WG:Advisory Board Meeting 2012-12-04

From phenoscape

Agenda

- 1. Introduction (Paula Mabee)
- 2. Project updates
 - Curation workflow (Hong Cui, Jim Balhoff)
 - Ontology development (Melissa Haendel, Peter Midford)
 - Phenotype curation (Alex Dececchi, Monte Westerfield)
 - Homology reasoning (Hilmar Lapp)
 - Semantic similarity (Todd Vision)
 - Outreach (Nizar Ibrahim and PM)
 - Knowledgebase development (JB)
- 3. Upcoming plans (Todd Vision)
- 4. Q&A and discussion (Board)

Minutes

Held by conference call (audio) and Adobe Connect (slideshow screen-shared).

- Advisory Board members present: John Day-Richter, Brian K. Hall, Cyndy Parr, Paul Schofield, Peter Vize, Alan Ruttenberg
- Phenoscapers present: Paula, Peter, Terry, Jim, Aaron, Hilmar, Todd, Alex, Melissa, Monte, Judy, Hong, Nizar
- Not present: David B., Paul S., Chris M., Wasila D.

1. Introduction (Paula Mabee)

- Self introductions of Advisory Board (AB) members (new advisory board member, Cyndy Parr) and new project team members (postdoc Prashanti Manda, to start Feb 2013)
- Recap of Phenoscape history, goals; Spring 2012 AB recommendations to be addressed here by project team members

2. Project updates

- Curation workflow NLP (Hong Cui)
 - (Schofield) How are you dealing with atomic terms (such as unossified) to turn this into an EQ expression to pull this apart on an automated basis? Some terms may only be able to be done through human curation.
 - Hong Indeed this is rather difficult. Taking cues from how humans curate these terms.
 - (Alan) A question is where such pulled apart information is kept? Typically you would have a lexicon to capture information about both automatically and humanly determined composite terms, and then the parsing code would use the lexicon as a resource.
 - Hong That is right. Keeping this knowledge in one place is important. We will develop a rule-bank like module to hold those heuristic rules, so if some rules

change, the algorthm/CharaParser will act accordingly.

• Curation workflow - Term broker (Jim Balhoff)

- (Day-Richter) Where to find documentation (more detailed technical specifications) about this? For example, use of backed and authentication technology.
 - Jim Doesn't exist not much that's worth pointing to will take this as an action item.
- (John) Interested in backend; will contact Jim via email

■ Ontology development - Anatomy (Melissa Haendel)

- (Parr) How large is the community around the ontology curation tool development.
 - Melissa Really just looking at 4 people for curating the phenoscape-ext.owl file, but UBERON community overall is much larger (~50 people submitting terms/editing the ontology)
- (Alan) Congrats on adopting OWL2/P4 successfully.

Ontology development - Vertebrate Taxonomy Ontology (Peter Midford)

- (Ruttenberg) Should make equivalence axioms rather than cross-references for NCBI terms that are trimmed out due to redundancy.
- (Parr) Let's talk later about ways to leverage EOL and Global Names, etc. for taxonomy ontology (include Jim)

Phenotype curation - Evolutionary Phenotypes (Alex Dececchi)

- (Ruttenberg) Are the PATO term additions available in one place? PATO may need some rethinking. Would be willing to look into for where our terms should best go.
 - Some on PATO tracker; some on the ORB (adding to PATO pending)

■ Phenotype curation - Model organisms (Monte Westerfield)

- (Ruttenberg) Some of the EQs looked like they were only partial for mappings of MP to EQ.
 - There are indeed some MP terms that are not completely described by their EQ mappings, though they should suffice for Phenoscape's purposes.
- (Schofield) How about issues with sets of digits? How far have you got with mapping HPO for limb phenotypes - this has proven to be very difficult. Is UBERON being used for all of these mappings?
 - There haven't been issues with sets of digits for MP. For HPO mapping to logical definitions, this is on the agenda for the LAMHDI project in the near future, but hasn't been tackled within Phenoscape yet.
 - Melissa-automate pipeline, uberon in combination with logical combination to decompose HPO. Have on LAMHDI agenda to help with HPO with logical definitions
 - Monte-we have embraced the human model organism, but besides what Melissa has discussed we have not addressed it very much

■ **Homology reasoning** (Hilmar Lapp)

- (Hall) Homologizing elements within a limb; are we homologizing between fore and hindlimb? How are we doing this, serial homology?
 - yes, we are distinguishing between phylogenetic and serial homology. Elements within a limb or between limbs of the same organism would, where appropriate, have serial homology asserted.
- (Ruttenberg) What does it mean that there needs to be an OWL model as well as one in RDF? I would think it one or the other?
 - For now the OWL model would be directly loaded into the (RDF-based) KB. It's still possible that the OWL will need to be transformed for better KB performance, but not expected as of now.

- **Semantic similarity** (Todd Vision)
 - (Schofield) Would appreciate more detail (perhaps before the April meeting) on the medianIC and the semantic similarity reasoning and scoring.
 - Todd will share this.
- Outreach (Nizar Ibrahim and PM)
 - (Hall) Are there plans for the coming year to attract more students through the outreach program?
 - Won't have more students at U Chicago due to equipment and resource constraints.
 Same at USD.
- Knowledgebase development (JB)
 - (Schofield) What is pre-reasoning?
 - We create fairly complicated OWL expressions, which can be difficult to query with a simple RDF reasoner. Therefore, create some hierarchies of subsuming class expressions in advance.

3. Upcoming plans (Todd Vision)

No comments/questions specific to this.

4. Questions from the Advisory Board/Discussion (Board)

- (Hall) Have we considered capturing data on gene regulation for model organisms rather than just gene expression? (noted in literature that gene regulation in medaka differs from zebrafish)
 - Monte This is question for MODs. Gene regulation and regulatory element data are beyond the resources of the MODS to capture or curate at the moment. Note that Gill Bejerano, one of the first people to develop automatic methods for finding regulatory elements in genomes, is attending the Feb Phenoscape workshop.
- (Schofield) Impressed with how the project has been addressing the feedback from the AB. Concern was primarily on "pushing the meat", and that's what seems to have a lot of forward movement. Utility is 90% of finding ways for sustainability.
- (Hall) Integrating the Model organisms and the annotations required a huge amount of work.
- (Day-Richter) How are we thinking about creating communities of users and incentivizing contribution. What measures have you taken so far. Have we received annotations from outside? Will send more questions by email.
 - Todd Having all curation centralized benefits quality as MODs show, but obviously isn't very scalable. We haven't tried to crowdsource annotations yet. The attribution of curators by their ORCIDs is the beginning of our thinking about nano publications, but going beyond that is still pie in the sky.
 - Paula We have people approach us wanting to curate data. However, curation process and adding terms to the ontologies is still rough for people who aren't well trained first. Not suitable yet for naive curators to jump in and curate.
- (Parr) Would encourage to think about where we are with respect to MODs and what that can tell us about scaling. Are our current methods really going to be working a few years down the road.

Action items

- Jim add system documentation for ontology broker (Alan)
- Alex, Paula: Send Alan R, the PATO term requests added + ORB request link
- Todd: share medianIC methodology/results with Paul
- Jim add documentation for OWL reasoning system prior to next advisory board meeting
- Terry: collect the cases that are a little more difficult -so we don't lose that experience (Alan)
- All: Try google hangouts! (John)

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