

Ca²⁺ channels: diversity of form and function

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The past year has seen some significant advances in our understanding of the structural and functional properties of neuronal voltage-gated Ca²⁺ channels. Molecular cloning and protein purification studies have identified structural components, and expression studies are beginning to define the biophysical and pharmacological properties of the cloned channels. A number of studies of native Ca²⁺ channels show that the concept of channel modulation includes gating by both voltage and ligands.

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Introduction

The rapid entry of Ca²⁺ into excitable cells is mediated by a heterogeneous class of molecule, the Ca²⁺ channel. In the nervous system, Ca²⁺ channels play roles in mediating a wide variety of events including the patterning of neuronal firing, neurotransmitter release, and gene expression. To date, four major types of Ca²⁺ channel have been described in neurons (called T, L, N and P) [1,2]. Low-threshold Ca²⁺ channels (T-type) are transiently activated by relatively small depolarizations from hyperpolarized holding potentials and are completely inactivated at positive potentials. The other three Ca²⁺ channel types are activated at more positive potentials (high threshold) and display diverse kinetic properties. A major distinguishing characteristic of the different high-threshold Ca²⁺ channels is their sensitivity to various pharmacological agents: L-type channels are sensitive to dihydropyridine (DHP) agonists and antagonists, N-type channels are blocked by the *Conus geographus* peptide toxin, ω -conotoxin GVIA (ω -CgTx), and P-type channels are blocked by a crude venom fraction from the funnel web spider *Agelenopsis aperta*. Much research has been directed towards understanding Ca²⁺ channel diversity and it has been proposed that the distinct Ca²⁺ channel types are selectively localized in neurons and that each type may make unique contributions to neuronal physiology (for examples, see [3,4]).

The molecular dissection of neuronal Ca²⁺ channels has been made possible by the previous biochemical and molecular characterization of the skeletal muscle L-type Ca²⁺ channel (reviewed in [5,6]). This channel is a heterooligomeric complex consisting of five subunits (α_1 , α_2 , β , γ and δ), with the α_2 - and δ -subunits being derived from the same gene and proteolytically cleaved *in vivo* [7,8]. In a number of test systems the α_1 -subunits of several cloned L-type channels have been shown to function as both the voltage sensor and Ca²⁺-selective pore [9–12]. This brief review will focus on recent studies ad-

ressing the molecular nature of neuronal Ca²⁺ channel diversity and on the modulation of Ca²⁺ channels by voltage and ligands.

Structural and functional diversity of cloned neuronal Ca²⁺ channels

While neuronal Ca²⁺ channel diversity is likely to be the result of a number of factors, including modulation by accessory subunits and second-messenger mechanisms (see below), it is now evident that a significant portion of Ca²⁺ channel heterogeneity is due to the expression of unique α_1 -subunits. Currently, molecular cloning has identified five primary types of α_1 -subunit and four of these are expressed in the mammalian nervous system (classes A, B, C and D), Table 1, [13]. The deduced amino acid sequences of the Ca²⁺ channel α_1 -subunits show an overall conservation in structure, and are evolutionarily related to voltage-gated Na⁺ and K⁺ channels (Fig.1).

In addition to the five major types of α_1 -subunit (Table 1), molecular cloning and polymerase chain reaction studies have revealed a previously unanticipated degree of Ca²⁺ channel heterogeneity. Within each of the class A–D α_1 -subunits, multiple isoforms are expressed. Of potential functional significance is the observation that in a number of instances the isoforms found within a given class are the result of a nearly precise substitution of putative transmembrane segments [14•,15–17]. Furthermore, the expression of some isoforms appears to be spatially regulated and may result in qualitative differences between Ca²⁺ channels in different cell types. In the case of the rat class C α_1 -subunit, the generation of distinct isoforms is clearly a result of alternate splicing from a single gene [14•]. Alternate splicing is also suggested, but not yet proven, to be responsible

Abbreviations

ω -CgTx— ω -conotoxin GVIA; DHP—dihydropyridine; G protein—GTP-binding protein.

Table 1. Properties of cloned full-length Ca ²⁺ channel α_1 -subunits.							
Class	cDNA	Amino acids	Predicted molecular mass (kD)	Channel properties	Distribution	Notes	References
Skeletal muscle (L-type)		1,873	212	High threshold, DHP-sensitive, slow activation in L cells	Skeletal muscle	Rate of activation and inactivation increased by β -subunit	[9,11,34 [•] ,35 [•] ,62]
A (P-type)	BI-1	2,273	257	High threshold, blocked by crude venom from <i>Agelenopsis aperta</i> , insensitive to nifedipine and ω -CgTx, 16pS single channel conductance	Brain, heart, pituitary, GH4C1, PC12, C-cells	In oocytes expression increased by skeletal muscle $\alpha_2\delta$ - and β -subunits High levels of expression in cerebellum Isoforms differ in carboxyl region	[19 ^{••} ,20]
	BI-2	2,424	273				
	rbA-I	2,212	252				
B	rbB-I	2,336	262	Not available	Brain, PC12, C cells	Antibodies against rbB-I immunoprecipitate brain ω -CgTx binding sites	[26 [•]]
C (L-type)	pCARD3	2,171	243	High-threshold, DHP-sensitive	Brain, heart, lung, pituitary, kidney, aorta, GH4C1, PC12, C cells	Distinct isoforms generated by splicing Modulated by β -, $\alpha_2\delta$ - and γ -subunits	[10,12] [14 [•] ,17] [32 [•] ,36 [•] ,37 [•]]
	pSCaL	2,166	242				
	rbC-I	2,140	240				
	rbC-II	2,143	240				
	VSm α_1	2,169	244				
D (L-type)	α_1D	2,161	245	High-threshold, DHP-sensitive, reversible block by 10–15 μ M ω -CgTx, little inactivation over 700 ms	Brain, heart, pituitary, pancreas, GH4C1, PC12, C cells	Expression in oocytes requires β -subunit	[18 ^{••} ,60,61]
	CACN4	2,181	248				
	RB α_1	1,634	187				

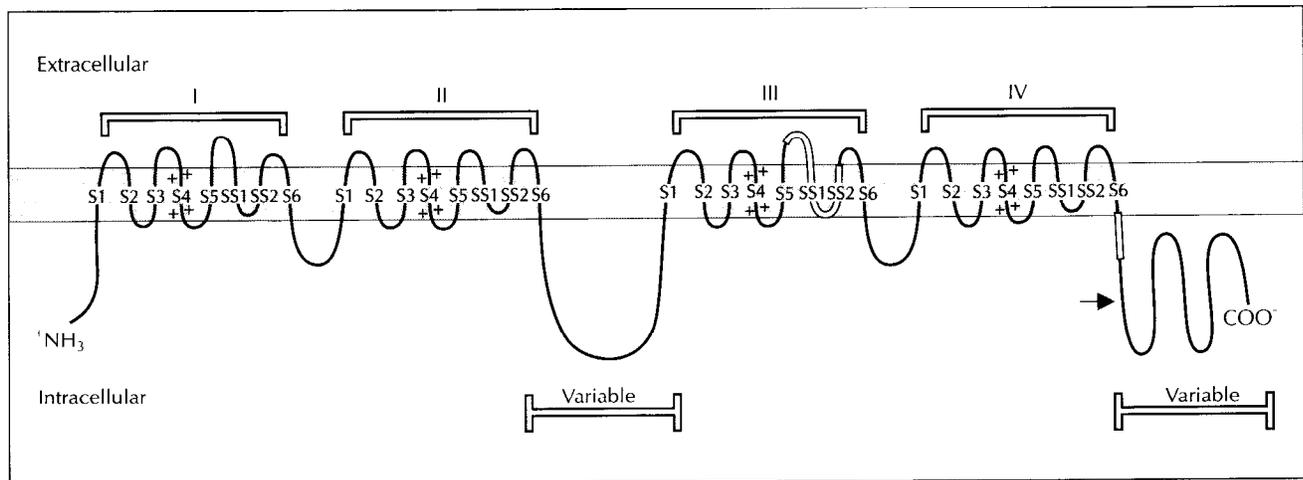


Fig. 1. The predicted transmembrane folding model of the Ca²⁺ channel α_1 -subunit. The five cloned types of Ca²⁺ channel α_1 -subunit are between 210 kD and 270 kD in predicted molecular mass and possess four internal repeated domains (I–IV) that are modelled to contain six α -helical transmembrane regions (S1–S6), including one (S4) that is positively charged and is thought to form part of the voltage sensor. By analogy to K⁺ channels the region separating segments S5 and S6 of each domain may contain two additional transmembrane segments (SS1 and SS2) that together form the pore of the channel [56]. The regions with the highest degree of amino acid identity among the five cloned types of α_1 -subunit are the four repeated domains and a short segment immediately flanking S6 of domain IV. The latter region, together with the loop between segments S5 and S6 in domain III, is implicated in the formation of the binding site for dihydropyridines in L-type Ca²⁺ channels (white regions) [57–59]. Compared with other cloned Ca²⁺ channel α_1 -subunits, the rat brain class D cDNA, RB α_1 [60], encodes a 187 kD protein truncated in the carboxyl terminus (arrow). The functional significance of this truncated α_1 -subunit remains unclear, however, as both a second rat brain class D clone (H Kim *et al.*: *Soc Neurosci Abstr* 1991, 17:772), and the human brain [18^{••}] and pancreas [61] class D homologues possess carboxyl segments that extend more than 500 amino acids past the end of RB α_1 .

for generating heterogeneity within the other classes of α_1 -subunits.

A further unexpected result from molecular studies is that two distinct classes of α_1 -subunit encode L-type

Ca²⁺ channels in neurons (Table 1). Overall, the class C and D α_1 -subunits are more closely related to each other (70–76% amino acid identity) than to the class A and B proteins (33–44%). The structural similarity between the class C and D proteins is also reflected in their functional properties. Antisense oligonucleotides against brain class C transcripts selectively block the expression of DHP-sensitive Ca²⁺ channels induced in *Xenopus* oocytes by rat heart RNA [14•]. Furthermore, microinjection of synthetic RNA derived from cardiac and lung class C α_1 -subunits into oocytes results in the expression of high-voltage threshold Ca²⁺ channels that are sensitive to both DHP agonists and antagonists [10,12]. Although the human brain class D α_1 -subunit alone is not functional in oocytes, co-injection of RNAs encoding the class D α_1 -subunit and a brain β -subunit results in the expression of high-threshold Ca²⁺ channels that are also DHP-sensitive [18••]. The class C and D Ca²⁺ channels appear to differ in their current-voltage relations as well as pharmacologically, as the class D channels are partially and reversibly blocked by 10–15 μ M ω -conotoxin [18••]. The class C and D Ca²⁺ channels are co-expressed in several cell lines (Table 1) and it will be particularly interesting to determine whether this co-localization extends to the mammalian nervous system, and if these distinct L-type Ca²⁺ channels make unique contributions to neuronal physiology.

Structurally, class A α_1 -subunits differ significantly from DHP-sensitive Ca²⁺ channels in two hydrophilic regions predicted to be cytoplasmic: the segment separating domains II and III, and the carboxy-terminal region (Fig.1) [19••,20]. Of particular interest is that the class A domain II–III segment is much larger than that of DHP-sensitive Ca²⁺ channels (~430 versus ~130 amino acids) and shows no detectable homology to other proteins. The analogous region of the skeletal muscle α_1 -subunit is involved in mediating excitation-contraction coupling [21], and it will be of interest to determine the functional role of this divergent region in Ca²⁺ channels expressed in the nervous system.

A significant development in the past year has been the demonstration that the rabbit brain class A α_1 -subunit encodes a high-threshold Ca²⁺ channel that is insensitive to both DHP antagonists and ω -CgTx, but is blocked by crude venom from the funnel web spider *Agelenopsis aperta* [19••]. Together with the finding that class A transcripts are highly abundant in the cerebellum [19••,20], these results suggest that class A Ca²⁺ channels correspond to the P-type Ca²⁺ channels first described by Llinas and coworkers [22]. The physiological properties of the rabbit class A Ca²⁺ channel expressed in *Xenopus* oocytes do not correlate exactly with those described for P-type channels in cerebellar Purkinje cells (for a discussion, see [23]). Two pieces of evidence suggest that P-type Ca²⁺ channels may be more widespread than originally thought: first, a high-threshold Ca²⁺ current that is not blocked by ω -CgTx or DHP antagonists has recently been described in a variety of central nervous system neurons [24••,25•]; and second, class A transcripts have been detected in the heart,

pituitary, and several neuronal and endocrine cell lines (Table 1).

Still actively being sought are cDNAs encoding T- and N-type Ca²⁺ channels. A possible candidate for N-type channels is the class B α_1 -subunit. Analogous to class A Ca²⁺ channels, the class B α_1 -subunit possesses a large hydrophilic segment linking domains II and III and also shares >80% amino acid identity in the four domains with class A α_1 -subunits [26•]. Interestingly, there is little detectable identity between the class A and B α_1 -subunits in either their domain II–III segments or in their carboxyl termini. Linking the class B protein to N-type channels are the results that antibodies directed against the class B α_1 -subunit immunoprecipitate radiolabelled brain ω -CgTx binding sites, and that class B transcripts are selectively localized to the nervous system [26•]. Direct evidence of the cloning of an N-type channel awaits the expression of the class B protein, however. At present, there are no leading molecular candidates for T-type Ca²⁺ channels. One possibility is that T-type channels are derived by alternate splicing from one of the classes of α_1 -subunit already cloned. Alternatively, T-type channels may be evolutionary quite distant from the high-threshold Ca²⁺ channels, and cloning may require the use of cell lines enriched for T-type channel expression and cloning strategies using alternative probes.

Subunit composition of neuronal Ca²⁺ channels

Neuronal N- and L-type Ca²⁺ channels are blocked with high-affinity by the *Agelenopsis aperta* peptide toxin ω -Aga-IIIa, indicating that they share some common structural features [27]. Furthermore, recent evidence suggests that neuronal N- and L-type channels are heterooligomeric complexes of roughly similar composition to the skeletal muscle DHP receptor. The purified brain ω -CgTx receptor is composed of a 230 kD α_1 -like protein that binds ω -CgTx and four smaller proteins of between 60 kD and 140 kD [28•]. Two of the smaller proteins are likely to be $\alpha_2\delta$ - and β -subunits as monoclonal antibodies against the skeletal muscle $\alpha_2\delta$ - [29,30] and β - [31•] subunits immunoprecipitate both brain DHP and ω -CgTx binding sites. Indeed, molecular cloning results from a number of laboratories have identified β - and $\alpha_2\delta$ -subunit isoforms that are expressed in the brain [18••,32•,33]. The existence of multiple brain β -subunit isoforms is also indicated by immunoprecipitation results that identify two distinct β -like subunits associated with the ω -CgTx receptor [31•].

Modulation of α_1 -subunits

A number of new studies are providing evidence as to the roles of the other subunits in affecting the functional properties of the α_1 -subunits. Perez-Reyes and co-workers [11] initially demonstrated that the skeletal muscle

α_1 -subunit transfected into mouse L cells resulted in the expression of Ca^{2+} currents that were DHP-sensitive, but which showed an uncharacteristically slow activation. Two independent studies have now shown that the rate of activation and inactivation of the skeletal muscle α_1 -subunit is dramatically increased by co-expression with the skeletal muscle β -subunit [34,35]. Co-expression of α_1 - and β -subunits also increases the number of DHP binding sites in transfected L cells. Surprisingly, further combining the $\alpha_2\delta$ - and γ -subunits seems to actually decrease the number of DHP binding sites found with α_1 and β together [35]. In *Xenopus* oocytes the level of expression of brain class A and cardiac class C α_1 -subunits is significantly increased by co-expression with $\alpha_2\delta$ - and β -subunits [19,32,36,37]. In addition, the $\alpha_2\delta$ -, β - and γ -subunits affect various functional properties of the class C α_1 -subunit, including changes in kinetics and the voltage-dependence of activation and inactivation [32,36,37]. No modulatory effects of the β - and $\alpha_2\delta$ -subunits on the class A channel have been reported [19]. While the modulatory effects of the other subunits seems to depend somewhat on the type of α_1 -subunit and the expression system used, it is clear that both the level of expression and the kinetic properties of Ca^{2+} channel α_1 -subunits are affected by the $\alpha_2\delta$ -, β - and γ -subunits.

Nomenclature

With the increasing number of full-length α_1 -subunit cDNAs isolated from various laboratories, it has become increasingly difficult to keep track of the names assigned to them. The situation becomes even more complex when the large number of partial cDNAs generated by RNA polymerase chain reaction analyses are considered. Some clones are named after the tissue from which the cDNAs were first derived, some are named alphabetically, and some are named numerically. With the more recent demonstration of specific isoforms of $\alpha_2\delta$ - and β -subunits, a further degree of complexity has been introduced into the nomenclature of Ca^{2+} channels. As future molecular cloning studies will undoubtedly define even further Ca^{2+} channel diversity, this may be an appropriate time to adopt a standardized Ca^{2+} channel nomenclature, perhaps analogous to that recently adopted for voltage-gated K^+ channels [38].

Both voltage and ligands can alter the kinetics of Ca^{2+} channels — facilitation

A subset of high-threshold Ca^{2+} channels exhibit facilitation—depolarizing pre-pulses enhance the amount of current evoked by subsequent depolarizations [39,40]. Facilitation involves a change in kinetic behavior of Ca^{2+}

channels such that the facilitated channel exhibits openings of longer duration accompanied by an increased probability of opening [41]. In contrast, in the absence of facilitation the channel is characterized by brief openings. For example, in chromaffin cells facilitation involves a change in the kinetic behavior of a 27 pS channel sensitive to DHPs [42]. It appears that more than one class of high-threshold Ca^{2+} channel may be subject to facilitation, however, as it is also exhibited by Ca^{2+} channels in rat sympathetic neurons sensitive to ω -CgTx [43].

The process underlying Ca^{2+} channel facilitation may be affected by receptor activation as well as by voltage. Neurotransmitters, acting through GTP-binding protein (G protein)-coupled receptors, have been shown to inhibit high-threshold Ca^{2+} currents in both sympathetic and dorsal root ganglion cells [44,45]. This inhibition appears to be due to a change in the voltage-dependence of the Ca^{2+} channel [46], and the agonist-induced inhibition can be overcome by depolarizing pre-pulses [47–49]. The available data suggest that under basal conditions the channels behave as if they had been facilitated by a depolarizing pre-pulse, i.e. they exhibit frequent long-duration openings. Agonist converts the channels to the mode in which openings of short duration predominate, and this can be overcome by depolarizing pre-pulses. Exactly the opposite situation prevails for those Ca^{2+} channels that are enhanced by agonists, i.e. the basal condition appears to be one in which the channels reside in the short-opening mode, and agonist converts them to a long-opening mode [50].

Because agonist-induced changes in Ca^{2+} channel behavior are relieved by depolarizing pre-pulses in a fashion essentially identical to that seen during the process of facilitation, it seems likely that the mode of Ca^{2+} channel behavior in any given cell is under constitutive control. An important question to be addressed is whether this phenomenon is a property of the Ca^{2+} channel complex or some as yet to be identified cellular process. A G-protein-dependent blocking molecule (either the G protein itself or a soluble intracellular messenger) coupled to Ca^{2+} channels has been proposed [46,51]. During differentiation of neuroblastoma cells there arises a tonic inhibition of an ω -CgTx-sensitive Ca^{2+} channel, and the effect appears to be mediated by a G protein [52]. Tonic modulation of Ca^{2+} channels provides a plausible explanation for both voltage- and agonist-mediated enhancement and diminution of Ca^{2+} channel activity.

From a functional point of view, facilitation may contribute to frequency-dependent Ca^{2+} entry in synaptic terminals. Ca^{2+} channels are facilitated not only by single long depolarizing pre-pulses, but also by a series of brief repetitive depolarizations. Moreover, the available data suggest that switching between two gating modes characterized by long- and short-duration openings is fundamental to modulation of Ca^{2+} channels. Dual control over this process by agonists and voltage provides a powerful and dynamic control over Ca^{2+} channel function in neurons.

An unexpected source of Ca²⁺-dependent inactivation of Ca²⁺ channels

It has long been known that voltage-gated Ca²⁺ channels can be inactivated by a rise in intracellular Ca²⁺. Classically, it has been thought that the source of such elevations in intracellular Ca²⁺ is the Ca²⁺ channel itself, an example of negative feedback. Using nystatin-perforated patch-clamp electrodes, which maintain the normal dynamics of intracellular Ca²⁺ levels, Kramer *et al.* [53•] have now shown that agonist-induced release of intracellular Ca²⁺, via production of inositol 1,4,5-trisphosphate, can also result in inhibition of voltage-gated Ca²⁺ channels. Because studies employing the whole-cell patch-clamp technique commonly buffer intracellular Ca²⁺, this phenomenon had been previously overlooked.

Low-threshold Ca²⁺ currents and action-potential generation

The central role of low-threshold Ca²⁺ channels (T-type) in the bursting behavior of neurons has been appreciated for nearly a decade [54]. Now, using a new technique called action-potential clamp, McCobb and Beam [55•] have shown that in dorsal root ganglion neurons the low-threshold Ca²⁺ current is the major Ca²⁺ current responsible for Ca²⁺ entry during a normal brief action potential, and is relatively insensitive to changes in action potential duration. In contrast, the contribution of high-threshold currents to Ca²⁺ entry increases markedly with action potential duration. These studies provide a new and intriguing glimpse into the functional roles of Ca²⁺ channels in neurons.

Perspective

A pressing objective of future molecular studies will be to correlate the functional properties of exogenously expressed Ca²⁺ channel cDNAs with those of Ca²⁺ channels described in neurons. The rather heterogeneous properties of Ca²⁺ channels found in the nervous system may make this task more difficult than it initially appears. In this regard, it may ultimately be necessary to compare the biophysical and pharmacological properties of Ca²⁺ channels found in defined neurons and neuronal cell lines with those of cDNAs isolated from the same cells. Together with the use of specific α_1 -subunit antibodies to determine the cellular and subcellular localization of cloned Ca²⁺ channels, these studies will help address questions concerning the contributions of individual Ca²⁺ channel subtypes to neuronal function and the mechanisms of processes such as facilitation.

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