Reference Genome Project: Plan for June-Dec 2010

| | June | July | August | September | October | November | December | | |
|---|---|--|--------|---|---------|-----------------------------|----------|--|--|
| PAINT (Ed, Suzi, Paul) | GO camp | Software improvements depending on how much staff we can get. Many suggestions for improvements are at: http://sourceforge.net/tracker/?group_id=184610&atid=1126623 Priorities have not yet been discussed. Paper describing PAINT | | | | | | | |
| Protein | | Version x of Panther should be available [date?] with 82 species | | | | | | | |
| Families | | This should roughly double the number of PAINT-generated ISS annotations | | | | | | | |
| MOD curation | | Wnt pathway primary annotation (Varsha) | | Transcription factors prime from Varsha's list) * Note that this is depended ontology WG – plan to fin progress is being made: http://wiki.geneontology.com/ | | | | | |
| Protein family annotation | | Mismatch repair tree annotation | | Protein N-linked glycosylation | | Wnt pathway tree annotation | | | |
| Curation report (Kara, Stan, Seth, Chris) | | | | | | | | | |
| Papers | | PAINT Phylogenomics: description of the rationale for doing annotations using phylogenetic trees; how we deal with inconsistencies in the tree; dealing with 'NOTs'; dealing with rapid divergence of sequences Reference Genome 'end product': annotations: annotation guidelines; show data from a few families (need to decide on the scope) compare results with other pipelines such automated ISO and automated triangulation; QC. | | | | | | | |
| New targets | Required groundwork: ontology review identify experts define the scope of the pathway to annotate: start- and endpoints, genes involved, PANTHER families; review and fix topology of PANTHER families get approval from ref genome group | | | | | | | | |