## Analyzing Nonlinear Dynamical Systems with Many Parameters

Adam Zheleznyak

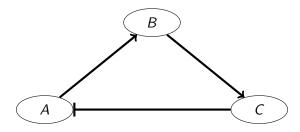
Konstantin Mischaikow Marcio Gameiro

DIMACS REU

June 1, 2020

## **Regulatory Networks**

Collection of species (DNA, RNA, proteins)



Dynamics of the system depend on parameters e.g.

- How much does A affect B?
- How much of A is needed to have an effect?
- How sensitive is the response to the amount of A?

We care about how different parameters may give different behaviors. However, the number of parameters can grow very large, which can make it difficult or impossible to analyze the dynamics. My mentors have developed methods and software that allow for a more efficient way to understand the overall dynamics of these kinds of biological systems.

Can create a coarser model by simplifying parameters, which can:

- Reduce the total number of parameters
- Make the model more "discrete" to allow for better computational techniques
- Allow the use of certain algebraic tools

The questions I will be exploring this summer:

How can the same techniques be used to create a general purpose tool for studying any nonlinear dynamical system with few variables and many parameters? How can we make sure this is efficient?

This work was carried out while I was a participant in the 2020 DIMACS REU program at Rutgers University, supported by NSF HDR TRIPODS award CCF-1934924.