

A Finite Memory Interacting Pólya Contagion Network and its Approximating Dynamical Systems

Somya Singh

Department of Mathematics and Statistics
Queen's University

In collaboration with **Fady Alajaji** and **Bahman Gharesifard** (Queen's)

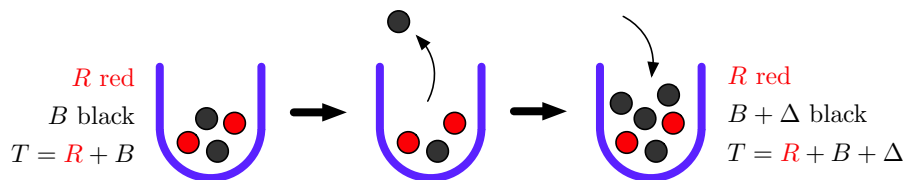
- 1 Introduction
- 2 Finite Memory Interacting Pólya urn Network
- 3 Stochastic Properties
- 4 Approximating Dynamical Systems
- 5 Simulation Results

A new model for spread of infection

Objective of the talk:

to use **a network of finite memory Pólya urns** to model the spread of infection in a population

Classical Pólya urn Model



At each time t , we add Δ **red** balls or Δ **black** balls if a red ball or a black ball is drawn respectively.

$$Z_t = \begin{cases} 1 & \text{if a \textbf{red} ball is drawn at time } t \\ 0 & \text{if a \textbf{black} ball is drawn at time } t. \end{cases}$$

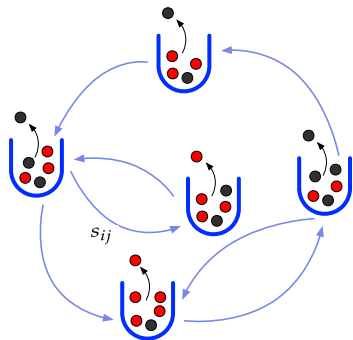
Z_t is the drawing random variable.

Interacting Pólya urn Network

Consider a **network of N Pólya urns**:

$$Z_{i,t} = \begin{cases} 1 & \text{w.p. } \sum_{j=1}^N s_{ij} U_{j,t-1} \\ 0 & \text{w.p. } 1 - \sum_{j=1}^N s_{ij} U_{j,t-1}, \end{cases}$$

where $U_{j,t-1}$ is the *ratio of red balls in urn j at time $t - 1$* .

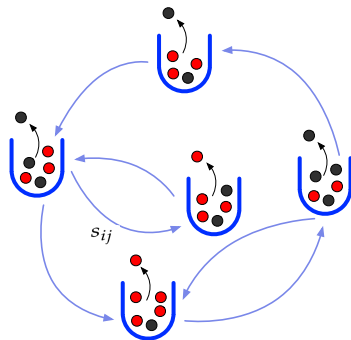


Interconnections prescribed by a **directed graph with a row-stochastic adjacency matrix** ($s_{ij} \geq 0$ and $\sum_{j=1}^N s_{ij} = 1$ for all $i, j \in \{1, 2, \dots, N\}$)

the interactions accounts for spatial infection among neighbours

Interacting Pólya urn Network

- At every time instant t , a ball is drawn for every urn.
- if a **red** ball is drawn for urn i at time t , we add $\Delta_{r,i}(t)$ red balls to the urn i at time t .
- if a **black** ball is drawn for urn i , we add $\Delta_{b,i}(t)$ black balls to urn i at time t .



The red balls account for the “infection” in urns and black balls account for the “healthiness” in urns.

Interacting Pólya urn Network

The draw variables $\{Z_{i,t}\}$'s are **conditionally independent given all past draws** in the Interacting Pólya urn Network i.e.,

$$P(Z_{1,t}, \dots, Z_{N,t} | \{Z_{1,k}\}_{k=1}^{t-1}, \dots, \{Z_{N,k}\}_{k=1}^{t-1}) = \prod_{i=1}^N P(Z_{i,t} | \{Z_{1,k}\}_{k=1}^{t-1}, \dots, \{Z_{N,k}\}_{k=1}^{t-1}).$$

We also can compute the **ratio of red balls** at each urn at a given time:

$$U_{i,t} = \frac{X_{i,t-1}}{X_{i,t}} U_{i,t-1} + \frac{Z_{i,t} \Delta_{r,i}(t)}{X_{i,t}},$$

where $X_{i,t} := T_i + \sum_{n=1}^t Z_{i,n} \Delta_{r,i}(n) + (1 - Z_{i,n}) \Delta_{b,i}(n)$ and T_i is the total number of balls in urn i at time $t = 0$.

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Some key observations

- An important characteristic of most reinforcement processes generated via urn models is that they are **non-Markovian**
- The composition of each urn at any given time **affects its composition at every time instant thereafter**
- This property is not realistic when modelling the spread of infection as **one should account for the possibility that infection is cured**

We consider interacting Pólya urn network with finite memory

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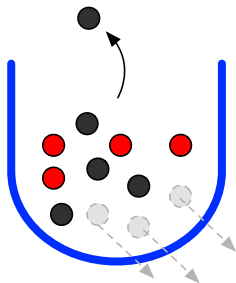
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Finite Memory

Interacting Pólya urn Network with N urns and **finite memory** $M \geq 1$ (denoted by IPCN(M, N)):

At time instant $t > M$, the reinforcing balls added at time $t - M$ are removed from the urn and hence have no effect on future draws.



Some parameters for IPCN(M, N)

Ratio of red balls in urn i at time t can be written as

$$U_{i,t} = \frac{\rho_i + \sum_{n=t-M+1}^t \delta_{r,i}(n)Z_{i,n}}{1 + \sum_{n=t-M+1}^t (\delta_{r,i}(n)Z_{i,n} + \delta_{b,i}(n)(1 - Z_{i,n}))}, \quad (1)$$

where the parameters defined below will be used later on:

$$\rho_i = \frac{R_i}{T_i}, \quad \delta_{r,i}(t) = \frac{\Delta_{r,i}(t)}{T_i}, \quad \delta_{b,i}(t) = \frac{\Delta_{b,i}(t)}{T_i}.$$

$U_{i,t}$ is the ratio of red balls in urn i at time t .

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Stochastic Properties

We define $Z_t = (Z_{1,t}, \dots, Z_{N,t})$

Note that $\{Z_t\}_{t=1}^{\infty}$ is a **time-varying M th order Markov chain**, and

$$\begin{aligned}
 & P[Z_{t+1} = a_{t+1} | Z_t = a_t, \dots, Z_{t-M+1} = a_{t-M+1}] \\
 &= \prod_{i=1}^N \left((2a_{i,t+1} - 1) \sum_{j=1}^N \frac{s_{ij} \left(\rho_i + \sum_{n=t-M+1}^t \delta_{r,i}(n) a_{i,n} \right)}{1 + \sum_{n=t-M+1}^t (\delta_{r,i}(n) a_{i,n} + \delta_{b,i}(n) (1 - a_{i,n}))} \right. \\
 & \quad \left. + (1 - a_{i,t+1}) \right) \tag{2}
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Even though we can compute the evaluations of this process using this master formula, analysis of this process is complex

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Even though we can compute the evaluations of this process using this master formula, analysis of this process is complex

Homogeneous Case

for all i and t :

$$R_i = R, T_i = T \text{ and depending on the value of draw variable at time } t, \\ \Delta_{r,i}(t) = \Delta > 0 \text{ or } \Delta_{b,i}(t) = \Delta > 0$$

Here $\{Z_t\}_{t=1}^{\infty}$ is a **time-invariant M th order Markov chain** and we can draw some conclusions about this process

We define $W_t := (Z_t, Z_{t+1}, \dots, Z_{t+M-1})$. Since $\{Z_t\}_{t=1}^{\infty}$ is an M th order Markov chain, $\{W_t\}_{t=1}^{\infty}$ forms a Markov chain of order one.

Lemma. For the homogeneous IPCN(M, N), the Markov process $\{W_t\}_{t=1}^{\infty}$ is irreducible and aperiodic.

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Lemma. For the homogeneous IPCN(M, N), the Markov process $\{W_t\}_{t=1}^{\infty}$ is irreducible and aperiodic.

Transition probability matrix for homogeneous case

For $M = 1$, the probability of going from state $a = (a_{1,t}, \dots, a_{N,t})$ to state $b = (b_{1,t+1}, \dots, b_{N,t+1})$ can be expressed as

$$q_{ab}^{(1,N)} := q_{ab}^{(1)} q_{ab}^{(2)} \cdots q_{ab}^{(N)}$$

where

$$q_{ab}^{(d)} = \begin{cases} \frac{\sigma + (1 - \sum_{k=1}^N s_{dk} a_{k,t}) \delta}{1 + \delta} & \text{if } b_{d,t+1} = 0 \\ \frac{\rho + \sum_{k=1}^N s_{dk} a_{k,t} \delta}{1 + \delta} & \text{if } b_{d,t+1} = 1, \end{cases} \quad (3)$$

where $d \in \{1, \dots, N\}$ and $\sigma = 1 - \rho$.

Transition probability matrix for homogeneous case

For $M > 1$, if the transition probability of going from state $a = ((a_{11}, a_{21}, \dots, a_{N1}), \dots, (a_{1M}, a_{2M}, \dots, a_{NM}))$ to state $b = ((b_{11}, b_{21}, \dots, b_{N1}), \dots, (b_{1M}, b_{2M}, \dots, b_{NM}))$ is nonzero, it is given by

$$q_{ab}^{(M,N)} := \tilde{q}_{ab}^{(1)} \tilde{q}_{ab}^{(2)} \dots \tilde{q}_{ab}^{(N)}$$

where

$$\tilde{q}_{ab}^{(d)} = \begin{cases} \frac{\sigma + \left(M - \sum_{i=1}^N s_{di} \left(\sum_{k=1}^M a_{ik} \right) \right) \delta}{1 + M\delta} & \text{if } b_{dM} = 0 \\ \frac{\rho + \left(\sum_{i=1}^N s_{di} \left(\sum_{k=1}^M a_{ik} \right) \right) \delta}{1 + M\delta} & \text{if } b_{dM} = 1, \end{cases} \quad (4)$$

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Approximation by dynamical systems

- The Markov chain $\{W_t\}_{t=1}^{\infty}$ for $IPCN(M, N)$ (not necessarily homogeneous from now on) has 2^{MN} states.
- Due to this exponential increase in the size of the transition probability matrix with the number of urns N and memory M , it is difficult to analytically solve for the stationary distribution in terms of the system parameters

This issue should is also present in the classical SIS/SIR model

We introduce a class of dynamical systems whose trajectory approximates the $P(Z_{i,t} = 1)$ at time t for any urn i in the network.

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Mean-field Approximations

At any given time t , we use the following **mean-field approximations**

- For $M = 1$,

$$P(Z_{1,t-1} = a_{11}, \dots, Z_{N,t-1} = a_{N1}) \approx \prod_{j=1}^N P(Z_{j,t-1} = a_{j1}). \quad (5)$$

- For $M > 1$,

$$\begin{aligned} P[\tilde{Z}_1 = \underline{a}_{1M}, \dots, \tilde{Z}_N = \underline{a}_{NM}] \\ \approx \prod_{j=1}^N \prod_{k=1}^M [a_{jk} P_j(t-k) + (1-a_{jk})(1-P_j(t-k))], \end{aligned} \quad (6)$$

where $\underline{a}_{iM} = (a_{i1}, a_{i2}, \dots, a_{iM})$, $\tilde{Z}_{i,t-1} = (Z_{i,t-1}, Z_{i,t-2}, \dots, Z_{i,t-M})$

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Approximation by dynamical systems: $M = 1$

We first define:

$$P_i(t) := P(Z_{i,t} = 1),$$

We have that

$$P_i(t) = \sum_{B_{N,1}} P(Z_{i,t} = 1 | Z_{1,t-1} = a_{11}, \dots, Z_{N,t-1} = a_{N1}) \quad (7)$$

$$\times P(Z_{1,t-1} = a_{11}, \dots, Z_{N,t-1} = a_{N1})$$

where

$$B_{N,1} := \{(a_{11}, a_{21}, \dots, a_{N1}) \mid a_{i,1} \in \{0, 1\} \text{ for } i \in \{1, 2, \dots, N\}\}. \quad (8)$$

Using *mean-field approximation*, we obtain

$$P_i(t) \approx \quad (9)$$

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Approximation by dynamical systems: $M = 1$

Theorem. For the IPCN(1, N) system, the infection vector given by $P(t) := [P_1(t), P_2(t), \dots, P_N(t)]^T$ satisfies the equation

$$P(t) \approx J_{N,1}P(t-1) + C_{N,1} \quad (10)$$

where $J_{N,1} \in \mathbb{R}^{N \times N}$, $C_{N,1} \in \mathbb{R}^{N \times 1}$ are matrices with respective entries:

$$[J_{N,1}]_{i \times j} = \frac{s_{ij}(\rho_j + \delta_{r,j})}{(1 + \delta_{r,j})} - \frac{s_{ij}\rho_j}{(1 + \delta_{b,j})} = s_{ij}(\beta_1^{(j)}(1) - \beta_1^{(j)}(0))$$

$$\text{and } [C_{N,1}]_{1 \times i} = \sum_{j=1}^N \frac{s_{ij}\rho_j}{(1 + \delta_{b,j})} = \sum_{j=1}^N s_{ij}\beta_1^{(j)}(0).$$

The approximating dynamical system for $M = 1$ is linear

Convergence properties: $M = 1$

Theorem. The approximating linear dynamical system for the IPCN(1, N) system given by (10) has a unique equilibrium point given by $P^* = (I - J_{N,1})^{-1}C_{N,1}$ and

$$\lim_{t \rightarrow \infty} P_i(t) = P_i^*$$

for all $i \in \{1, \dots, N\}$.

In the homogeneous case, we obtain an expected result:

Corollary. For a homogeneous IPCN(1, N) system, the equilibrium of (10) is given by $P^* = \rho \mathbf{1}_N$, where $\mathbf{1}_N$ is vector of ones of size N and ρ is the ratio of red balls in the urns.

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Approximation by dynamical systems: $M > 1$

Theorem. For the IPCN(M, N) system, the approximating dynamical system can be written as

$$P_i(t) \approx \sum_{j=1}^N s_{ij} \beta_M^{(j)}(0) + \sum_{j=1}^N \sum_{n=1}^M \left[\left(\sum_{k=0}^n \left((-1)^{n-k} \binom{n}{k} s_{ij} \beta_M^{(j)}(k) \right) \right) \times \left(\sum_{\substack{(d_1, \dots, d_n) \\ \in H_{n,M}}} P_j(t - d_1) \cdots P_j(t - d_n) \right) \right],$$

where

$$H_{n,M} := \{(d_1, d_2, \dots, d_n) \mid d_i \in \{1, \dots, M\}, d_i \neq d_j \forall i, j \in \{1, \dots, n\}\}.$$

The approximating dynamical system for $M > 1$ is highly nonlinear

Linearization for the approximation dynamical systems: $M > 1$

The dynamical system that we have obtained for $M > 1$ are complex to analyze

we further linearize the dynamical systems for the case $M > 1$

Corollary. The linear part of the dynamical system is given by

$$P_i(t) \approx \sum_{j=1}^N s_{ij} \beta_M^{(j)}(0) + \sum_{j=1}^N \sum_{k=1}^M s_{ij} \left(\beta_M^{(j)}(1) - \beta_M^{(j)}(0) \right) P_j(t - k). \quad (11)$$

We now discuss the quality of these approximations

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Simulation Results

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- **For the IPCN(M, N) system:** we plot the average empirical sum at time t , which is given by

$$\frac{1}{N} \sum_{i=1}^N I_t(i) \quad \text{where} \quad I_t(i) = \frac{1}{t} \sum_{n=1}^t Z_{i,n}.$$

For each plot, the average empirical sum is computed 100 times and the mean value is plotted against time.

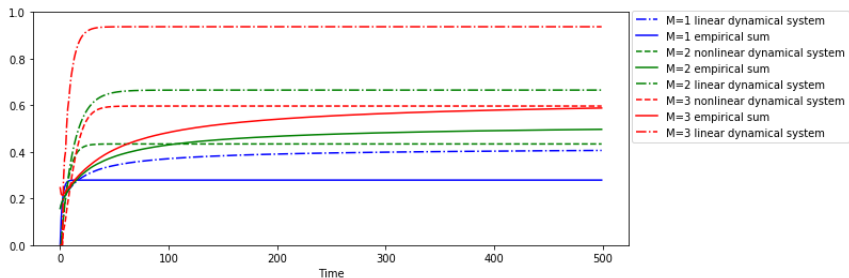
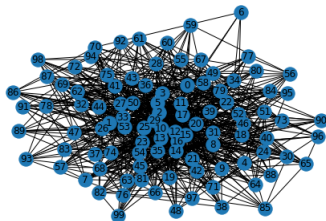
- **For the dynamical systems:** we plot the average infection rate at time t , which is given by

$$\frac{1}{N} \sum_{i=1}^N P_i(t).$$

Highly non-Homogeneous Interacting urns

A 100 node Barabási-Albert network

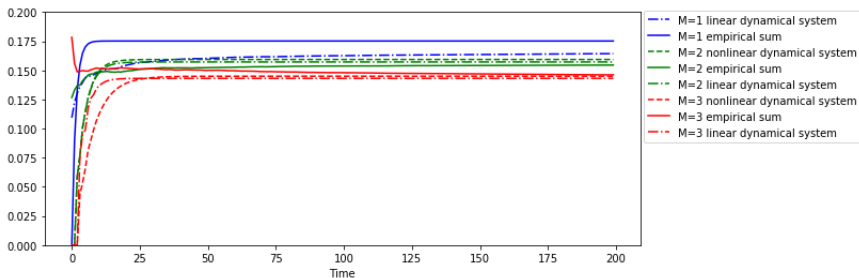
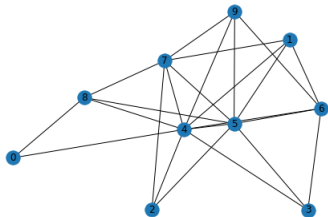
- *The nonlinear approximating dynamical system provides a good approximation and its quality increases with M .*
- *The linearization can perform poorly when the network is highly non-Homogeneous.*



Nearly Homogeneous Interacting urns

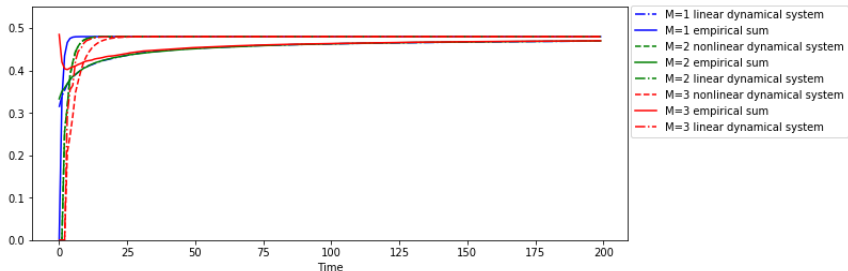
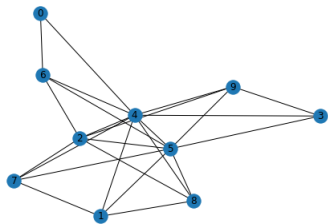
A 10 node network

- *Similar to highly non-Homogeneous case, the nonlinear approximating dynamical system provides a good approximation and its quality increases with M .*
- *The linearization performs well when the network is near homogeneous.*



Homogeneous Interacting urns

For the Homogeneous case, the empirical sum and average infection rate (approximated using dynamical systems) converge to ρ irrespective of the memory of the system.



Conclusions and future directions

- A finite memory interacting Pólya urn network, $\text{IPCN}(M, N)$
- Some stochastic properties $\text{IPCN}(M, N)$
- Approximating dynamical systems for the $\text{IPCN}(M, N)$

Future directions: Many avenues including

- Stability analysis for the approximating non-linear dynamical systems.
- Comparison of $\text{IPCN}(M, N)$ with SIS and SIR models.

Thank You!