

Signaling through BK_{Ca} channels - a view by functional proteomics

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Cellular and fast intercellular signaling mainly occurs through proteins integral to the plasma membrane such as ion channels and G protein-coupled receptors. The respective signalling processes are highly specific and often tightly restricted in time and space - functional characteristics that result from integration of channels and receptors into 'macromolecular complexes' (termed signaling supercomplexes or microdomains). So far, however, membrane protein-associated signalling supercomplexes have largely escaped molecular analysis predominantly for technical reasons arising from their poor solubility and low stability.

Our group has developed a novel approach for isolation of such signaling supercomplexes from plasma membranes and identification and analyses of their individual protein constituents.

In this presentation I will show application of our *functional proteomics* to Ca²⁺ and voltage-activated potassium channels of the BK-type (BK_{Ca}). These channels are fundamental modulators of signaling in the brain by contributing to action potential repolarization, mediating the fast phase of afterhyperpolarization, controlling dendritic Ca²⁺ spikes and establishing a feedback loop between membrane potential and cytosolic Ca²⁺ that regulates the release of hormones and transmitters. I will discuss organisation and working of BK_{Ca} channel complexes and show how they reconstitute 'Ca²⁺ nano-domains' where Ca²⁺ influx through the co-assembled voltage-gated Ca²⁺ channels (Cav) activates BK_{Ca}. Moreover, I will discuss how other constituents fine-tune the operation of BK_{Ca} supercomplexes and thus promote the specificity of their signaling required in different types of cells and distinct subcellular compartments.

Literature

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