# **Progress Report**

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### Goals:

- 1. Implement Bayesian Weighting Scheme on Toy Network. This was part of the larger project to compare different weighting schemes for interactomes. This implementation will later be used on a larger network with actual biological data.
- 2. Write up/formalise math behind another weighting scheme. This follows directly from the project idea as well. This will eventually lead to another implementation to use for comparison.
- 3. Read Papers about interactome weighting, directionality, gene function enrichment. This helps to find more interactomes and learn about their weighting schemes.
- 4. Find one application network to illustrate different weighting schemes. Finding a particular interactome to apply different weighting schemes will help ascertain how helpful the different schemes are based on existing biological knowledge of the actual signalling pathways and the interactome as a whole.

## What Was Accomplished

- 1. Implemented Bayesian Weighting Scheme on Toy Network and posted to Graph Space.
- 2. Implemented a small portion of the scoring portion of the HIPPIE interactome
- 3. Read a few papers on HIPPIE interactome and Human Base.
- 4. Finished Bayesian Weighting Write Up Posted to Blog.

### **Problems**

- 1. The implementation was too slow and needs to sped up in order to apply to bigger real-world networks.
- 2. Failed to fully understand Parameter Selection and obtain working knowledge of Logistic Regression for HIPPIE Interactome Paper

### **Next Steps**

- 1. Change Bayesian Implementation to be faster
- 2. Follow up on Statistics resources to understand statistics concepts in HIPPIE Interactome but also come up with a list of descriptive statistics that might be useful to try to calculate for to compare weighting scheme implementations.