

## Short Note on the Role of RNA in Stress Granules

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### DESCRIPTION

Hormones and neurotransmitters activate the  $G\alpha_q$ /phospholipase  $C\beta_1$  (PLC $\beta_1$ ) signaling system eliciting cellular calcium responses. Beside the plasma layer, PLC $\beta_1$  has a cytosolic population that binds stress granule related proteins preventing their collection, and activation of  $G\alpha_q$  shifts PLC $\beta_1$  to the membrane releasing bound proteins and advancing the development of stress granules. The cellular impact of stress granules shaped upon routine  $G\alpha_q$  protein signaling is obscure. Here, we have characterized Ago<sub>2</sub> stress granules formed in response to neurotransmitter activation in cultured PC<sub>12</sub> cells. We observe these stress granules have a distinct protein composition, and unlike stress granules formed upon heat shock, contain just two mRNA transcripts, chromogranin B, which is engaged with secretory capacity, and ATP synthase 5f1b, which is expected for ATP synthesis. Our investigations show a surprising pathway where  $G\alpha_q$ /PLC $\beta$  manages the interpretation of specific proteins.

Binding of extracellular ligands like acetylcholine, serotonin and histamine, to their particular G protein coupled receptor will actuate  $G\alpha_q$ , one of the four significant G proteins pathways.  $G\alpha_q$ , thusly, actuates phospholipase  $C\beta$ , which catalyzes the hydrolysis of the signaling lipid phosphoinositide 4,5 bisphosphate leading an expansion in intracellular calcium. Alongside this significant layer work, PLC $\beta_1$  has been found to have an abnormal cytosolic population that ties to the advertiser of RNA-induced silencing, C<sub>3</sub>PO, as well as a several proteins involved with pressure granules formation. Stress granules are ended ribosomal edifices that safeguard mRNAs under stress conditions, for example, arsenite treatment, heat/cold shock and osmotic stress. Proteins that bind spot PLC $\beta_1$  include  $\epsilon$ FI5A, Polyadenylate Binding Protein (PABPC1) and Ago<sub>2</sub>. Ago<sub>2</sub> is

moreover the nuclease component of the RNA-induced silencing complex (RISC) that degrades mRNA with the assistance of C<sub>3</sub>PO. Whenever Ago<sub>2</sub>-bound mRNA matches impeccably with a bound miR, Ago<sub>2</sub> transitions to an active conformation to hydrolyze the mRNA. In any case, assuming matching is defective, it will frame a slowed down complex resulting in stress granules.

Our new study showed that reducing the cytosolic PLC $\beta_1$  population increases the number and size of particles containing Ago<sub>2</sub> along with the pressure granule makers Polyadenylate-binding protein 1(PABPC1) and G3BP1. Restricting among PLC $\beta_1$  and stress granule proteins assists keep them with scattering, while initiation of  $G\alpha_q$  promotes relocation of cytosolic PLC $\beta_1$  to the plasma film, advancing arrival of bound proteins and the development of stress granules. This system proposes that  $G\alpha_q$  might be associated with protein translation through cytosolic PLC $\beta_1$ .

We have characterized the composition of Ago<sub>2</sub> stress granules formed in response to neurotransmitter activation in differentiated PC<sub>12</sub>, and contrasted these with traditional stress responses. PC<sub>12</sub> cells have a large endogenous articulation of  $G\alpha_q$  and PLC $\beta_1$ , and although not neuronal in origin, when treated with nerve growth factor, the cells take on neuronal morphology and secrete particles mimicking synaptic vesicles. We find that  $G\alpha_q$  activation produces pressure granules that have an unmistakable protein composition when contrasted with different stress. Likewise, not at all like heat shock that contain different mRNA and miRs,  $G\alpha_q$  stress granules contain just two major mRNA transcripts. Our studies show a connection between physiological G protein activation and protein translation.

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