

MOLECULE PAGE

Galpha s

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Version 1.0, June 10, 2011 [DOI: 10.3998/panc.2011.19]

Gene symbols: [GNAS/Gnas](#)

1. General Information

G α_s is the catalytic subunit of one of the first heterotrimeric G proteins identified and is among the best characterized. It becomes activated upon binding GTP and then activates adenylyl cyclase (AC) leading to the production of cyclic AMP (2,5). Because of the widespread use of cAMP as a signaling molecule the activating pathway plays a role in many functions including development, muscle contraction, learning and memory and endocrine and exocrine secretion. G α_s is expressed in almost all cells and is activated by beta adrenergic, dopamine, H₂, secretin, VIP, TSH, LH and GLP-1 receptors among others. These receptors are all of the 7 transmembrane domain family and when liganded act as a guanine nucleotide exchange factor (GEF) for G α_s and accelerate the release of GDP. The α_s subunit can then bind GTP, dissociate from its associated $\beta\gamma$ complex and activate all known isoforms of AC. G α_s has an intrinsic rate of deactivation by hydrolyzing the bound GTP. Whether this can be accelerated by its effector acting as a GTPase activating protein (GAP) is

unclear. The GDP bound form can also be activated by aluminum fluoride (5).

G α_s is coded in humans and mice by a gene with 13 exons that can generate multiple gene products through 4 alternative promoters and first exons (21). Two of these are G α_s which is broadly expressed and an N terminal extended version XL α_s expressed primarily in neuroendocrine tissues. The gene is imprinted in a tissue specific manner with G α_s being expressed primarily from the maternal allele. The protein G α_s is a 45 KDa protein which is postranslationally modified by N-terminal palmitoylation which targets it to the plasma membrane. G α_s is alternatively spliced in exon 3 to produce a long and short form differing by 14 amino acids (2) which function similarly. G α_s is also located on intracellular membranes and may play a role in membrane trafficking. This may involve additional effectors beyond AC.

The structure of G α_s has been solved both while binding GTP γ S or GDP plus AIF and in combination with its effector AC (17,18). Like other G α subunits, G α_s is made up of a ~220 amino acid Ras like GTPase domain which

includes the sites for GTP binding and effector interaction and a ~120 amino acid alpha helical domain that helps form a pocket for guanine nucleotides. Binding of GTP leads to structural movement in several switch regions. Mutations in specific amino acids in these regions can lead to permanent activation or inactivation. Cholera toxin (CT) catalyzes the ADP ribosylation of Arg 201 leading to constitutive activation of $G\alpha_s$ by blocking the GTPase turn off mechanism. An activating mutation ($G\alpha_s$ Q227L) exists similar to Q to L mutations in other G proteins.

Mutations in $G\alpha_s$ have been linked to a number of diseases (8,12). Inactivating mutations in $G\alpha_s$ are associated with the inherited disorder, Albright's Hereditary Osteodystrophy or pseudohyperparathyroidism with the resulting syndrome affected by whether the mutation is on the maternal or paternal allele. Activating mutations are associated with pituitary or thyroid adenomas and the McCune-Albright syndrome. Some of these mutations have been reproduced with mouse models (21). Whole body knockout of $G\alpha_s$ is embryonic lethal and heterozygotes show reduced viability. Floxed $G\alpha_s$ mice have been generated and used in a tissue specific manner by the Weinstein group to delete $G\alpha_s$.

2. Specific Function in the Pancreas

$G\alpha_s$ has been observed by immunohistochemistry to be expressed at high levels in mouse islet beta cells and at lower levels in surrounding acini (22). Within acini, $G\alpha_s$ is localized to the plasma membrane and to a lesser extent intracellularly in the Golgi region (3,9). Isolated rat zymogen granule membranes were reported to contain $G\alpha_s$ by Western blotting in one study (10) but not to be present in another study (9). ADP ribosylation studies have labeled multiple forms of the protein in response to CT in acinar cell membranes (14) and in AR42J cells (7). It has also been identified in rat parotid gland membranes (1,20). We are

not aware of similar studies on pancreatic ductal epithelium or in stellate cells.

Functional studies of $G\alpha_s$ have mainly used CT to activate the G protein. CT increases AC activity in pancreatic membranes (6,19), increases cAMP in dissociated acini (4,11,15) and slightly increases in amylase secretion (4,11,15). In the perfused cat and rat pancreas, CT stimulates bicarbonate rich fluid secretion (6,16).

Activation of $G\alpha_s$ can also be carried out by overexpression of $G\alpha_s$ (Q227L) mutant (the long form) either by plasmid or adenoviral vector (13) which increases cyclic AMP in acini. This overexpression of active $G\alpha_s$ did not affect the activation of RhoA or Rac1 and did not affect acinar morphology.

The importance of $G\alpha_s$ can also be assessed by tissue specific knockout studies. Xie et al used a Pdx1-Cre to delete $G\alpha_s$ throughout the pancreas (23). Most of the findings were due to effects on the islets similar to earlier study with Beta cell deletion using Rat insulin-Cre (22) but in addition the pancreas weight was larger than normal and exocrine histology was stated to be altered. More definitive analysis will require deleting in acinar or duct cells independent of the islets.

3. Tools to Study $G\alpha_s$

a. cDNA clones

cDNA clones for human GNAS are available from the [Missouri S & T cDNA Resource Center](#) for both the short and long forms of human GNAS including wild type, Q to L activating mutations, and an internal Glu-Glu epitope tagged version.

b. Antibodies

[Biocompare](#) lists 45 commercially available $G\alpha_s$ antibodies. We are unable to provide a recommendation of which ones work.

c. Viral Vectors

A constitutively active G α_s Q227L mutant has been prepared and used by us in mouse pancreatic acini (13).

d. Mouse lines

Whole body gene deletion is embryonic lethal. A G α_s with floxed exon 1 has been constructed by the laboratory of Lee Weinstein and used to delete G α_s in osteoblasts, liver, kidney, and islets (21).

4. References

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