GO Editorial Office Report — March 2006

1. GO Staff

Full-time staff unchanged: Midori Harris, Jane Lomax, Amelia Ireland, Jennifer Clark. From January–July 2006 we have a visiting student, Erika Feltrin (see below).

2. Annotation Progress

Not applicable

3. Methods of Annotation

Not applicable

4. Ontology development

Summary table:

	March31, 2005	March 23, 2006	New	%change
function	6949	7399	450	+6.5
process	9263	10517	1254	+13.5
component	1484	1688	204	+13.7
total	17696	19694	1908	+10.8

Note: the table does not include obsolete terms (987 in March 2005; 1000 in March 2006), whereas reports at previous meetings have included obsoletes in the summary table.

Content highlights:

- Draft of cellular component ontology with complete is_a paths (see poster)
- Progress towards integrating cell type ontology (CL) terms into GO
 - changes to GO term names (as well as CL terms) to reconcile GO and CL
 - reviewing OBOL output of proposed cross-product terms
- New effort to improve transporter terms in molecular function (see poster)
- Content meeting in November 2005 led to proposals in immunology (with lots of work form Alex Diehl and the immunology experts who participated), self/

- non-self processes, interaction between organisms, and cell death. All are under consideration; also see poster on two-taxon annotation.
- Many additions and changes to improve MetaCyc and GO consistency (includes metacyc2go mapping)
- Several development topics, e.g. neurogenesis, metamorphosis, embryonic morphogenesis, etc.
- Additions and changes in cellular component for consistency with MetaCyc component ontology

5. Publications/Tutorials/Presentations with substantial GO component

Publications

- Clark JI, Brooksbank C, Lomax J. 2005. It's all GO for plant scientists. Plant Physiol. 138(3): 1268–1279.
- Harris MA, Lomax J, Ireland A, and Clark JI. 2005. The Gene Ontology project. In Encyclopedia of Genetics, Genomics, Proteomics and Bioinformatics, Part 4, Bioinformatics. S. Subramanian (section ed.), Wiley & Sons, Inc., New York.
- Lomax J, The Gene Ontology Consortium. 2005. Get ready to GO! A biologist's guide to the Gene Ontology. *Brief. Bioinform.* **6:** 298-304.
- The Gene Ontology Consortium. 2006. The Gene Ontology Project in 2006. *Nucleic Acids Res.* **34**: D322–D326.

Tutorials

- HGMP Microarray Workshop, Cambridge, UK. Jane Lomax & Amelia Ireland, March 2005
- One day course on GO Bioinformatics Masters, Instituto Gulbenkian de Ciencia, Lisbon, Portugal. Amelia Ireland, May 2005
- One-day workshop on GO, Plant & Animal Genomes XIV, San Diego, CA. Jen Clark, John Day-Richter, Harold Drabkin & Rama Balakrishnan; January 2006
- GO tutorial on annotation and microarray analysis at MUGEN course, Milan. Jane Lomax, February 2006
- Introduction to GO and AmiGO walkthrough. Computing Workshop, European Fission Yeast Meeting, Hinxton, UK. Midori harris & Jane Lomax, March 2006

Presentations

 The Gene Ontology. Wellcome Trust Advanced Course, Cambridge, UK. Jane Lomax, June 2005.

- GOing forward: Refining the content of the Gene Ontology. ISMB 2005, Detroit, MI. Midori Harris, June 2005.
- The Gene Ontology. MetaboMeeting 1.0, Cambridge, UK. Jane Lomax, July 2005.
- Does your species show in GO? Xenopus tropicalis Annotation Jamboree, Hinxton, UK. Jen Clark, July 2005.
- Does your species show in GO? 1st European Farm Animal Functional Genomics Workshop, Edinburgh, UK. Jen Clark, September 2005.
- Developing and Using the Gene Ontology. GENIA workshop on Text mining, Ontologies and Natural Language Processing in Biomedicine, manchester, UK. Midori Harris, March 2006.

also several posters (e.g. ISMB, Biocurator, etc.)

6. Other Highlights

- · New interest groups: neurobiology, farm animals
- mips2go & multifun2go mappings updated; work on mapping IMG to GO
- Progress in GO annotation outreach; about 110 groups contacted
- GO user survey conducted, and results analyzed
- New, improved GO bibliography page with search and configurable display (and update script); also, now up to 799 entries; new categories proposed: use of GO in clinical applications; use of GO in network modeling and analysis; use of GO in comparative genomics and evolutionary analysis; use of GO to support predictions (i.e. predicting something other than GO annotations)
- · Restyled web pages
- · Additions to documentation for specific content topics
- OBO-Edit working group formed, with John Day-Richter and several MOD curators (see poster)
- Participation in AmiGO working group
- Visitors:
 - Mikel Egaña Aranguren, University of Manchester (Ph.D. student), September 2005. Work examining compositional structure of GO terms, GO relationships, OBO /OWL conversion (follow-up to GONG project), work with Chris Mungall on GO /CL integration.
 - Erika Feltrin, University of Padova, Italy (Ph.D. student), January–July 2006. Work on ontology development in neurobiology, annotation of UniProt entries (e.g. glutamate receptors); will probably also work on some signaling pathways in GO, and will look at relevant bits of Reactome.
- GO Users Meeting at MGED8, Bergen, Norway



