

The p-bodies and
reinitiate translation
(11). p-bodies
increase



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The inositol phosphate family is an important family of signalling molecules that was discovered in the 1920s and is conserved from yeast to humans. All the members of this family are derived from myo-inositol and are involved in cytoplasmic and nuclear signalling.

The hydroxyl groups of the inositol ring can be phosphorylated in a combinatorial manner. Inositol hexakisphosphate (IP₆) is the most abundant inositol phosphate and is generated from IP₅ by inositol pentakisphosphate 2 kinase (IPPK) (1). In mammals, IP₆ further gives rise to high energy pyrophosphate molecules (2), IP₇ and IP₈, by the action of IP₆ kinases (IP₆K1, IP₆K2 and IP₆K3) and IP₇ kinases (PPIP5K1 and PPIP5K2). IP₆K1 is implicated in a number of cellular processes in mice such as vesicular trafficking (3), insulin secretion (4, 5), male fertility (5), neutrophil phagocytosis (6), blood clotting (7) and DNA repair (8, 9).

IP₆K1 modulates protein functions by mainly three mechanisms: a) IP₇ synthesized by IP₆K1 can pyrophosphorylate proteins, b) IP₇ can directly bind to proteins, and c) IP₆K1, independent of its catalytic activity, can also directly bind to proteins and modulate their functions. Processing bodies (P-bodies) are small membrane-less cytoplasmic foci that are involved in mRNA degradation (10) and storage (11). mRNA in the cytoplasm is targeted to degradation by either exosome mediated 3'-5' degradation or 5'-3' exoribonuclease mediated degradation (12). mRNA deadenylation precedes both these decay pathways (13). All the enzymes involved in 5'-3' degradation pathway are localized to P-bodies (14-19). However, not all mRNA that is localized to P-bodies is targeted for degradation as they can also escape P-bodies and reinitiate translation (11). P-bodies increase in size

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when mRNA decay is inhibited and reduce in size and number if mRNA deadenylation is blocked or if one of the scaffolding proteins is missing (10). P-bodies are involved in mRNA surveillance (20), gene silencing (21), degradation of mRNA containing AU rich elements (22) and translational repression.

P-bodies are dynamic and motile entities that form when the abundance of stalled mRNA increases in the cytoplasm (23). This implies that the size and number of P-bodies depend on the amount of mRNA undergoing de-capping. To enter P-bodies mRNA has to be released from polysomes. Translation inhibiting drugs like cycloheximide that block mRNA on polysomes reduce the number of P-bodies drastically whereas drugs like puromycin that block translation by releasing the mRNA from polysomes induce more P-bodies (10).

The proteins present in P-bodies are generally also diffused in the cytoplasm and there is dynamic flow of proteins in and out of P-bodies. Hence, changes in P-body number and size per cell is related to the cell cycle phase, proliferation status of the cell and availability of nutrients (24). *Ip6k1*^{-/-} male mice are infertile (5). Our lab has shown that IP6K1 protein is indispensable for the differentiation of round spermatids and is required for regulating the assembly of chromatoid bodies during spermiogenesis (Malla et al., unpublished).

A chromatoid body is a dense structure in the cytoplasm of male germ cells and is mainly composed of RNA and RNA binding proteins and is thus a type of RNP granule, similar to P-bodies. It was observed that these chromatoid

bodies are depleted in male germ cells of *Ip6k1*^{-/-} knockout mice and *Ip6k1*^{-/-} mouse embryonic fibroblasts (MEFs) have diminished P-bodies compared to *Ip6k1*^{+/+} MEFs.