Impact of gating modulation in Ca_v1.3 L-type calcium channels

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Ca_v1.3 L-type channels control inner hair cell (IHC) sensory and sinoatrial node (SAN) function, and excitability in central neurons by means of their low-voltage activation and inactivation properties. In SAN cells Ca_v1.3 inward calcium current (l_{ca}) inactivates rapidly whereas in IHCs inactivation is slow. A candidate suggested in slowing Ca_v1.3 channel inactivation is the presynaptically located ribbon-synapse protein RIM that is expressed in immature IHCs in presynaptic compartments also expressing Ca_v1.3 channels. Ca_v1.3 channel gating is also modulated by an intramolecular C-terminal mechanism. This mechanism was elicited during analysis of human C-terminal splice variants that differ in the length of their C-terminus and that modulates the channel's negative activation range and slows calcium-dependent inactivation.

Voltage-gated L-type calcium channels (LTCCs) form the Ca_v1 channel family, which comprises the isoforms Ca, 1.1, Ca, 1.2, Ca_v1.3 and Ca_v1.4. Ca_v1 channels are well-known pharmacotherapeutic targets of Ca2+ channel blockers such as dihydropyridines. Among the Ca_v1 family, Ca_v1.3 L-type channels physiologically control inner hair cell (IHC) sensory and sinoatrial node (SAN) function, as well as excitability in central neurons by means of their peculiar low-voltage activation and differential inactivation properties.1 A lot of our knowledge about Ca_v1.3 channels comes from knock-out mouse models $(Ca_v 1.3^{-/-})$.¹⁻³ $Ca_v 1.3^{-/-}$ mice are deaf and suffer from sinoatrial node dysfunction.^{1,2} The deafness is due to the complete absence of L-type Ca²⁺ currents (I_{Ca}) in IHCs and outer hair cells of the cochlea^{1,4} where Ca_v1.3-mediated L-type currents comprise about 90% of the calcium current. $Ca_v 1.3^{-/-}$ mice also exhibit an arrhythmic and bradycardic heart beat. This phenotype is due to an intrinsic defect in the SAN present at low heart rates.¹

 $Ca_v 1.3$ channels can conduct calcium inward current in the operating range of IHCs and SAN cells (which is between -60 and -40 mV) due to their negative activation range.^{1,5-7} In neurons, $Ca_v 1.3$ channels shape neuronal firing as for example in striatal medium spiny neurons⁸ and contribute to dendritic calcium oscillations in Dopamine (DA)-releasing neurons of the *substantia nigra* pars compacta (SNc).⁹ Ca²⁺ ions entering SNc

DA neurons through LTCCs elevate cellular vulnerability to toxins used to create animal models of Parkinson's disease (PD).¹⁰ In animal models of PD, block of Ca_v1.3 channels appears to underlie neuroprotective therapeutic effects of dihydropyridine (DHP) LTCC blockers¹⁰ and DHPs ameliorate the development of L-DOPA-induced dyskinesias.¹¹ In a retrospective human study, long-term use of calcium channel blockers was associated with a significantly reduced risk of a PD diagnosis.¹² In this context it is important to note, that DHPs act as state-dependent blockers that need the channel's inactivated state and are thereby very likely to show activity-dependent potency.

The typical low activation threshold is intrinsic to Ca, 1.3 channels; however, their inactivation properties vary in a wide range and seem to be differentially modulated. Whereas Ca, 1.3 I_{Ca} inactivates rapidly in SAN cells^{1,13} its inactivation is slow in IHCs.1 CaM-like Ca2+ binding proteins (CaBPs) have been shown to eliminate calcium-dependent inactivation (CDI) of a short rat Ca, 1.3 channel isoform by competing CaM binding to the channel's C-terminus^{14,15} and also Ca₄B2 was recently reported to slightly affect CDI in IHCs,16 but voltage-dependent inactivation remained largely unaltered. A candidate suggested in slowing Ca, 1.3 channel inactivation was a presynaptically located ribbon-synapse protein called Rab3-interacting molecule (RIM) that is co-localized with Ca, 1.3 in the same presynaptic compartments of IHCs.17 In tsA-201 cells, RIM proteins inhibit Ca, 1.3 inactivation by slowing both CDI and VDI and induce a non-inactivating current component typical for Ca, 1.3 currents in IHCs.¹⁷ The modulatory effects of RIM are mediated via its binding to the Ca_b-subunit of the Ca²⁺ channel complex (Fig. 1). Because RIM mRNA is detected in the organ of Corti in IHC preparations before the onset of hearing, RIM proteins might therefore partly account for the slow inactivation of Ca, 1.3 IHC currents at least in an early developmental stage.

Furthermore, $Ca_v 1.3$ channel gating is also controlled by an intramolecular protein-protein interaction in the channel's C-terminus.¹⁸ An intrinsic C-terminal modulator (CTM) controls both activation as well as inactivation properties via binding of the distal $Ca_v 1.3$ C-terminus to a more proximal domain containing the EF-hand, pre-IQ- and IQ-motif, and a regulatory domain right after the IQ-motif (Fig. 1) that has also been identified to be important in $Ca_v 1.2$ channels.¹⁹ Most interestingly, alternative splicing generates $Ca_v 1.3 \alpha 1$ -subunits with long or short C-termini in various tissues¹⁸ and might thereby enable tight control of channel gating. In tsA-201 cells, the absence of

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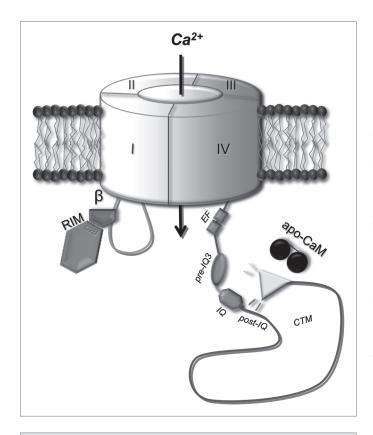


Figure 1. Cartoon of a proposed model for differential Ca_v1.3 gating modulation. The Ca_v1.3 channel is given as transmembrane pore forming α 1-subunit in grey. RIM protein (pink) via its C2B domain interacts with the Ca_vB subunit (magenta), which in turn binds the I–II loop (light blue). The C-terminus (dark blue) contains the CDI maschinery (comprising the EF hand, pre-IQ and IQ domains) and a post-IQ domain that interacts with the distal C-terminus forming the CTM in Ca_v1.3 channels. Modulation of the channel's CDI is suggested a competitive mechanism in which the Ca_v1.3 CTM competes with apoCaM for binding near the channel IQ domain.^{18,23}

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the CTM in a human short splice variant led to a lower activation range, a negative shift of the voltage-dependence of inactivation as well as more pronounced CDI of the channel compared to the long variant.¹⁸ The physiological implications of the Ca_v1.3 CTM are still ambiguous. The CTM in long Ca_v1.3 channels may be suitable for longer lasting Ca²⁺ signals triggered by stronger depolarization inducing CREB phosphorylation and synaptic plasticity.²⁰ Differences in the CTM inactivation pattern of long and short Ca_v1.3 channels (e.g., due a different extend of accumulation in their inactivated state) could though underlie different shapes or firing rates of action potentials as observed in different types of neurons.²¹ In terms of pharmacological intervention the Ca_v1.3 CTM should also have a strong impact on the efficiency of DHP block because this correlates with the amount of channels inactivated.^{5,6}

Based on the emerging role of Ca, 1.3 for normal and potential pathological cellular function, the discovery of the Ca_v1.3 CTM also raises a question about its potential as an alternative concept for pharmacological modulation of Ca_v1.3 channels. Potential Ca_v1.3 selective drugs may be envisaged to have disadvantages, such as slowing of the heartbeat, as seen in Ca₁1.3^{-/-} mice¹ and adaptive mechanisms seem not be able to restore pace-making in cardiac cells. The pathophysiological consequences observed in mice lacking function Ca_v1.3 channels about ten years ago are meanwhile also reported in two consanguineous Pakistani deafness families that also show severely impaired SAN function due to a splice variant specific Ca_v1.3 loss-of-function mutation.²² Interference with only the Ca, 1.3 CTM interaction should be beneficial to alter cellular excitability by switching form the long to short channel gating mode in avoidance of complete channel block. Such shift in Ca_v1.3 gating properties within a limited range could provide also a novel strategy for therapeutic Ca2+ channel modulation that avoids complete state-dependent inhibition of these channels, but could nevertheless induce functional changes to obtain the desired pharmacological effects.

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