REVIEW



Voltage-gated sodium currents in cerebellar Purkinje neurons: functional and molecular diversity

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Abstract

Purkinje neurons, the sole output of the cerebellar cortex, deliver GABA-mediated inhibition to the deep cerebellar nuclei. To subserve this critical function, Purkinje neurons fire repetitively, and at high frequencies, features that have been linked to the unique properties of the voltage-gated sodium (Nav) channels expressed. In addition to the rapidly activating and inactivating, or transient, component of the Nav current (I_{NaT}) present in many types of central and peripheral neurons, Purkinje neurons, also expresses persistent (I_{NaP}) and resurgent (I_{NaR}) Nav currents. Considerable progress has been made in detailing the biophysical properties and identifying the molecular determinants of these discrete Nav current components, as well as defining their roles in the regulation of Purkinje neuron excitability. Here, we review this important work and highlight the remaining questions about the molecular mechanisms controlling the expression and the functioning of Nav currents in Purkinje neurons. We also discuss the impact of the dynamic regulation of Nav currents on the functioning of individual Purkinje neurons and cerebellar circuits.

Keywords Conductance · Accessory subunit · Cerebellum · Intrinsic excitability

Introduction

In most mammalian central neurons, voltage-gated sodium (Nav) currents underlie the generation and propagation of action potentials and regulate repetitive firing rates [1, 2]. The detailed time- and voltage-dependent properties of the Nav currents, in concert with the many other voltage- and non-voltage-gated currents expressed, underlie the generation of cell-type-specific differences in the waveforms of individual action potentials and the rates and patterns of repetitive firing [1, 2]. In cerebellar Purkinje neurons, which fire repetitively at high frequency in vivo [3] and in vitro [4], independent of synaptic inputs [4, 5], for example, three Nav current components have been distinguished: (1) a rapidly activating and inactivating, i.e., transient, component, I_{NaT} ; (2) a non-inactivating, persistent component, I_{NaP} [6];

and (3) a resurgent component, I_{NaR} [5]. These Nav current components all contribute to shape the waveforms of action potentials and control the repetitive firing rates of Purkinje neurons [5, 7, 8]. Deletion or mutation of *Scn1a* or *Scn8a*, which encode the Nav channel pore-forming (α) subunits Nav1.1 or Nav1.6, respectively, attenuates Nav currents and repetitive firing rates in Purkinje neurons, the sole output neurons of the cerebellar cortex, and impairs motor performance [9–12]. Interestingly, modulation of the voltage-dependent properties of Nav currents in Purkinje neurons also disrupts high-frequency firing and affects motor coordination and balance [13].

Here, we review the time- and voltage-dependent properties of the Nav currents expressed in cerebellar Purkinje neurons and explore the roles of the three Nav current components, I_{NaT} , I_{NaP} , and I_{NaR} , in shaping the firing properties and the functioning of these cells. We also discuss recent studies focused on identifying the molecular determinants of native Nav currents in Purkinje neurons and the mechanisms contributing to I_{NaT} , I_{NaP} , and I_{NaR} gating. Finally, we discuss upstream mechanisms that may contribute to Nav current modulation and important, unanswered questions about the regulation of Nav channel expression, properties, and functioning in cerebellar Purkinje neurons.

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Purkinje neurons are the sole output of the cerebellar cortex

In pioneering work conducted during the 1960s, Eccles, Llinás, and Sasaki directly linked the cerebellum to motor function and mapped the physiological underpinnings of the cerebellar circuit [14–21]. Purkinje neurons function as the final destination of sensory information processing in the cerebellar cortex (Fig. 1) and the relay of information from the cerebellar cortex to downstream targets [22]. Sensory information enters the cerebellum via mossy fibers (Fig. 1), originating from over two-dozen brain and spinal cord nuclei [22], that synapse on and excite granule neurons [14]. The granule neurons, in turn, project parallel fiber axon bundles into the molecular layer of the cerebellar cortex (Fig. 1), where they make excitatory synapses on the large dendritic trees of Purkinje neurons [15]. The axons of Purkinje neurons exit the cerebellar cortex (Fig. 1) and provide GABA-mediated inhibition to neurons in the deep cerebellar nuclei [23-25]. In addition to granule cell inputs, Purkinje neurons also receive excitatory synaptic climbing fiber inputs from neurons in the inferior olivary nucleus [16] and local GABAergic



Fig. 1 Schematic of the cerebellar circuit shown in sagittal and coronal planes. Climbing fibers, originating in the inferior olive, form excitatory synapses on the dendrites of Purkinje neurons (green). Mossy fibers, which originate in multiple hindbrain and spinal cord nuclei, form excitatory synapses on granule neurons (red) in the cerebellar cortex. The granule cells, in turn, project parallel fiber axons into the molecular layer and form excitatory synapses on the dendrites of Purkinje neurons. Stellate and basket cells (blue), in contrast, form inhibitory synapses on Purkinje neurons, the sole output of the cerebellar cortex, which project inhibitory axons to the deep cerebellar nuclei

inhibitory inputs from cerebellar basket [26] and stellate [17] cells (Fig. 1).

The interplay of inhibitory and excitatory synaptic inputs modulates the repetitive firing rates and can cause brief pauses in the firing of individual Purkinje neurons [19, 27], as well as synchronize the repetitive firing of adjacent Purkinje neurons [28, 29]. The synchronized firing of Purkinje neurons is important in relaying spike timing information to neurons in the deep cerebellar nuclei [30], and is evident during cerebellar-related motor behaviors [31, 32]. In the absence of synaptic inputs, however, Purkinje cells continue to fire action potentials spontaneously and repetitively at high frequencies [4, 5]. This sustained, highfrequency repetitive firing, which depends on the unique intrinsic membrane properties of Purkinje neurons, is vital to the functioning of the cerebellar cortex. Indeed, alterations in the rates and patterns of repetitive firing of Purkinje neurons impact balance and motor performance [9-13].

Nav currents underlie high-frequency repetitive firing in cerebellar Purkinje neurons

In 1980, Llinas and Sugimori demonstrated that the highfrequency repetitive firing of action potentials observed in Purkinje neurons in acute cerebellar slices requires subthreshold Na⁺-dependent, tetrodotoxin (TTX)-sensitive membrane depolarizations [6, 33]. Combining voltageclamp and action potential-clamp recordings on isolated Purkinje neurons, Raman and Bean detailed the time- and voltage-dependent properties of the critical subthreshold currents underlying spontaneous action potential firing, providing new insights into the roles of these currents in shaping the waveforms of individual action potentials and in controlling the rates and patterns of repetitive firing [5, 7,]34]. Importantly, these experiments revealed that the intrinsic, high-frequency repetitive firing of action potentials in cerebellar Purkinje neurons is not dependent on calcium currents (I_{Ca}) or hyperpolarization-activated cation currents (I_{h}) [7], conductance pathways often associated with pacemaker activity [35-37]. Instead, TTX-sensitive Nav conductances were identified as the primary source of subthreshold membrane depolarizations, both during the action potential and the inter-spike interval, and to be critical in controlling the repetitive firing of Purkinje neurons [5, 7]. Indeed, spontaneous, high-frequency repetitive firing persists in isolated Purkinje neurons with pharmacological inhibition of I_{Ca} and $I_{\rm h}$, but not in the presence of TTX [7].

Action potential-clamp recordings also revealed that the TTX-sensitive inward Nav currents are activated in Purkinje neurons at subthreshold membrane potentials during the upstroke of individual action potentials, as well as during the inter-spike interval [7]. Combining action potential-clamp and voltage-clamp recordings, Carter and Bean [38] went on to examine $I_{\rm Na}$ availability during action potentials and during inter-spike intervals in Purkinje neurons firing at different rates. To quantify $I_{\rm Na}$ as a function of time and voltage, they applied steady-state depolarizing voltage steps at different times during action potentials [38]. These experiments revealed that Purkinje neurons continue to fire at > 100 Hz even with ~75% of the Nav channels inactivated. In Purkinje neurons firing at 300 Hz, $I_{\rm Na}$ availability fell to 15% of its (original) maximal value [38].

Multiple Nav current components drive and fine tune repetitive firing in Purkinje neurons

Steady-state voltage-clamp recordings revealed that the critical Nav conductance in cerebellar Purkinje neurons included a rapidly activating and inactivating, i.e., a transient Nav current component, I_{NaT} , and a non-inactivating, i.e., persistent Nav current component, I_{NaP} [5, 7, 9, 39], as illustrated in Fig. 2a. In mouse Purkinje neurons, I_{NaP} , which is observed at voltages as negative as -80 mV, was shown to contribute to the regulation of action potential thresholds and the amplification of dendritic excitatory and inhibitory synaptic potentials [40]. Voltage-clamp experiments also revealed that the decay of I_{NaT} in cerebellar Purkinje neurons follows a bi-exponential time course [9]. At room temperature, the fast component of I_{NaT} decay inactivates with a time constant of ~1 ms, and the slow component of

 I_{NaT} decay inactivates with a time constant of ~8 ms [9]. Recovery of the fast inactivating component of I_{NaT} (from inactivation) is best described by a single exponential; the time constant of recovery at -40 mV (and at -80 mV) at room temperature, for example, is 3.8 ms [5, 41]. This rate is considerably faster than the rate of I_{NaT} recovery observed in other mammalian central neurons. In hippocampal CA1 pyramidal cells, for example, the time constant of I_{NaT} recovery from inactivation at -70 mV at room temperature was reported to be 9 ms [42].

Voltage-clamp analyses also led to the discovery of a distinct "resurgent" Nav current component, I_{NaR} , in Purkinje neurons, revealed on membrane repolarizations following depolarizing voltage steps [5, 43], as illustrated in Fig. 2b. I_{NaR} , therefore, would be expected to provide depolarizing, inward (Na⁺) current during the falling phase of action potentials and inter-spike intervals in Purkinje neurons [8, 41]. Action potential-clamp recordings from Purkinje neurons indeed revealed an increase in TTX-sensitive inward Nav current during the action potential downstroke [7, 38, 44, 45].

To explore the functional role(s) of the individual Nav current components in the regulation of high-frequency repetitive firing in Purkinje neurons, Khaliq et al. [8] constructed a conductance-based Purkinje neuron model based on the experimentally determined properties of the three Nav current components, I_{NaT} , I_{NaP} , and I_{NaR} , together with the three voltage-gated potassium (Kv) currents, two voltagegated calcium (Cav) currents, and the calcium-activated potassium current, identified in these cells [7–9, 34]. The Nav currents were simulated using a Markov model [34]





Fig. 2 Three Nav current components, I_{NaT} , I_{NaP} , and I_{NaR} , resolved in steady-state voltage-clamp experiments. In the example voltageclamp paradigm illustrated, I_{NaT} and I_{NaP} , are revealed during the depolarizing voltage step to +10 mV (**a**), and I_{NaR} is observed on membrane repolarization (from +10 mV) to -40 mV (**b**). In both

the left and right panels, the waveform of the evoked Nav currents is shown below the voltage-clamp paradigm, and schematics of Nav channel gating states are presented; the intrinsic channel inactivation gate is shown in red and the open-channel blocking particle is depicted in blue

that included an 'open blocked' state to simulate I_{NaR} . This conductance-based model allowed manipulation of I_{NaR} and I_{NaT} densities and revealed that decreasing I_{NaR} reduced repetitive firing rates in model cells [8]. Using this model [34] to simulate I_{NaR} , Ransdell et al. [41] demonstrated that (positively) scaling the magnitude of I_{NaR} in dynamic-clamp recordings [46, 47] increased instantaneous and evoked firing rates in Purkinje neurons in acute cerebellar slices.

Nav channel gating in cerebellar Purkinje neurons

Each Nav channel pore-forming (α) subunit contains four homologous domains (domains I-IV), each with six transmembrane segments (S1-S6) that are linked by cytosolic peptides (Fig. 3a) [48]. The S5 and S6 transmembrane domains contribute to the formation of the Na⁺-selective pore (Fig. 3b). The voltage-sensing, S4, segment in each domain has multiple positively charged residues, repeated at three-residue intervals, followed by two hydrophobic amino acids [48, 49]. The positively charged amino acid residues in the S4 domain respond to changes in membrane voltage and move (outward) on membrane depolarization [50]. This charge movement opens the Nav channel pore, allowing Na⁺ influx. Following opening, Nav channels inactivate, mediated by the cytosolic DIII-DIV linker peptide, specifically by three residues in the DIII-DIV linker that form the hydrophobic (IFM) motif [51], terminating Na⁺ influx. The IFM motif binds to a site in the channel pore that is only made available after the outward movement (opening) of the DIV S4 segment [52–54]. Importantly, this means that Nav channel inactivation, although tightly linked to voltage-dependent Nav channel activation [55], is a voltage-independent process [52, 56].

Interestingly, the S4 voltage sensors in each of the four domains of an Nav channel α subunit have different voltage sensitivities [53]. Because Nav channel inactivation requires the movement or activation of the domain IV voltage sensor [54], and Nav channels conduct Na⁺ after activation of domains I, II, and III [57, 58], I_{NaP} can arise at membrane potentials that activate the domain I, II, and III voltage sensors if domain IV remains deactivated. Importantly, these gating mechanisms indicate that differential regulation of the voltage sensitivities of the domain I–IV voltage sensors may regulate the proportion/density and voltage dependence of I_{NaP} and I_{NaR} .

The "resurgent" component of the Nav current in Purkinje neurons, I_{NaR} , is evident on membrane repolarization following brief depolarizations to positive test potentials (Fig. 2b) [5, 43]. In Purkinje neurons, I_{NaR} , peaks at -35 to -45 mV, activates with a time constant of 5-6 ms and inactivates with a time constant of $\sim 20 \text{ ms} [5, 41, 43]$. Importantly, the fact that I_{NaR} is not revealed instantaneously on membrane repolarization, but rather takes time to activate, reveals that it (I_{NaR}) does not reflect I_{NaP} tail currents. Rather, I_{NaR} must be mediated by newly opened or recovered Nav channels. In Purkinje neurons, prolonged (40 ms), moderate depolarizations (e.g., to -30 mV) result in I_{NaR} (at -40 mV) that is much smaller than I_{NaR} (also measured at -40 mV) after a short (5 ms) depolarizing step to more positive (e.g., + 30 mV) potentials [34]. Using (both) voltage-clamp protocols mimicking these paradigms on the same cell, Raman and Bean [34] showed that I_{NaT} recovery paralleled I_{NaR} , i.e., I_{NaT} recovery was greater when measured after a hyperpolarizing voltage step producing a large I_{NaR} , than following steps yielding little I_{NaR} . This result has several important



Fig. 3 Schematic of the membrane topology of Nav α and β subunits. a Nav α subunits have four homologous (DI–DIV) domains, each of which has six transmembrane segments; the voltage-sensing S4 segment in each domain is red and the two (S5 and S6) transmembrane domains contributing to the pore are brown. The inactivation gate (grey) in the DIII–DIV linker and the Nav channel transmembrane accessory (β) subunits, Nav β 1–Nav β 4, are also shown. Nav β 2 and

Nav $\beta4$ (green) form covalent disulfide bonds with Nav α subunits, whereas Nav $\beta1$ and Nav $\beta3$ (blue) interact non-covalently with Nav α subunits. Binding sites for additional Nav channel interacting proteins, including AnkyrinG (purple) in the DII–DIII cytosolic linker and iFGF14 (blue) in the C-terminus, are also indicated. **b** The 24 transmembrane spanning segments of a single Nav α subunit assemble to form an Na⁺ selective pore (brown)

implications. First, because the magnitude of I_{NaR} was dependent on the duration and potential of the pre-step, with shorter, more depolarized potentials resulting in larger I_{NaR} , Raman and Bean reasoned that there were two competing Nav channel inactivation mechanisms [34]. Specifically, they suggested that Nav channels undergo conventional inactivation (mediated by the cytosolic DIII–DIV linker) at more hyperpolarized potentials and a novel form of Nav channel inactivation at more depolarized potentials.

This novel mechanism of Nav channel inactivation was referred to as "open-channel block", i.e., activated Nav channels opened at depolarized membrane potentials were blocked by a particle distinct from the DIII–DIV linker. Grieco et al. [59] specifically suggested that a blocking particle, not part of the Nav α subunit, competes with conventional inactivation, occludes the Nav channel pore during strong depolarizations, and is expelled from the pore on membrane repolarization allowing 'resurgent' Na⁺ influx (before the Nav channels enter a conventional inactivated state). In the open-channel block mechanism, Nav channels, or at least the subset of Nav channels that undergo openchannel block, forego conventional inactivation allowing faster recovery of I_{NaT} during hyperpolarization [34].

The proposed mechanism for I_{NaR} generation requires the presence/functioning of a particle that competes with conventional inactivation and occludes the Nav channel pore at depolarized potentials. The Nav channel β 4 subunit (Nav β 4), which binds covalently to Nav α subunits [60, 61] and has a short (20 residues) intracellular C-terminus, which includes several positively charged amino acids, in addition to a phenylalanine residue, was suggested by Lewis and Raman to function as an endogenous "blocking" particle [62]. In a landmark study, Grieco et al. [63] demonstrated that I_{NaR} , measured in membrane patches excised from Purkinje neurons, was abolished when the intracellular face of the membrane was exposed to proteases specific for cleavage at positively charged (trypsin) or aromatic (chymotrypsin) residues. In addition, following protease treatment, I_{NaR} was rescued by intracellular application of a synthetic peptide, referred to as $\beta 4_{peptide}$, homologous to the Nav $\beta 4$ C-terminus [63]. Interestingly, although co-expression of Nav α subunits with Nav_{β4} in heterologous cells results in Nav currents that lack I_{NaR} [64–66], intracellular applications of the $\beta 4_{peptide}$ to HEK-293 cells expressing Nav1.1, 1.4, 1.5, or 1.7 produce robust I_{NaR} [65–68]. It has also been reported that shRNA-mediated 'knockdown' of Navβ4 reduced or eliminated I_{NaR} in cerebellar granule neurons [69], as well as in dorsal root ganglion neurons [70]. More recently, however, it was shown that targeted deletion of Nav β 4 (Scn4b^{-/-}) reduces, but does not abolish I_{NaR} in striatal medium spiny neurons [71] or in cerebellar Purkinje neurons [41]. In fact, I_{NaR} amplitudes/densities in $Scn4b^{-/-}$ Purkinje neurons were ~ 50% of wild-type I_{NaR} levels, and in addition,

the time- and voltage-dependent properties of the residual I_{NaR} were indistinguishable from wild-type I_{NaR} [41]. These results clearly indicate that Nav β 4-independent mechanisms contribute to the generation of native I_{NaR} .

An alternative, or perhaps additional, mechanism for I_{NaR} generation is that there is a range of membrane voltages and/ or a distinct subset of Nav channels in which the rate of Nav channel deactivation is slower than the rate of channel recoverv from inactivation [41, 43]. If Nav channel closing (deactivation), for example, is slower than recovery from inactivation at membrane potentials between -50 and -30 mV, Na⁺ influx through recovered Nav channels will occur. This mechanism was first suggested by the results of experiments demonstrating that β -toxins from *Centruroides* scorpions [72] slow the kinetics of deactivation of heterologously expressed Nav1.6 channels in HEK-293 cells [73, 74]. The change in deactivation kinetics resulted in the generation of resurgent Nav currents in cells exposed to the toxin [73, 74]. It has also been reported that exposure of rat cerebellar Purkinje neurons to Cn2 toxin, a β -toxin from *Centruroides* noxius, produced an additional Na⁺-mediated resurgent current with voltage-dependent activation properties distinct from native I_{NaR} [73]. It remains to be determined if there are intrinsic mechanisms in Purkinje neurons, or other neuronal cell types, which similarly modulate Nav current deactivation kinetics to produce resurgent Nav currents.

A fascinating aspect of I_{NaR} in Purkinje neurons that has not been thoroughly explored to date is that, at negative potentials, the magnitude and the rate of decay of I_{NaR} parallel the magnitude and the rate of decay of the slow component of I_{NaT} inactivation. Representative recordings from an isolated Purkinje neuron demonstrate that the magnitude and decay phase of I_{NaR} at -40 mV (from a depolarized membrane potential) are indistinguishable from the slow component of I_{NaT} inactivation evoked directly on membrane depolarization to -40 mV from a hyperpolarized membrane potential (Fig. 4). Noting this property, Raman and Bean proposed that, rather than competing with conventional inactivation, Nav channels responsible for I_{NaR} may undergo open-channel block at all voltages that activate I_{NaT} [34].

Nav channel alpha and accessory subunits in the functioning of Purkinje neurons

Transcripts encoding the Nav1.6 (*SCN8A/Scn8a*) and Nav1.1 (*SCN1A/Scn1a*) pore-forming α subunits are robustly expressed in cerebellar Purkinje neurons [75]. In situ hybridization studies have provided conflicting results [76–78] regarding the expression of *Scn2a*, which encodes the Nav1.2 α subunit, in cerebellar Purkinje neurons. In addition, although readily detected in cerebellar granule neurons, the Nav1.2 protein was not detected on Purkinje neuron



Fig. 4 Decay phases of I_{NaT} and I_{NaR} . In the panel on the left, Nav current waveforms recorded from an isolated neonatal Purkinje neuron during two different voltage-clamp paradigms are shown; the voltage-clamp paradigms are illustrated above the current records. In the first case, membrane depolarization to -40 mV from a holding potential (HP) of -90 mV (red) evoked I_{NaT} (red); note the fast (red, open arrow) and slow (red, filled arrow) components of inactivation of I_{NaT} . Membrane repolarization to -40 mV following a brief (5 ms) strong (+10 mV) depolarization from the same HP revealed I_{NaR} (black). Note that the time course of I_{NaT} decay (black, filled arrow) are indistinguishable. The currents are shown on expanded amplitude and time scales in the records shown in the panel on the right

somata, dendrites and axons [12]. In contrast, robust anti-Nav1.6 and anti-Nav1.1 protein labeling has been reported in Purkinje neurons [79, 80], and the Nav1.6 and Nav1.1 proteins are differentially localized. Anti-Nav1.1 labeling. for example, was observed on the cell bodies and dendrites of Purkinje neurons [79] and, in addition, was detected in the segment of the axon proximal to the soma [81]. In contrast, anti-Nav1.6 staining was reported to be localized to the axon initial segment (AIS) [80, 81]. Focal application of TTX (to block Nav currents) or β -pompilidotoxin (to increase I_{NaR}) [82] revealed that action potentials in Purkinje neurons are initiated in the distal portion of the AIS, although when AIS Nav channels were blocked, somatic Nav channels in Purkinje neurons were sufficient to generate repetitive firing [83]. Simultaneous extracellular recordings from Purkinje neuron somata and different sites along the axon [84], in combination with the results of antibody labeling [80, 81], suggest that Nav1.6 plays a critical role in the initiation of action potentials in the AIS and the regulation of repetitive firing in Purkinje neurons.

In studies designed to explore directly the contributions of individual Nav α subunits to the regulation of action potential generation and repetitive firing in Purkinje neurons, the functional consequences of the targeted disruption of the genes encoding individual Nav α subunits were examined. The first of these studies used the naturally occurring *Scn8a^{med}* mutant mouse [85], which results in the functional 'knockout' of *Scn8a* and loss of the Nav1.6 protein [9, 85, 86]. As might have been expected, loss of Nav1.6 resulted in attenuation of repetitive firing in Purkinie neurons. Recordings from cells isolated from 14- to 19-day-old mice revealed spontaneous firing rates reduced to 9 Hz in Scn8a^{med}, compared with 35 Hz in wild type, Purkinje neurons [8]. In addition, peak I_{NaT} densities in Scn8A^{med} Purkinje neurons were ~ 65% of wild-type levels [9]. Interestingly, I_{NaP} and I_{NaR} densities were reduced to 30 and 10–20%, respectively, of wild-type levels in Scn8A^{med} Purkinje neurons [9]. Targeted disruption of Scn1a and loss of Nav1.1 also reportedly reduced spontaneous, repetitive firing rates in acutely isolated postnatal day 13-14 Purkinje neurons to 45 Hz, compared with 69 Hz in wild-type cells [12]. Loss of Nav1.1 resulted in I_{NaT} , I_{NaP} , and I_{NaR} densities that were 42, 41, and 31% of those measured in wild-type Purkinje neurons, respectively.

Taken together, these observations suggest that both Nav1.1 and Nav1.6 contribute to the high rates of repetitive firing in Purkinje neurons and animals lacking either of these subunits have severe deficits in balance and motor coordination [10, 12]. In addition, both the Nav1.1 and Nav1.6 α subunits contribute to I_{NaT} , I_{NaP} , and I_{NaR} , although loss of Nav1.6 disproportionately affects I_{NaR} and I_{NaP} . Using the Cre–lox system to eliminate *Scn8a* expression selectively in Purkinje *or* in cerebellar granule neurons, Levin and colleagues demonstrated that loss of Nav1.6 in granule neurons resulted in only minor behavioral defects, whereas mice lacking Nav1.6 in Purkinje neurons exhibited ataxia, tremors, and impaired coordination [11].

It is important to note that the biophysical properties of native I_{NaP} and I_{NaR} in Purkinje neurons are distinct from those of heterologously expressed Nav1.1- or Nav1.6-encoded Nav currents. Heterologous expression of Nav1.1, for example, revealed persistent Nav current amplitudes corresponding to 2–13% of I_{NaT} [87]. It has been estimated that Nav1.1-mediated I_{NaP} in cerebellar Purkinje neurons is ~ 1.4% of Nav1.1-mediated I_{NaT} [12]. Even more striking is the fact that *neither* Nav1.6 *nor* Nav1.1 produces resurgent Nav currents in heterologous expression systems [64, 65].

Accumulated evidence suggests that native neuronal Nav channels, like many other types of ion channels, function in macromolecular protein complexes comprising the pore-forming α subunits and a complement of accessory and regulatory proteins [88–90]. Various Nav channel accessory proteins have been linked to the regulation of Nav channel trafficking, localization, and gating [65, 91–93], increasing the complexities by which Nav channel regulation can influence intrinsic neuronal excitability, action potential waveforms, and repetitive firing. Clear roles for the intracellular fibroblast growth factors (iFGF11–14), also referred to as fibroblast homologous factors (FHFs) [13, 81, 94, 95], and the Nav channel β subunits (Nav β 1–4) [41, 43, 63] in the regulation of Purkinje neuron Nav channels, action potential

waveforms, and repetitive firing patterns, for example, have been provided. *FGF14* is the locus of mutations in spinal cerebellar ataxia type 27 (SCA27) [96], an autosomal dominant disorder characterized by progressive ataxia and cognitive decline [96]. iFGFs are unique from the canonical FGFs in that they lack a signal sequence for secretion and they do not activate any known FGF receptors [97, 98]. Rather, the iFGFs function intracellularly, binding to the C-terminus of Nav α subunits/channels [99–101].

iFGF14 is robustly expressed in Purkinje neurons and localizes with Nav1.6 at the AIS [81, 102]. iFGF14 null $(Fgf14^{-/-})$ mice display severe deficits in balance and motor coordination [103] and the vast majority of $Fgf14^{-/-}$ Purkinje neurons are quiescent and lack the ability to fire repetitively, even in response to depolarizing current injections [95]. Interestingly, shRNA-mediated 'knockdown' of iFGF14 in adult Purkinje neurons also resulted in deficits in balance and motor coordination and markedly attenuated firing in Purkinje neurons [13]. Additional experiments revealed that attenuated firing in $Fgf14^{-/-}$ Purkinje neurons could be attributed to a hyperpolarizing shift in the voltagedependence of steady-state inactivation of I_{NaT} and that firing could be rescued by membrane hyperpolarizations to membrane potentials that allow inactivated Nav channels to recover. It has also been reported that shRNA-mediated 'knockdown' of iFGF14 in neonatal Purkinje neurons in culture results in reduced I_{NaR} [94]. Loss of iFGF14, however, does not measurably affect I_{NaR} in acutely dissociated Purkinje neurons (unpublished). Additional iFGFs may also serve important roles in the regulation of Nav currents and the excitability of Purkinje neurons. Goldfarb and colleagues [102] demonstrated, for example, that animals lacking both iFGF12 and iFGF14 (Fgf12^{-/-}, Fgf14^{-/-}) display significantly worse motor defects than $Fgf14^{-/-}$ animals and that granule neuron excitability in $Fgf12^{-/-}$, $Fgf14^{-/-}$ mice was markedly attenuated. The role of iFGF12 in Purkinje neurons, however, has not been explored to date.

Transcripts (*Scn1b–Scn4b*) encoding Nav β 1–4 are expressed in developing cerebellar Purkinje neurons [41, 61, 93, 104, 105], although *Scn3b* (Nav β 3) is undetectable in mature Purkinje neurons [105]. As discussed above, Nav β 4 contributes to I_{NaR} and repetitive firing rates are attenuated in adult *Scn4b^{-/-}* Purkinje neurons. In marked contrast, repetitive firing rates and properties in neonatal (P14–P15) *Scn4b^{-/-}* and wild-type Purkinje neurons are indistinguishable [41], suggesting that the functional effects of Nav β 4 on the repetitive firing rates of Purkinje neurons are developmentally regulated. Adult *Scn4b^{-/-}* animals also have deficits in balance and motor coordination [41], although not as severe as those observed in *Fgf14^{-/-}* animals [13, 103].

In contrast with expectations, voltage-clamp experiments revealed that I_{NaR} is readily detected in $Scn4b^{-/-}$ Purkinje neurons and that the mean amplitude of the current is ~ 50%

of the magnitude of I_{NaR} in wild-type cells [41]. These observations demonstrate that additional, perhaps compensatory, mechanisms contribute to the generation of I_{NaR} in Purkinje neurons. Although the critical, alternate mechanism(s) remain to be identified, there are several possibilities. As mentioned above, it has been reported that shRNA-mediated 'knockdown' of iFGFf14 in neonatal Purkinje neurons in dissociated cell culture resulted in reduced I_{NaR} [94]. In addition, I_{NaR} has been reported to be attenuated (compared with wild-type levels) in neonatal $Scn1b^{-/-}$ granule neurons in culture [93]. Importantly, no changes in I_{NaP} or I_{NaT} densities or properties were observed in Scn1b^{-/-} granule neurons [93], indicating specificity for I_{NaR} . In contrast, Nav β 2 (Scn2b) more closely resembles Nav_{β4} than Nav_{β1} and includes multiple positively charged residues in the C-terminus [62]. However, relative I_{NaR} (to I_{NaT}) amplitudes are similar in $Scn2b^{-/-}$ and wild-type Purkinje neurons [63]. The effects of the additional loss of Nav β 1 or Nav β 2 with Nav β 4 on Nav currents in Purkinje neurons have not been investigated to date.

Teasing out the additional molecular determinants of I_{NaR} and determining if there are physiologically relevant mechanisms distinct from open-channel block of Nav channels that contribute to I_{NaR} will be important in efforts to provide a detailed understanding of the diverse roles of Nav currents in controlling the physiology and functioning of cerebellar Purkinje neurons.

Summary, open questions, and future directions

The discovery and detailed characterization of the biophysical properties of I_{NaP} [6] and I_{NaR} [5] have provided important insights into the contributions of these Nav current components, together with I_{NaT} , to regulate spontaneous, high-frequency, repetitive firing in Purkinje neurons [8, 9, 38, 41, 44]. Cerebellar Purkinje neurons are also being utilized to uncover the important contributions of distinct Nav channel accessory proteins on the properties of the Nav currents and on the regulation of action potential waveforms and repetitive firing properties [13, 63, 94, 106]. However, this work is still in its infancy. At present, for example, the only Nav channel accessory proteins shown to contribute to the regulation of the excitability and the functioning of cerebellar Purkinje neurons are iFGF14 and Nav β 4. In addition to the contribution of the many other Nav channel accessory proteins that are expressed in Purkinje neurons (including Nav β 1, Nav β 2, iFGF11, iFGF12, and iFGF13), we should also consider the possibility that multiple proteins are involved and that this could result in non-linear contributions to membrane excitability. Binding of calmodulin on the C-terminus of Nav1.4, which is expressed primarily in skeletal smooth muscle, for example, has been reported to affect channel permeability [107]. Over-expression of calmodulin has also been shown to attenuate Nav1.6-mediated I_{NaP} in Purkinje neurons in vitro [106]. It is certainly possible that calmodulin and iFGF14 (or other iFGFs), both of which bind to the C-terminus of Nav α subunits (Fig. 3), act competitively or synergistically to modulate Nav channel gating properties. Interestingly, mice with global deletion of Fgf12 do not exhibit changes in motor coordination or granule neuron excitability, whereas loss of Fgf14 caused deficits in both animal coordination and granule neuron firing [102]. The combined deletion of *Fgf14* and *Fgf12*, however, resulted in a more severe deficit in both these phenotypes than Fgf14 deletion alone [102]. These observations suggest that the expression/functioning of these two Nav channel accessory proteins may be linked in ways that are physiologically relevant. An important area for future research will be to explore the functional relation-

ships among the various Nav channel accessory subunits (Fig. 3) and to define their roles in the dynamic regulation of Purkinje neuronal excitability.

Beyond the combined and/or non-linear impact of Nav channel accessory subunits, there are also compelling recent studies that have identified potential upstream regulators that promote the binding of iFGF14 to Nav α subunits. These include glycogen synthase kinase 3 (GSKIII) [108, 109] and, more recently, casein kinase II (CKII) [110], which is a priming kinase for GSKIII. Inhibition of either of these kinases in hippocampal neurons disrupts iFGF14:Nav a subunit complexes, specifically Nav1.6- and Nav1.2-channel complexes [108, 110], and alters Nav channel localization and membrane excitability [110]. While GSKIII is required during neuronal development and actively regulates neuronal proliferation and migration [111–113], it also functions in the regulation of synaptic transmission [114]. GSKIII is detected in both pre-synaptic and post-synaptic neuronal compartments [115], and interestingly, GSKII has been found to participate in both long-term synaptic depression [116] and potentiation [117]. The finding that GSKIII also functions in the regulation of iFGF14:Nav α subunit complexes may signify a mechanism by which network-level activity regulates or modulates the intrinsic excitability of individual neurons. Although this hypothesis has not been tested at the network level, Purkinje neurons and the cerebellar circuit are an ideal model system to examine these processes.

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