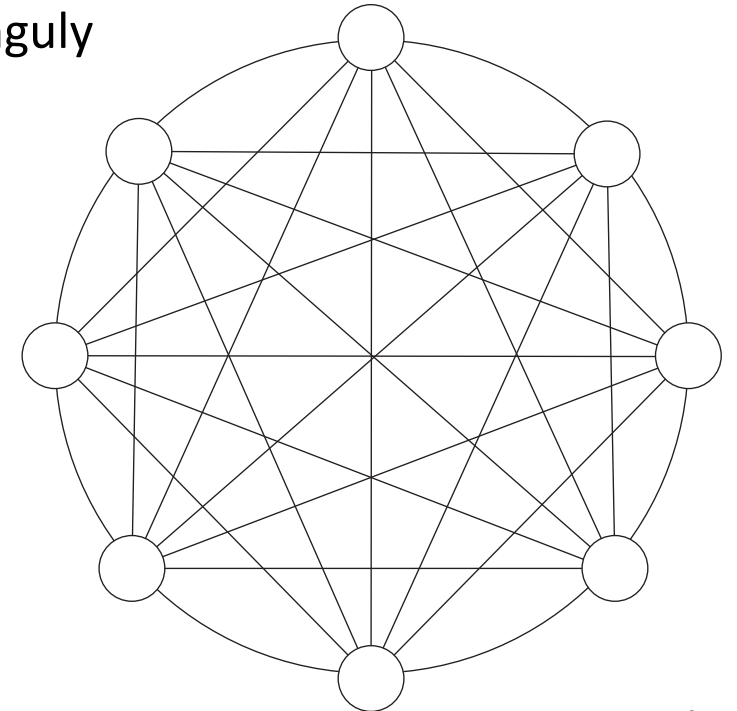
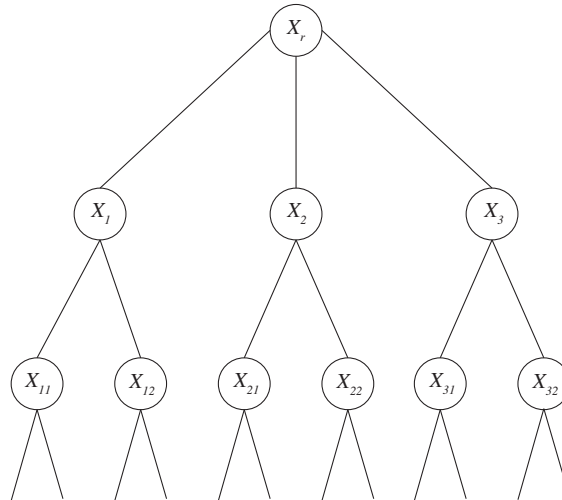
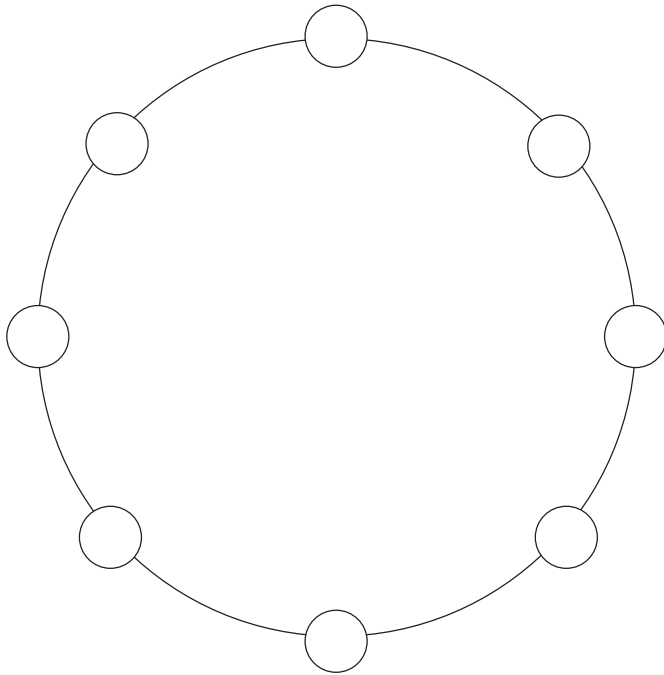
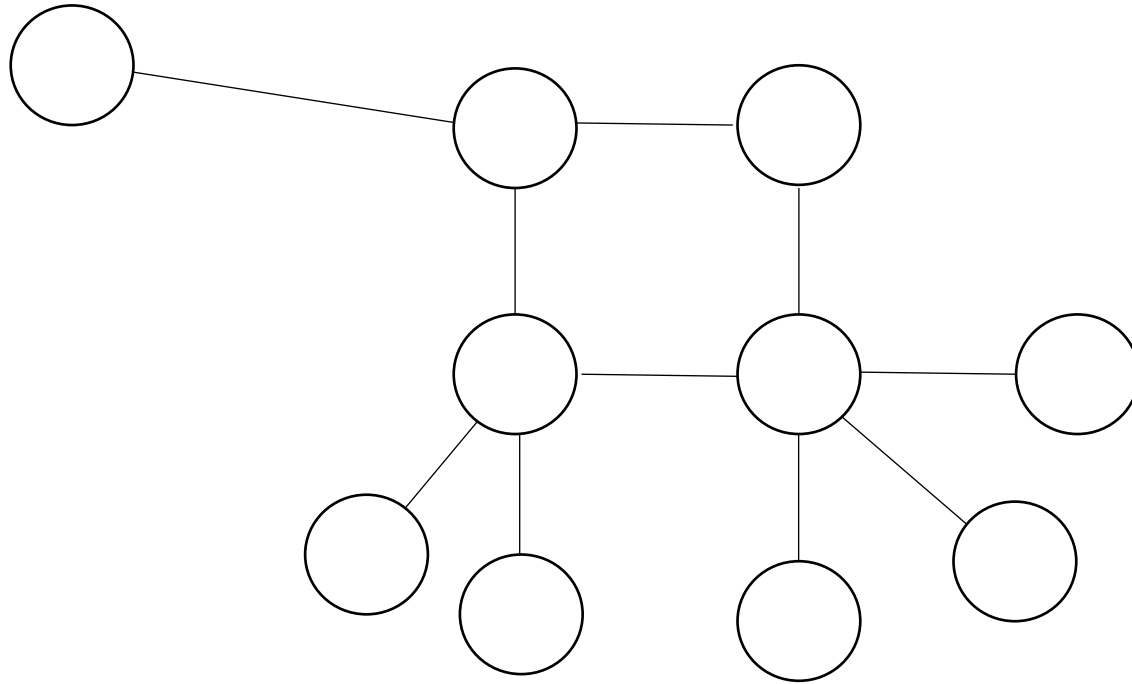


Interacting Particle Systems and Efficient Approximations for Large Sparse Graphs

Senior Thesis Supervisor: Kavita Ramanan
Graduate Student Mentor: Ankan Ganguly
Presenter: Mitchell Wortsman

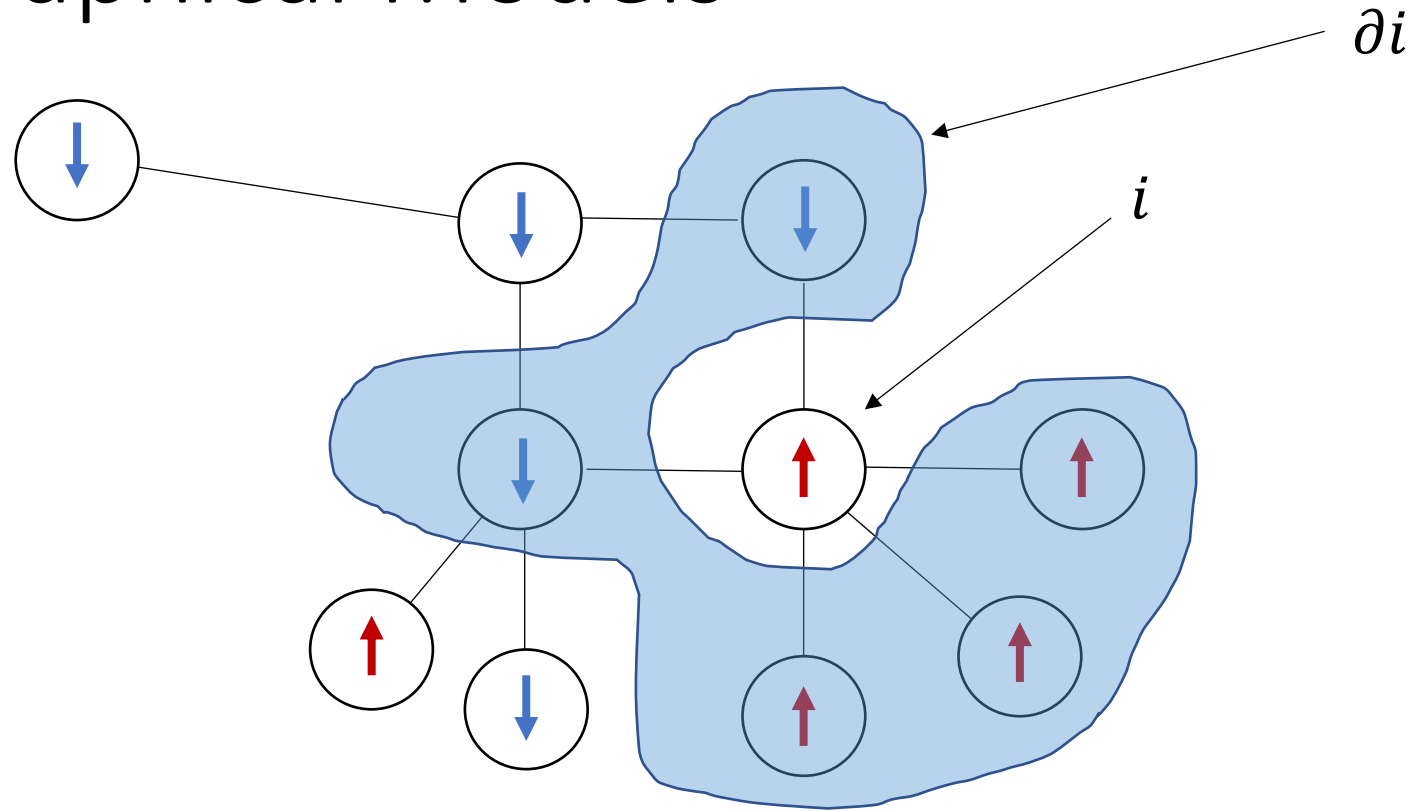


Step 1: Graphs



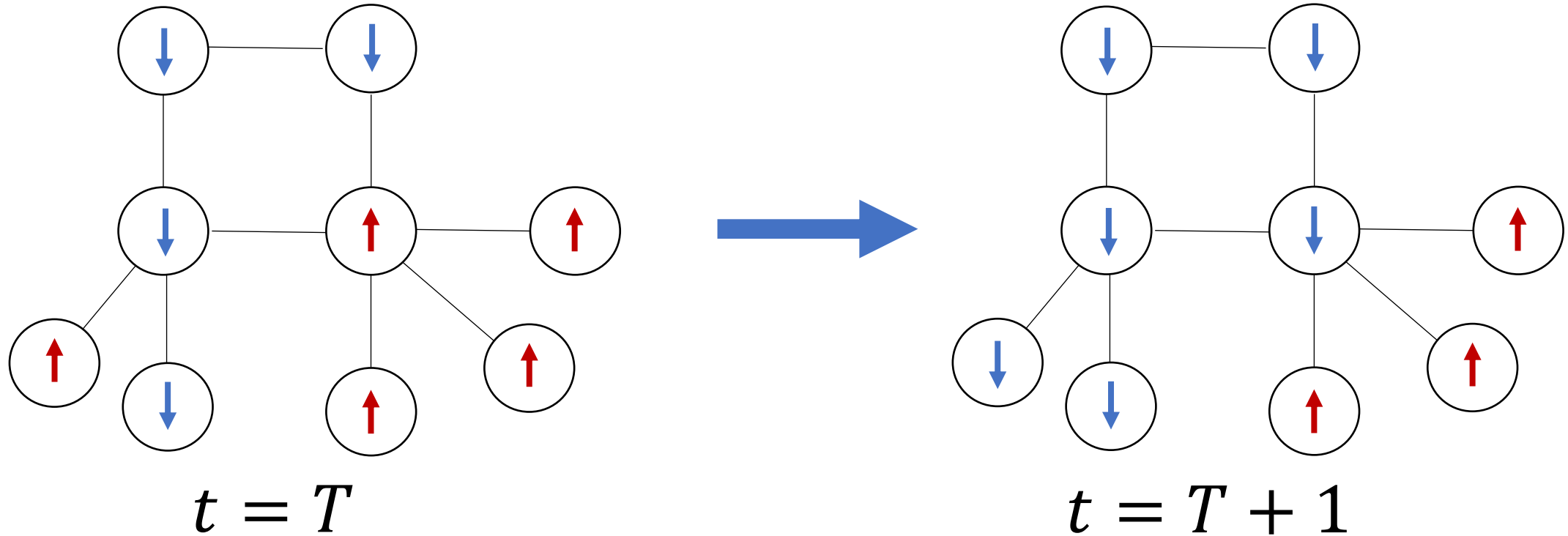
- $G = (V, E)$
 - V is the set of particles.
 - E is the set of connections.

Step 2: Graphical Models



- Each particle has a value.
- The probability that particle i has value x_i depends only on the state of its neighbors.

Step 3: Interacting Particle Systems



- Each particle has a state which evolves through time.
- Particle i evolves based only on the current state of its neighbors (and some independent noise).

Step 4: Formally...

- The state of particle i at time t is given by $X_i(t)$
- The random vector $X(t) = (X_0(t), X_1(t), \dots)$ represents the state of **all** particle at time t .
- The system satisfies the following properties:

1. The Markov Property

$$\begin{aligned}\mathbb{P}(X(t+1) = \vec{x} \mid X(0), \dots, X(t)) \\ = \mathbb{P}(X(t+1) = \vec{x} \mid X(t))\end{aligned}$$

- If we know the present, knowing the past does not give us any more information about the future.

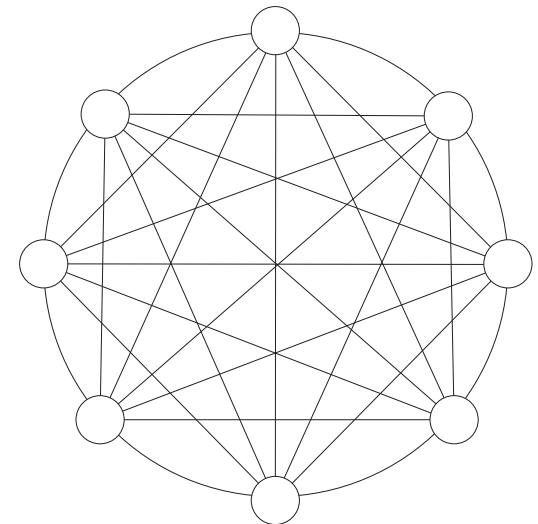
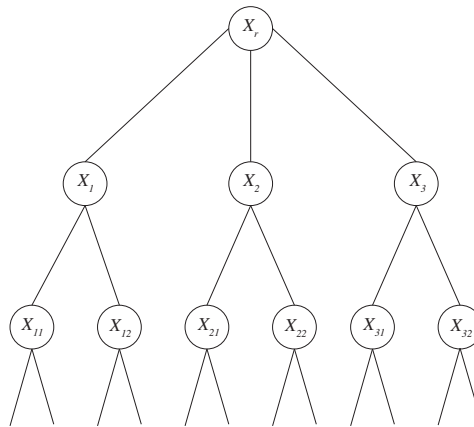
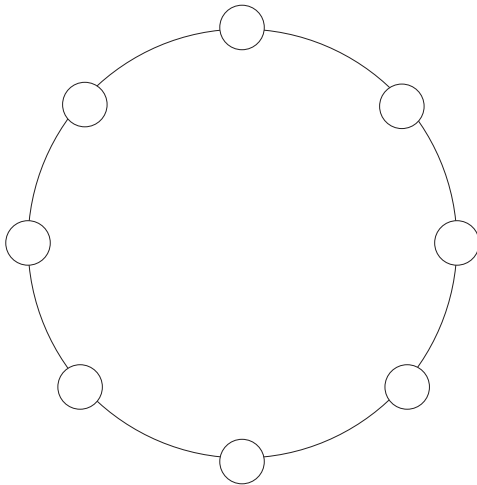
2. The Local Interactions Property

$$\begin{aligned}\mathbb{P}(X_i(t+1) = v \mid X(t)) \\ = f(v, X_i(t), X_{\partial i}(t))\end{aligned}$$

- Particle i evolves based only on the current state of its neighbors.

Step 5: The Goal

- Characterize the dynamics of a typical particle.
- Particle dynamics?
 - What is the probability that particle i is in state x_i at time t ?
- Typical particle?
 - Assume that graph is symmetric and so all particles are identical.



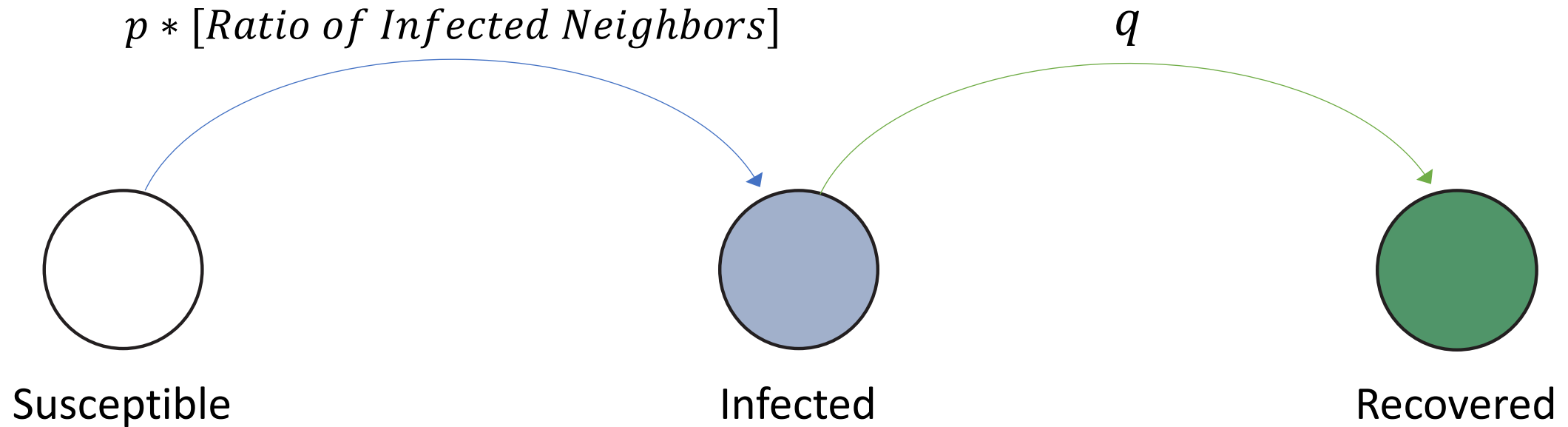
Step 6: Why is this a challenge?

- **We can simulate the full system for a small number of particles N , but what about as N grows large and is perhaps infinite?**
- **The standard Mean Field Approximation works very well, but only for *dense* graphs.**

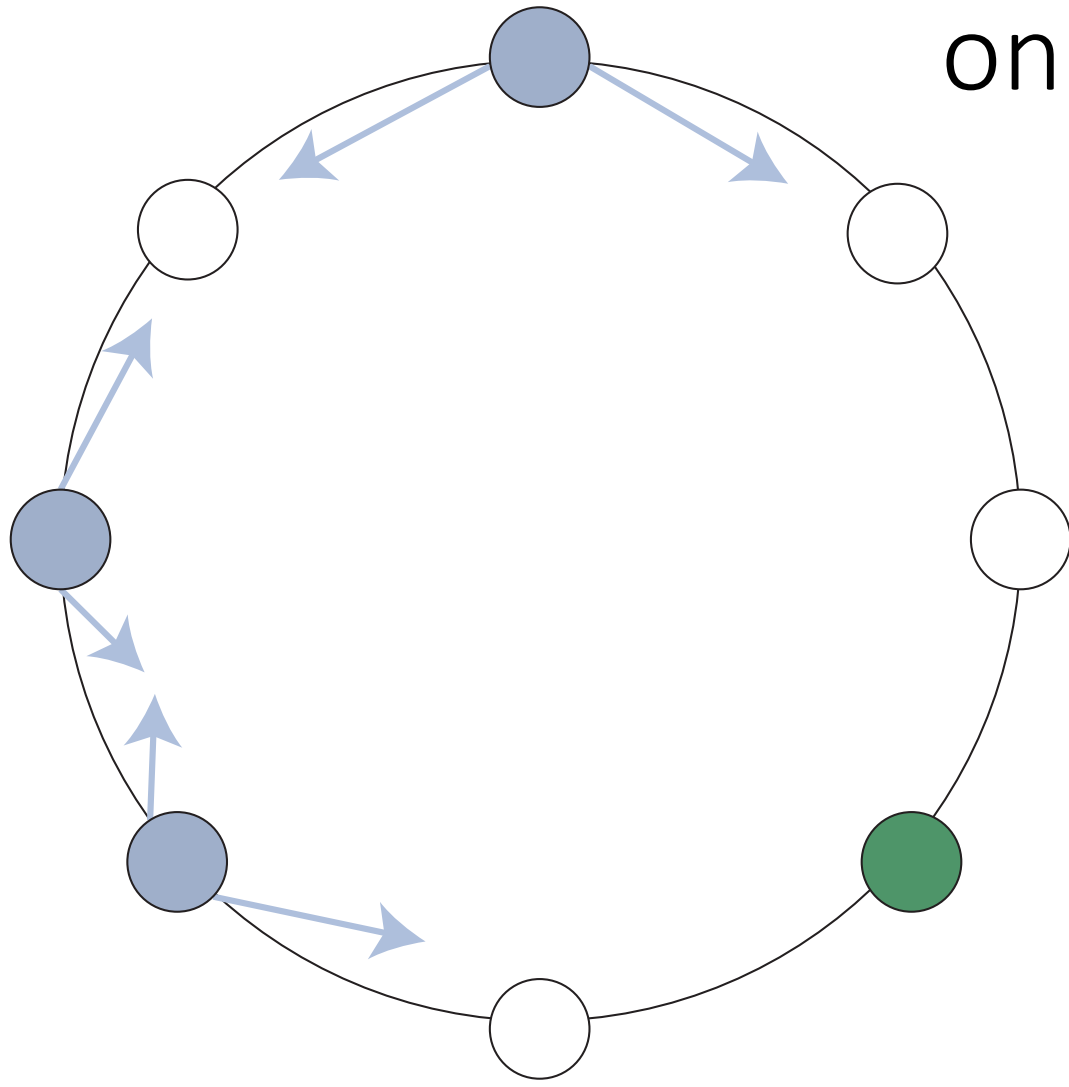
Step 7: Why are we interested?

- Epidemiology
- Statistical Physics
- Load Balancing
- Information Theory
- Genetics
- Machine Learning

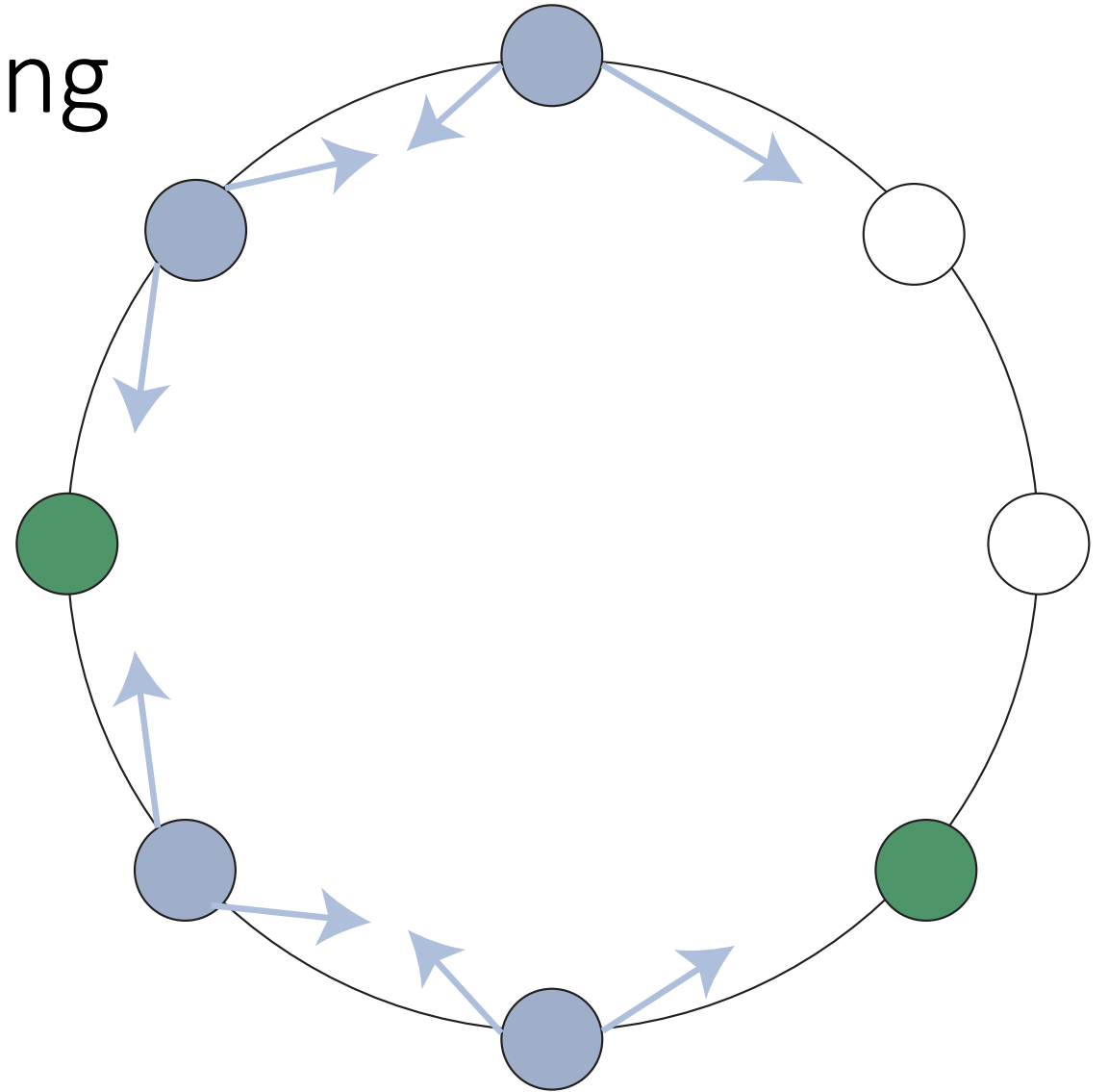
Susceptible-Infected-Recovered (SIR) Process



SIR Process on a Ring



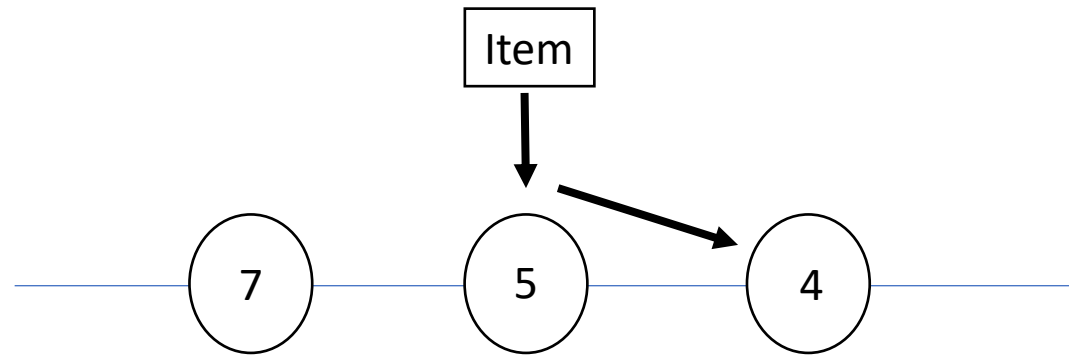
$t = T$



$t = T + 1$

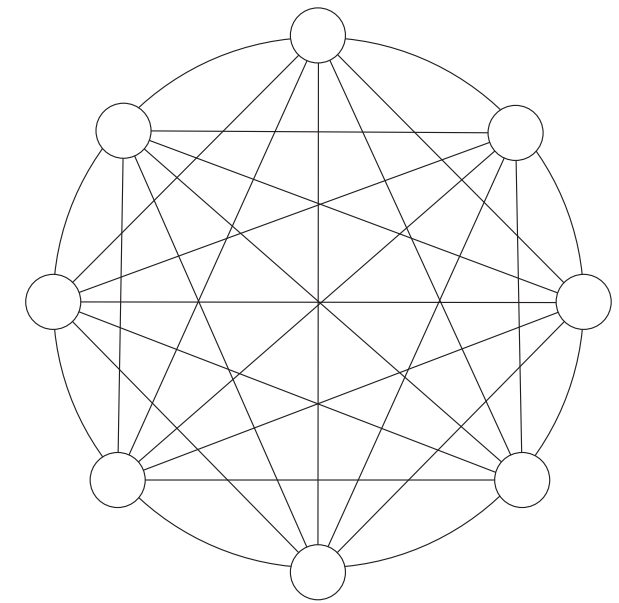
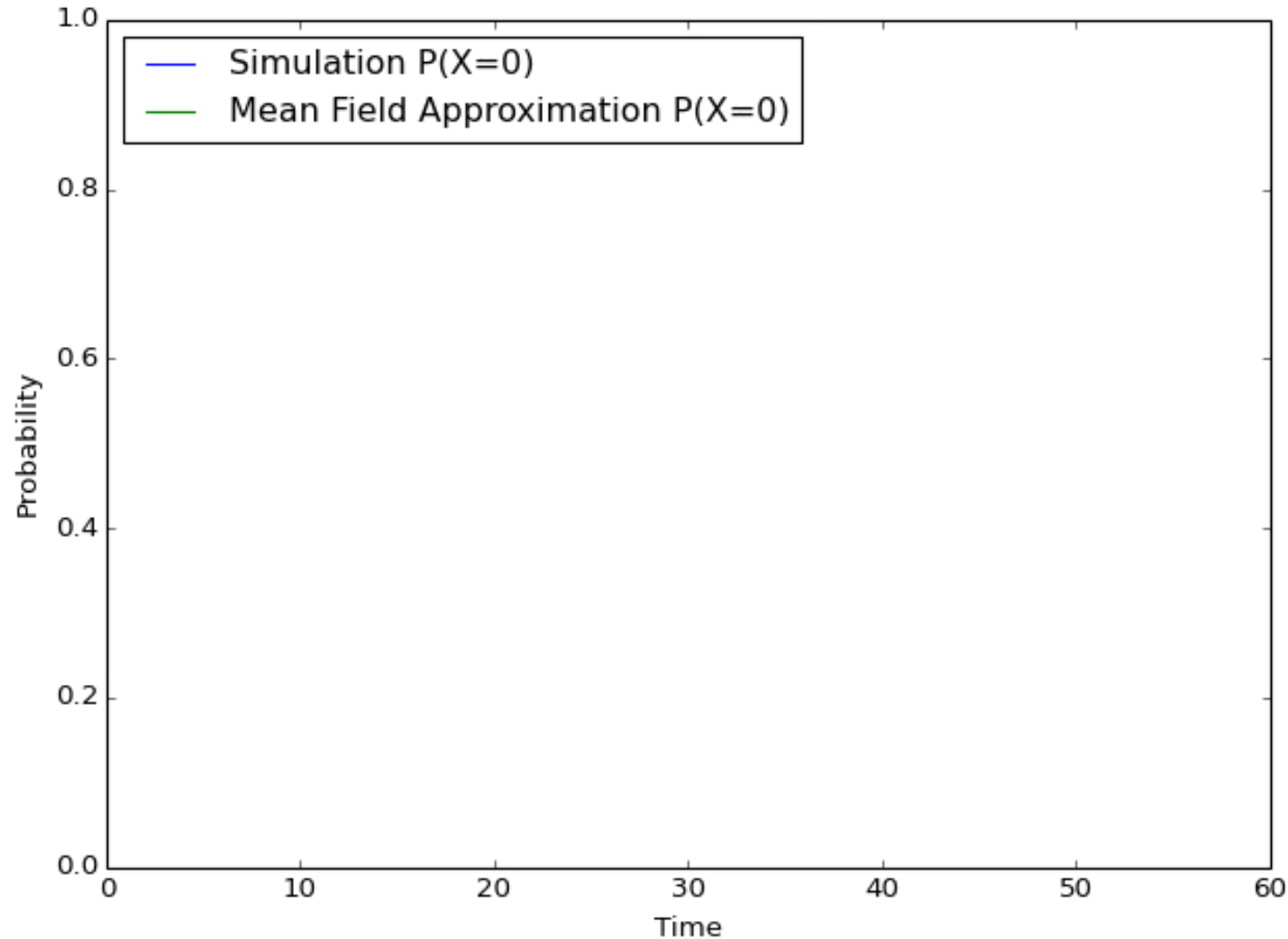
Load Balancing

- The state of each particle is the number of items in a queue.
- When an item arrives at queue i , it is routed to the shortest queue among i and its neighbors.



An item incident at the queue with 5 items
will be routed to the queue with 4 items.

The SIR Process on a Complete Graph

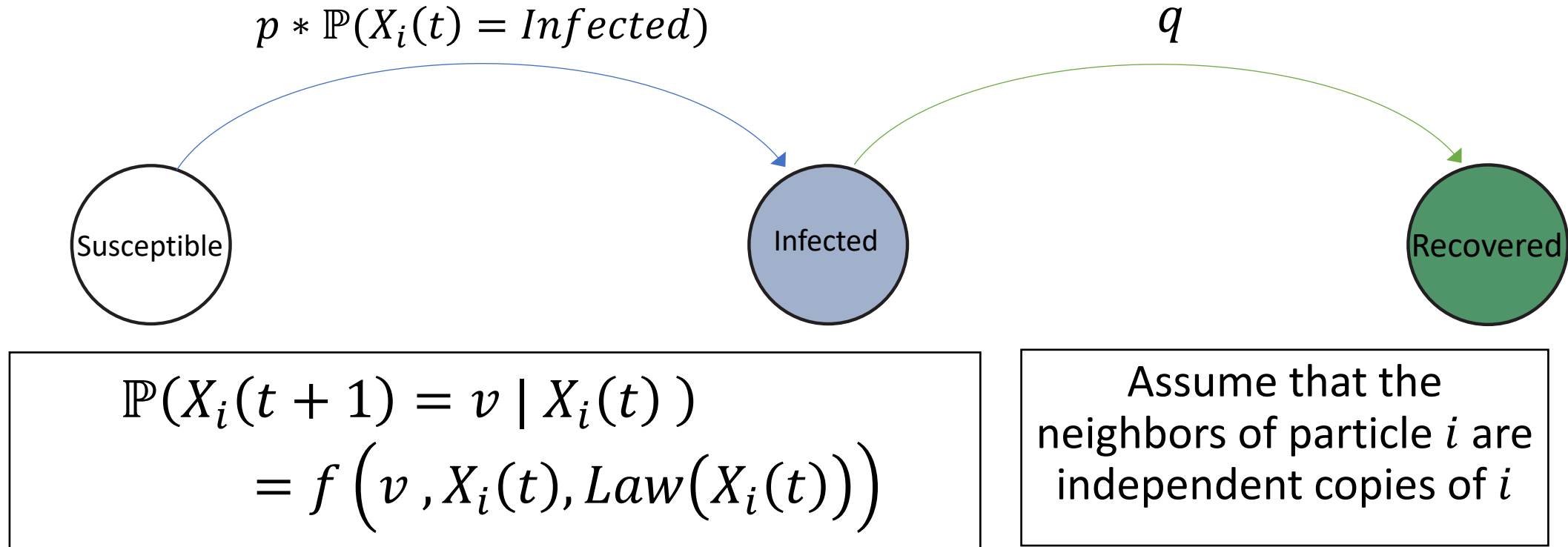


Interaction Network

$n = 50$

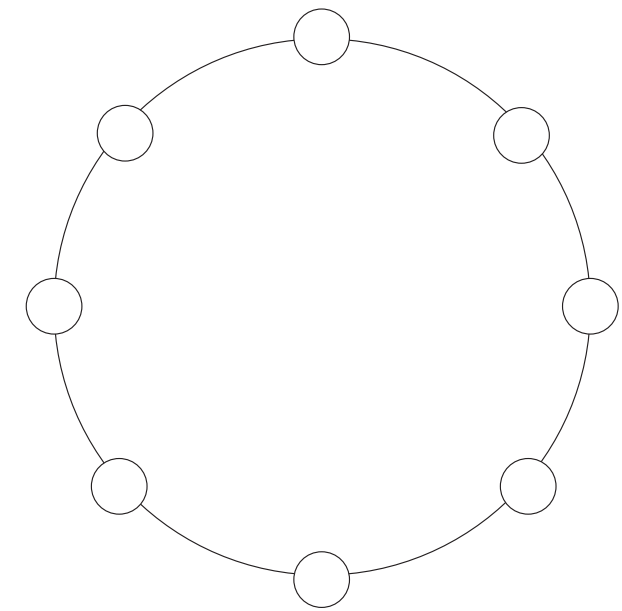
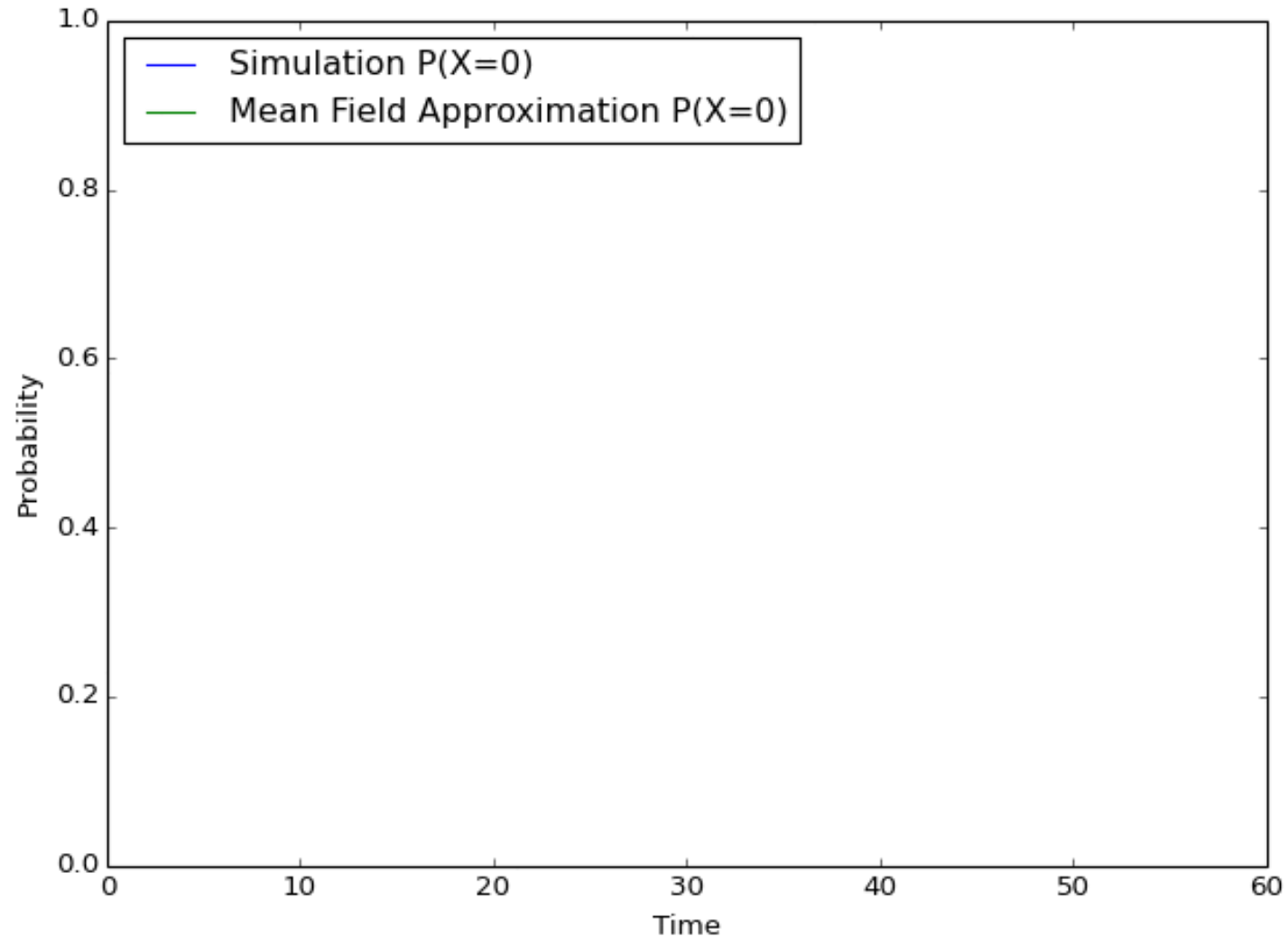
The Mean Field Approximation

Mean Field Approximation



- This approximation is **exact** for the complete graph as the population size tends to infinity (Oelschläger, 1984).
- Recently been shown to be asymptotically accurate for sequences of dense graphs whose degree goes to infinity (Bhamidi, Budhiraja, Wu, 2017).
- In many applications the mean-field approximation is used for any graph.

The SIR Process on a Ring

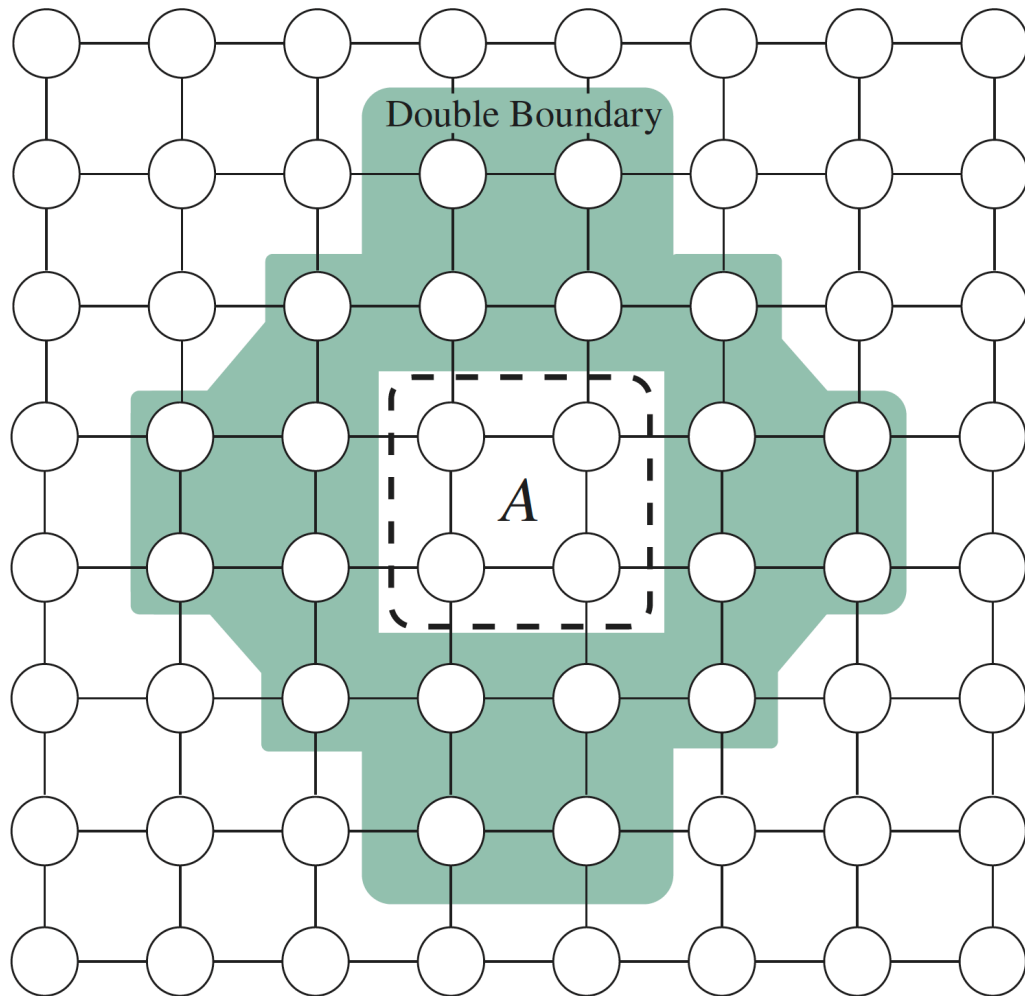


Interaction Network
 $n = 50$

Step 8: Our Local Recursions

- **We build on recent theoretical results by Lackner, Ramanathan, and Wu (2018) to construct a novel '*Local Recursion*'.**

Dynamic Conditional Independence (Lacker, Ramanan, Wu 2018)

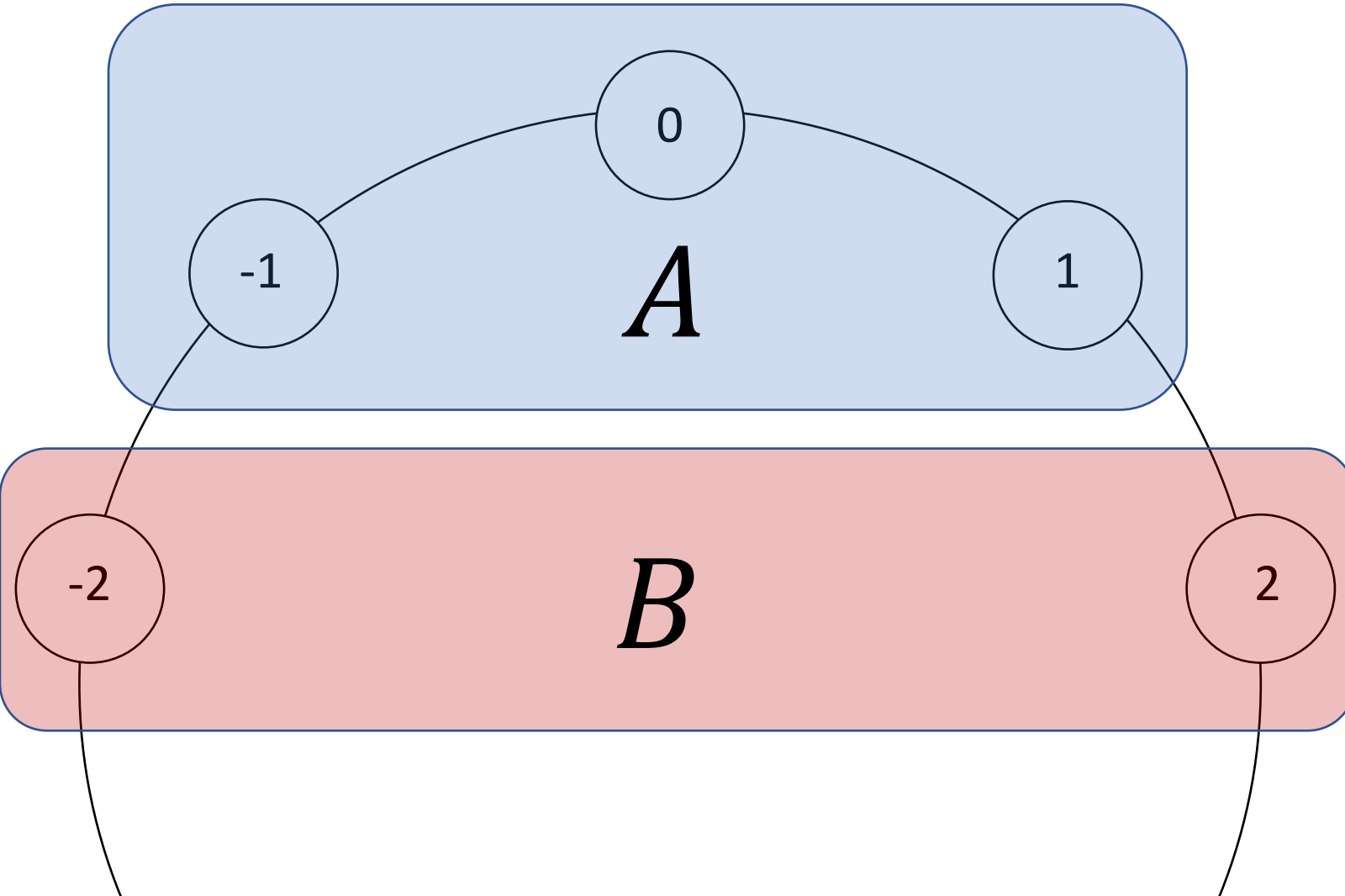


- **Notation**

- *The Full Trajectory:* $X_A^T = (X_i(t))_{i \in A, t \leq T}$
- *The Double Boundary:* $D = \partial^2 A$

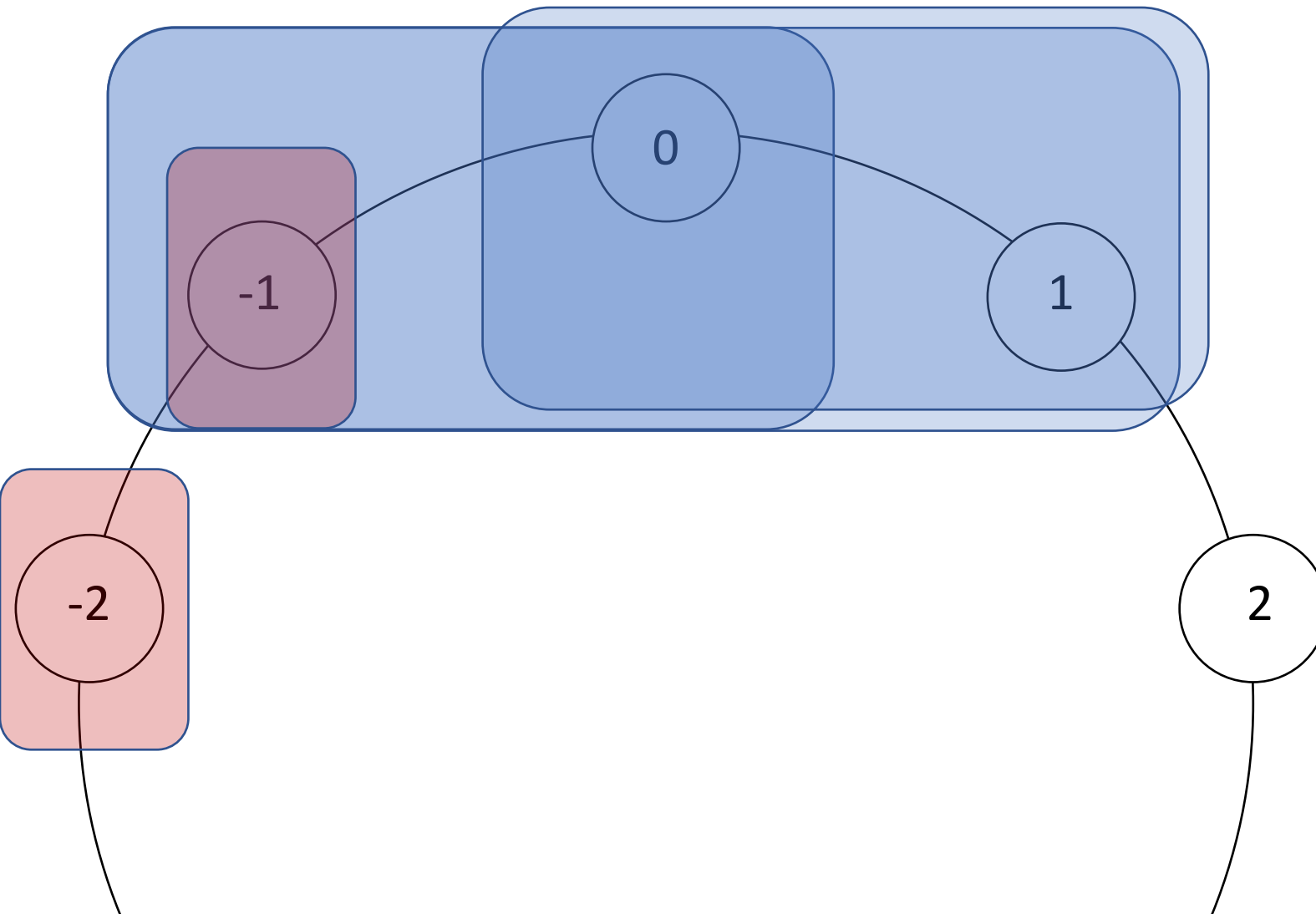
$$X_A^T \perp X_{V \setminus (D \cup A)}^T \mid X_D^T$$

Our Local Recursion



- Consider local region A .
- We would like a *self-contained* equation for A .
- Assume that we know the joint distribution of local region A up to time t .
- To compute the joint distribution of region A up to time $t + 1$ we need to know the distribution of region B at time t (conditioned on the region A).

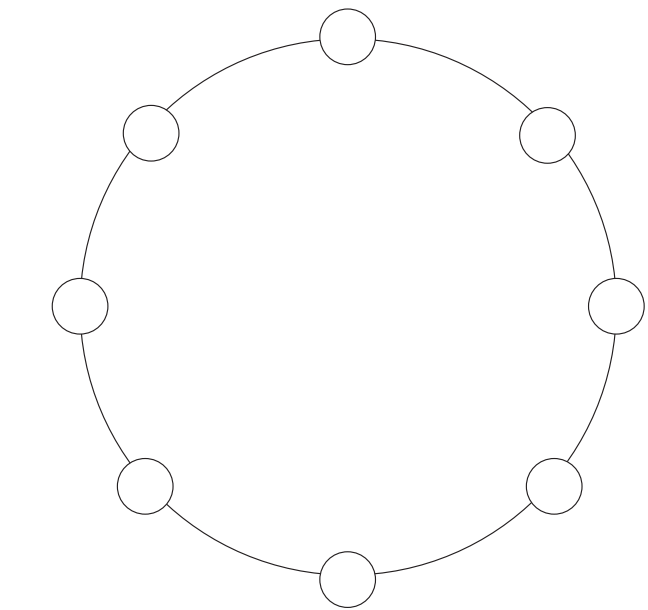
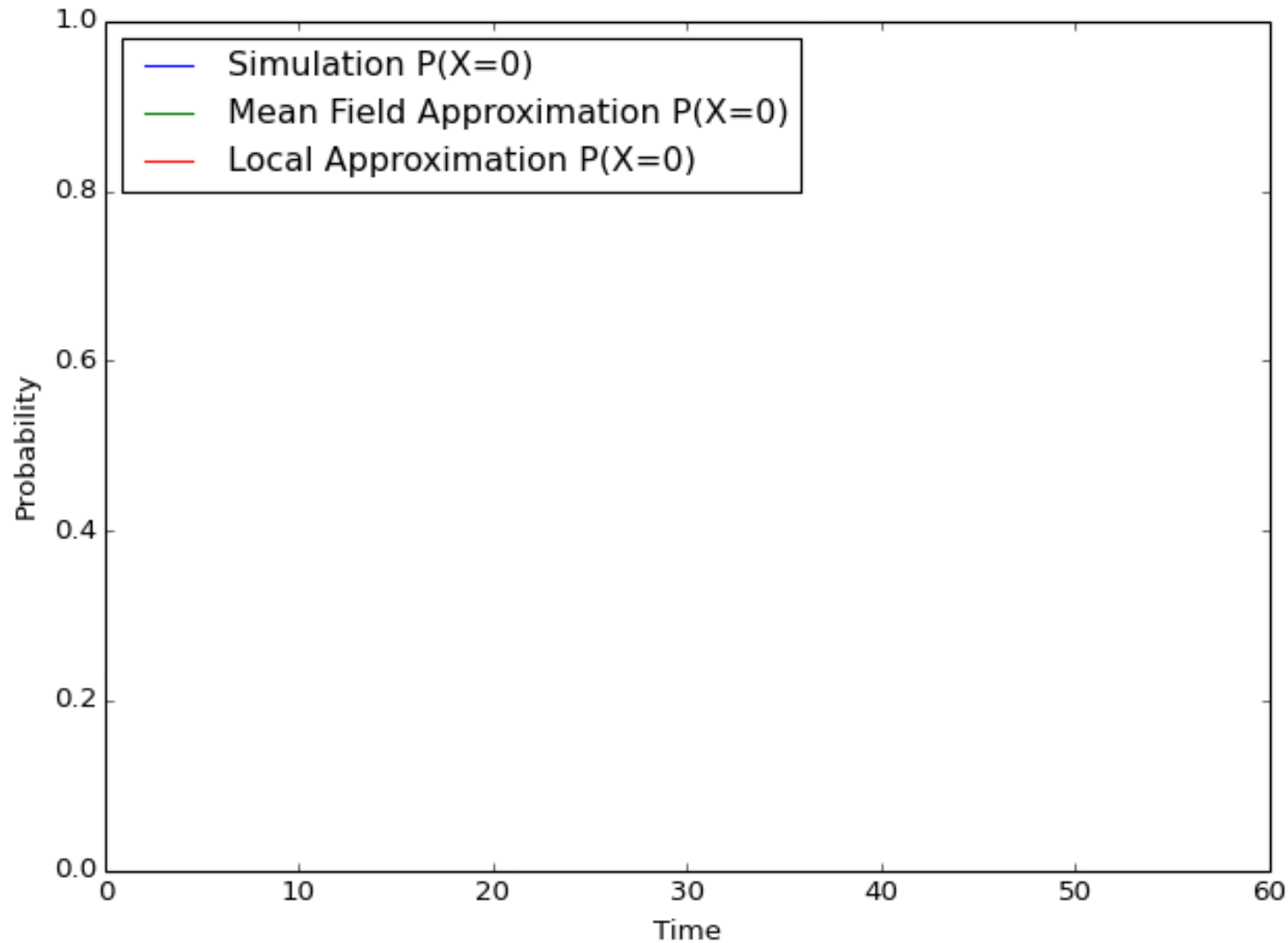
Our Local Recursion (2)



- CDB = Conditional Independence Given the Double Boundary.
- By CDB, the state of particles -2 and 2 at time t are independent given the trajectory of A .
- First we consider particle -2.
- By CDB, conditioning on A is equivalent to conditioning on the trajectory of -1,0.
- Since all particles are identical, this is equal in distribution to the state of particle -1 at time t conditioned on the trajectory of 0,1.
- Which we know by our inductive assumption.

How well does our local approximation work?

The SIR Process on a Ring (with our local approximation)



Interaction Network
 $n = 50$

Local Recursions: Time Complexity

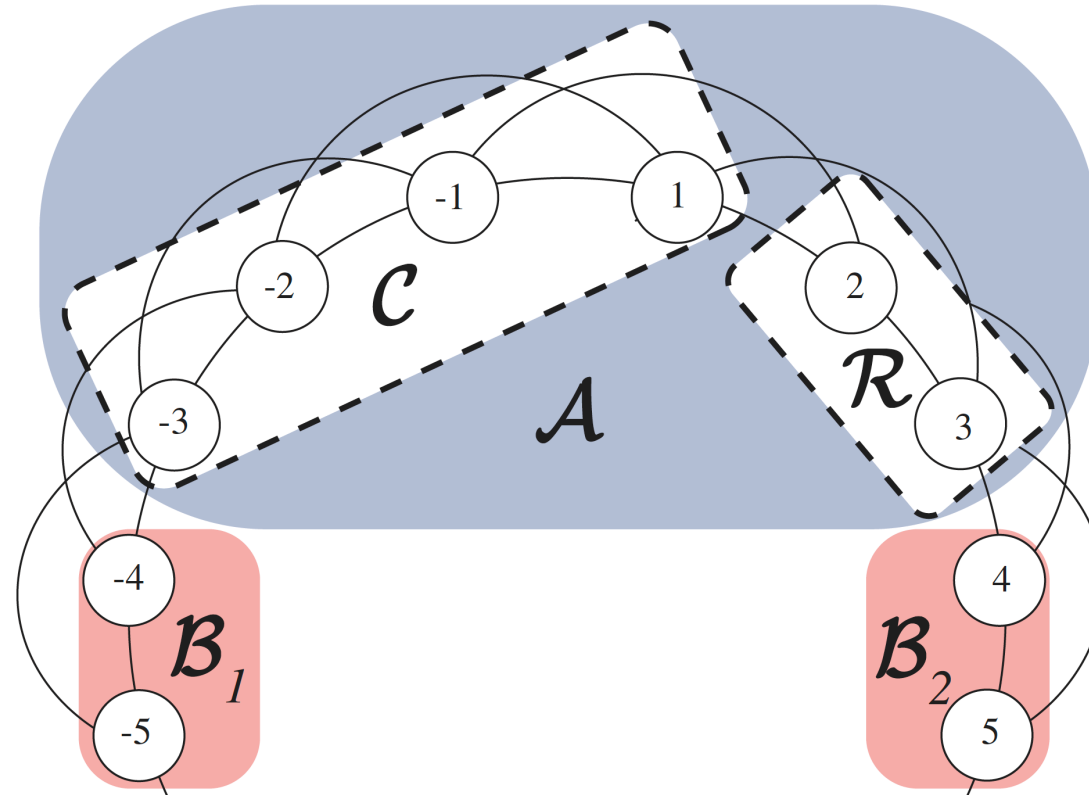
- As an approximation to the exact recursions, we may consider only the last τ time steps.
 - We call this **The τ Approximation**.

Other Existing Approximations

- Dynamic Cavity Method
 - Only exact for trees.
 - Requires consideration of the dynamics of all particles in the graph (slow).
 - Not clear how to implement for continuous state space.
 - No fast approximation.
- Moment closure methods (i.e. Pair Approximation):
 - Makes a huge assumption (that the time marginals satisfy the spatial Markov property).
 - For many models (e.g. SIR) this approximation is very inaccurate (Gast, 15).

Generalizations to Other Graphs

- Our method generalizes to non locally-tree-like graphs.
 - For example, the graph which arises in Load Balancing on a ring.



Summary

In this presentation we have...

- Defined Interacting Particle Systems and given examples.
- Discussed existing approximations for understanding the behavior of these systems (Mean Field) and highlighted cases where they are not applicable.
- Discussed recent theory which reveals a new conditional independence structure.
- Building off this recent theory we have demonstrated the effectiveness of our new Local Recursions.

Questions?