



GO On, Tell Us What You Think: Community Input into the Gene Ontology

Jennifer I. Clark, Amelia Ireland, Jane Lomax, Midori A. Harris and the Gene Ontology Consortium.
EMBL Outstation - European Bioinformatics Institute, Hinxton, Cambridge CB10 1SD, UK

We welcome **questions, requests** and **ideas** from our users.

I need terms to cover mesoderm development.

How do I browse the GO annotations?

How do I get a version of GO that includes only terms relevant to prokaryote research?

How do I annotate the function of a gene product that acts as part of a complex?

Participate in GO's development

Submit your requests and comments online

geneontology.sf.net

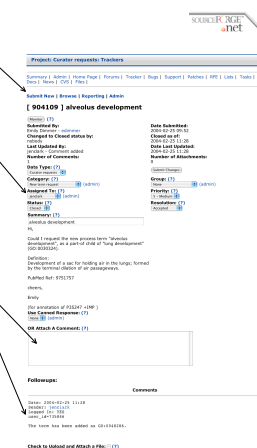
All development issues in the Gene Ontology are discussed on a **publicly accessible** internet discussion forum. Anyone may comment on each item, to ask **questions**, contribute **information** or suggest **solutions** to problems.

You can submit requests for:

- **Software improvements** - AmiGO + DAG-Edit
 - *Would it be possible to export/save the history file as text so it can be used to generate log files?*
- **New or better GO terms**
 - *I'd like terms to cover mesoderm development.*
 - *The definition of GO:0046530 does not apply to the species I work on.*
- **Help with GO annotation issues**
 - *Is antibody cross-reactivity annotated with the ISS evidence code?*
 - *How is the 'NOT' qualifier used?*
- **GO integrity checks**
- **GO website improvements**
 - *Please can you add more detailed information on the synonym syntax in the OBO file format?*

What happens when I submit a request?

1. Click **'Submit New'** to make a request.
2. The request is taken up by a named **curator** who will take responsibility for the request throughout the process.
3. Possible solutions are discussed and anyone may comment. Read other people's comments lower on the page.
4. Each time a comment is added an **e-mail update** is sent to everyone involved.
5. A mutually acceptable **solution** and implementation date are **proposed**.
6. If no problems are found then the plan will be **implemented** by the curator.



Can I meet Consortium members and GO users in person?

Yes: attend a **GO Users Meeting**. These meetings are open to anyone interested in the GO project, and provide opportunities for GO users and developers of GO-related analysis and visualization tools to share their work with each other and with GO Consortium members.

Next Meeting: **October 14, 2004;**
Northwestern University, Chicago, IL, USA.
http://dictybase.org/GOmeeting/GO_Users_Meet.html

Can I ask questions by e-mail?

The consortium has moderated e-mail lists for questions on:

- **database** issues (go-database@mail.fruitfly.org),
- **content** issues (go@geneontology.org),
- **annotation** issues (annotation-request@geneontology.org)
- **announcements** (lower traffic - gofriends@geneontology.org)

Lists are archived, and all but the GO-database archive can also be searched using Google. To subscribe see www.geneontology.org/GO.contacts.html.

Contact GO

www.geneontology.org