figure 7.** Colocalization of CAPS2, NT3, and BDNF in cerebellar cultures at 14 DIV. *A–C*, Confocal fluorescent images of cerebellar neurons immunostained with the anti-CAPS2 (*A*, green) and anti-NT-3 (*B*, red) antibodies. *D–F*, Confocal fluorescent images of cerebellar neurons immunostained with the anti-CAPS2 (*D*, green) and anti-BDNF (*E*, red) antibodies. *C, F*, Merged images between *A, B* and *D, E*, respectively. Pu, Purkinje cell. Scale bars, 10  $\mu$ m.



**Figure 8.** Enhanced release of NT-3 and BDNF by exogenously expressed CAPS2 from PC12 cells. PC12 cells were transfected with the CAPS2 constructs (wt, C2 + PH,  $\Delta$ C2 + PH; see Materials and Methods) together with NT-3 (*A*) or BDNF (*B*), and the amounts of NT-3 (*A*) and BDNF (*B*) released into the culture media with or without 50 mM KCl stimulation were analyzed by ELISA. Data points represent the average of values obtained from three independent experiments  $\pm$  SD  $^{**}p < 0.01$ .

### Overexpression of CAPS2 promotes depolarization-induced neurotrophin release activity and Purkinje cell survival in cerebellar cultures

We next investigated CAPS2-mediated neurotrophin release from cultured granule cells. We exogenously overexpressed the CAPS2 in 7 DIV granule cells using the adenoviral vector gene delivery system. Granule cells at 24 hr after infection (8 DIV) with the adenoviral vector alone (Ad-mock) had immunoreactivity to the endogenous CAPS2 in the soma and neurites (Fig. 9*A*) and showed an increase in NT-3 immunoreactivity in the culture

media by 50 mM KCl stimulation (Fig. 9*C*). By infecting with the recombinant adenovirus vector carrying the CAPS2 (Ad-CAPS2), the CAPS2 immunoreactivity was augmented over the cells and observed as a punctate pattern on the neurites (Fig. 9*B*). In Figure 9*C*, the NT-3 immunoreactivity in the media in response to the KCl stimulation was approximately twofold higher in the Ad-CAPS2-infected cells than that in the Ad-mock-infected cells. Without high-KCl stimulation, exogenous CAPS2 did not enhance the immunoreactivity. These results reveal that CAPS2 is involved in depolarization-dependent NT-3 secretion from granule cells. Secreted BDNF was at undetectable levels in the media of 7, 14, and 21 DIV cultures (data not shown), which was probably attributable to the low concentration of released BDNF and the sensitivity of our ELISA system.

The cultures infected with the adenovirus vectors at 3 DIV were analyzed at 11 d after infection (14 DIV). As shown in Figure 9*D*, the number of calbindin-immunopositive Purkinje cells was increased approximately twofold in the Ad-CAPS2-infected cultures compared with that in the Ad-mock-infected cultures. Overexpression of functional CAPS2 probably increases the amount of neurotrophins released, the frequency of their release under normal culture conditions, or both without any stimulation such as depolarization, leading to the enhancement of their functional effect in the cultures. This possibility is supported by a previous study that showed that both BDNF and NT-3 promote Purkinje cell survival in dissociated cultures (Larkfors et al., 1996; Morrison and Mason, 1998). On the contrary, the number of MAP2-immunopositive neurons, mostly granule cells, showed only a slight increase (Fig. 9*D*).

### Discussion

The molecular basis of neurotrophin release from neurons is poorly understood regardless of the functional importance of neurotrophins in many cellular aspects of the nervous system. In the present study, we identified the presynaptic protein CAPS2, which is predominantly localized

to the PF terminals of developing granule cells and associated with vesicles containing CGB, NT-3, and BDNF, and demonstrated that overexpressed CAPS2 has a promoting activity of depolarization-induced NT-3 and BDNF release as well as of Purkinje cell survival. These results indicate that CAPS2 is a molecule involved in NT-3 and BDNF release from granule cells. Moreover, the expression profiles characteristic of postnatal cerebellar development and the promoting activity of cell survival suggest that CAPS2 has a specific role in both granule cell development

and Purkinje cell development, which are modulated by both NT-3 and BDNF (Lindholm et al., 1993; Segal et al., 1995; Lindholm et al., 1997; Schwartz et al., 1997; Doughty et al., 1998; Bates et al., 1999).

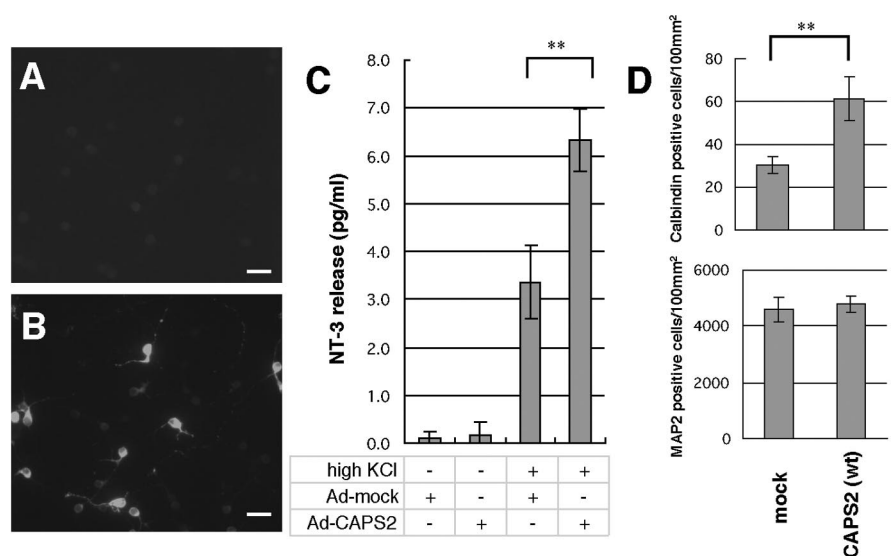
### CAPS2 is involved in activity-dependent neurotrophin release from neurons

We showed that CAPS2 and CAPS1 are differentially distributed in mouse cerebellum; the former predominates at the PF terminals in the ML, whereas the latter is preferentially located at the glomeruli in the IGL, indicating their differential cellular function in the cerebellum. CAPS1 is known to regulate  $Ca^{2+}$ -dependent release of [ $^3H$ ]norepinephrine from the preloaded PC12 cells (Ann et al., 1997) and rat brain synaptosomes (Berwin et al., 1998; Tandon et al., 1998). CAPS2 might control the release of such catecholamines. However, cerebellar granule cells express neither tyrosine hydroxylase, the first enzymatic step in catecholamine synthesis (Austin et al., 1992; Takada et al., 1993), nor dopamine- $\beta$ -hydroxylase (Verney et al., 1982), indicating that CAPS2 is involved in the release of substances, except for catecholamines, in the vesicles of PF terminals. We then showed that exogenously expressed CAPS2 promotes the NT-3 and BDNF release from PC12 cells and NT-3 release from granule cells in a depolarization-dependent manner. Substances released by CAPS1 in cerebellar glomeruli remain elusive.

BDNF and NT-3 are released in an activity-dependent manner (Goodman et al., 1996; Heymach et al., 1996; Kohara et al., 2001) and regulate their own secretion (Canossa et al., 1997; Kruttgen et al., 1998). However, little research into the molecules involved in the neurotrophin release from neurons has been reported (Canossa et al., 2001; Balkowiec and Katz, 2002; Wang et al., 2002). Thus, the present finding provides a clue to elucidating the underlying molecular basis.

### CAPS2-associated vesicles within the PF fiber terminals are distinct from synaptic vesicles

We found CAPS2 in vesicular and membranous structures in the PF terminals. The purified CAPS2 vesicles contain BDNF and NT-3, an SG marker CGB, but did not contain another SG marker, CGA, and SV markers Syt I and II and VAMP-2, indicating that the CAPS2 vesicles are completely separated from typical SVs. CGA is a key regulator of dense-core SG biosynthesis in PC12 cells, but CGB is not (Corradi et al., 1996; Kim et al., 2001). Our electron micrographs of PF terminals (Fig. 5A–C) revealed no apparent dense-core-like vesicles as previously reported (Carter et al., 2002), indicating that CAPS2 vesicles of the PF terminals are probably different from typical dense-core vesicles. This feature in the molecular composition and electron microscopic view is likely consistent with the fact that CGB does not play a direct role in dense-core SG biogenesis in PC12 cells (Corradi et al., 1996; Kim et al., 2001). On the contrary, the presence of NT-3 and BDNF in large dense-core SGs has been reported in the retinotectal terminals of chicken embryos (Wang et al., 2002) and the axonal terminals of rat dorsal root ganglions (Michael et al.,



**Figure 9.** Cellular localization of exogenously overexpressed CAPS2 and its promoting effect on NT-3 release and Purkinje cell survival. *A, B*, CAPS2-immunofluorescent images of cerebellar granule cells (8 DIV) infected with Ad-mock (*A*) and Ad-CAPS2 (*B*). Scale bars, 10  $\mu$ m. *C*, ELISA of NT-3 levels released into the conditioned media from the cultures (8 DIV) infected with Ad-mock and Ad-CAPS2 with or without 50 mM KCl stimulation. *D*, Number of Purkinje cells and granule cells in the cultures infected with Ad-mock and Ad-CAPS2. At 11 d after infection (14 DIV), the numbers of calbindin-immunopositive Purkinje cells and MAP-2-positive granule cells were counted. Data points represent the average of values obtained from four independent experiments  $\pm$  SD  $**p < 0.01$ .

1997), respectively. Cerebellar granule cells, although they express BDNF and NT-3, do not express CGA mRNAs (Mahata et al., 1991). Thus, the biochemical properties and electron microscopic view of vesicles containing neurotrophins appear different from cell type to cell type.

In SV exocytosis, a variety of molecules are involved in a series of stages (Jahn and Sudhof, 1999; Chen and Scheller, 2001; Rettig and Neher, 2002). On the other hand, the detailed molecular mechanism underlying SG exocytosis remains to be studied (Haynes et al., 2001; Voets et al., 2001a,b; Fukuda et al., 2002b; Shin et al., 2002). CAPS2 contains a predicted C2 domain, as does CAPS1, and enhances the NT-3 and BDNF release by stimulation with high KCl, which generally elicits membrane depolarization leading to  $Ca^{2+}$  influx. How CAPS2 functions in the vesicle exocytosis remains to be studied. In addition, to understand its underlying molecular machinery, CAPS2-related molecules require elucidation.

### A role of CAPS2-mediated neurotrophin release in cerebellar development

During postnatal development, the mouse cerebellum undergoes a series of magnificent cellular events to accomplish the formation of a functional circuit and architecture. Granule cells are generated by vigorous cell proliferation in the EGL to reach an immense cell number of approximately half the neurons in the brain, extend their PF axons, and migrate downward to form the IGL. Purkinje cells undergo robust outgrowth of dendrites and form elaborate arborization with numerous synapses with extending PFs of granule cells. During this period, neurotrophins act as important modulators for differentiation and survival of cerebellar neurons (Lindholm et al., 1993, 1997; Segal et al., 1995; Schwartz et al., 1997; Doughty et al., 1998; Bates et al., 1999). Among them, NT-3 and BDNF are closely related to the interaction between granule cells and Purkinje cells. Mice genetically deficient in the genes of these neurotrophins showed abnormal



cerebellar morphology (Schwartz et al., 1997; Bates et al., 1999). We showed that the exogenous CAPS2 overexpression promotes the Purkinje cell survival as the result of an increase in the neurotrophin release activity from granule cells. In our culture system (7–10 Purkinje cells per  $10^5$  plating cells), Purkinje cells were more susceptible to cell death than granule cells, and their survival primarily depended on the numbers of granule cells or astrocytes (Yuzaki and Mikoshiba, 1992). Effects on granule cell viability, neuronal differentiation, including neurite outgrowth and synapse formation, or both would be expected to be seen by use of appropriate systems such as granule cell cultures at low cell density or in low-KCl medium and genetically manipulated animal models.

In contrast to the predominant localization of CAPS2 to the PF terminals, CAPS1 distributes specifically in glomerular rosettes in the cerebellum and is widespread in other brain areas. The present study suggests that CAPS1 and CAPS2 share their roles in modulation of cerebellar development and function, although what kinds of substances, including, e.g., neurotrophins and neuropeptide, are released by CAPS1 should be examined, and CAPS2 is specifically associated with the cerebellar postnatal development during which tens of millions of neurons undergo vigorous differentiation events. Moreover, CAPS2-mediated BDNF release may be involved in synaptic plasticity even in the fully developed cerebellar circuitry, because BDNF has a modulatory role in the short-term plasticity of PF → Purkinje cell synapses (Carter et al., 2002).

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