

Gene Ontology (GO) Extensions

A. Gene Summary Page GO Display

Gene Ontology Annotations for *ctnnA*

Molecular Function

beta-catenin binding **binds** *aarA*
beta-catenin binding **binds** Q02248
actin filament binding

Biological Process

mitotic cytokinesis **genetically interacts with** *mhcA*
protein secretion
sorocarp stalk development
culmination involved in sorocarp development
actin filament bundle assembly
positive regulation of cytoskeleton organization
centrosome localization
Golgi localization
establishment or maintenance of bipolar cell polarity
cellular protein localization **localizes** *dcsA* **during** sorocarp stalk development
cellular protein localization **localizes** *rgaA*, *ctxA*, *mhcA* **during** sorocarp stalk development
sorocarp stalk morphogenesis

Cellular Component

cell cortex
cell-cell junction
basal cortex **during** epithelial cell development **during** sorocarp stalk development

Annotation Extensions: Recently, the GO consortium developed annotation extensions, to add deeper information such as ‘under what conditions’ or ‘during which developmental stage’, a process, activity or localization occurs. This results in annotations that form sentence-like structures, providing a more complete answer to biological questions.

Left: GO annotations displayed on the Gene Summary page. This example depicts a gene, which contains **manual** annotations in each of the three GO aspects. When manual annotations are available, only those are shown on this page. See below for all GO annotations. The relations of the annotation extensions (such as **binds**, **during** **regulates**, **at**) are marked in black and bold.

B. GO Page Display

Gene Ontology Annotations for *p2xA*

| All GO | Manual GO | Experimental GO | Electronic GO | | |
|---|-----------|----------------------|------------------------------------|------------|--------|
| Biological Process | | | | | |
| GO term + Extension | Evidence | With | Reference | Date | Source |
| ATP-gated ion channel activity | IMP | | Sivaramakrishnan & Fountain (2012) | 21-02-2014 | DDB |
| calcium ion transmembrane transport requires ATP | IGI | p2xA p2xC,p2xD,p2xE | Sivaramakrishnan & Fountain (2012) | 21-02-2014 | DDB |
| cation transport | IBA | | Gaudet et. al (2010) | 01-03-2011 | GOC |
| cation transmembrane transport | IEA | UniProtKB-KW:KW-0406 | GO_REF:0000037 | 25-04-2015 | SPKW |
| cellular hypotonic response | IGI | p2xA,p2xC'p2xD,p2xE | Ludlow et. al (2009) | 14-03-2014 | DDB |
| ion transport | IDA | | Ludlow et. al (2009) | 10-03-2010 | DDB |
| negative regulation of GTPase activity regulates rab11A | IDA | | Parkinson et. al (2013) | 22-04-2014 | DDB |
| positive regulation of vesicle fusion at plasma membrane | IDA | | Parkinson et. al (2013) | 22-04-2014 | DDB |
| regulation of calcium-mediated signaling | IGI | p2xA,p2xC'p2xD,p2xE | Ludlow et. al (2009) | 14-03-2014 | DDB |
| response to ATP | IEA | InterPro:IPR001429 | GO_REF:0000002 | 25-04-2015 | IPRO |

Display of all annotations on the GO page. Only the Biological Process aspect is shown. At the top are tabs that allow easy filtering of annotations based on evidence code classes. This view shows a mixture of experimental annotations (IMP, IGI, IDA), manual annotations (all of the above plus IBA), and electronic annotations (IEA). GO terms and annotation extensions are found in the first column. In general, annotation extensions are associated with experimental annotations.