Application for KRMP

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Project start date: December 18, 2017 Project end date: June 30, 2018

Title of project: Modeling effects of viral infection on cellular fate using probabilistic Boolean networks

Abstract: To replicate inside of a host cell, viruses must make changes to cellular signal transduction networks. These changes are accomplished in a myriad of ways. For example, changing the expression patterns of genes/gene products. In this study, we will attempt to model these changes by implementing probabilistic Boolean networks. A probabilistic Boolean network is a Boolean network in which each node in the network has the potential to have multiple rules that govern them. We will look at a few properties of interest for this study including fixed points/cyclic attractors, effects on the stability of the network, and how the flow of information is changed in these networks.

Purposes of the project:

- 1. Identify the best format for network data to be used for simulations.
- 2. Review the mathematical algorithm for finding the determinative power of nodes, fixed points/ cyclic attractors, and determining the stability of the network. Develop appropriate analytic models.
- 3. Prepare Matlab/python codes to be used with actual network data. Fine tune the codes along the way, and possibly add new codes as needed.
- 4. Generate random PBN data in suitable format and run simulations. This is the most timeconsuming part of the project.
- 5. Collect results for all network simulations.
- 6. Data analysis of the results. This task may involve further aspects that we cannot foresee now and that are dependent on the actual network properties.
- 7. Writing the KRMP report.

Timetable:

Month	Purpose items
December 2017 – January 2018	1, 2, 3, 4, 5
January - March 2018	4,5,6
April - May 2018	6
June 2018	7
July 2018-forward	Preparing results for publication.