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FGF14 regulates presynaptic Ca²⁺ channels and synaptic transmission

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SUMMARY

Fibroblast growth factor homologous factors (FHF) are not growth factors, but instead bind to voltage-gated Na⁺ channels (Na_v) and regulate their function. Mutations in *FGF14*, a FHF that is the locus for spinocerebellar ataxia 27 (*SCA27*) are believed pathogenic because of a dominant-negative reduction of Na_v currents in cerebellar granule cells. Here, we demonstrate that FGF14 also regulates members of the presynaptic Ca_v2 Ca²⁺ channel family. Knockdown of FGF14 in granule cells reduced Ca²⁺ currents and diminished vesicular recycling, a marker for presynaptic Ca²⁺ influx. As a consequence, excitatory post-synaptic currents (EPSCs) at the granule cell to Purkinje cell synapse were markedly diminished. Expression of the *SCA27*-causing *FGF14* mutant in granule cells exerted a dominant negative reduction in Ca²⁺ currents, vesicular recycling, and the resultant EPSCs in Purkinje cells. Thus, FHF are multimodal, regulating several discrete neuronal signaling events. *SCA27* likely results at least in part from dysregulation of Ca²⁺ channel function.

Introduction

The four members of the family of fibroblast growth factor homologous factors (FHF), a subset of the fibroblast growth factors (FGF), have received increasing attention for their unanticipated modulation of voltage-gated Na⁺ (Na_v) channels and regulation of neuronal excitability. Although FHF acquired their name because of their homology to FGFs (Smallwood et al., 1996), several defining features set FHF (FGF11-FGF14) apart from other FGFs. Most notably, lacking a signal sequence, FHF are not secreted (Smallwood et al., 1996) and do not appear capable of functioning as growth factors (Olsen et al., 2003). Physiologic roles for FHF remained ill-defined until a confluence of experiments identified FHF as modulators of Na_v channels and regulators of neuronal signaling, and genetic data pinpointed *FGF14* as the locus for spinocerebellar ataxia 27 (*SCA27*).

Focus on FHF regulation of neuronal excitability began when *Fgf14*^{-/-} mice showed ataxia (Wang et al., 2002), providing a basis for exploring the implications of a linkage analysis that identified a F150S missense mutation in “b” splice variant of *FGF14* (FGF14b^{F150S};

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termed FGF14^{F145S} in some studies that used numbering based on the alternatively spliced FGF14a variant) as the etiology for the autosomal dominant SCA27 in an extended Dutch family (van Swieten et al., 2003). The specific mechanism(s) by which the human *FGF14* mutant or knockout of *Fgf14* in mouse affected neuronal signaling and led to ataxia were subsequently explained by the discovery of FGF12 as a binding partner for the C-terminus of Nav1.9 via a yeast two-hybrid strategy (Liu et al., 2001). Several studies then showed that FHF, through their interaction with Nav C-termini, can modulate Na⁺ channel currents (Liu et al., 2003; Lou et al., 2005). Moreover, the SCA27 missense mutant FGF14b^{F150S}, when expressed in cultured hippocampal neurons, decreased Nav channel currents and depressed neuronal excitability in a dominant negative manner (Laezza et al., 2007). Ataxia phenotypes have also been associated with a frame-shift mutation causing early termination of FGF14 or a chromosomal translocation that disrupted *FGF14* (Dalski et al., 2005; Misceo et al., 2009). Some experiments hint that FGF14 can regulate other neuronal processes, however, such as synaptic transmission in hippocampal neurons (Xiao et al., 2007). How FGF14 contributes to these other signaling pathways are not known, and whether mechanisms other than Na⁺ channel dysfunction contribute to the ataxia phenotype has not been examined.

Here, we focused on the role of FGF14 at the cerebellar granule cell to Purkinje cell synapse. We discovered that FGF14 in granule cells is a potent regulator of P/Q-type Cav2.1 Ca²⁺ channels, the dominant presynaptic Ca²⁺ channels in granule cells, and synaptic transmission. Further, the SCA27-causing mutant FGF14 impaired Cav channels in cerebellar neurons and affected synaptic transmission at the granule cell to Purkinje cell synapse. Thus, FGF14 is a regulator of multiple ionic currents and the pathogenic effects of mutant FGF14 are likely mediated by dysregulation of both Ca²⁺ channels and Na⁺ channels.

Results

FGF14 affects Cav2 channels in cerebellar neurons

Although FGF14 regulates Nav channel currents in granule cells (Goldfarb et al., 2007), we suspected that additional mechanisms might contribute to the SCA27 phenotype based on our observation that most SCA-associated channelopathies or the related episodic ataxias result from perturbed Ca²⁺ channel function (Shakkottai and Paulson, 2009) and with the observation that loss-of-function mutations in Cav2.1 underlie the ataxia phenotype in *tottering* mice (Fletcher et al., 1996). We therefore first asked whether FGF14 affected voltage-gated Ca²⁺ channels in granule cells within mixed primary rat cerebellar cultures containing both Purkinje cells and granule cells, which could be readily distinguished by their characteristic morphology and membrane capacitance (granule cell, 8.8 ± 0.2 pF, n=105; Purkinje cell, 15.4 ± 0.6 pF, n=50). To assess the effects of FGF14 on voltage-gated Ca²⁺ channels in granule cells, we recorded Ca²⁺ currents after FGF14 knockdown by shRNA.

A ramp protocol with 10 mM Ba²⁺ as the charge carrier, as shown in Figure 1A, revealed that FGF14 knockdown significantly reduced Ca²⁺ channel currents (summary data normalized to cell capacitance are shown in Figure 1B for FGF14 shRNA, scrambled shRNA, and a GFP transfection control). Peak inward Ba²⁺ currents were also elicited with a step protocol in which currents were evoked by a 500 ms voltage step from -80 mV to -10 mV, revealing that FGF14 knockdown significantly reduced Ca²⁺ channel current density (Figure 1C–D). We next assessed whether FGF14 affected influx through Ca²⁺ channels in response to an action potential waveform, by isolating the Cd²⁺-sensitive integrated current. We found that knockdown of endogenous FGF14 by shRNA significantly decreased the integrated current compared to transfection with GFP or scrambled shRNA (Figure 1E–F).

The efficacy of the FGF14 shRNA construct was validated in hippocampal neurons, which (because they also express the related FGF13) allowed us to test specificity. Figure S1A shows effective knockdown of the axon initial segment (AIS)-enriched FGF14 by shRNA, but not by a scrambled control. FGF13, however, was unaffected (Figure S1B). The FGF14 shRNA, but not the scrambled control, also reduced FGF14 protein expressed in HEK293T cells (Figure S1C). After confirming efficacy and specificity of the FGF14 shRNA, we transfected it (or a scrambled control shRNA or GFP) into granule cells and recorded Ca^{2+} currents.

Since 95% of the somatodendritic Ca^{2+} current in granule cells is carried by $\text{Ca}_v2.1$ P/Q-type channels (Jun et al., 1999), as well as a similar fraction of presynaptic Ca^{2+} influx at the granule cell to Purkinje cell synapse (Mintz et al., 1995), we tested whether FGF14 specifically affected $\text{Ca}_v2.1$ P/Q Ca^{2+} channel currents in a heterologous expression system. The $\text{Ca}_v2.1$ pore-forming α_{1A} subunit along with the accessory β_2b and $\alpha_2\delta$ subunits were expressed in HEK293T cells and currents were evoked by step depolarizations with 10 mM Ba^{2+} as charge carrier (Figure 2A). The current-voltage (I-V) relationship shows that co-expression of FGF14b increased current density over a broad range of voltages (Figure 2B), without affecting kinetics of activation or steady-state inactivation (Figure S2A–B, Table 1). The integrated inward current through $\text{Ca}_v2.1$ in response to an action potential waveform also increased with FGF14 co-expression (Figure 2C–D). Co-expression of FGF14b also increased current density of the other major presynaptic Ca^{2+} channel, the $\text{Ca}_v2.2$ N-type channel (Figure 2E). In contrast, FGF14b did not affect the $\text{Ca}_v1.2$ L-type or $\text{Ca}_v2.3$ R-type channels, which are predominantly somatodendritic in location (Figure 2F–G). Together, these results showed that endogenous FGF14 affected granule cell Ca^{2+} channels and that FGF14 was capable of regulating the presynaptic $\text{Ca}_v2.1$ and $\text{Ca}_v2.2$ Ca^{2+} channels.

Because FGF14 did not affect $\text{Ca}_v2.1$ kinetics of activation or steady-state inactivation, we suspected that a major mechanism by which FGF14 increased current density was by increasing the number of channels at the plasma membrane. In HEK293T cells expressing $\text{Ca}_v2.1$ channels, we therefore measured the gating charge, with and without FGF14 co-expression, as a means to assess the number of channels at the cell surface. We isolated the gating charge by depolarizing the cell from a holding potential of -80 mV to the reversal potential (determined individually for each cell, average $+43.2$ mV \pm 0.9 mV, $n=27$) and eliminated any remaining ionic current by blocking channels with Cd^{2+} (300 μM). Figure 3A–B shows that co-expression of FGF14b increased gating charge by $\sim 35\%$ compared to the GFP-only control, suggesting that FGF14b does increase the number of channels at the plasma membrane. This increase in gating charge could be attributed solely to the transfected $\text{Ca}_v2.1$ channels, and not to FGF14b effects upon endogenous ionic currents in HEK293T cells because transfection of FGF14b in the absence of $\text{Ca}_v2.1$ did not increase gating charge compared to cells transfected with GFP only (2.2 ± 0.2 fC/pF, $n=12$; and 2.0 ± 0.3 fC/pF, $n=10$, respectively; $p>0.05$).

Presynaptic FGF14 regulates baseline transmission at the granule cell to Purkinje-cell synapse

Having established that FGF14 affects $\text{Ca}_v2.1$ channels, the predominant presynaptic Ca^{2+} channels at the granule cell to Purkinje cell synapse and since FGF14 is abundant in granule cell axons (Wang et al., 2002), we explored whether FGF14 affected synaptic transmission at the granule cell to Purkinje cell synapse. We used paired recordings to identify a granule cell to Purkinje cell synapse in which only the presynaptic granule cell was transfected with FGF14 shRNA, a scrambled control shRNA or GFP (Figure 4A). We then evoked unitary excitatory post-synaptic currents (EPSCs) in the presence of 20 μM (–) bicuculline. Knockdown of FGF14 in granule cells reduced the EPSC amplitude in Purkinje cells by more than 80% compared to EPSCs recorded from pairs in which the granule cell was

transfected with GFP only or with the control scrambled shRNA (Figure 4B–C). Because FGF14 is also present in Purkinje cell somata (Shakkottai et al., 2009), we also checked whether FGF14 exerted postsynaptic effects in Purkinje cells by examining glutamate- and GABA-evoked responses after knocking down endogenous FGF14 by shRNA. The glutamate- and GABA-evoked currents were obtained by directly application of drugs to Purkinje cells. We found that neither glutamate-nor GABA-evoked inward currents were altered compared to currents from Purkinje cells transfected with a scrambled control shRNA or GFP only (Figure S3A–B), suggesting FGF14 had no effect on post synaptic responses.

FGF14 regulates vesicular turnover and short-term synaptic plasticity

We hypothesized that FGF14 knockdown in granule cells reduced EPSCs at the granule cell to Purkinje cell synapse because of a diminished presynaptic Ca^{2+} current through $\text{Ca}_v2.1$ channels, consistent with the reduced $\text{Ca}_v2.1$ Ca^{2+} current seen in Figure 2. We therefore tested whether FGF14 might affect presynaptic Ca^{2+} influx in granule cells. We assessed this indirectly by two different means. First, we measured synaptic vesicular turnover after a 90 sec depolarization with 90 mM KCl. We measured turnover of synaptic vesicles with FM 4–64. The styryl FM dyes become trapped in vesicles that underwent endocytosis following synaptic activity (Ryan et al., 1993), providing a measure of vesicular turnover that is closely correlated with the amount of presynaptic Ca^{2+} influx (Evans and Cousin, 2007; Yamashita, 2012). We transfected the mixed cultures with shRNA to knockdown FGF14 or a scrambled control shRNA and quantified FM 4–64 uptake in granule cells after a 90 sec depolarization by 90 mM KCl. The styryl dye was included only during the 90 sec depolarization, after which the neurons were immediately washed then fixed. In these sparsely transfected cultures, we then followed a GFP-positive granule cell axon until it synapsed with an untransfected Purkinje cell and then counted the number of puncta positive for FM 4–64 and GFP fluorescence (Figure 5A). In all cases, quantification was performed with experimenter blinded to the identity of the transfection. Knockdown of FGF14 reduced the number of recycled punctae by 54 % compared to scrambled control shRNA (Figure 5B–C). Second, we assessed short term plasticity. Pre-synaptic Ca^{2+} influx plays an essential role in neurotransmitter release in central nervous system synapses, and also contributes to short-term synaptic plasticity. A reduction in presynaptic Ca^{2+} influx is predicted to increase the paired pulse ratio (PPR) in response to two closely spaced stimuli (Zucker and Regehr, 2002). In the presence of 20 μM (–) bicuculline, we measured the PPR in response to two stimuli at 10 Hz (Figure 5D–E) and found that when the granule cell expressed GFP only or the scrambled control shRNA, the PPR was near unity (1.03 ± 0.04 , $n=16$; and 1.05 ± 0.06 , $n=10$, respectively). Knockdown of FGF14, however, significantly increased the PPR (1.45 ± 0.11 , $n=27$), consistent with an effect on presynaptic Ca^{2+} entry. Together with the reduction in vesicular turnover after FGF14 knockdown, these data provided corroborating evidence that presynaptic Ca^{2+} influx was reduced by FGF14 knockdown.

The SCA27-causing FGF14 mutant affects synaptic transmission in a dominant negative manner

We next addressed the effects of the FGF14b^{F150S} mutant associated with SCA27. We chose the “b” splice variant for study, since it is the most abundantly expressed in brain (Wang et al., 2000). First, we examined whether FGF14b^{F150S} affected voltage-gated Ca^{2+} channel currents (with 10 mM Ba^{2+} as a charge carrier) by step or ramp depolarization in granule cells. As seen in Figure 6A–D, transfection of FGF14b^{F150S} significantly reduced Ca^{2+} channel current density. This decrease in Ca^{2+} current was also observed in response to an action potential waveform (Figure 6E–F). In contrast, overexpression of FGF14b^{WT} markedly increased Ca^{2+} currents in response to a step depolarization, ramp protocol, or an

action potential waveform. Since FGF14 increased Ca^{2+} currents above control, these data suggest that the effects of endogenous FGF14 are not saturated (Figure 6A–F).

We next examined whether the FGF14b^{F150S} mutant affected presynaptic Ca^{2+} influx in granule cells by using FM 4–64 labeled vesicular recycling as an indicator. Compared to control, the SCA27 mutant reduced vesicular recycling (Figure 7A–C). In contrast, overexpression of FGF14b^{WT} increased vesicular recycling (Figure 7A–C), consistent with the observed effects on total Ca^{2+} current. Finally, we used paired recordings to measure EPSCs in Purkinje cells after stimulation of a granule cell transfected with FGF14b^{F150S}, allowing us to measure directly the effect of FGF14b^{F150S} on synaptic transmission. As seen in Figure 7D–E, expression of FGF14b^{F150S} in a granule cell exerted a dominant negative effect and markedly reduced the EPSCs amplitude in Purkinje cell by 63 %. Overexpression of FGF14b^{WT}, in contrast, markedly increased the EPSCs by 317 %, which also suggests that the effects of endogenous FGF14 were not saturating.

Discussion

Understanding of the physiological roles of FIEFs has greatly evolved since their initial discovery (Smallwood et al., 1996). Recognition of FIEFs as Na_V channel binding partners and modulators, demonstration that *Fgf14*^{-/-} mice displayed ataxia, identification of *FGF14* as the locus for SCA27, and demonstration that FIEFs cannot activate FGF receptors redirected most attention to the influence of FIEFs intracellularly and specifically focused attention on their roles in neuronal excitability (Liu et al., 2001; Liu et al., 2003; Olsen et al., 2003; van Swieten et al., 2003; Wang et al., 2002). Further focus on modulation of Na_V channels followed the demonstration that the SCA27 mutant version of FGF14 acted as a dominant negative to suppress Na_V currents and excitability in hippocampal neurons (Laezza et al., 2007) and that granule cells from *Fgf14*^{-/-};*Fgf12*^{-/-} mice displayed a deficit in intrinsic excitability and altered Na_V channel inactivation properties (Goldfarb et al., 2007). The ataxia phenotype common to *Fgf14*^{-/-} mice and patients with the dominant negative FGF14b^{F150S} mutation has thus been suspected to result from Na_V channel dysfunction in granule cells or Purkinje cells (Goldfarb et al., 2007; Shakkottai et al., 2009).

Nevertheless, several observations suggest that FIEFs might possess capabilities beyond Na_V channel regulation. For example, CA1 synapses in *Fgf14*^{-/-} mice have fewer total and docked vesicles (Xiao et al., 2007) and *Fgf13* knockdown in *Xenopus* oocytes influences neuronal development by affecting bone morphogenetic protein receptor activation of the MEK5-ERK5 pathway (Nishimoto and Nishida, 2007). Moreover, certain FIEFs localize to the nucleus (Munoz-Sanjuan et al., 2000), a subcellular location in which FIEFs are unlikely to influence Na_V channels, and FIEFs appear to have binding partners other than Na_V channels, such as kinase scaffolds (Schoorlemmer and Goldfarb, 2001). In this context, our demonstration that FGF14 regulates $\text{Ca}_V2.1$ channel currents and synaptic transmission at the granule cell to Purkinje cell synapse adds an additional dimension to FIEF function. How FGF14 affects these processes is unclear, but we suspect that mechanism differs from how FIEFs affect Na_V channels, for which there is direct binding with the Na_V C terminus (Wang et al., 2012). In contrast, we were unable to detect direct interactions between FGF14 and Ca_V2 channels by either co-immunoprecipitation or recombinant protein binding studies targeting intracellular domains of Ca_V2 channels or their auxiliary subunits. Consistent with our observations, FGF14 was not annotated as a component of the Ca_V2 -anchored proteome in a recent analysis (Muller et al., 2010). Thus, we hypothesize that the FGF14 acts indirectly, or possibly transiently, to increase the number of Ca_V2 channels at the plasma membrane.

Importantly, our data provide a novel mechanism by which FGF14 could affect synaptic transmission at a granule cell to Purkinje cell synapse, thereby offering new insight into the etiology of the disease phenotypes in humans with SCA27 and the ataxia phenotype in *Fgf14*^{-/-} mice. As we showed, the mutant FGF14b^{F150S} acts as a dominant negative, reducing Ca²⁺ channel currents in granule cells similar to the effect of shRNA knockdown of endogenous FGF14. The dominant-negative effect of FGF14b^{F150S} on Ca²⁺ currents in granule cells and the consequent reduction in EPSCs in Purkinje cells are both consistent with the observation that the mutant also acts as a dominant-negative to reduce Na⁺ channel currents in hippocampal neurons (Laezza et al., 2007). Our data measuring gating charge point to a potential role for FGF14 in trafficking Ca²⁺ channels to, or regulating removal from, the plasma membrane. As such, FHF's may have a broader role in trafficking ion channels, since we recently demonstrated that endogenous FGF13 increases the number of cell surface Na_v1.5 Na⁺ channels in cardiomyocytes (Wang et al., 2011). On the other hand, our data also underline the concept that individual FHF's confer channel-specific regulatory effects. For example, although FGF14 regulated Ca_v2.1 and Ca_v2.2, the major presynaptic Ca²⁺ channels, FGF14 did not affect currents through Ca_v1.2 or Ca_v2.3, which are mainly restricted to the somatodendritic compartments. Such data argue against the possibility that FGF14 mediates its effects via interaction with a Ca²⁺ channel auxiliary protein, which would be common to all Ca²⁺ channels, but could suggest that FGF14 controls a regulator, such as a kinase, that has effects specific to individual types of Ca²⁺ channels.

The FGF14b^{F150S}-induced reduction in Ca²⁺ channel currents fits well with the observation that most SCA-associated channelopathies or the related episodic ataxias result from perturbed Ca²⁺ channel function (Shakkottai and Paulson, 2009) and with the observation that loss-of-function mutations in Ca_v2.1 underlie the ataxia phenotype in *tottering* mice (Fletcher et al., 1996). Thus, the previously identified actions of FGF14 on granule cell Na_v currents (Goldfarb et al., 2007), together with its effects upon granule cell Ca_v currents and the consequences for Purkinje cell EPSCs described herein, suggest that altered Purkinje cell output from the cerebellum in *FGF14* loss-of-function or dominant negative mutations derives from multiple mechanisms. Along with the previously demonstrated FGF14b^{F150S}-dependent reduction in Na_v currents and channel availability (Laezza et al., 2007), these data suggest that FGF14 is truly multimodal and the ataxia phenotype caused by mutant FGF14 results from several independent mechanisms.

Experimental Procedures

Molecular Biology

Mouse FGF14b or FGF14b^{F150S} were cloned into pIRES2-AcGFP1. The F150S mutation in FGF14b was generated with QuickChange (Agilent). Constructs were sequenced in both directions. A cDNA for Ca_v2.1 (Kraus et al., 1998) was kindly provided by A. Lee (U. Iowa) by permission of J. Striessnig (U. Innsbruck). A cDNA for Ca_v2.3 (Bannister et al., 2004) was kindly provided by Brett Adams (Utah State). The cDNAs for Ca_v1.2, β_{2b} and α_{2δ} were previously described (Wang et al., 2007). Hairpins targeted to FGF14 were designed with Invitrogen's RNAi Designer. The sequences were synthesized via Integrated DNA Technologies and subsequently cloned into pLVTHM (Addgene). Neurons were transfected with the different constructs followed by immunocytochemical staining to determine the efficacy and specificity of knockdown. The most effective shRNA has the sequence 5'

CGCGTGGAGGCAAACAGTCAACAAGTGCATTCAAGAGATGCACTTGTTGACTG
GTTTGC CTCCTTTTTTAT - 3' and was used for the experiments described in this work. A scrambled shRNA that exhibits no significant homology to genes in rodent genomes was used as a control. This scrambled shRNA has been previously described (Wang et al., 2011).

Primary Cerebellar Culture and transfection

Primary dissociated cerebellar cultures were prepared using minor modifications of a previously described procedure for preparation of hippocampal cultures (Wang et al., 2007). Briefly, the cerebellum cortex was dissected on ice from P0-P1 male or female Wistar rat pups, digested with 0.25% trypsin for 10 min at 37°C with Dulbecco's Modified Eagle's Medium (DMEM, Sigma), and dissociated into single cells by gentle trituration. The cells were seeded onto coverslips coated with 50 µg/ml poly-D-lysine (Sigma) and 25 µg/ml laminin (Sigma) at a density of $2.5\text{--}3.0 \times 10^5$ cells/coverslip (12 × 12 mm coverslip) in DMEM supplemented with 10% heat-inactivated fetal bovine serum (FBS). The cells were maintained in a humidified incubator in 5% CO₂ at 37°C. After 15–16h, the medium was replaced with Basal Medium Eagle (BME, Sigma) supplemented with 2% B27 (Invitrogen), 1% or 5% FBS, 25 µM uridine, 70 µM 5-fluorodeoxyuridine, and 20 mM KCl. After 5–7 days *in vitro* (DIV) culture, the neurons were transiently transfected with 1 µg plasmid DNA per coverslip with calcium phosphate, as described previously (Wang et al., 2007). Experiments were carried out 7–12 days after transfection.

HEK 293T Cell Culture and Transfection

Human embryonic kidney (HEK) 293T cells were maintained in DMEM containing 10% FBS at 37°C in a 5% CO₂ incubator. The cells were plated in 60-mm tissue culture dishes, grown to 65–75% confluency and transfected with Lipofectamine 2000 (Invitrogen) in serum-reduced medium (Opti-MEM; Invitrogen) following the manufacturer's instructions. The total amount of cDNA used per dish was 8 µg, which included 3 µg of Ca_v2.1, Ca_v2.2, Ca_v2.3, or Ca_v1.2 subunits (α_{1A} , α_{1B} , α_{1E} , or α_{1C} , respectively); 2.2 µg of β_2b ; 1.8 µg of $\alpha_2\delta$; 2 µg of the empty pIRES2-acGFP1 vector or FGF14b in 5 ml transfection medium. After 24h transfection, the cells were re-plated on coverslips coated with 50 µg/ml poly-D-lysine (Sigma) at a low density for recording.

Electrophysiological Recordings

Whole cell voltage-clamp recordings were obtained from cultured cerebellar granule cells and Purkinje cells 7–12 days after transfection. The granule cells and Purkinje cells were identified based on their size and morphology. For paired recording experiments, whole cell recordings used an Axopatch 200A and 200B amplifier (Axon Instruments, Inc., Union City, CA), the signal was filtered at 2 kHz to 5 kHz bandwidth. The data acquisition was performed using a DigiData 1322A (Axon Instruments, Inc) digitizer and stored on a personal computer running pClamp software, version 10. For EPSCs recording, patch pipettes with 5–6 MΩ resistances were filled with internal solution containing (in mM): 120 K-gluconate, 10 KCl, 5 MgCl₂, 0.6 EGTA, 5 HEPES, 10 phosphocreatine, 50 U/ml creatine-phosphokinase, 2 Mg-ATP, and 0.2 GTP, pH 7.3 with KOH (290–300 mOsm). The external solution contained (in mM): 135 NaCl, 2 CaCl₂, 1 MgCl₂, 5 KCl, 10 glucose, and 5 HEPES, pH 7.3 with NaOH (300–310 mOsm). To block GABA_A receptor activity, (–) bicuculline 20 µM was added to external solution. A brief 5 mV hyperpolarizing step was performed at the end of each sweep to monitor series resistance, capacitance, and input (leak) resistance throughout the experiment. Cells were rejected from analysis if the series resistance changed by >15–20%. The neuronal Ca²⁺ currents were recorded using an EPC 10 USB patch amplifier (HEKA Elektronik, Lambrecht/Pfalz, Germany). The signal was filtered at 2.9 Hz and digitized at 20 Hz. To record currents through voltage-gated Ca²⁺ channels, we used a bath solution containing (in mM): 124 NaCl, 20 TEA-Cl, 1 MgCl₂, 10 BaCl₂, 5 HEPES, and 10 glucose, pH 7.3 with NaOH (300–310 mOsm). The internal solution contained (in mM): 115 CsCl, 20 TEA-Cl, 1 CaCl₂, 2 MgCl₂, 10 HEPES, and 2 Mg-ATP, pH 7.3 with CsOH (290–300 mOsm). To block Na⁺ currents, 1 µM Tetrodotoxin (TTX) was supplemented in external solution.

Whole cell voltage-clamp recordings were obtained from HEK293T cells at room temperature 2–3 days after transfection using external and internal solutions as described above. The liquid junction potential and series resistance for these recordings were not corrected, and cells were discarded if series resistance was more than 10 M Ω .

Protocols and data analysis

Data analysis was performed using PatchMaster, FitMaster, and Clampfit 10.2 software. All averaged data presented the mean \pm SEM. Statistical significance was determined using Student's *t*- or one-way ANOVA tests. In whole-cell voltage-clamp mode, the unitary EPSCs were obtained by stimulating a transfected presynaptic granule cell and recording from a neighboring untransfected postsynaptic Purkinje cell 7–12 days after transfection (Figure 4A). To elicit unitary EPSCs, the two cells were both held at -70 mV membrane potential, and a 20 ms depolarization pulse was delivered to the granule cell. The EPSC amplitude was determined by an average of 5–10 EPSCs from each cell. The neuronal Ca²⁺ current was obtained by 500 ms step depolarization from a holding potential of -80 mV to -10 mV or by ramp from a holding potential of -80 mV to $+50$ mV in 1 sec. The current amplitude was normalized to each cell's capacitance.

Recombinant HEK293T cells were voltage-clamped at a holding potential (V_h) of -80 mV, and Ca_v2.1, Ca_v2.2, Ca_v2.3, or Ca_v1.2 current I_{Ca} was elicited by depolarizing pulses of 300 ms from -80 mV to $+60$ mV (in 10-mV increments). Peak I_{Ca} amplitude during the test pulse was divided by the corresponding cell capacitance to obtain a measure of current density (pA/pF). Current density-to-voltage relationships were plotted (I - V relationship). Ca_v2.1 currents were also evoked by an action potential waveform command. The action potential waveform (APW) was modified from a previously reported study (Borst et al., 1995). In brief, the APW began at -80 mV and peaked at $+33$ mV with a 2-ms half-amplitude duration. The maximal rising and falling slopes were $+127$ V/s and -52 V/s, respectively. Ca²⁺ channel currents were defined as the CdCl₂ (300 μ M) sensitive fraction. Gating charge was obtained by depolarization for holding potential (-80 mV) to the reversal potential (40 – 50 mV, determined individually for each cell) for 20 ms.

FM dye staining was performed on cultured cerebellar neurons. On day 6 (DIV), the neurons were transfected with pIRES2-AcGFP1 (Control), scrambled control shRNA, FGF14 shRNA, FGF14b^{WT}, or FGF14b^{F150S}. FM4-64 (Molecular Probes) dye loading was performed 9 days after transfection (15 DIV). The neurons were first washed in HBS (in mM) 139 NaCl, 2.5 KCl, 10 HEPES, 10 glucose, 2 CaCl₂, and 1.3 MgCl₂, pH 7.3 (300 mOsm), then depolarized for 90 s at room temperature with 90 mM KCl solution (in mM: 48.5 NaCl, 90 KCl, 10 glucose, 10 HEPES, 2 CaCl₂, 1.3 MgCl₂, 0.05 APV, and 0.02 DNQX) containing 15 μ M FM4-64. The neurons were immediately washed for 2 min with a calcium-free HBS solution containing 0.1 mM Advasep-7 (Sigma) and then rinsed 3 times with HBS solution to remove all nonspecific membrane bound FM4-64. Finally, the cells were fixed for 10 min with 4% paraformaldehyde/4% sucrose.

Imaging was performed with a Zeiss LSM 510 confocal microscope using an oil immersion 40X objective. GFP and FM4-64 dye were excited at 488 and 543 nm, respectively. All images were collected at 1024 \times 1024 pixel resolution. Nerve terminals of transfected neurons were identified by tracing the GFP-positive axons. Synaptic punctae incorporation along a traced axonal process was identified at a region away from the cell body. For quantification, the experimenter was blinded to the identity of the transfected plasmid. A 45 μ m \times 45 μ m square region of interest was selected and the numbers of puncta were calculated using NIH ImageJ software. Particles <0.3 μ m in diameter were excluded from analyses.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

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Highlights

FGF14 regulates presynaptic CaV2.1 at the granule cell to Purkinje cell synapse

FGF14 regulates presynaptic Ca²⁺ influx at the granule cell to Purkinje cell synapse

FGF14 regulates synaptic transmission at the granule cell to Purkinje cell synapse

Mutant FGF14 affects all these processes adversely

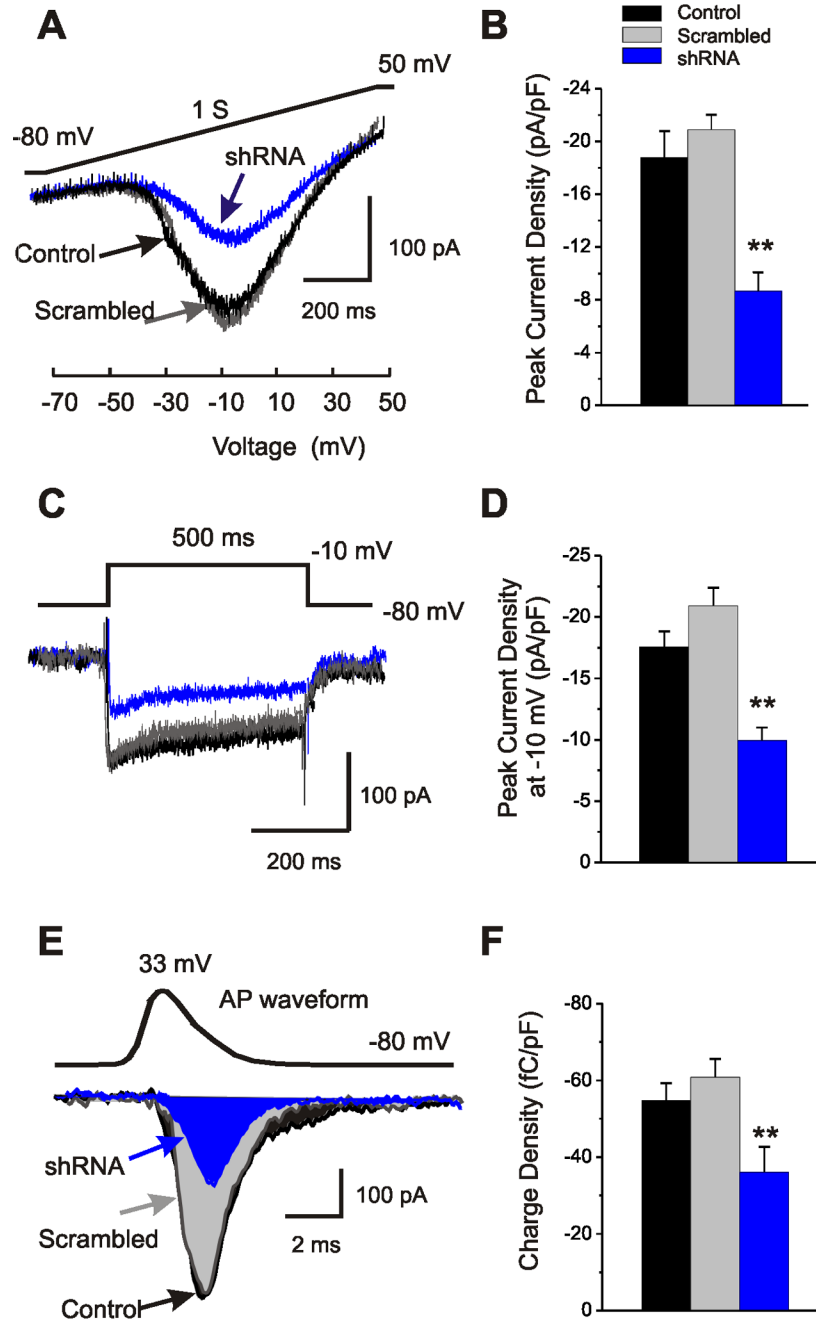


Figure 1. Endogenous FGF14 regulates Ca²⁺ channel currents in granule cells

A. Example Ca²⁺ channel current traces (using Ba²⁺ as the charge carrier) recorded from a cerebellar granule cell transfected with GFP-control (black), scrambled control shRNA (gray), or FGF14 shRNA (blue). The currents were evoked by a ramp protocol from a holding potential of -80 mV to 50 mV in 1 sec. **B.** Summary data from granule cells expressing GFP control (n=17), scrambled control shRNA (n=10), or FGF14 shRNA (n=12). **C.** Example Ca²⁺ channel current traces recorded from a cerebellar granule cell transfected with GFP-control (black), scrambled control shRNA (gray), or FGF14 shRNA (blue). The currents were evoked by a step protocol from a holding potential of -80 mV to

–10 mV in 500 ms. **D.** Summary data from granule cells expressing GFP control (n=22), scrambled control shRNA (n=11), or FGF14 shRNA (n=10). **E.** Representative Cd²⁺-sensitive Ba²⁺ currents evoked by a single action potential waveform (APW, top) command recorded from granule cells transfected with GFP control (black), scrambled control shRNA (gray), or FGF14 shRNA (blue). The integrated current (Q) is colored with black, gray, or blue. **F.** Summary data of the integrated current (Q) normalized to each cell capacitance. Summary results were obtained from granule cells expressing GFP control (n=25), scrambled control shRNA (n=11), or FGF14 shRNA (n=10). **p<0.01 versus Control.

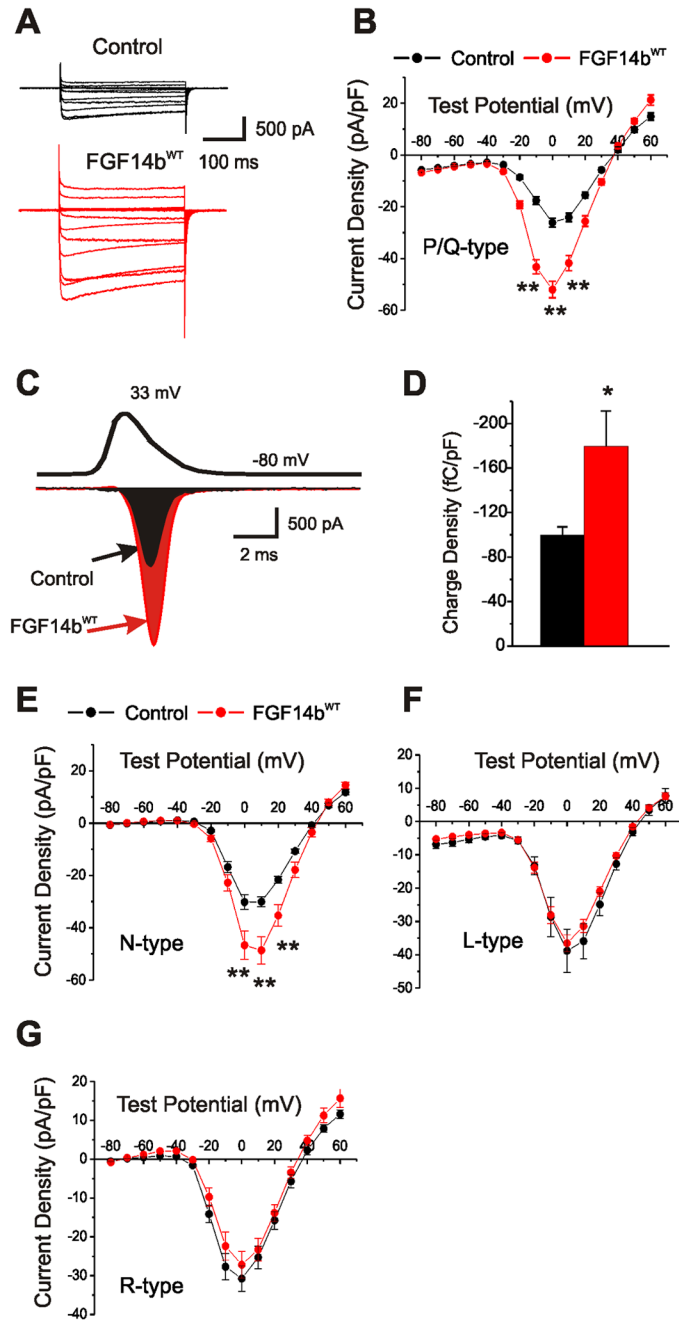


Figure 2. FGF14 modulates Ca_v2.1 and Ca_v2.2 channels

A. Example Ca²⁺ channel current traces (using Ba²⁺ as the charge carrier) recorded from HEK293T cells in which Ca_v2.1 channels were co-expressed with GFP control (black) or FGF14^{WT} (red). The currents were elicited by depolarizing pulses of 300 ms from -80 mV to +60 mV (in 10-mV increments). **B.** Current-voltage relationships (normalized to cell capacitance) for cells in which Ca_v2.1 was co-transfected with GFP control (black) or FGF14^{WT} (red). **C.** Representative Cd²⁺-sensitive Ba²⁺ currents evoked by a single action potential waveform (APW, top) command from HEK293T cells in which Ca_v2.1 channels were co-transfected with GFP control (black) or FGF14^{WT} (red). The integrated current (Q)

is colored with black or red. **D.** Summary data of the integrated current (Q) normalized to cell capacitance for each cell expressing GFP (n=9) or FGF14^{WT} (n=9). **E-G.** Current-voltage relationships (normalized to cell capacitance) for cells in which Ca_v2.2 (E), Ca_v1.2 (F) or Ca_v2.3 (G) were co-transfected with GFP (black) or FGF14^{WT} (red). The current amplitude values were divided by the capacitance of each cell to obtain current density (pA/pF). *p<0.05, **p<0.01 versus Control.

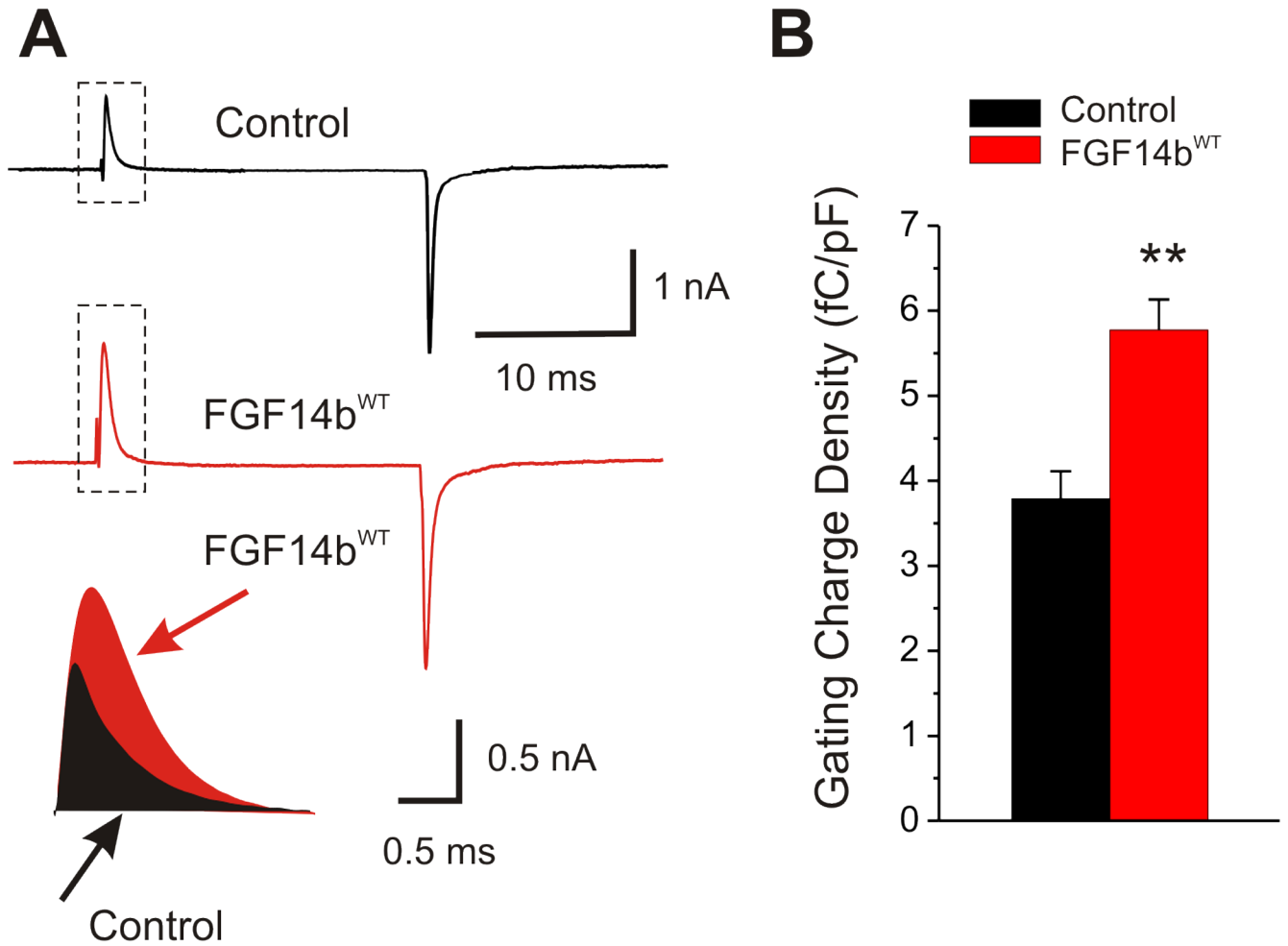


Figure 3. FGF14 increases Ca²⁺ gating charge in Ca_v2.1 channels

A. Example Ca²⁺ channel current traces (using Ba²⁺ as the charge carrier) recorded from HEK293T cells in which Ca_v2.1 channels were co-expressed with GFP control (black) or FGF14^{WT} (red). The example traces were elicited by a 20 ms step from the holding potential (−80 mV) to the reversal potential. Magnification of the integrated gating currents is shown below. **B.** Summary data of the integrated current (Q) normalized to each cell capacitance for cells expressing Ca_v2.1 or GFP control (n=14) or FGF14^{WT} (n=13). **p<0.01 versus Control.

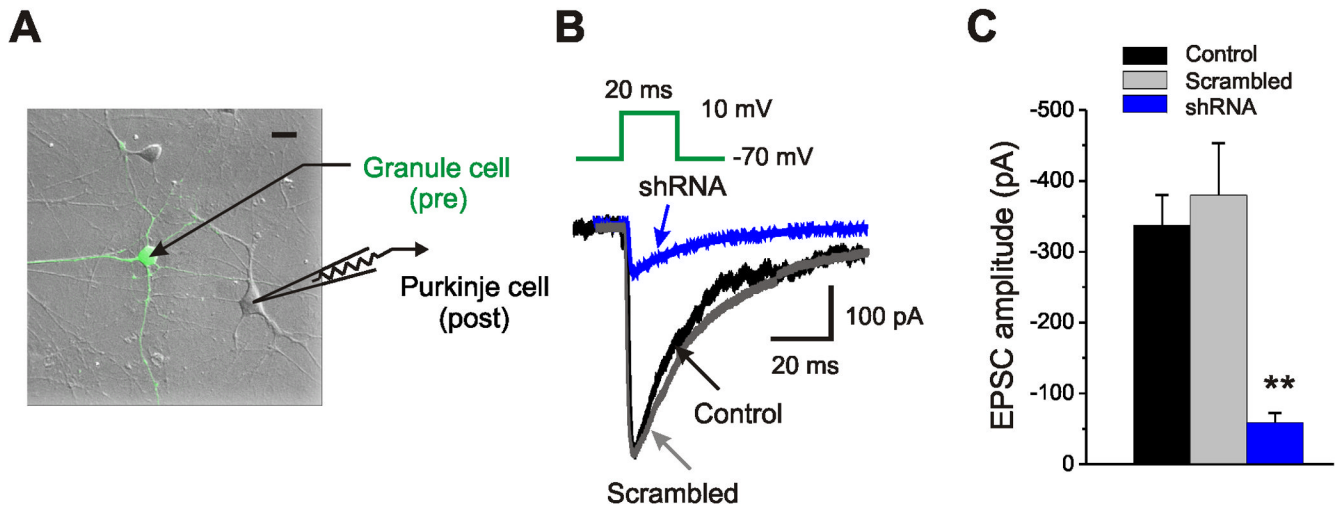


Figure 4. Endogenous FGF14 regulates synaptic transmission at the granule cell to Purkinje cell synapse

A. Evoked EPSCs in an untransfected Purkinje cell were elicited by a 20 ms depolarization of a transfected granule cell from a holding potential of -70 mV to -10 mV. **B.** Representative EPSC traces recorded from Purkinje cells in which the presynaptic granule cell was transfected with GFP control (black), scrambled control shRNA (gray), or FGF14 shRNA (blue). **C.** Averaged amplitude of unitary Purkinje cell EPSCs when the presynaptic granule cells expressed GFP control ($n=14$), scrambled control shRNA ($n=25$), or FGF14 shRNA ($n=20$). $**p<0.01$ versus Control.

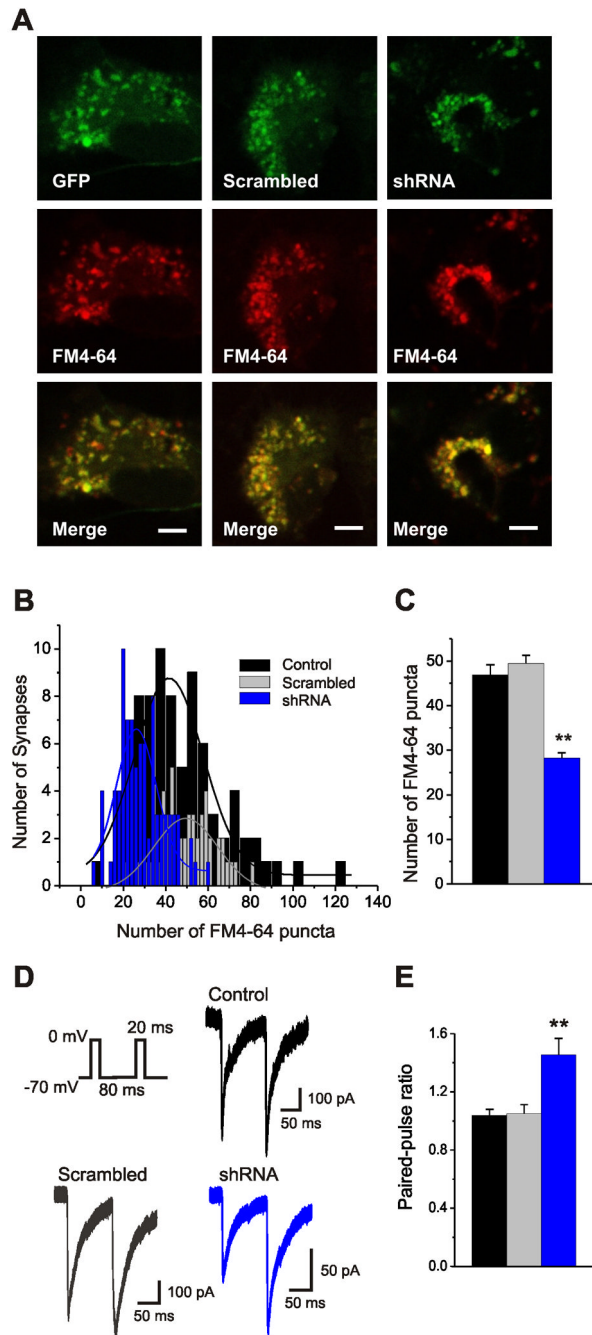


Figure 5. Endogenous FGF14 regulates vesicular recycling and short-term synaptic plasticity

A. Confocal images from cultured cerebellar neurons expressing GFP (left), scrambled control shRNA (middle), or FGF14 shRNA (right) that were loaded with FM4-64 by a 90 sec depolarization using 90 mM KCl. Scale bar, 5 μ m. **B.** The distribution of FM4-64 puncta per synapse within a $45 \times 45 \mu\text{m}^2$ region of interest (ROI) in neurons transfected with control GFP (black, n=81), scrambled control shRNA (gray, n=49), or FGF14 shRNA (blue, n=87). The data for each group were fit to a Gaussian distribution. **C.** Average puncta per $45 \times 45 \mu\text{m}^2$ ROI for neurons over-expressing control GFP, scrambled control shRNA, or FGF14 shRNA. **D.** Representative EPSCs evoked by a paired-pulse protocol (indicated in inset; 80

ms interstimulus interval) when the presynaptic granule cells expressed GFP control (black), scrambled control shRNA (gray), or FGF14 shRNA (blue). **E.** Averaged paired-pulse ratio (amplitude of the second EPSC divided by the amplitude of the first EPSC). ** $p < 0.01$ versus Control.

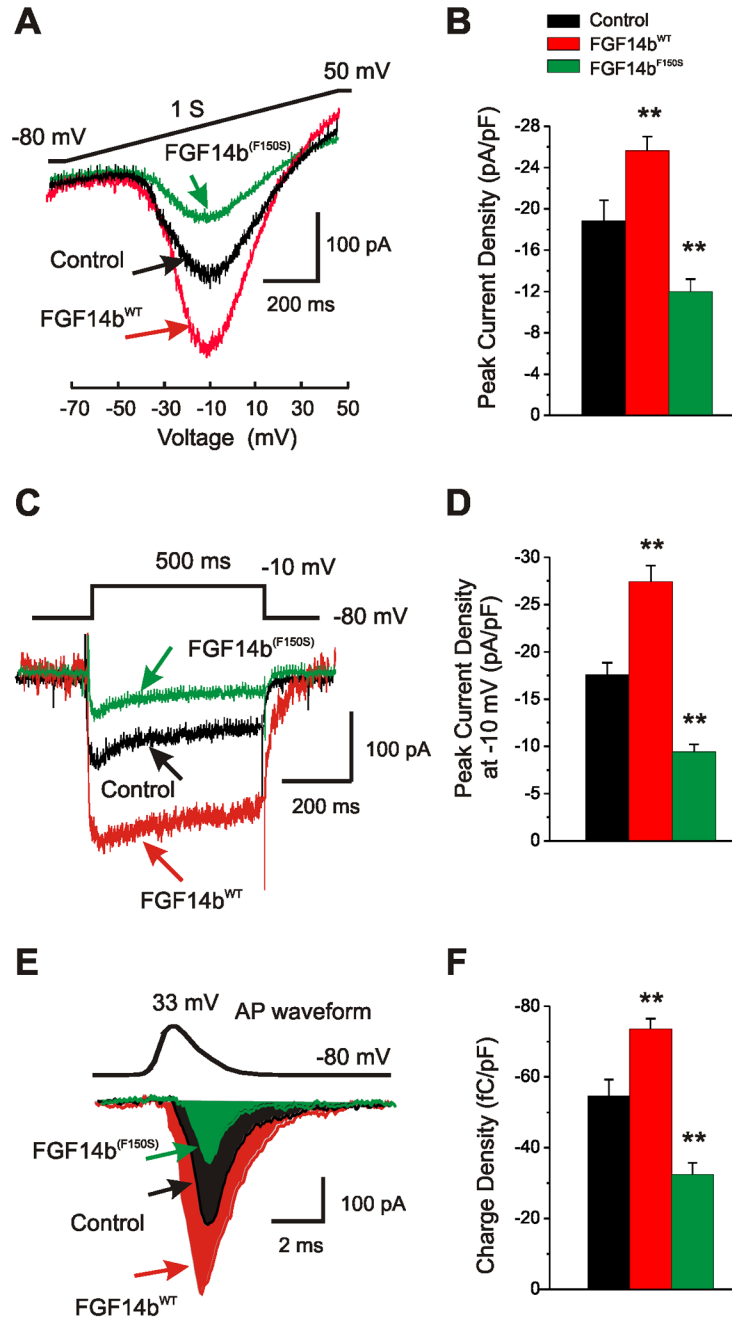


Figure 6. The SCA27 FGF14 mutant reduces granule Ca²⁺ currents

A. Example Ca²⁺ current traces (using Ba²⁺ as the charge carrier) recorded from a cerebellar granule cell transfected with GFP-control (black), FGF14^{WT} (red), or FGF14b^{F150S} (green). The currents were evoked by a ramp protocol from a holding potential of -80 mV to 50 mV in 1 sec. **B.** Summary data from granule cells expressing GFP control (n=17), FGF14^{WT} (n=25), or FGF14b^{F150S} (n=16). **C.** Example Ca²⁺ channel current traces recorded from a cerebellar granule cell transfected with GFP-control (black), FGF14^{WT} (red), or FGF14b^{F150S} (green). The currents were evoked by a step protocol from a holding potential of -80 mV to -10 mV in 500 ms. **D.** Summary data from granule cells expressing GFP

control (n=22), FGF14^{WT} (n=37), or FGF14b^{F150S} (n=13). **E.** Representative Cd²⁺-sensitive Ba²⁺ currents evoked by a single action potential waveform (APW, top) command recorded from granule cells transfected with GFP control (black), scrambled control shRNA (gray), or FGF14 shRNA (blue). The integrated current (Q) is colored with black, red, or green. **F.** Summary data of the integrated current (Q) normalized to each cell capacitance. Summary results were obtained from granule cells expressing GFP control (n=25), FGF14^{WT} (n=28), or FGF14b^{F150S} (n=16). **p<0.01 versus Control.

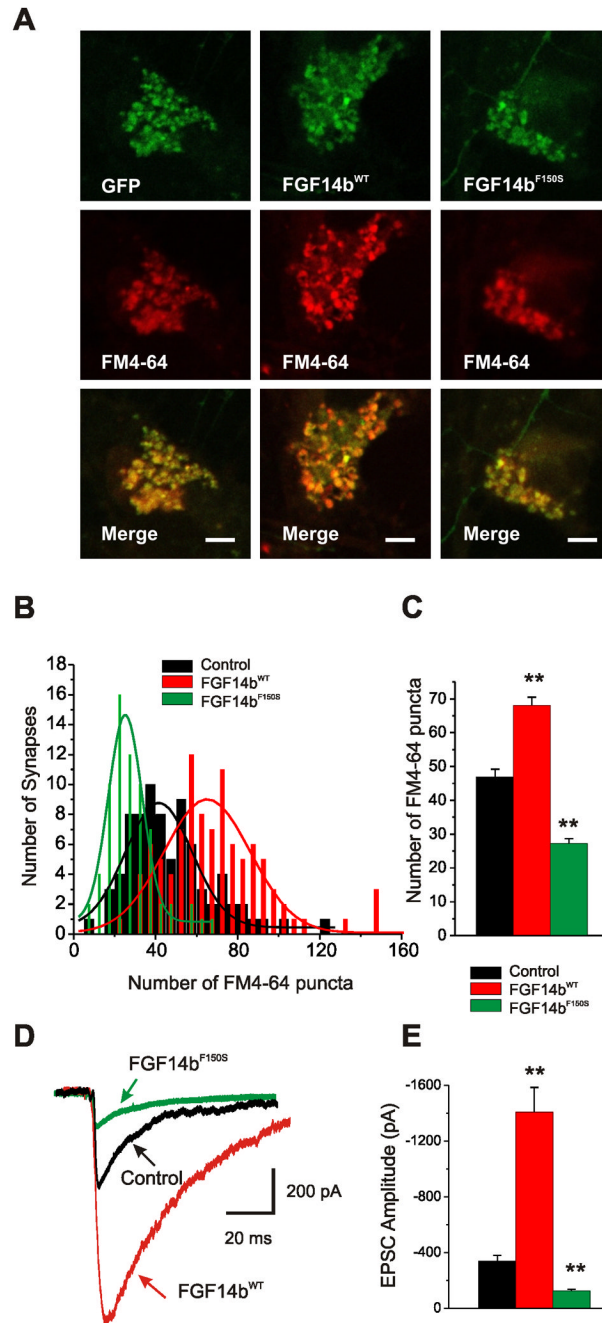


Figure 7. The SCA27 FGF14 mutant in granule cells reduces presynaptic Ca^{2+} influx and EPSCs at a granule cell to Purkinje cell synapse

A. Confocal images from cultured cerebellar neurons expressing GFP (left), FGF14^{WT} (middle), or FGF14b^{F150S} (right) that were loaded with FM4-64 by a 90 sec depolarization using 90 mM KCl. Scale bar, 5 μm . **B.** The distribution of FM4-64 puncta per synapse within a 45 \times 45 μm^2 ROI in neurons transfected with GFP (black, n=81), FGF14^{WT} (red, n=98), or FGF14b^{F150S} (green, n=69). The data for each group were fit to a Gaussian distribution. **C.** Averaged puncta number per 45 \times 45 μm^2 ROI for neurons over-expressing control GFP (black, n=81), FGF14^{WT} (red, n=98), or FGF14b^{F150S} (green, n=69). **D.** Representative EPSC traces recorded from Purkinje cells evoked by a 20 ms depolarization

of the granule cell from a holding potential of -70 mV to -10 mV in which the presynaptic granule cell was transfected with GFP (black), FGF14^{WT} (red), or FGF14b^{F150S} (green). **E.** Averaged amplitude of Purkinje cell EPSCs when the presynaptic granule cell expressed GFP control (n=14), FGF14^{WT} (n=19), or FGF14b^{F150S} (n=10). **p<0.01 versus Control.

Table 1Summary electrophysiology data for Ca_v2.1 and Ca_v2.2 channel activation and inactivation in HEK293 cells

		V _{1/2} (mV)	K	n
Activation				
Ca_v2.1	Control	-11.5 ± 0.9	5.9 ± 0.2	31
	FGF14 ^{WT}	-12.5 ± 0.5	5.4 ± 0.2	44
Ca_v2.2	Control	-4.8 ± 0.8	5.6 ± 0.2	24
	FGF14 ^{WT}	-5.2 ± 0.8	5.5 ± 0.2	25
Inactivation				
Ca_v2.1	Control	-19.3 ± 0.8	17.5 ± 1.4	23
	FGF14 ^{WT}	-18.2 ± 2.3	15.1 ± 0.6	20
Ca_v2.2	Control	-18.4 ± 1.5	14.8 ± 1.3	15
	FGF14 ^{WT}	-20.0 ± 1.5	14.1 ± 0.6	14

Activation kinetics were obtained from fits with a Boltzmann equation of the form: $G = G_{\max} / [1 + \exp((V - V_{1/2})/k)]$, where G_{\max} is the extrapolated maximum Ca²⁺ conductance, V is the test voltage, $V_{1/2}$ is the half-activation voltage, and k is the slope factor. Steady-state inactivation were obtained from fits with a Boltzmann relationship, $I/I_{\max} = (1 + \exp((V - V_{1/2})/k))^{-1}$.