

Resolving Phospholipase C Regulation

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Phospholipase C β (PLC β) increases intracellular calcium in response to diverse hormones, regulating numerous processes including cell proliferation and survival. Dysregulation of its expression or activity contributes to heart disease, cancer, and other pathophysiological conditions. PLC β interacts with the cytoplasmic leaflet of the plasma membrane, where it hydrolyzes phosphatidylinositol-4,5-bisphosphate (PIP₂) to produce second messengers that increase intracellular Ca²⁺ and activate protein kinase C. PLC β exists in an autoinhibited state that is relieved via direct interactions with the heterotrimeric G protein subunits, G α_q and G $\beta\gamma$, and the membrane, coupling lipase activity to stimulation of G protein-coupled receptors. Prior structural and functional studies have revealed how G α_q binds to and activates PLC β , but where G $\beta\gamma$ binds to PLC β to regulate its activity is controversial. Using cryo-electron microscopy, we have determined a series of reconstructions of the G $\beta\gamma$ -PLC β complex, allowing identification of the G $\beta\gamma$ binding surface, as well as an unexpected role for G $\beta\gamma$ as a potential scaffold for PLC β , wherein a pre-activated complex is maintained at the membrane.