

Modeling Restrained Epidemic Routing on Complex Networks

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July 16, 2019

Outline

Introduction

Epidemic Routing

Analysis

Numerical Examples

Conclusion

Background: DTN Routing and Approaches for Realizing Efficiency Delivery

- ▶ DTN (Delay/Disruption Tolerant Networking) routing
 - ▶ **Store-carry-and-forward** message routing
 - ▶ Common performance metrics
 - ▶ Message delivery delay, delivery ratio, throughput
- ▶ Approaches for realizing efficient delivery
 - ▶ Increase the chance of delivery
 - ▶ **Message replication**
 - ▶ Message coding
 - ▶ Avoid congestion/resource starvation
 - ▶ (Utility-based) selection of relaying node
 - ▶ Message replication suppression with ACK
 - ▶ Data compression
 - ▶ Message expiration with TTL (Time-To-Live)

Background: Issues in Concurrent Message Transfer and Broadcasting ACK

- ▶ **Epidemic routing**
 - ▶ Disseminate message replicas through network
 - ▶ A large number of message replicas generated
- ▶ When the amount of workload is **small**...
 - ▶ Can achieve **near-optimal** performance
- ▶ When the amount of workload is **large**...
 - ▶ May result in performance degradation due to many message replicas

Related Works: Modeling (Biological) Virus Dissemination (1/2)

- ▶ **Biological virus** dissemination modeling (1920's–)
 - ▶ Node states
 - ▶ SI (Susceptible Infected) model
 - ▶ SIS (Susceptible Infected Susceptible) model
 - ▶ SIR (Susceptible Infected Recovered/Removed) model
 - ▶ Contact among nodes
 - ▶ **Fully-fixed** (identical nodes, identical contact rates)
 - ▶ Graph (**contact relationship** graph)

Related Works: Modeling (Biological) Virus Dissemination (2/2)

- ▶ **Biological virus** dissemination modeling (1920's–)
 - ▶ Model state
 - ▶ Macro model: the number of nodes in every state (e.g., N_S, N_I)
 - ▶ Micro model: state of every node (e.g., s_1, \dots, s_N)
 - ▶ Model description
 - ▶ Discrete (Markov chain): exact but for small systems
 - ▶ Continuous (differential equations): approximate but can be **large-scale**

Related Works: Epidemic Routing Modeling

- ▶ Mapping from **virus epidemic** to **epidemic routing**
 - ▶ What to be infected
 - ▶ Human being, animals
→ Nodes (terminals)
 - ▶ What infects
 - ▶ Biological (single) virus
→ (**Multiple**) messages
 - ▶ Recovery from infection
 - ▶ Natural recovery
→ **Message discard by TTL or ACK**
 - ▶ Objective
 - ▶ Virus elimination
→ **Rapid/reliable/low-cost message delivery**

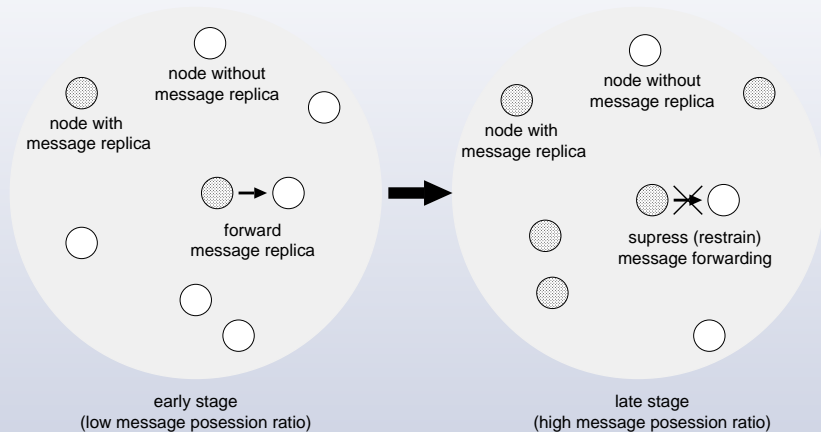
Related Works: Epidemic Routing Modeling

- ▶ Restrained epidemic routing
 - ▶ Objective: Fast message delivery under concurrent messages routing
 - ▶ Idea: **Intentionally refrain** message replication at a later stage
- ▶ Development of epidemic modeling on complex networks
 - ▶ Describe dynamics of all nodes
 - State space explosion
 - ▶ Homogeneous node assumption
 - Loss of graph structure
 - ▶ **Describe dynamics of node classes**
 - Reduction in state space
 - ▶ **DBMF (Degree-Based Mean Field)** approximation (2002)

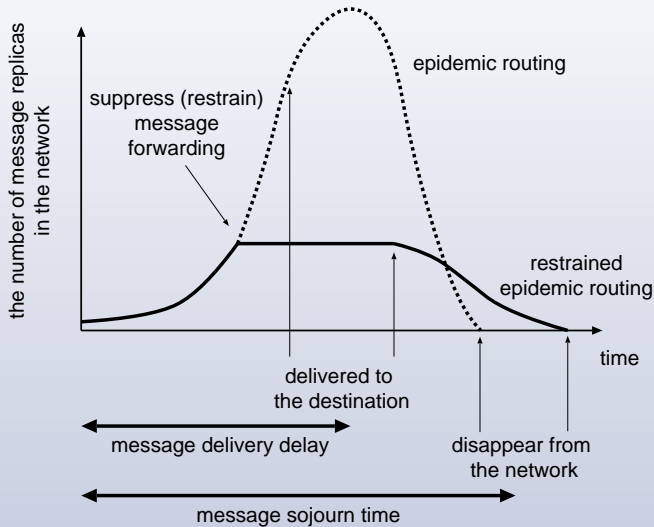
Objective: Modeling Epidemic Routing in Complex Networks

- ▶ Describe the dynamics of **restrained epidemic routing**
- ▶ Using **DBMF (Degree-Based Mean Field)** approximation
 - ▶ Clarify the impact of node contact relationship (complex network) on message delivery

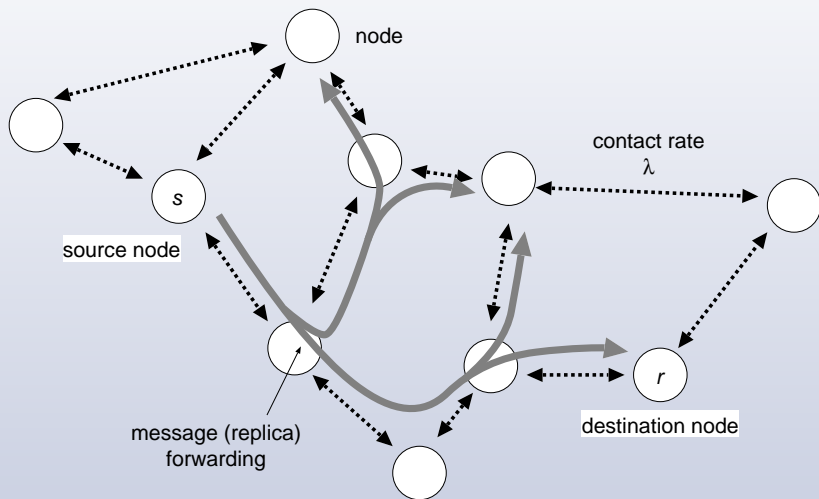
Restrained Epidemic Routing: Overview



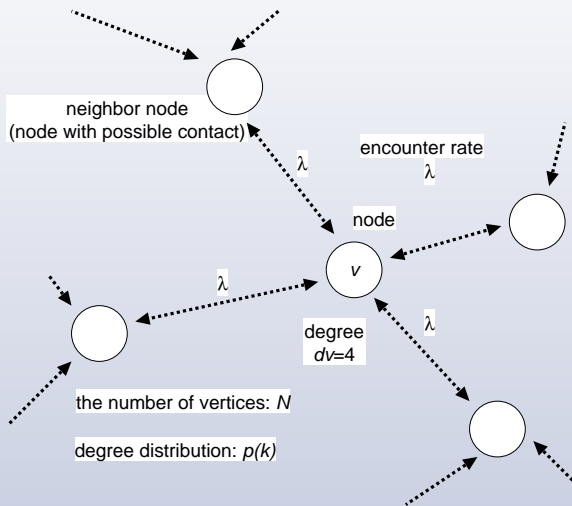
Restrained Epidemic Routing: Comparison with Normal Epidemic Routing



Analytic Model: Message Delivery from Source Node to Destination Node



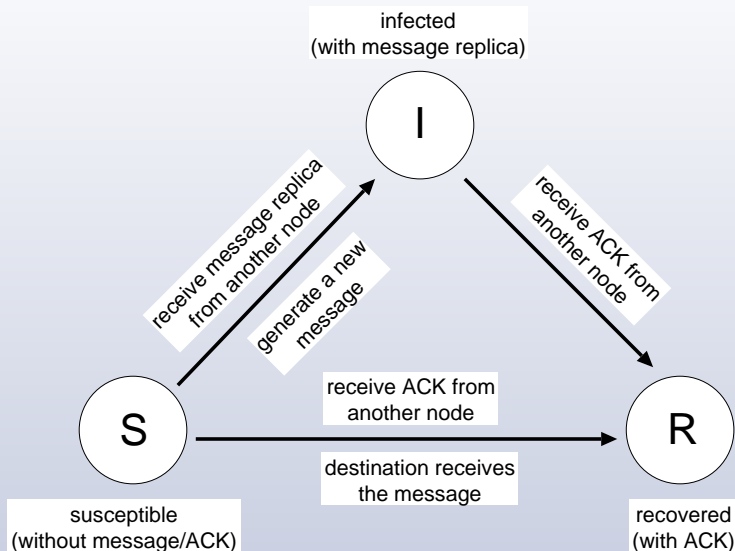
Analytic Model: Representing Node Contacts as Undirected Graph



Assumptions

- ▶ Restrained epidemic routing with broadcast ACK
- ▶ N nodes
- ▶ Message is generated at a source node at $t = 0$
- ▶ Contact duration follows Poisson distribution with mean λ
- ▶ Degree distribution of contact relationship: $P(k)$

Analytic Model: Mapping to SIR Model



Analysis: Initial State to Message Restraint

- ▶ Initial state

$$\rho_k^I(0) = \begin{cases} \frac{1}{NP(k)} & k = d_s \\ 0 & \text{otherwise} \end{cases} \quad (1)$$

$$\rho_k^R(0) = 0 \quad (2)$$

$$\rho_k^S(0) = 0 \quad (3)$$

- ▶ Dynamics of the number of infected nodes in class k

$$\frac{d\rho_k^I(t)}{dt} = \lambda k \rho_k^S(t) \Gamma_k(t) \quad (4)$$

$$\Gamma_k(t) = \sum_{k'} P(k'|k) \rho_{k'}^I(t) \quad (5)$$

Analysis: Message Restraint to Message Delivery

- ▶ The number of infected nodes does not change until delivery completion

$$\frac{d\rho_k^I(t)}{dt} = 0 \quad (6)$$

$$\frac{d\rho_k^R(t)}{dt} = 0 \quad (7)$$

$$\frac{d\rho_k^S(t)}{dt} = 0 \quad (8)$$

- ▶ From message restraint to message delivery

$$t_2 - t_1 = \frac{1}{\lambda d_r \Gamma_{d_r}(t_1)} \quad (9)$$

Analysis: Message Delivery to Broadcast ACK Dissemination

- ▶ When message is delivered

$$\rho_k^I(t_2) = \rho_k^I(t_1) \quad (10)$$

$$\rho_k^R(t_2) = \begin{cases} \frac{1}{NP(d_r)} & k = d_r \\ 0 & \text{otherwise} \end{cases} \quad (11)$$

$$\rho_k^S(t_2) = 1 - \left(\rho_k^I(t_2) + \rho_k^R(t_2) \right) \quad (12)$$

- ▶ Message replica reduction with broadcast ACK

$$\frac{d\rho_k^R(t)}{dt} = \lambda k (1 - \rho_k^R(t)) \Omega_k(t) \quad (13)$$

$$\Omega_k(t) = \sum_{k'} P(k'|k) \rho_{k'}^R(t) \quad (14)$$

Numerical Examples: Three Types of Degree Distributions

► Poisson

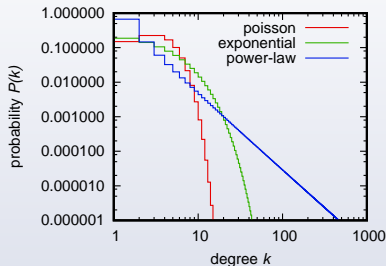
$$P(k) = e^{-\bar{k}} \frac{\bar{k}^k}{k!} \quad (15)$$

► Exponential

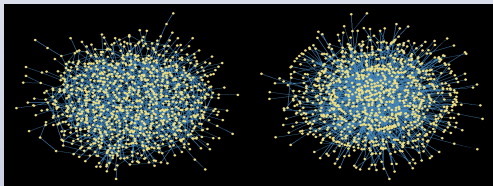
$$P(k) = (1 - e^{-\mu}) e^{-\mu k} \quad (16)$$

► Power-law

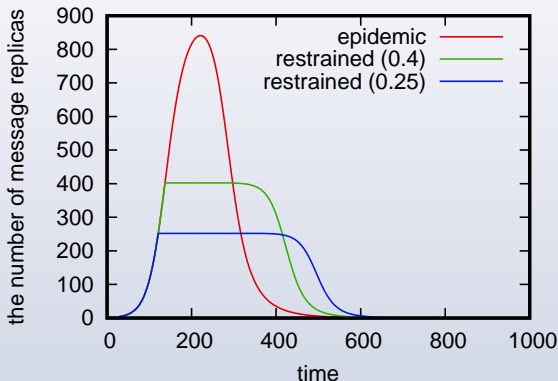
$$P(k) = \frac{k^{-\alpha}}{\zeta(\alpha)} \quad (17)$$



average degree $\bar{k} = 3$

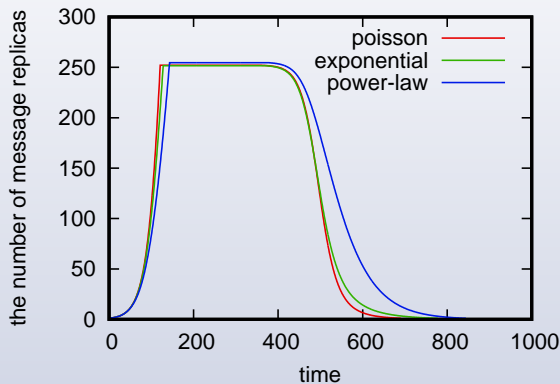


Numerical Example: Evolution of the Number of Message Replicas



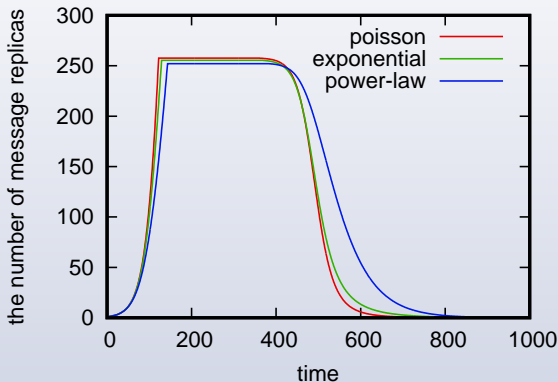
1,000 nodes, degree distribution: Poisson, source and destination degree: 1, contact rate: $1/60$

Numerical Examples: Effect of Contact Rate on Message Delivery



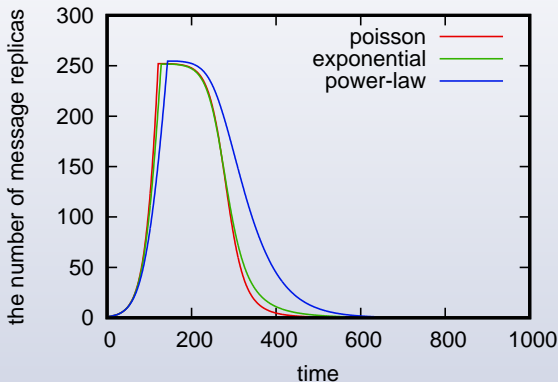
1,000 nodes, source and destination node degree: 1, contact rate: $1/60$, p_T : 0.25

Numerical Examples: Effect of Contact Relationship on Message Delivery



1,000 nodes, source node degree: 10, destination node degree: 1, contact rate: $1/60$, p_T : 0.25

Numerical Examples: Effect of Contact Relationship on Message Delivery



1,000 nodes, source node degree: 1, destination node degree: 10, contact rate: $1/60$, p_T : 0.25

Conclusion

- ▶ Model **restrained epidemic routing**
 - ▶ Contact relationship is given by **a complex network**
 - ▶ Describe the dynamics of a single message routing
- ▶ Derive **average message delivery delay** and **average message sojourn time**
- ▶ Investigate the impact of contact relationship on message delivery
 - ▶ When contact relationship graph is **power-law**
 - ▶ **Message delivery delay is larger** than non-power-law cases

Future Works

- ▶ Derive exact solutions of average message delivery/sojourn times under **specific** degree distributions
- ▶ Modeling resource contention under **multiple concurrent messages routing**
- ▶ Designing a DTN routing mechanism utilizing our analysis