

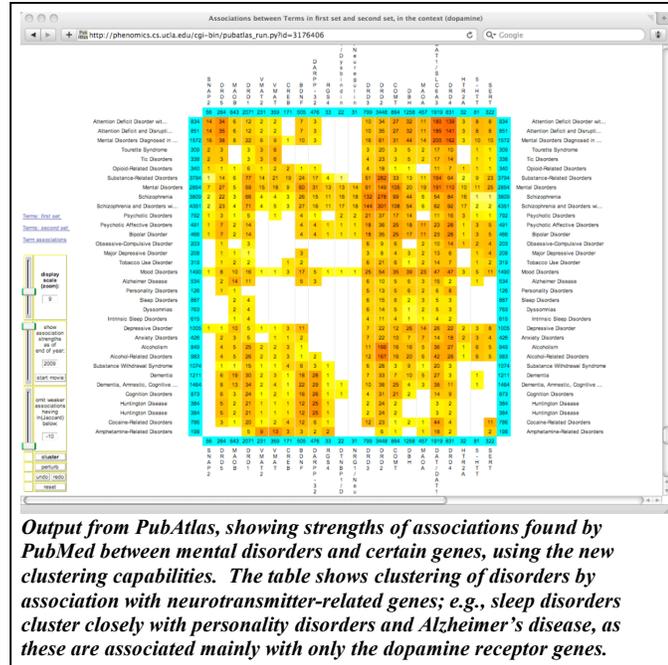
The Hypothesis Web
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<http://www.hypothesisweb.org>

Project Goals

The central goal of the Hypothesis Web project within the Consortium for Neuropsychiatric Phenomics is to aid in the development of interdisciplinary hypotheses spanning multiple disciplines of neuroscience. The Hypothesis Web itself is a software platform that permits collaborative formulation of complex scientific hypotheses, so that:

- hypotheses are the focus of collaboration.
- literature, data, and annotations are combined within a single system.
- interdisciplinary research groups work together to develop a web site for a given hypothesis that organize a group's accumulated evidence.
- resulting hypothesis web sites can be shared, modularized, and published.

The Hypothesis Web is to be used for collaborative development of hypotheses within the other CNP projects, particularly as regards the development of phenomics as a discipline. This is an important part of the effort because understanding the neuropsychiatric syndromes under study at the CNP require transdisciplinary collaborations.



Progress over the period June 2008 – May 2009

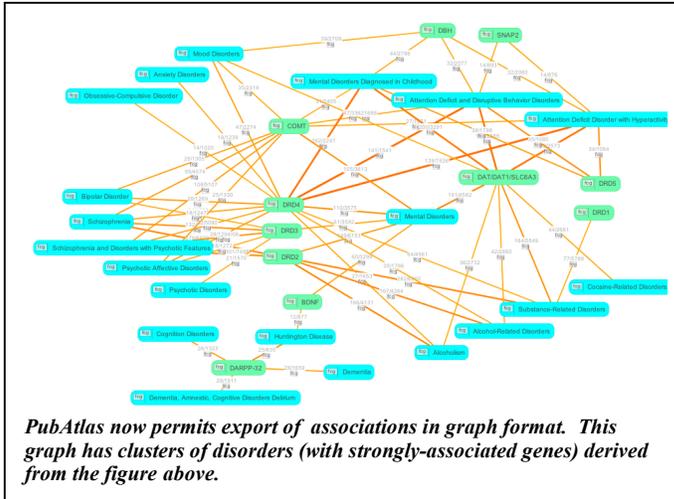
We have made substantial progress for the Hypothesis Web effort in the following areas:

- Expansion of *PubAtlas*, a tool (and website) that permits high-level views of the PubMed literature, giving perspective on concepts and their associations.
- Initial deployment of *HyPoint*, a tool (and website) for constructing multi-level graph representations of hypotheses, with links to the scientific literature, and exploration of hypotheses related to CNP projects.
- Development and refinement of multiple tools for text mining and literature mining, including management of controlled vocabularies (lexica) used in *PubAtlas* and *HyPoint*.
- Formalization of roles for informatics to play in phenomics.

These software components are building blocks of the Hypothesis Web, facilitating collaborative development of hypotheses, and in managing associations between multiple levels of transdisciplinary science.

PubAtlas

Over this period we extended *PubAtlas* (<http://www.pubatlas.org>), the web service we initiated last year. *PubAtlas* provides a visual interface to PubMed, and can be viewed as a “BLAST for PubMed”, extending the PubMed interface to give a high-level view. The main extension was a new architecture that stores PubMed information in a central knowledge base, permitting many new services to be added over time. We extended *PubAtlas* to support more forms of association analysis, and in particular included features for client-side interactive biclustering, allowing users to regroup the data in ways that makes it easier to spot patterns of interest. We also included services for downloading results in different formats (including spreadsheet and lexicon), and saving the state of a *PubAtlas* interaction for later analysis. This allows embedding of links to *PubAtlas* results to be included directly in other documents; for example http://phenomics.cs.ucla.edu/cgi-bin/pubatlas_run.py?term_collection_1=CNP_people&intersection_threshold=10 gives a dynamic window on collaborations within CNP. This particular link also shows how *PubAtlas* makes it possible to construct “semi-automated review papers” that summarize the literature relative to a given vocabulary – while remaining both thorough (covering all publications in PubMed) and always up-to-date.



Several services were added to provide alternative kinds of literature map – particularly graph layout maps using the VUE hierarchical graph editor, giving it capabilities of the earlier PubGraph system. For example, the graph figure here shows a graph literature map produced by PubAtlas for the genotype/phenotype associations in the figure above. Edge thickness and color intensities reflect the degree of co-occurrence, as in PubAtlas. Other literature map formats and analytical tools are planned.

One application of PubAtlas has repeatedly generated interest – mapping of associations between researchers and other sets of topics (including other sets of people). PubAtlas now includes lexica for all large neuroscience groups at UCLA, and we plan to extend it

with more services specifically for formulating queries about researchers.

We presented PubAtlas at the March 2009 *AMIA Summit on Translational Bioinformatics*, and updated the PubAtlas site. In late April 2009 we will be running on a larger multicore server to handle increasing traffic.

HyPoint

Over this period we developed an initial version of HyPoint (<http://www.hypoint.org>), a new web service that provides a means for translating a high-level specification into a hypothesis graph. This version is a derivative of PubAtlas – it takes a lexicon of concepts and associations, and derives a VUE graph representing them. A prototype graph developed for the Gpr6 hypothesis developed within the CNP is shown in the figure below.

Literature Mining

We have made progress in representation of semantic knowledge used in CNP hypotheses. Specifically we have results in *lexicon cleaning, assertion extraction, query extraction, and document retrieval*.

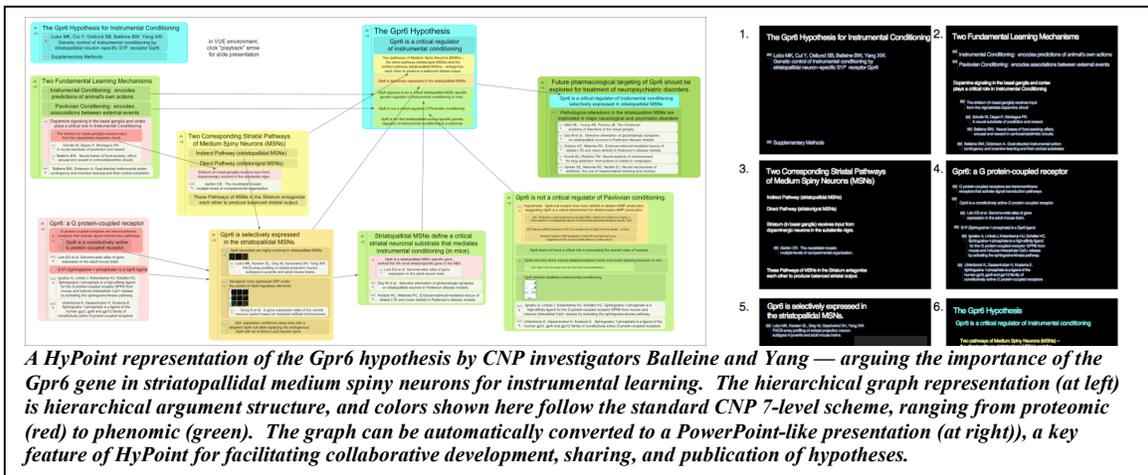
Lexicon Cleaning

We have developed a program that helps clean cognition lexicon, starting from an initial list of over 2,000 *cognition* candidate terms obtained as frequent terms from textbooks and other online resources. To further clean the lexica, we used a stemming tool to group syntactically similar terms together, and then sort them based on their common prefixes or suffixes. For example, family terms can be grouped such as

attention, auditory attention, central attention, divided attention, focus of attention, focused attention, involuntary attention, passive attention, selective attention, spatial attention, sustained attention

and

attention, attention capacity, attention span, attention shifting, attentional blink, attentional effort, attentional focusing, attentional resources, attentional state, attentional subsystems.



In this way, irrelevant terms can be easily detected and pruned by domain experts. For example in the first group only "attention" and "sustained attention" are kept; furthermore this grouping technique can be used as an aid in constructing concept hierarchies.

Assertion Extraction

We built a knowledge based tool to extract assertions. Given two or more concept terms, the assertion extractor is able to find all sentences (assertions) that mention these concept terms (or their synonyms) in a given knowledge base (e.g. a specified set of scientific documents). For example, for the concept terms "dopamine" and "working memory" and a set of PDF documents by CNP researchers, we obtained the following results from the assertion extractor:

Working memory function is strongly dependent upon dopaminergic function in PFC [29, 30]

Dopamine D1 receptors are involved in promoting stability of active representations in working memory [31, 32]

D2 receptors are involved in updating or resetting of working memory neural ensembles [33]

This tool is particularly useful for researchers who want to construct complex hypotheses.

Query Expansion

Query expansion can be used to relax users' queries so as to retrieve more documents. It can also serve as a tool for constructing a knowledge base of concepts. For a given concept term we are able to find conceptually related concepts by text mining a domain specific corpus. For example, we obtained the following list of the top 5 terms related to the concept term "dopamine" in cognition phenotypes and neural system:

Cognition:

1. [memory consolidation]
2. [insight]
3. [cognition]
4. [motor control]
5. [nondeclarative memory]

Neuroanatomy:

1. [archipallium, striatum]
2. [nucleus accumbens]
3. [frontal cortex, frontal lobe]
4. [prefrontal cortex]
5. [nucleus caudatus, caudate nucleus]

We are now working towards expanding queries containing multiple concepts (e.g., "dopamine" and "working memory"). Clearly, the additional concept term "working memory" will alter the expansion of "dopamine" with different emphasis.

Document Retrieval

We have built a search engine for document retrieval. Given an expanded query, we can retrieve relevant scientific report, and sort them based on their relevance to the query. Document retrieval is critical to the hypothesis web project because the content of the relevant scientific document that supports the hypothesis. Compared with the assertion extractor, document retrieval is more advanced in that it does not require query terms to appear in the same sentence. Instead, the relevance of each document to the query is decided by their term frequency and inverse document frequency (TF-IDF) weights. We have evaluated the precision for complex queries such as "response inhibition AND prefrontal", and obtained good verification of the search results from domain experts.

Plans

Over the next period we plan to consolidate the Hypothesis Web platform. This requires both a more complete ontology and new mechanisms for development of hypotheses. All tools described here (PubAtlas, HyPoint, and the Literature Mining tools) will also be extended in work with other CNP projects on representation of hypotheses.

Publications

Bilder RM, Sabb FW, Cannon TD, London ED, Jentsch JD, Parker DS, Poldrack RA, Evans C, Freimer NB. Phenomics: The systematic study of phenotypes on a genome-wide scale. *Neuroscience*. Epub 2009 Jan 20.

D.S. Parker, W.W. Chu, F.W. Sabb, A.W. Toga, R.M. Bilder, Literature Mapping with PubAtlas -- extending PubMed with a 'BLASTing' interface Proc AMIA Summit on Translational Bioinformatics, San Francisco, March 2009.

Parker DS, Poldrack RA, Sabb FW, Bilder RM, Quantologies, under review at *BMC Bioinformatics Journal*.