

# Calmodulin Interactions with Voltage-Gated Sodium-Channels

Subjects: Cell Biology

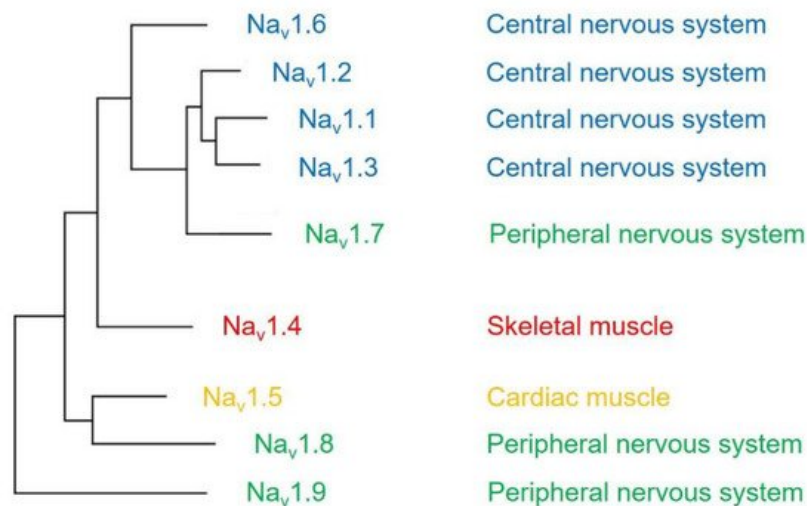
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Calmodulin (CaM) is a small protein that acts as a ubiquitous signal transducer and regulates neuronal plasticity, muscle contraction, and immune response. It interacts with ion channels and plays regulatory roles in cellular electrophysiology. CaM modulates the voltage-gated sodium channel gating process, alters sodium current density, and regulates sodium channel protein trafficking and expression.

Keywords: calmodulin ; voltage-gated sodium channel ; IQ domain ; sodium channel isoform ; genetic mutation

## 1. Introduction

The voltage-gated sodium channel (Na<sup>v</sup>) plays a vital role in the generation and propagation of action potential in excitable cells such as neurons and cardiac myocytes <sup>[1]</sup>. The family of voltage-gated sodium channels has nine members named Na<sup>v</sup> 1.1 through Na<sup>v</sup> 1.9 <sup>[2]</sup>. Among them, Na<sup>v</sup> 1.1, Na<sup>v</sup> 1.2, Na<sup>v</sup> 1.3, and Na<sup>v</sup> 1.6 are predominantly expressed in neurons of the central nervous system <sup>[3][4]</sup>. Three isoforms (Na<sup>v</sup> 1.7, Na<sup>v</sup> 1.8, and Na<sup>v</sup> 1.9) are widely expressed in neurons of the peripheral nervous system, such as dorsal root ganglia (DRG) neurons <sup>[5][6][7]</sup>. The isoform Na<sup>v</sup> 1.4 is responsible for upstroke of the action potential in skeletal muscle <sup>[8]</sup>, and Na<sup>v</sup> 1.5 is a cardiac-specific isoform known as cardiac sodium channel ( **Figure 1** ) <sup>[9]</sup>.



**Figure 1.** A phylogenetic tree and tissue distribution of voltage-gated sodium channels. Isoforms predominantly expressed in the central nervous system (blue), peripheral nervous system (green), skeletal muscle (red), and cardiac muscle (orange) are highlighted.

An essential property of voltage-gated sodium channels is “inactivation”, which prevents the reopening of the channel until complete recovery <sup>[10]</sup>. The inactivation regulates the frequency of action potential firing initiated by sodium channels in excitable cells. It reduces the breakdown of ionic gradients and cell death.

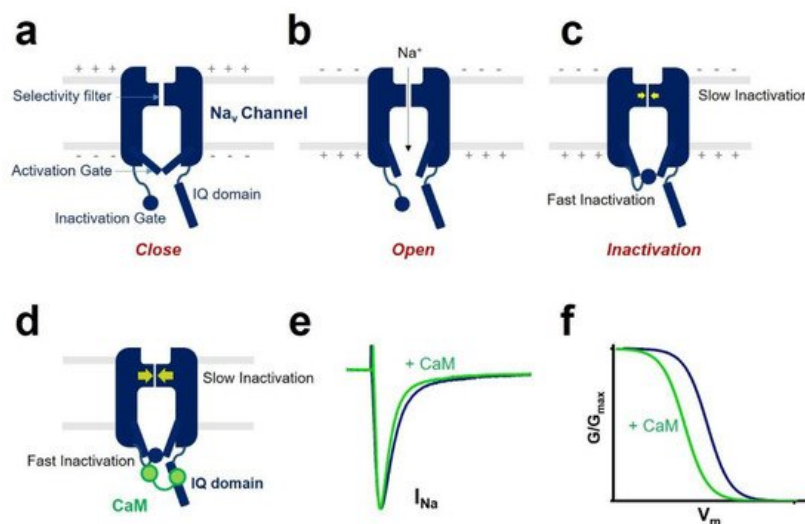
All voltage-gated sodium channels contain a calmodulin (CaM)-binding IQ domain necessary for the channel inactivation <sup>[11]</sup>. CaM is a small protein expressed in all eukaryotic cells <sup>[12]</sup>. It acts as a ubiquitous signal transducer and regulates essential processes such as neuronal plasticity, muscle contraction, and immune response <sup>[13]</sup>. CaM has two globular domains (i.e., N-terminal lobe and C-terminal lobe) connected by a linker. Each lobe contains two EF-hand motifs binding to Ca<sup>2+</sup> <sup>[14]</sup>. In the Ca<sup>2+</sup>-free state, the EF-hands are collapsed in a compact configuration. When Ca<sup>2+</sup> is bound to CaM, a conformational change occurs in the protein and rearranges the structural information of CaM.

The CaM-binding IQ domain is located within the C-terminal domain of the voltage-gated sodium channel. It contains around 25 residues with two highly conserved amino acids, isoleucine (I) and glutamine (Q), in the middle of the motif. The residues in the IQ domain form a seven-turn  $\alpha$ -helix binding to CaM in a  $\text{Ca}^{2+}$ -independent manner. In addition to the IQ domain, a globular domain in the C-terminus of voltage-gated sodium channel consists of EF hand-like (EFL) motifs that interact with CaM in the absence of  $\text{Ca}^{2+}$  [15][16], whereas an intracellular loop connecting domains III and IV (III-IV linker) in some isoforms of the sodium channel is shown to interact with CaM in the presence of  $\text{Ca}^{2+}$  [17].

## 2. CaM Regulation of Voltage-Gated Sodium Channel

CaM binds to the voltage-gated sodium channel and modulates the sodium channel gating process [15], alters sodium current density [18], and regulates sodium channel protein trafficking and expression [19].

One of the CaM modulations of voltage-gated sodium channel gating is inactivation ( **Figure 2** ). Voltage-gated sodium channel has two different types of inactivation, known as fast inactivation and slow inactivation. Fast inactivation occurs by occlusion of the intracellular pore by an “inactivation gate” formed by the cytoplasmic III-IV linker [10][20]. Slow inactivation is related to conformational rearrangements of the selectivity filter of the channel [21][22]. CaM mediates the fast inactivation by an interaction between III-IV linker and C-terminal domain of the voltage-gated sodium channel [23]. It enhances the slow inactivation by inducing a hyperpolarized shift of voltage dependence of inactivation and reducing the channel's availability. CaM interaction with the sodium channel affects channel inactivation kinetics. It strengthens the development of inactivation and slows the recovery from inactivated state of the sodium channel.



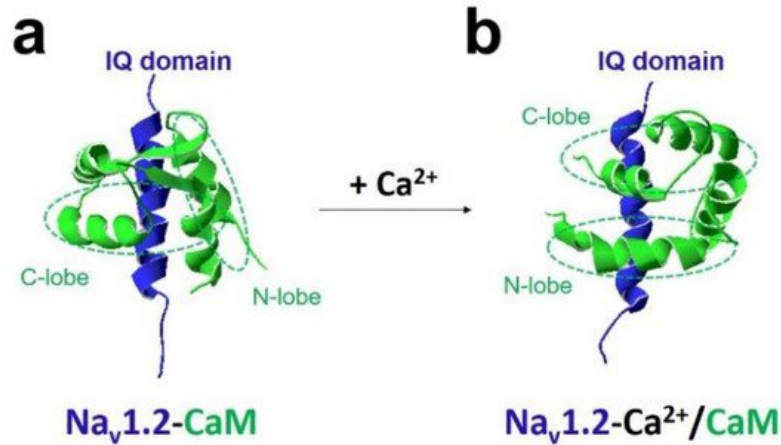
**Figure 2.** CaM regulation of sodium channel function by interaction with the IQ domain. (a-c) Voltage-gated sodium channel ( $\text{Na}_v$ ) has three states known as “close” (a), “open” (b), and “inactivation” (c). Several functional parts (inactivation gate, activation gate, and selectivity filter) are involved in channel gating. The “+” and “-” represent charge separation across the membrane. Differences in charges on opposite sides of the cellular membrane generate the membrane potential. At rest, the activation gate is closed (a). When the membrane is depolarized, the activation gate opens and allows  $\text{Na}^+$  to enter (b). At the same time, depolarization of the membrane causes the inactivation gate to close and induces the fast inactivation. The channel also shows slow inactivation related to conformational rearrangements of the selectivity filter (highlighted in yellow arrows in c). (d-f) CaM binds to the voltage-gated sodium channel IQ domain and modulates the sodium channel inactivation. CaM affects channel's fast inactivation kinetics (e) by an interaction between the inactivation gate formed by cytoplasmic III-IV linker and the C-terminal domain of the channel (d). CaM enhances the slow inactivation (highlighted in bigger yellow arrows in d). It induces a hyperpolarized shift of voltage dependence of inactivation and reduces the channel's availability (f).

In some isoforms of the voltage-gated sodium channels, CaM influences the activation process. It was reported that co-expression of CaM and  $\text{Na}_v 1.1$  produced a hyperpolarized shift of the voltage dependence of activation of the  $\text{Na}_v 1.1$  sodium channel [24].

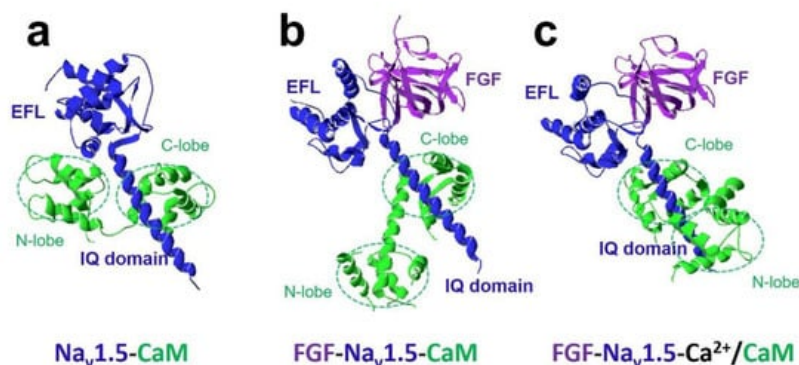
Moreover, CaM can modulate sodium channel current density [18]. Studies reported that CaM regulation of sodium current density is  $\text{Ca}^{2+}$ -dependent [24][25]. CaM does not affect sodium current amplitude in the absence of  $\text{Ca}^{2+}$ . In contrast, it increases the current density in the presence of a high concentration of  $\text{Ca}^{2+}$ .

### 3. Modulators Involved in CaM Regulation of Sodium Channel

The most crucial regulator is  $\text{Ca}^{2+}$  [11][26]. In the presence of  $\text{Ca}^{2+}$ ,  $\text{Ca}^{2+}$ /CaM ( $\text{Ca}^{2+}$ -binding form of CaM) has a different regulation of sodium channel function compared with apo-CaM ( $\text{Ca}^{2+}$ -free CaM), and  $\text{Ca}^{2+}$  binding to CaM induces rearrangements between the voltage-gated sodium channel and CaM lobes. In the absence of  $\text{Ca}^{2+}$ , the N-lobe of CaM does not target the IQ domain (Figure 3 a and Figure 4 a,b). In the presence of  $\text{Ca}^{2+}$ , the structure of the N-lobe is rearranged by  $\text{Ca}^{2+}$ . It generates an allosterically conformational change binding to the distal IQ domain of the channel (Figure 3 b and Figure 4 c). However, the C-lobe of CaM can bind to the voltage-gated sodium channel IQ domain in a  $\text{Ca}^{2+}$ -independent manner.



**Figure 3.**  $\text{Ca}^{2+}$  modulates the interaction between CaM and Na<sub>v</sub>1.2 IQ domain. The NMR structures showed Na<sub>v</sub>1.2 IQ domain (blue) in complex with CaM (green) in the absence (a) or presence (b) of  $\text{Ca}^{2+}$ . Two lobes (N-lobe and C-lobe) of CaM are indicated. C-lobe of CaM bound to the Na<sub>v</sub>1.2 channel IQ domain independent of  $\text{Ca}^{2+}$  concentration. In the absence of  $\text{Ca}^{2+}$ , the N-lobe of CaM did not interact with the IQ domain (PDB: 2KXW) (a). When binding to  $\text{Ca}^{2+}$ , the N-lobe of CaM was induced to interact with the distal IQ domain of Na<sub>v</sub>1.2 channel (PDB: 2M5E) (b).



**Figure 4.** FGF and  $\text{Ca}^{2+}$  modulate the interaction between CaM and Na<sub>v</sub>1.5 IQ domain. (a) The structure of the complex of the C-terminal domain of Na<sub>v</sub>1.5 with CaM showed C-lobe of CaM bound to the IQ domain, and N-lobe of CaM interacted with EFL (PDB: 4OVN). (b) Crystal structure showed the interactions between the C-terminal domain of Na<sub>v</sub>1.5, FGF, and CaM (PDB: 4DCK). When FGF bound to the channel, the N-lobe of CaM was rearranged. It did not make contact with EFL and was free to interact with other parts of the channel complex. The N-lobe of CaM likely interacted with the inactivation gate of the sodium channel and modulated the channel's inactivation. (c) In a high concentration of  $\text{Ca}^{2+}$ , the N-lobe of CaM was induced to interact with the distal IQ domain of Na<sub>v</sub>1.5 channel (PDB: 4JQ0).

The FHF2s were shown to form a complex with CaM and modulate CaM regulation of Na<sub>v</sub>1.5 function [27]. A recent study discovered fibroblast growth factor homologous factor FGF13 (fibroblast growth factor 13, also known as FHF2) tuned arrhythmogenic late sodium currents in CaM binding-deficient channels in cardiac myocytes [27]. The crystal structures in Figure 4 showed that FGF and  $\text{Ca}^{2+}$  mediated the interaction between CaM and Na<sub>v</sub>1.5 IQ domain. It indicated that the N-lobe of CaM interacted with the EFL of the channel in the absence of FGF. However, when FGF was bound to the channel, the N-lobe of CaM was rearranged. It did not make contact with the EFL and was free to interact with other parts of the channel complex. The N-lobe of CaM likely interacted with the III-IV linker of the sodium channel and modulated the channel's inactivation.

In addition, CaMKII has been reported to influence CaM interaction with the sodium channel. One study found CaMKII-mediated phosphorylation of the Na v 1.1 sodium channel IQ domain increased CaM binding affinity to the channel [28]. Three phosphorylation sites (T1909, S1918, and T1934) were identified in the Na v 1.1 IQ domain. Other studies demonstrated that CaMKII phosphorylated Na v 1.5 at residue S571 to decrease channel availability. The CaMKII phosphorylation site S571 is located at the intracellular DI-DII loop of the Na v 1.5 sodium channel, and mutation at position S571 abolished the hyperpolarized shift of the voltage dependence of inactivation produced by CaMKII phosphorylation. The effects of CaMKII-mediated phosphorylation of IQ domain on sodium currents have not been characterized. CaMKII-dependent IQ domain phosphorylation might confer normal sodium currents important for the function of cells where the Na v channels are expressed. The CaMKII-mediated effect seems to be isoform-specific. CaMKII phosphorylation enhances CaM affinity for sodium channel Na v 1.1 but not for Na v 1.2 [28][29].

With these partners, CaM regulates voltage-gated sodium channel function by altering channel gating, sodium current density, and expression of the sodium channel protein. Studies have shown that CaM has isoform-specific modulation of voltage-gated sodium channel function [30][31][32][33].

## 4. Interaction between CaM and Sodium Channel Isoforms

Na v 1.4 is the predominant subtype of sodium channel initiating skeletal muscle action potential [8]. Na v 1.4 mutations associated with periodic paralysis disorders affect skeletal muscle excitability [34].

The sodium channel Na v 1.5 IQ domain mutations associated with arrhythmia reduce CaM binding affinity. In HEK293 cells, overexpression of CaM attenuates late sodium currents caused by Na v 1.5 IQ domain mutations [35]. CaM binding to Na v 1.5 channel IQ domain also modifies late sodium currents in cardiac myocytes [27]. Study in transgenic mouse models showed FHF-tuned cardiac late sodium currents in ventricular myocytes. FHF-tuned late sodium current of the Na v 1.5 channel IQ/AA mutation (substitution of two conserved residues isoleucine (I) and glutamine (Q) with double alanines (AA) in the IQ domain), IQ/AA mutation has been reported to disrupt CaM binding to the IQ domain of voltage-gated sodium channels.

Studies in guinea-pig ventricular myocytes revealed CaM enhances the cardiac sodium current density [36]; this is different from the result that CaM does not influence on sodium current amplitude when co-expressing with Na v 1.5 channel in HEK293 cells [35]. It indicates that some environmental factors in cardiac tissues are involved in CaM regulation of Na v function.

To date, there is no study investigating the interaction between CaM and Na v 1.9 channel. The Na v 1.9 IQ domain shares high sequence homology with other sodium channels ( **Figure 5** ). It remains to be determined if CaM regulation of Na v 1.9 channel plays a role in channel function in sensory neurons.

Channel	Gene				★ ★					
Na <sub>v</sub> 1.1	SCN1A	1912	RKQEEVSAVI	I	QRAYRRHLLKRTVKQA	1938				
Na <sub>v</sub> 1.2	SCN2A	1902	RKQEEVSAIII	I	QRAYRRYLLKQKVKV	1928				
Na <sub>v</sub> 1.3	SCN3A	1897	RKQEEVSAII	I	QRNFRCYLLKQRLKNI	1923				
Na <sub>v</sub> 1.4	SCN4A	1724	RKHEEVCAIKI	I	QRAYRRHLLQSRMKQA	1750				
Na <sub>v</sub> 1.5	SCN5A	1898	RKHEEVSA	MVI	QRAFRHLLQ	1924				
Na <sub>v</sub> 1.6	SCN8A	1892	RKQEEVSAVVL	Q	RAYRGHLARRGFICK	1918				
Na <sub>v</sub> 1.7	SCN9A	1886	RKQEDVSATVI	Q	RAYRRYRLQNVKNI	1912				
Na <sub>v</sub> 1.8	SCN10A	1848	WKQEDISATVI	Q	KAYRSYVLHRS	1874				
Na <sub>v</sub> 1.9	SCN11A	1730	RKEEERGA	II	QKAFKYMVKVTKGDQ	1756				
										★ ★

**Figure 5.** Alignment of voltage-gated sodium channel family IQ domain and CaM-binding IQ domain mutations. Sodium channel mutations in the IQ domain associated with human diseases are highlighted in yellow. The conserved residues Ile (I) and Glu (Q) are indicated by stars.

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