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RIM modulates Ca_VI.3 Ca²⁺ channels

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Calcium channel β subunits (Ca_Vβs) are essential cytoplasmic components of voltage-gated calcium channels (VGCCs) affecting their gating and targeting. Ca_Vβs bind with high affinity to the cytoplasmic loop between transmembrane segments I and II of the α 1 subunit (loop-I-II). To identify new proteins that modulate VGCCs by interaction with Ca_Vβs we performed a yeast two-hybrid screen using Ca_vβ2a as bait. Screening of a human fetal brain cDNA library identified a C-terminal fragment of RIM1α (Rab3-interacting molecule) which contains a highly conserved C2B domain as potential interaction partner. To proof the interaction between RIM and Ca_vβs we developed a protein targeting assay in tsA-201 cells heterologously expressing the loop-I-II of Ca_v1.3 channels with diverse $Ca_V\beta$ subunits. The $Ca_V1.3$ -loop-I-II was transported to the plasma membrane and co-targeted all Ca_Vβ subunits indicating that the $Ca_V1.3$ -loop-I-II and the $Ca_V\beta$ subunits formed a functional complex. The C-terminal fragment of RIM1α or the full-length form of RIM2β exhibited a cytoplasmic distribution but when coexpressed with Ca_Vβs in presence of the Ca_V1.3-loop-I-II both were co-localized at the plasma membrane. Using qualitative RT-PCR analysis we detected various RIM isoforms in the total organ of Corti and RIM2 α in cochlea inner hair cells (IHCs) at an early developmental stage, before the onset of hearing. As RIM is a presynaptic active zone protein involved in Ca²⁺-induced neurotransmitter release, we asked the question whether the association of RIM with Ca_V1.3 could account for the slow Ca_V1.3 cur-

rent inactivation seen in IHCs. In whole-cell patch-clamp analysis of tsA-201 cells using 15 mM Ca2+ as charge carrier the C2B domain containing fragments of RIM1α and RIM2α caused a significant depolarizing shift of the activation-curve of Ca_V1.3 (7–12 mV) and slowed the inactivation of both Ca^{2+} and Ba^{2+} currents (p < 0.05) albeit to a lesser extent as found in native IHCs. To investigate if a slowly inactivating Ca_v1.3 spliceform (1b) could contribute to this effect we examined its expression with RT-PCR analysis. However, we did not detect Ca_V1.3(1b) transcripts in the total organ of Corti at the same developmental stage as we found RIM. Taken together these data showed that indeed RIM modulated Ca_V1.3 Ca²⁺ channels. However, we assume that a mixture of diverse proteins and/or Ca_v1.3 splice variants probably accounts for the slow current inactivation of these channels in native IHCs.

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