Ablation of glutamate receptor GluRδ2 in adult Purkinje cells causes multiple innervation of climbing fibers by inducing aberrant invasion to parallel fiber innervation territory.
Ablation of Glutamate Receptor GluR82 in Adult Purkinje Cells Causes Multiple Innervation of Climbing Fibers by Inducing Aberrant Invasion to Parallel Fiber Innervation Territory

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Glutamate receptor GluR82 is exclusively expressed in Purkinje cells (PCs) from early development and plays key roles in parallel fiber (PF) synapse formation, elimination of surplus climbing fibers (CFs), long-term depression, motor coordination, and motor learning. To address its role in adulthood, we previously developed a mouse model of drug-induced GluR82 ablation in adult PCs (Takeuchi et al., 2005). In that study, we demonstrated an essential role to maintain the connectivity of PF–PC synapses, based on the observation that both mismatching of presynaptic and postsynaptic specializations and disconnection of PF–PC synapses are progressively increased after GluR82 ablation. Here, we pursued its role for CF wiring in adult cerebellum. In parallel with the disconnection of PF–PC synapses, ascending CF branches exhibited distal extension to innervate distal dendrites of the target and neighboring PCs. Furthermore, transverse CF branches, a short motile collateral rarely forming synapses in wild-type animals, displayed aberrant mediolateral extension to innervate distal dendrites of neighboring and remote PCs. Consequently, many PCs were wired by single main CF and other surplus CFs innervating a small part of distal dendrites. Electrophysiological recording further revealed that surplus CF-EPSCs characterized with slow rise time and small amplitude emerged after GluR82 ablation, and increased progressively both in number and amplitude. Therefore, GluR82 is essential for maintaining CF monoinnervation in adult cerebellum by suppressing aberrant invasion of CF branches to the territory of PF innervation. Thus, GluR82 fuels heterosynaptic competition and gives PFs the competitive advantages over CFs throughout the animal’s life.

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Introduction
Cerebellar Purkinje cells (PCs) receive two distinct excitatory afferents (Palay and Chan-Palay, 1974). Each PC receives 10^5–10^6 inputs from parallel fibers (PFs) at distal dendrites, whereas each PF forms one or two synapses to individual PCs (Napper and Harvey, 1988). In contrast, each PC is innervated by a single climbing fiber (CF) that originates from the inferior olive and forms hundreds of synapses by twisting around proximal dendrites. This PC circuitry is established through heterosynaptic competition between PFs and CFs and homosynaptic competition among multiple CFs (Mariani et al., 1977; Crepel, 1982; Hashimoto et al., 2009). This competitive wiring is regulated by various signaling molecules expressed in cellular elements constituting PC synapses, including metabotropic glutamate receptor mGluR1, P/Q-type Ca^{2+} channels, glutamate receptor GluR82 (GluD2), and Cbln1 (or precerebellin) (Kano et al., 1995, 1997, 1998; Kashiwabuchi et al., 1995; Offermanns et al., 1997; Ichikawa et al., 2002; Miyazaki et al., 2004; Hirai et al., 2005).

Of these, GluR82 regulates heterosynaptic competition to the advantage of PF innervation (Watanabe, 2008; Yuzaki, 2009). GluR82 is dominantly expressed in PCs (Araki et al., 1993; Lomeli et al., 1993) and localized on spines contacting PFs (Takayama et al., 1995; Landsend et al., 1997). Targeted disruption of GluR82 gene in mice causes mismatching between presynaptic and postsynaptic specializations and disconnection at PF–PC synapses (Guastavino et al., 1990; Kashiwabuchi et al., 1995; Kurihara et al., 1997; Lalouette et al., 2001). The N-terminal domain of GluR82 has been demonstrated to mediate PF–PC synaptogenesis by interacting neurexin through Cbln1 (Uemura et al., 2007, 2010; Kakegawa et al., 2008, 2009; Uemura and
Mishina, 2008; Torashima et al., 2009; Matsuda et al., 2010). The loss of GluR2 further affects CF innervation; CFs extend distally to take over free spines of the target and neighboring PCs, causing multiple CF innervation (Hashimoto et al., 2001; Ichikawa et al., 2002). Intriguingly, mutant mice lacking the Ca₉.2,1, a pore-forming subunit of P/Q-type Ca²⁺ channels, display almost reciprocal phenotypes that PF territory expands, and CF territory regresses, down to PC somata and basal dendrites (Miyazaki et al., 2004). Therefore, the construction of functional PC circuits stands on competitive equilibrium among afferents prefermented by distinct molecular mechanisms.

Heterosynaptic competition is still active in adult cerebellum, as surgical lesion to olivocerebellar projections or pharmacological blockade of corticofugal activities alters innervation territories of PFs and CFs reciprocally (Sotelo et al., 1975; Rossi et al., 1991a; Cesa et al., 2005). To address the molecular mechanisms that maintain functional synaptic wiring in adult cerebellum, we previously developed a mouse model of CrePR/loxP-mediated GluR2 ablation in adult PCs (Takeuchi et al., 2005). In that study, we demonstrated that GluR2 plays an essential role to maintain the connectivity of PC–PC synapses. Here, we addressed that GluR2 is also indispensable to maintain CF monoinnervation. After induction of GluR2 ablation, CF innervation extended distally and mediolaterally in the cerebellar cortex, giving rise to aberrant surplus branches that caused multiple innervation.

Materials and Methods

Animals. We used littersmates derived from crossing of GluR2flox/flox and GluR2flox/CrePR mice on the pure C57BL/6 genetic background, as reported previously (Takeuchi et al., 2005). GluR2flox/CrePR mice were produced by crossing a target mouse line carrying the GluR2 gene flanked by loxP sequences with a Cre mouse line carrying the Cre recombinase-progesterone receptor (CrePR) fusion protein under the control of the GluR2 gene promoter. All animal experiments were performed according to the guidelines for the care and use of laboratory animals of the Hokkaido University School of Medicine. The GluR2floxc allele was identified by PCR using primers 5′-AGCAACATCATACTCCCAAAGAAC-3′ (FD2P2) and 5′-ATTCCGTGGCCAAGACGACAGAACAA-3′ (FD2P4). The GluR2CrePR allele was identified by PCR using the CreP1 and CreP2 primers (Tsujita et al., 1999). For induction of PC-specific gene recombination, 11B-[(dimethylamino)phenyl]-17β-hydroxy-17α-propynyl)estr-4-9-dien-3-one (RU-486) (Sigma-Aldrich), suspended at a concentration of 50 mg/ml in water containing 0.25% (v/v) carboxymethyl cellulose and 0.5% (v/v) Tween 80, was injected intraperitoneally at a dose of 1 mg/g body weight RU-486 at postnatal day 42 (P42) to P45 for 2 consecutive days (Takeuchi et al., 2005). We also used global or null-type GluR2 knockout mice, which were produced as a Cre knock-in mouse line D2CreN under the control of the GluR2 gene promoter (GluR2Cre/C0) using the C57BL/6 ES cell line RENKA (Mishina and Sakimura, 2007). Details of the GluR2Cre/C0 mouse will be described elsewhere.

In each morphological analysis, we analyzed three GluR2floxFlox and three GluR2floxC0 as control and mutant mice, respectively. Under deep pentobarbital anesthesia, mice were perfused transcardially with 4% paraformaldehyde in 0.1 M sodium phosphate buffer, pH 7.2. After deep pentobarbital anesthesia, mice were perfused transcardially with the same fixative and processed for preparation of parasagittal or horizontal microslicer sections (50 μm in thickness) were prepared with an ultramicrotome (Ultracut; Leica), and photographs were taken with an H-7100 electron microscope (Hitachi).

Electrophysiology. Whole-cell recordings were made from visually identified PCs using an upright microscope (BX51WI; Olympus) at 31°C. Patch pipettes had resistance of 2–4 MΩ when filled with an internal solution composed of the following (in mM): 60 CsCl, 10 Cs-n-gluconate, 20 TEA (tetraethylammonium)-Cl, 20 BAPTA, 4 MgCl₂, 4 ATP, 0.4 GTP, and 30 HEPES, pH 7.3, adjusted with CsOH. The pipette access resistance was compensated by 80%. The composition of the standard bathing solution was as follows (in mM): 125 NaCl, 2.5 KCl, 2 CaCl₂, 1 MgSO₄, 1.25 NaH₂PO₄, 26 NaHCO₃, and 20 glucose, bubbled with 95% O₂ and 5% CO₂. Bicuculline methochloride (10 μM; Tocris Bioscience) was always added to the bath solution for blocking inhibitory synaptic transmission. Ionic currents were recorded with a patch-clamp amplifier (EPC10; HEKA). The signals were filtered at 2 kHz and digitized at 20 kHz. On-line data acquisition and off-line data analysis were performed previously (Araki et al., 1993; Miyazaki et al., 2003; Miura et al., 2006). For GluR2 immunofluorescence, sections were digested with 1 mg/ml pepsin (Dako) in PBS, pH 7.4/0.2N HCl for 3 min at 37°C in a water bath. All immunohistochemical incubations were done at room temperature in a free-floating state. For immunofluorescence, cerebellar sections were incubated with 10% normal donkey serum for 20 min, a mixture of primary antibodies overnight, and a mixture of Alexa Fluor 488-, indocarbocyanine (Cy3)-, and indodicarbocyanine (Cy5)-labeled species-specific secondary antibodies for 2 h at a dilution of 1:200 (Invitrogen; Jackson ImmunoResearch). Images were taken with a confocal laser-scanning microscope (FV1000; Olympus) or with a fluorescence microscope (AX-70; Olympus) equipped with a digital camera (DP70; Olympus), and analyzed with MetaMorph software (Molecular Devices). The reach of CFs was evaluated by the distance from the base of the molecular layer to the tips of VGluT2-positive CF terminals relative to the total vertical height of the molecular layer.

Anterograde tracer labeling. Under anesthesia with chloral hydrate (350 mg/kg body weight, i.p.), a glass pipette (G-1.2; Narishige) filled with 2–3 μl of 10% solution of biotinylated dextran amine (BDA) (3000 molecular weight; Invitrogen) or dextran Alexa 594 (DA-594) (Invitrogen) in PBS was inserted stereotaxically to the inferior olive by the dorsal approach. Tracers were injected by air pressure at 20 psi with 5 s intervals for 1 min (Pneumatic Picompump; World Precision Instruments). After 4 days of survival, mice were anesthetized and fixed by transcardial perfusion. For bright-field light microscopy, BDA-labeled CFs were visualized by overnight incubation with peroxidase-labeled streptavidin (Nichirei) and visualized in dark using 3′,3′-diaminobenzidine (DAB) and cobalt. For combined labeling by tracer and immunofluorescence, DA-594-labeled microslicer sections were incubated with a mixture of calbindin and VGluT2 antibodies followed by incubation with fluorescent secondary antibodies for 2 h. Images of double or triple labeling were taken with a confocal laser-scanning microscope. For quantitative analyses of transverse CF collaterals, six optical sections taken in the z-axis plane at an interval of 1 μm were stacked into a single image, and >15 images were obtained from each mouse. We measured the number of transverse collateral branches originating from ascending CF branches; the incidence was calculated as the number at the center 100 μm of the parent ascending branches. We counted the number of VGluT2-positive terminals per 100 μm of transverse branches. Quantitative analyses were performed with MetaMorph software, and all data were described as the mean ± SEM. Statistical significance was evaluated by Mann–Whitney U test. Statistical significance was assumed when p < 0.05.

For immunoelectron microscopy, BDA-labeled microslicer sections were incubated with VGluT2 antibody diluted with Tris-buffered saline containing 1% bovine serum albumin and 0.004% saponin. Sections were incubated with 1.4 nm gold particle-conjugated anti-guinea pig antibody (1:200; Nanogold; Nanoprobes) for 3 h. Immunogold for VGluT2 was silver-enhanced with an HQ-silver enhance kit (Nanoprobes), whereas BDA was detected by overnight incubation with peroxidase-labeled streptavidin (Nichirei) and visualized with DAB. Sections were postfixed with 1% osmium tetroxide for 15 min, dehydrated in graded alcohols, and embedded in Epon 812. Ultrathin sections (70 nm in thickness) were prepared with an ultramicrotome (Ultratcut; Leica), and photographs were taken with an H-7100 electron microscope (Hitachi).
performed using PULSE software (HEKA). Stimulation pipettes (5–10 μm tip diameter) were filled with the standard saline and used to apply square pulses for focal stimulation (duration, 0.1 ms; amplitude, 0–90 V). Decay time constant of CF-mediated EPSC (CF-EPSC) was measured by fitting the EPSC decay with single exponential (LLano et al., 1991). Passive membrane properties were calculated based on the equivalent circuit model proposed by LLano et al. (1991). The current in response to hyperpolarizing voltage steps from −70 to −80 mV was biphasic and fitted with two exponentials. Both time constants for the fast (τ1) and slow (τ2) components were not significantly different between control and mutant mice at any time points examined (supplemental Table S1, available at www.jneurosci.org as supplemental material). Statistical significance in frequency distribution was evaluated by χ² test or Mann–Whitney U test. Statistical significance was assumed when p < 0.05. For linear regression analyses, Pearson’s product moment correlation coefficient (r²) was calculated. Results

The global GluR2-knock-out mice display two major anatomical phenotypes in terms of CF–PC wiring: distal extension of CF territory and multiple CF innervation by surplus innervation to distal dendrites of CF territory and multiple CF innervation (Kashiwabuchi et al., 1995; Ichikawa et al., 2001). The current in response to hyperpolarizing voltage steps from −70 to −80 mV was biphasic and fitted with two exponentials. Both time constants for the fast (τ1) and slow (τ2) components were not significantly different between control and mutant mice at any time points examined (supplemental Table S1, available at www.jneurosci.org as supplemental material). Statistical significance in frequency distribution was evaluated by χ² test or Mann–Whitney U test. Statistical significance was assumed when p < 0.05. For linear regression analyses, Pearson’s product moment correlation coefficient (r²) was calculated. Rotarod test. Control (n = 15) and mutant (n = 11) mice were used for motor behavioral test at 8, 12, 16, 20, 24, and 28 weeks after RU-486 administration. Mice were housed individually and were handled for ~1 min a day for 7–10 d before behavioral tests. An animal was placed on the rod (3.2 cm diameter) rotating at 15 rpm (RRAC-3002; O’Hara), and given three trials with 45–60 min intertrial intervals. The latency to fall (retention time) was measured with cutoff time of 5 min. Rotarod test was performed in a blind fashion. Statistical significance was evaluated by ANOVA with repeated measures followed by Student’s t test. Statistical significance was assumed when p < 0.05.

Distal extension of CF innervation is induced after GluR2 ablation

Using parasagittal cerebellar sections, we applied double immunofluorescence for calbindin and VGluT2, markers for PCs and CF terminals, respectively (Fig. 1A–D). At all time points examined, CF terminals in control mice climbed up along proximal or shaft dendrites until their transition to distal dendrites or spiny branchlets (Fig. 1A, E), suggesting little influence, if any, on dendritic morphology and CF innervation by RU-486 administration per se. Reflecting the territorized innervation, CF terminals were distributed within basal four-fifths of the molecular layer in control mice. In mutant mice at 2 weeks after RU-486 administration, the distribution of CF terminals was comparable with that in control mice (Fig. 1B). At later time points, however, CF terminals were distributed closer to the pial surface (Fig. 1C, D). At high magnifications, distal dendrites of mutant PCs were stunted and studded with shrunken spines (Fig. 1F). Furthermore, CF terminals in mutant mice were not confined to proximal shaft dendrites, but also associated with the atrophied distal dendrites (Fig. 1F). These phenotypic alterations were confirmed by measuring the thickness of the molecular layer (Fig. 1G) and the height of VGluT2-positive CF terminals (Fig. 1H). The mean thickness of
the molecular layer decreased progressively (Fig. 1G), whereas the mean relative height of CF terminals in molecular layer increased progressively (Fig. 1H), both showing significant differences at 8, 16, and 24 weeks (p < 0.00001 for each comparison, U test). These results suggest that, after induction of GluR2 ablation, CF territory extends distally along atrophied distal dendrites.

After RU-486 administration, the ablation of GluR2 protein in mutant mice proceeds unevenly among PC populations (Takeuchi et al., 2005). We examined the relationship between distal extension of CF territory and GluR2 ablation by triple immunofluorescence for calbindin (blue), VGluT2 (red), and GluR2 (green) at 12 weeks after RU-486 administration (Fig. 2A–E). In this analysis, we used horizontal cerebellar sections, in which PC dendrites are seen as single straight bars and GluR2 is detected, if expressed, on the surface of dendritic bars. In control mice, the reach of VGluT2-positive CF terminals up to basal four-fifths of the molecular layer was confirmed in horizontal sections (mean ± SEM, 77.2 ± 0.5% of the molecular layer; n = 54) (Fig. 2A, B, E). In mutant mice, the relative reach was significantly extended along dendritic bars lacking GluR2 (88.2 ± 0.4%; n = 38), compared with that along dendritic bars retaining GluR2 (75.2 ± 0.8%; n = 38; p < 0.00001, U test) (Fig. 2C–E). Thus, distal extension of CF innervation selectively occurs along GluR2-ablated PC dendrites.
GluR62 ablation induces multiple innervation by ascending CF branches

Then we tested whether the distally extended ascending branches caused multiple innervation. To this end, we applied triple fluorescent labeling for calbindin (blue), VGluT2 (green), and anterograde tracer DA-594 (red) to distinguish anatomical forms of CF innervation (Fig. 3). In this analysis, PC dendrites that were associated with CF terminals either double-labeled for DA594 and VGluT2 or single-labeled for VGluT2 only were judged to be monoinnervated (supplemental Fig. S1A, available at www.jneurosci.org as supplemental material), whereas those associated with double- and single-labeled CF terminals simultaneously were to be multiply innervated (supplemental Fig. S1B, available at www.jneurosci.org as supplemental material).

In parasagittal cerebellar sections, monoinnervation pattern was overwhelming in control mice at each time point examined (Fig. 3A) and in mutant mice at 2 weeks after RU-486 administration (Fig. 3B). At 8 weeks and thereafter, mutant mice frequently displayed anatomical patterns of multiple innervation (Fig. 3C,D). For example, PC dendrites, marked as PCD-a, were mainly innervated by ascending branches of tracer-labeled CF-a (Fig. 3C) or tracer-unlabeled CF-a (Fig. 3D), respectively. These dendrites received additional innervation by tracer-unlabeled CF-b (Fig. 3C, green arrows) or tracer-labeled CF-b (Fig. 3D, red arrows), respectively, thus demonstrating the innervation by multiple CFs. Predominant monoinnervation in control mice and predominant multiple innervation in mutant mice were also confirmed using horizontal cerebellar sections at 16 weeks after RU-486 administration (supplemental Fig. S2, available at www.jneurosci.org as supplemental material). In particular, alternate innervation of single dendritic bars by tracer-labeled CF-a and tracer-unlabeled CF-b was readily captured in mutant mice (supplemental Fig. S2C,D, available at www.jneurosci.org as supplemental material). In general, terminals of additional CFs were few in number and intermingled with those formed by main CFs at 8 weeks (Fig. 3C), whereas additional CFs often covered significant portions of PC dendrites and tended to be segregated from main CFs within the same dendrites at 16 and 24 weeks after RU-486 administration (Fig. 3D; supplemental Fig. S2C,D, available at www.jneurosci.org as supplemental material). Thus, the ablation of GluR62 induces aberrant wiring of ascending CF branches to dendrites of nearby PCs, thereby causes multiple innervation, and this is exacerbated with time after RU-486 administration.

Aberrant mediolateral elongation and terminal differentiation of transverse CF branches

We also examined the transverse branch of CFs. This branch is a thin motile collateral, which originates from the parental ascending branches, extends mediolaterally in the horizontal and transverse cerebellar planes, and rarely forms conventional synapses in wild-type rodents (Rossi et al., 1991a; Sugihara et al., 1999; Nishiyama et al., 2007). In horizontal cerebellar sections, anterograde tracer BDA clearly visualized vertical ladders of ascending CF branches subjected to dextran Alexa-594 tracer labeling for CFs (DA-594; red) and immunofluorescence for calbindin (calb; blue) and VGluT2 (green) in GluR62 flox/flox at 2 weeks after RU-486 administration (A) and GluR62 flox/CrePR mice at 2 weeks (B), 8 weeks (C), and 24 weeks (D) after RU-486 administration. Immunoreaction for calbindin is pseudocolored in blue (A1–D1) or brown (A2–D2, A3–D3). CF-a indicates the CF mainly innervating a PC dendrite of interest (PCD-a). In GluR62 flox/CrePR mice at 8 and 24 weeks, PCD-a is multiply innervated by both main CF-a and additional CF-b (arrows). Scale bars, 10 μm.
branches in both types of mice (Fig. 4A–E). In such middle-power magnification of bright-field images, transverse branches were too short and thin to be identified at all time points in control mice and at 2 weeks in mutant mice (Fig. 4A, B). By contrast, a few transverse branches were clearly discerned to elongate in mutant mice at 8 weeks (Fig. 4C, arrows). At 16 and 24 weeks, transverse branches were further increased in number and length, particularly, in the superficial half of the molecular layer (Fig. 4D, E, arrows).

Morphological changes of transverse branches were examined more in detail by triple fluorescent labeling (Fig. 4F–O). In control mice, DA-594-labeled transverse branches were uniformly thin in caliber and rarely formed VGluT2-positive terminals at each time point examined (Fig. 4F, G, arrowheads). In mutant mice, transverse branches were immunonegative to VGluT2 at 2 weeks (Fig. 4H, I, arrowheads), but differentiated a few VGluT2-positive terminals at 8 weeks (Fig. 4J, K, arrows). These ectopic terminals were formed more around dendritic bars lacking GluRΔ2 (6.1 ± 0.6 per 100 μm of transverse branches; n = 29) than around those retaining GluRΔ2 (2.6 ± 0.7; n = 31; p < 0.00001, U test) (Fig. 2F, G), indicating that terminal differentiation on transverse branches was preferentially induced at around GluRΔ2-ablated PC dendrites. In parallel with marked elongation of transverse branches, VGluT2-positive terminals were greatly increased in number at 16 and 24 weeks (Fig. 4L–O, arrows).

Indeed, the mean number of VGluT2-positive terminals per 100 μm of transverse branches increased strikingly from 8 weeks to 16 and 24 weeks, showing significant differences between control and mutant mice at 8 weeks (p < 0.01, U test) and at 16 and 24 weeks (p < 0.00001 for each) (Fig. 5A). We also measured the mean number of transverse branches per 100 μm of ascending branches, and found significant increase in mutant mice at 24 weeks after RU-486 administration (Fig. 5B) (p < 0.00001). Then, we plotted the number of VGluT2-positive terminals on a given transverse branch against its relative vertical height in the molecular layer (Fig. 5C–F). In both types of mice, the majority of transverse branches originated and ran in the superficial one-half of the molecular layer [control mice, 74.5% (143 of 192 branches) and 76.7% (145 of 189 branches) at 8 and 24 weeks, respectively; mutant mice, 70.0% (147 of 210 branches) and 70.4% (162 of 230 branches) at 8 and 24 weeks, respectively]. Therefore, GluRΔ2 ablation has induced marked mediolateral elongation and extensive terminal differentiation in transverse CF branches, particularly, at the superficial molecular layer.

Multiple CF innervation is also caused by transverse CF branches

Whether the aberrant transformation of transverse branches also caused multiple innervation was examined using mutant mice at 16 weeks after RU-486 administration. Figure 6, A and B, shows a typical example, in which a DA-594-labeled transverse branch CF-b (red) formed VGluT2-positive terminals, some of which attached to a distal dendrite PCD-a of a neighboring PC (arrowheads). This PCD-a, however, was innervated by many terminals of a DA-594-unlabeled ascending branch CF-a (green or light blue), thus representing multiple innervation. In general, such ectopic terminals of transverse CF branches were few in number and intermingled with those of main ascending CFs. In parasag-
nals of a BDA-labeled/VGlut2-labeled ascending CF-a (Fig. 6E, H). In the adjacent sections, its distal portion was further innervated by a single small terminal of a BDA-unlabeled/VGlut2-labeled CF-b (Fig. 6E2,F). Because the trajectory of CF-b was perpendicular to that of the innervating dendrite, CF-b is likely a transverse branch originating from an ascending CF branch in the neighbor. Therefore, the ablation of Glur82 in adult PCs induces multiple innervation by aberrant wiring of transverse CF branches.

Together, these anatomical results demonstrate that the loss of Glur82 in adult PCs does permit CFS to extend distally and mediolaterally, and frequently causes multiple innervation to neighboring and remote PCs.

**CF phenotypes in Glur82^floxFloxCrePR mice mimic those in global Glur82 knock-out**

The above results indicate that the two major anatomical phenotypes of global Glur82-knock-out mice are reproduced in Glur82^floxFloxCrePR mice. In addition, we confirmed in the present study that multiple CF innervation caused by both ascending and transverse branches was also observed in global Glur82-knock-out mice (supplemental Fig. S3, available at www.jneurosci.org as supplemental material).

As Glur82^floxFloxCrePR mice at 24 weeks after RU-486 administration correspond to 30 weeks of age, we further compared CF phenotypes using the age-matched global Glur82-knock-out mice (supplemental Fig. S4, available at www.jneurosci.org as supplemental material). The number of VGlut2-positive terminals per 100 μm of transverse branches was abnormally high in global Glur82-knock-out mice at 30 weeks of age (supplemental Fig. S4A, available at www.jneurosci.org as supplemental material), although the number was significantly lower than that in Glur82^floxFloxCrePR mice (p < 0.00001, U test). However, the mean number of transverse branches per 100 μm of ascending branches was comparable between age-matched Glur82^floxFloxCrePR and global Glur82-knock-out mice (supplemental Fig. S4B, available at www.jneurosci.org as supplemental material) (p = 0.4250, U test). When plotting the terminal number of a given transverse branch against its relative vertical height in the molecular layer, 69.6% (135 of 194) of transverse branches were distributed in the superficial one-half of the molecular layer in global Glur82-knock-out mice at 30 weeks of age (supplemental Fig. S4C, available at www.jneurosci.org as supplemental material), showing distribution plots quite similar to age-matched Glur82^floxFloxCrePR mice (Fig. 5F). From these similarities, CF phenotypes by Glur82 ablation in adult PCs mimic those by global Glur82 knock-out in many respects.

**Multiple CF innervation increases progressively after RU administration**

These genotypic differences were further examined electrophysiologically. Using parasagittal cerebellar slices, PCs were recorded in the whole-cell configuration, and CFS were stimulated with a glass pipette placed in the granular layer near the recorded PCs (Konnerth et al., 1990; Hashimoto et al., 2001). Individual CF-EPSC was clearly discerned, because it was elicited in an all-or-none fashion, and showed prominent paired-pulse depression (Konnerth et al., 1990; Hashimoto et al., 2001). Individual CF-EPSC was clearly discerned, because it was elicited in an all-or-none fashion, and showed prominent paired-pulse depression (Konnerth et al., 1990; Hashimoto et al., 2001). Individual CF-EPSC was clearly discerned, because it was elicited in an all-or-none fashion, and showed prominent paired-pulse depression (Konnerth et al., 1990; Hashimoto et al., 2001). Individual CF-EPSC was clearly discerned, because it was elicited in an all-or-none fashion, and showed prominent paired-pulse depression (Konnerth et al., 1990; Hashimoto et al., 2001). Individual CF-EPSC was clearly discerned, because it was elicited in an all-or-none fashion, and showed prominent paired-pulse depression (Konnerth et al., 1990; Hashimoto et al., 2001).
ined: 75.0% (27 of 36 PCs) at 2 weeks, 78.1% (25 of 32) at 8 weeks, and 75.6% (28 of 37) at 24 weeks after RU-486 administration (Fig. 7A–C, left traces; D–F, white bars), showing no significant differences between time points examined \( (p/H_{11005} = 0.41 \text{ between 2 and 8 weeks}, \quad p/H_{11021} = 0.90 \text{ between 8 and 24 weeks}, \quad \chi^2 \text{ test}) \). Thus, most PCs in control mice are innervated by single CFs. In mutant mice, the percentage of monoinnervated PCs was 73.0% (27 of 37) at 2 weeks, but decreased to 8.3% (3 of 36) at 8 weeks, and to 2.7% (1 of 37) at 24 weeks (Fig. 7A–C, right traces; D–F, black bars). Between control and mutant mice, the frequency distribution of CF-EPSC step numbers showed no significant difference at 2 weeks \( (p = 0.53, \chi^2 \text{ test}) \) but differed significantly at 8 and 24 weeks \( (p < 0.0001 \text{ for each comparison}) \). Significant difference in the frequency distribution was also found in mutant mice between 2 and 8 weeks, and between 8 and 24 weeks \( (p < 0.0001 \text{ for each comparison}, \chi^2 \text{ test}) \). Accordingly, the mean number of

Figure 6. Multiple CF innervation caused by aberrant wiring of transverse CF branches. A–D, Triple fluorescent labeling for anterograde tracer DA-594 (red), calbindin (blue), and VGluT2 (green) in horizontal (A, B) and parasagittal (C, D) cerebellar sections of GluR2\textsuperscript{flox/CrePR} mice at 16 weeks after RU-486 administration. The boxed regions in A and C are enlarged in B and D, respectively. CF-a represents tracer-unlabeled CFs mainly innervating PC dendrites of interest (PCD-a), whereas CF-b represents tracer-labeled transverse branches additionally innervating the PCD-a (red arrowheads). E–H, Double-labeling serial electron microscopy for VGluT2 (metal particles) and anterograde tracer BDA (diffuse precipitates) in GluR2\textsuperscript{flox/CrePR} mice at 24 weeks after RU-486 administration. The boxed regions in E2 and E3 are enlarged in F–H. PCD-a pseudocolored in green is a typical distal dendrite of PCs, but has numerous free spines (E, arrowheads; G, FS), thus indicating GluR2-ablated dendrite. Note that tracer-unlabeled CF-b (F) and tracer-labeled CF-a (H) both innervate the PCD-a. The asterisks indicate spines contacting to terminals CF-a or CF-b. The small arrowheads in G indicate the edges of the postsynaptic density of free spine. The black arrows in F–H indicate the neck of spines protruding from PCD-a. PFs, Parallel fiber axons. Scale bars: A, C, 20 μm; B1, D1, 5 μm; E, 2 μm; F–H, 500 nm.
CF-EPSC steps was progressively increased in mutant PCs (1.3 ± 0.5, 3.3 ± 1.9, and 8.2 ± 3.6 CFs at 2, 8, and 24 weeks, respectively), whereas it remained stable in control PCs (1.3 ± 0.4, 1.3 ± 0.5, and 1.3 ± 0.6 CFs at 2, 8, and 24 weeks, respectively). These results indicate that multiple CF innervation progressively increased in mutant PCs.

**Surplus CF-EPSCs with small amplitudes and slow kinetics emerge after GluR62 ablation**

In both types of mice, CF-EPSCs in multiply innervated PCs were composed of a single main step having the largest amplitude and one or more surplus step(s) with much smaller amplitude (supplemental Fig. S5, available at www.jneurosci.org as supplemental material). The main CF-EPSCs recorded from multiply innervated PCs and CF-EPSCs from monoinnervated PCs showed no significant differences in terms of the amplitude or kinetics (data not shown), therefore they were pooled together as “main CF-EPSCs,” and their amplitudes and kinetics were compared with those of “surplus CF-EPSCs.” These main CF-EPSCs displayed a clear paired-pulse depression to a similar extent in mutant and control mice at any time points examined (supplemental Fig. S7A, available at www.jneurosci.org as supplemental material).

In control mice, both main and surplus CF-EPSCs were distributed as a single population with a peak around 0.5 ms, and their amplitude and 10–90% rise time were similar among various time points after RU-486 administration (Fig. 8A, C, E; supplemental Fig. S5A, C, E; available at www.jneurosci.org as supplemental material). In mutant mice at 2 weeks, the distributions of amplitude and rise time of CF-EPSCs were comparable with those in control mice (Fig. 8B; supplemental Fig. S5B, available at www.jneurosci.org as supplemental material). In mutant mice at 8 and 24 weeks, however, the rise time segregated into two distinct populations (Fig. 8D, F; supplemental Fig. S5D, F, available at www.jneurosci.org as supplemental material): The fast population (<0.7 ms, mean SD for CF-EPSCs from control mice) consisted mostly of the main CFs and centered around 0.5 ms (Fig. 8D, F; supplemental Fig. S5D, F, available at www.jneurosci.org as supplemental material). However, the slow population (>0.7 ms) consisted entirely of surplus CFs peaked around 2–3 ms (3.0 ± 1.6 ms at 8 weeks; n = 75) (Fig. 8D; supplemental Fig. S5D, available at www.jneurosci.org as supplemental material) (2.2 ± 1.0 ms at 24 weeks; n = 246) (Fig. 8F; supplemental Fig. S5F, available at www.jneurosci.org as supplemental material). Importantly, when those slow CF-EPSCs were omitted, the frequency distribution of fast CF-EPSCs was comparable between control and mutant mice at each time point (Fig. 8G–I), indicating that the CF input generating slow EPSCs was selectively recruited to mutant PCs. Surplus slow CF-EPSCs were also increased in amplitude between 8 and 24 weeks (19.7 ± 14.7 pA, n = 69; 61.0 ± 86.5 pA, n = 259; mean ± SD at 8 and 24 weeks, respectively; p < 0.0001, U test) (supplemental Fig. S6, available at www.jneurosci.org as supplemental material). At 24 weeks, in addition to slow CF-EPSCs with small amplitude, those with large amplitude (>200 pA) were occasionally observed (supplemental Fig. S6B, available at www.jneurosci.org as supplemental material). At each time point, there was no significant correlation between the rise time and amplitude of slow CF-EPSCs [Pearson’s correlation coefficients (r2) = 0.04 and 0.03 at 8 and 24 weeks, respectively] (supplemental Fig. S6, available at www.jneurosci.org as supplemental material).

In parallel with the increase of surplus slow CF-EPSC, the 10–90% rise time and decay time constant of the main CF-EPSCs became slower. At 2 weeks, the 10–90% rise time of the main CF from mutant mice was similar to that from control mice (0.4 ± 0.1 ms, n = 30; 0.4 ± 0.1 ms, n = 27; mean ± SD from control and mutant mice, respectively; p = 0.71, U test) (Fig. 8A, B; supplemental Figs. S5A, B, S7B, available at www.jneurosci.org as supplemental material), and 8 weeks (0.5 ± 0.1 ms, n = 31; 0.5 ± 0.1 ms, n = 31; from control and mutant mice, respectively; p = 0.71) (Fig. 8C, D; supplemental Figs. S5C, D, S7B, available at
However, it was significantly slower in mutant mice than in control mice at 24 weeks (0.5 ± 0.1 ms, n = 38; 0.6 ± 0.2 ms, n = 33; from control and mutant mice, respectively; p < 0.001) (Fig. 8E,F; supplemental Figs. S5E,F, S7B, available at www.jneurosci.org as supplemental material). The decay time constant of the mutant main CF was similar to that of control at 2 weeks (5.6 ± 1.6 ms, n = 30; 6.1 ± 1.4 ms, n = 28; mean ± SD from control and mutant mice, respectively; p = 0.25, U test); however, it became significantly slower than that of control one thereafter (5.5 ± 1.1 ms, n = 29; 7.4 ± 2.3 ms, n = 31; at 8 weeks, from control and mutant mice, respectively; 4.8 ± 1.1 ms, n = 29; 6.8 ± 2.3 ms, n = 32; at 24 weeks, from control and mutant mice, respectively; p < 0.0001 for each comparison) (supplemental Fig. S7C, available at www.jneurosci.org as supplemental material).

The slow kinetics of CF-EPSCs might be attributable to the distal extension of CF territory (Hashimoto et al., 2001; Uemura et al., 2007); however, EPSC waveforms can also be affected by other factors, including the passive membrane properties and dendritic arborization (Hashimoto et al., 2001; Roth and Hausser, 2001). Therefore, we next examined the passive membrane properties of PCs. Both time constants for the fast (τ1) and slow (τ2) components were not significantly different between control and mutant PCs at any time points examined (supplemental Table S1, available at www.jneurosci.org as supplemental material). Then we calculated several parameters representing passive membrane properties and found that there was no significant difference between the two strains at 2 and 8 weeks. However, in mutant mice at 24 weeks, the lumped dendritic capacitance (C2) became smaller, and lumped resistance (R3) became larger than control mice (supplemental Table S1, available at www.jneurosci.org as supplemental material). These results suggest that, at 24 weeks, the average total membrane area of the mutant PC dendrites is considerably smaller than that of control. This is consistent with the morphological data that the thickness of the mutant molecular layer decreases progressively (Fig. 1G). Therefore, the slow rise and decay time constants in mutant PCs cannot be explained by the altered passive membrane properties. Together, these results collectively demonstrate that GluR2 ablation in adult PCs induces aberrant innervation of distal dendrites by both main and surplus CFs, which may lead to the generation of numerous slow EPSCs with small amplitude and the slowing of main CF-EPSC kinetics.

Figure 8. CFs with slow rise time increase after induction of GluR2 ablation. A–F, Frequency histograms showing the number of the 10–90% rise time of CF-EPSCs. At 8 and 24 weeks, CF-EPSCs are segregated into fast and slow populations in GluR2^floxFlox^CrePR mice. Rise times of main (A, C, E, white bars; B, D, F, black bars) and surplus (A–F, gray bars) CFs are overlaid. G–I, The frequency distribution of the number of CF-EPSCs with fast rise time only (<0.7 ms). Note no apparent differences between GluR2^floxFlox^ (white bars) and GluR2^floxFlox^CrePR (black bars) PCs.
Motor discoordination progresses slowly after GluR62 ablation

Finally, changes in motor performance were assessed (Fig. 9). When the footprint pattern was examined, mutant mice at 8 weeks after RU-486 administration showed no ataxic gait and could walk along a straight line. However, mutant mice at 30 weeks were unable to walk straight and took tottering steps (Fig. 9A). Such motor discoordination was not observed in RU-486-treated control mice even at 30 weeks. To evaluate more quantitatively, we performed a rotating rod test to compare the retention time on the rotating rod at 15 rpm. Control and mutant mice performed similarly at 8 weeks after RU-486 administration (Fig. 9B). However, the retention time in mutant mice gradually decreased thereafter with statistical differences being first apparent at 16 weeks and further exacerbated thereafter (Student’s t test, p < 0.01 at 16 weeks, and p < 0.001 at 24 and 28 weeks). There was a significant difference between the two genotypes (ANOVA with repeated measures, genotype effect, F(1,24) = 12.0, p = 0.002). Thus, motor discoordination is manifested slowly but progressively after RU-486 administration.

Discussion

Using a conditional mouse model of GluR62 ablation in adult PCs, we have demonstrated that GluR62 plays an essential role in maintaining the mode of CF innervation that has been established during normal cerebellar development.

GluR62 suppresses distal extension and innervation by ascending CF branch

In GluR62^{flox/CrePR} mice, relative height of CF terminals in the molecular layer started to increase at 8 weeks after RU-486 administration and further progressed until 24 weeks, when numerous VGluT2-positive CF terminals were distributed along atrophied distal dendrites (Fig. 1G,H). In horizontal cerebellar sections, this change was reflected as extended ladders of ascending CF branches along straight dendritic bars (Fig. 4). Importantly, distal extension of ascending branches occurred selectively along GluR62-ablated dendrites (Fig. 2A–E), and VGluT2-positive terminals were differentiated on them (Fig. 1E,F). Therefore, GluR62 suppresses aberrant distal extension and innervation by ascending CF branches in adult cerebellum.

This suppressive action to CF innervation should be an indirect role of GluR62. First, CF synapses in adult PCs lack GluR62 (Takayama et al., 1995, 1996; Landsend et al., 1997; Zhao et al., 1998). Second, spines on distal dendrites are innervated exclusively by PFs in adult wild-type animals (Palay and Chan-Palay, 1974; Napper and Harvey, 1988), and the loss of GluR62 generates free spines on distal dendrites (Ichikawa et al., 2002). Third, viral transfer of GluR62 to adult GluR62-knock-out mice rapidly restores PF synapse formation in transfected PCs (Kakegawa et al., 2008, 2009). These results suggest that the primary role of GluR62 is to control and consolidate the connectivity of PF–PC synapses, which then suppresses aberrant extension and innervation by CFs. The present finding eventually highlights that GluR62 actively fuels heterosynaptic competition in adult cerebellum and gives PFs a competitive advantage over CFs. This molecular function will underlie rapid reinnervation of PC spines by sprouting PFs after surgical lesion to PFs (Chen and Hillman, 1982) and may also account for proximal expansion of PF innervation after surgical denervation of CFs (Sotelo et al., 1975) or CF regression after activity blockade by TTX (Bravin et al., 1999; Morando et al., 2001; Cesa et al., 2003, 2007).

Then, the countermechanisms should be working in adult cerebellum to balance the heterosynaptic competition. In this regard, P/Q-type Ca^{2+} channel may be one of the likely candidates, because it exerts such action during cerebellar development (Miyazaki et al., 2004). Whatever the counter mechanisms, it will ensure rapid reinnervation of adult PCs by surviving CFs after subtotal lesion of the inferior olive (Rossi et al., 1991a,b).

GluR62 suppresses aberrant mediolateral wiring by transverse CF branch

GluR62 ablation also drastically changed transverse CF branches, i.e., aberrant mediolateral elongation and progressive terminal differentiation (Figs. 4, 5). Ectopic terminals on transverse branches contacted PC dendrites at variable distances from parenter ascending branches (Fig. 4F–O). Because transverse branches in wild-type rodents rarely form synapses on PCs (Rossi et al., 1991a; Sugihara et al., 1999; Nishiyama et al., 2007), our finding indicates that transverse branches retain synapse-forming ability, but this ability is potently suppressed by mechanisms, including GluR62. Because these aberrant transverse branches were preferentially distributed in the superficial molecular layer (Fig. 5E,F), where PF–PC synapses are the main synaptic constituent, and also because ectopic CF terminals did innervate distal dendrites of GluR62-ablated PCs (Figs. 1F, 6), the emergence of free spines on distal dendrites likely induce aberrant mediolateral and distal extension of CF innervation.

Mediolateral extension of CF wiring will have another impact on the cerebellar physiology. Longitudinal Olivocerebellar micro-
zones have been identified in the cerebellar cortex, based on high synchrony of complex spike activity (Llinás and Sasaki, 1989; Sasaki et al., 1989; Sugihara et al., 1993; Lang et al., 1999; Fukuda et al., 2001) and Ca$^{2+}$ spikes (Ozden et al., 2008; Mukamel et al., 2009; Schultz et al., 2009). Each microzone is ~500 μm in width, stable across behavioral states, and has sharp boundary with the neighboring microzones (Mukamel et al., 2009). This synchrony is based on electrical coupling of nearby olivary neurons through dendrodendritic gap junction (Llinás et al., 1974; Sotelo et al., 1986; Angaut and Sotelo, 1989; de Zeeuw et al., 1990) and topographical projection from given subnuclei of the inferior olive to specific longitudinal cortical zones (Sugihara et al., 1999, 2001). Axons derived from single olivary neurons project rostrocaudally to a narrow band (8 μm in width) within a single lobule or across multiple lobules, but do not project mediolaterally (Sugihara et al., 1999, 2001). Therefore, very frequent mediolateral wiring by transverse branches will broaden and even disrupt functional organization of the olivocerebellar microzone system in GluR2-ablated cerebellum. Considering that GluR2-deficient PCs produce abnormal oscillating action potentials at ~10 Hz because of enhanced CF activities and are thought to relate to involuntary spontaneous eye movement with characteristic 10 Hz oscillation (Yoshida et al., 2004), the aberrant mediolateral wiring could be one such anatomical basis for anomalous oscillatory activities and severe motor deficits in this mutant.

**GluR2 is essential to maintain CF monoinnervation**

Aberrant distal expansion and mediolateral wiring eventually produced an enormous number of surplus branches that innervated many PCs around the main target, leading to frequent occurrence of multiple innervation (Figs. 3, 6). This anatomical phenotype was in good agreement with electrophysiological data. The number of CF-EPSC steps in GluR2<sup>2flox/CrePR</sup> mice was 2 weeks after RU-486 administration was comparable with that in GluR2<sup>2flox/flox</sup> mice (Fig. 7D). However, the number of CF-EPSC steps was significantly increased at 8 weeks (Fig. 7E), and ~90% of PCs exhibited more than four steps at 24 weeks (Fig. 7F). Most surplus CF-EPSCs had small amplitude and slow kinetics (Figs. 7, 8; supplemental Fig. S5, available at www.jneurosci.org as supplemental material). These surplus CF-EPSCs with slow kinetics increased in both number and amplitude, whereas those with fast kinetics were unchanged (Fig. 8G–I; supplemental Figs. S5, S6, available at www.jneurosci.org as supplemental material). Considering that CF-EPSCs are strongly attenuated and slowed by dendritic filtering (Hashimoto et al., 2001; Roth and Hauser, 2001), slow CF-EPSCs with small amplitude likely correspond to aberrant wiring formed by ascending and transverse CF branches onto distal dendrites.

In contrast to GluR2<sup>2flox/CrePR</sup> mice, a substantial fraction of surplus CF-EPSCs in global GluR2<sup>2flox</sup>-knock-out mice displays fast as well as slow kinetics (Hashimoto et al., 2001). It is during the second postnatal week that PF synapses are formed enormously onto spines of growing PC dendrites in wild-type rodents (Woodward et al., 1971; Sotelo, 1978; Takács and Hámori, 1994), and that free spines increase drastically in global GluR2<sup>2flox</sup>-knock-out mice (Kurihara et al., 1997). Furthermore, it is also during this period that a single “winner” CF translocates to PC dendrites (Hashimoto et al., 2009). Since PF synapse activity is required to eliminate surplus CFs in normal cerebellar development (Bravin et al., 1995; Kakizawa et al., 2000), severe impairment during the period of active PF–PC synaptogenesis in global GluR2<sup>2flox</sup>-knock-out mice may allow surplus CFs with fast kinetics to survive into adulthood. Furthermore, free spines generated on growing PC dendrites might help surplus CFs to translocate from soma to proximal dendrites. By contrast, the elimination of surplus CFs and dendritic translocation of a single winner CF have completed well before the onset of GluR2<sup>2flox/CrePR</sup> mice. Therefore, unoccupied postsynaptic substrates are only available at distal dendrites in GluR2<sup>2flox/CrePR</sup> mice, resulting in more purely distal type of multiple innervation than global GluR2<sup>2flox</sup>-knock-out mice. Considering that the density of VGluT2-positive terminals on transverse branches was significantly higher in GluR2<sup>2flox/CrePR</sup> mice than global GluR2<sup>2flox</sup>-knock-out mice (supplemental Fig. S4A, available at www.jneurosci.org as supplemental material), some compensatory mechanisms might work less in adulthood than during development.

**Temporal relationship of molecular, synaptic, and behavioral deficits**

Together with our previous study using GluR2<sup>2flox/CrePR</sup> mice (Takeuchi et al., 2005), temporal relationships of molecular abnormality, synaptic wiring abnormality, and behavioral deficit can be discussed. After RU-486 administration, the percentage of PCs expressing GluR2 mRNA was reduced to one-half level at 2 weeks and 20% at 4 weeks. Cerebellar contents of GluR2 protein were reduced to one-half level at 4 weeks, 21% at 8 weeks, and <10% by 24 weeks (Takeuchi et al., 2005). Considering that RU-486 was only administered at 2 consecutive days during P42–P45, this protracted time course should represent high stability of GluR2 mRNA and protein rather than low efficiency of gene recombination.

Synaptic wiring abnormalities, including mismatched PF synapses, free spines, extended CF territory, and multiple CF innervation, all became evident by 8 weeks after RU-486 administration, when cerebellar GluR2 contents were reduced to 21% of the control level. The abnormalities of PF–PC synapses correlated inversely with synaptic GluR2 protein densities (Takeuchi et al., 2005) and the aberrant CF innervation occurred against GluR2<sup>2flox/CrePR</sup> PC dendrites. These results indicate that defects of synaptic wiring proceed in a manner highly sensitive to GluR2 loss. In comparison, motor coordination was not apparent at 8 weeks and gradually deteriorated thereafter (Fig. 9). Considering severe abnormalities in synaptic wiring at 16–24 weeks after RU-486 administration, our findings suggest that animals come to manifest cerebellar symptoms and disorders after considerable accumulation of wiring defects.

**References**


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