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Biophysical Characteristics of Mg²⁺ Blockades in Ion Channels and Their Physiological Roles

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Ionic currents through many different cation channels are specifically reduced by internal and/or external Mg²⁺ within a concentration range of physiological relevance. Although there are many ways for a divalent cation to reduce channel currents, the current blockade by directly binding to a conduction pore has been most well studied. Since the binding of Mg²⁺ to a site (or sites) within transmembrane voltage-drop affects the fundamental characteristics of an ion channel, biophysical mechanism and physiological significance of such a blockade have been under intensive investigation.

We have been studied the blockade of two different channels by Mg2+. Ionic currents of cyclic nucleotide-gated (CNG) channels are reduced by sub-millimolar concentration of intra- and extracellular Mg²⁺. In an earlier study, we identified a glutamte (Glu³⁶³) residue within the pore-forming region of the channel as the high-affinity binding site of extracellular Mg²⁺. We have further studied the electrophysiological and structural features of the site. In addition, we constructed several point-mutations of CNG channel at S6 region which is likely to compose of the internal vestibule of CNG channels to reveal the location of the second Mg2+-binding site exposed to internal side of the membrane. In a recent study, we accidentally observed that the current-voltage (I-V) relationship of another channel, small conductance Ca2+-activated K+ channel (SKCa channel), got rectified as the concentration of internal Mg²⁺was raised. The apparent affinity for Mg²⁺ varied as a function of membrane voltage and intracellular Ca²⁺ concentration, and extracellular K⁺ altered the voltage-dependence as well as the apparent affinities of Mg^{2+} from intracellular side. From the results, we tentatively concluded that the inwardly rectifying I–V relationship of SK_{Ca} channels are due to the voltage–dependent blockade of intracellular Mg^{2+} and that the Mg^{2+} -binding site is located within the ion–conducting pathway. We are currently localizing the Mg^{2+} -binding site within the putative pore–forming region of SK channels using site–directed mutagenesis and investigating the functional significance of the I–V rectification of SK_{Ca} channels.