

# Summary of Action Items from GO Consortium meeting: 23-24th September 2007

## Summary of Action Items from Sunday 23<sup>rd</sup> September

1. Tutorial on wiki discipline (assigned to: Jim Hu).
2. Look at and comment on outstanding items regarding the Regulates relationship: [http://gocwiki.geneontology.org/index.php/Part\\_3](http://gocwiki.geneontology.org/index.php/Part_3) (assigned to: ALL)
3. Check whether there should be a relationship between pigment metabolic process and pigmentation (assigned to: David H)
4. A reference to the annotation SOP pages should go in next newsletter. (assigned to: Jen D)
5. Add a link from outreach to annotation SOPs (assigned to: Jen D)
6. Investigate why terms requests aren't coming in, do we need things we need to do to make it easier? Regarding the SF tracker list and annotation list - who are on these lists and do other people need to be on those lists? (assigned to: Jen D)
7. Michael has sent URL for getting a ISSN for the GO Newsletter to Eurie, who needs to act on this. (assigned to: Eurie H)
8. e-mail Ben if you are not getting a gp2protein check for your database (assigned to: ALL)
9. Reactome annotations should be available from GO by the next GO Consortium meeting. Add new evidence code EXP for 1:1 Reactome to literature, add all other Reactome with TAS to Reactome source. (assigned to: Reactome, Chris M, Alex D, Jen D and Ruth L)
10. Convert Reactome complex terms to GO terms (assigned to: Reactome, Chris M, Alex D, Jen D and Ruth L)
11. Jen D to do a pilot project with a minimal set of terms, as an experiment and bring back results for next GO meeting (assigned to: Jen D)
12. Make difficult sensu terms organism specific (biologist intuitive) (i.e plant vacuole, fungal vacuole). However GO definitions will still be designed to be formal, not depending on species to define the term (assigned to: David H)

## Summary of Action Items from Monday 24<sup>th</sup> September

13. David Hill to organize Webex meeting to ensure all editors understand what they need to do when inputting cross-product information. (Assigned to: David H)
14. OBO file renaming. Wiki: [http://gocwiki.geneontology.org/index.php/Versioning\\_Proposal](http://gocwiki.geneontology.org/index.php/Versioning_Proposal) On the best practices page:

[http://gocwiki.geneontology.org/index.php/Best\\_Practises](http://gocwiki.geneontology.org/index.php/Best_Practises) (assigned to: Chris M and John D-R)

15. work on specification needed for new Amigo features (assigned to: Midori H)
16. Deploy the part that created SF items based on a friendly webform, and would like to see an OBO format in the SF item (assigned to: David H, Midori H, Seth C)
17. ORB: Make link to how to make a perfect GO term from the term request tool (assigned to: Seth C)
18. link to GOOSE from front page (assigned to: Amelia I) DONE
19. write Gene Association file filter script to split files into two – manual and electronic annotations (assigned to: Mike C)
20. Create more advanced interface to download custom gene association files by versioning (assigned to: Chris M)
21. Predictive Activities. Collaborations with external groups. Reports into next GOC meeting as to these kinds of activities. (assigned to: ALL)
22. Finalizing proposed evidence code documentation – abbreviated version on web pages and more detailed on GOC Wiki should be created – separate documentation for users and curators (assigned to: Evidence code working group)
23. Querying communities on awareness of evidence codes – do they know what they are, what do they can use them for? Also regarding the proposal of expanding evidence codes – we could then get a feel for what would be of benefit? (assigned to: Outreach group and Rama)
24. Evidence working group to make new evidence code hierarchy in the context of what has been discussed this afternoon (assigned to Evidence code working group)
25. Evidence code Revise evidence code documentation so that a mutation in only one gene can only be IMP (protein localization IGI example) (assigned to: Evidence code working group)
26. Curators should check whether you have used IGI in this manner and update annotations (assigned to: ALL curators)
27. Provide an evidence code proposal in the context of what was discussed in the Evidence code session on the Monday afternoon (assigned to: Rama , Sue R, Michelle G-G)
28. Update evidence code decision tree in response to today's discussion on evidence code usage (Jen D and Evidence code working group)