

# Monte-Carlo Estimation of s-t Reliability in Acyclic Networks

Rico Zenklusen

Institute for Operations Research, D-MATH, ETH Zurich

`rico.zenklusen@ifor.math.ethz.ch`

Joint work with Marco Laumanns

European Conference on Complex Systems, Dresden, 1 Oct 2007



# Outline

## 1 Introduction

- The  $s$ - $t$  reliability problem
- Motivation: spreading processes on networks
- Complexity results

## 2 New algorithm for $s$ - $t$ reliability estimation in DAGs

- The method
- Worst-case bounds on the running time
- Computational results

## 3 Conclusions

# Outline

## 1 Introduction

- The  $s$ - $t$  reliability problem
- Motivation: spreading processes on networks
- Complexity results

## 2 New algorithm for $s$ - $t$ reliability estimation in DAGs

- The method
- Worst-case bounds on the running time
- Computational results

## 3 Conclusions

## The s-t reliability problem

In classical network reliability problems we consider a graph whose **edges fail independently** of each other with some given probability.

Given is a network  $G = (V, E, p)$  with

- $V$ : set of vertices
- $E$ : set of edges (directed, undirected or mixed)
- $p(e)$ : failure probability of edge  $e$  ( $\forall e \in E$ )

and  $s, t \in V$  two special terminals.

### Definition (s-t reliability)

The s-t reliability  $\text{REL}_{s,t}(G)$  is the **probability** that the graph obtained from  $G$  after edge failures contains a **path from s to t**.

# Outline

## 1 Introduction

- The  $s$ - $t$  reliability problem
- **Motivation: spreading processes on networks**
- Complexity results

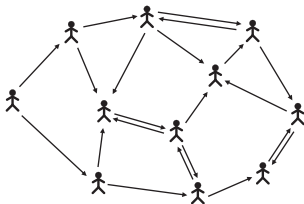
## 2 New algorithm for $s$ - $t$ reliability estimation in DAGs

- The method
- Worst-case bounds on the running time
- Computational results

## 3 Conclusions

## SIR disease spreading model

- Continuous time model for the spread of a disease over the social network of a fixed population.

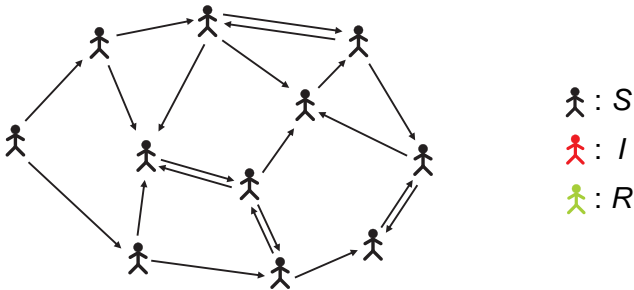


- At every time each individual is in one of the states  $S$ ,  $I$  or  $R$ .



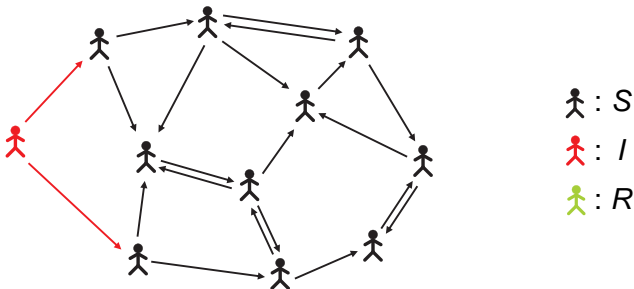
## SIR model (example simulation)

Every node has an associated recovery time and every edge an associated infection rate.



## SIR model (example simulation)

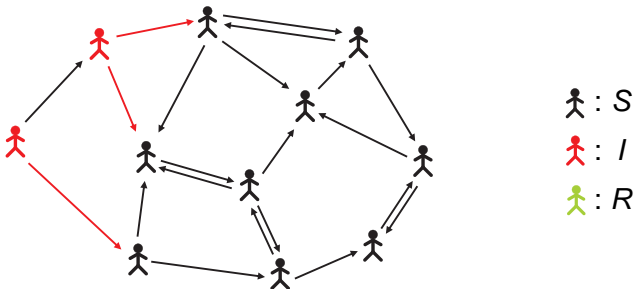
Every node has an associated recovery time and every edge an associated infection rate.



Initial outbreak of a disease.

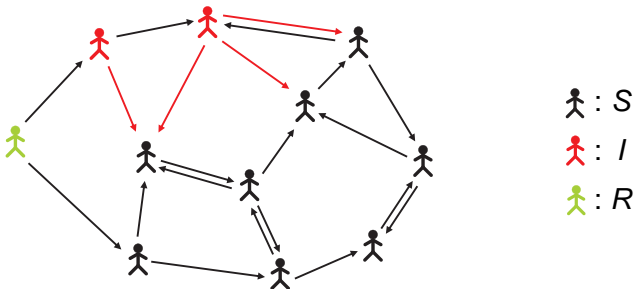
## SIR model (example simulation)

Every node has an associated recovery time and every edge an associated infection rate.



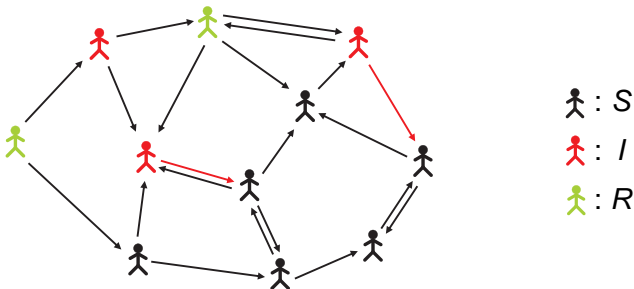
## SIR model (example simulation)

Every node has an associated recovery time and every edge an associated infection rate.



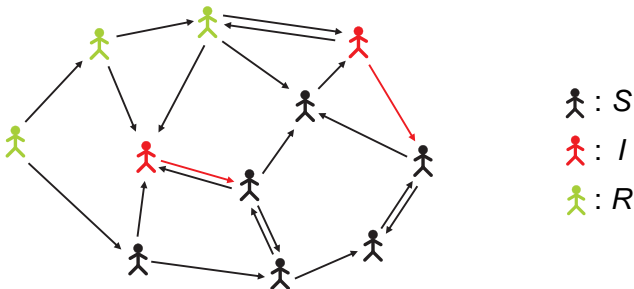
## SIR model (example simulation)

Every node has an associated recovery time and every edge an associated infection rate.



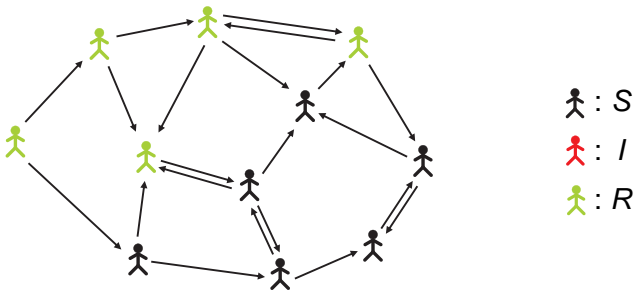
## SIR model (example simulation)

Every node has an associated recovery time and every edge an associated infection rate.



## SIR model (example simulation)

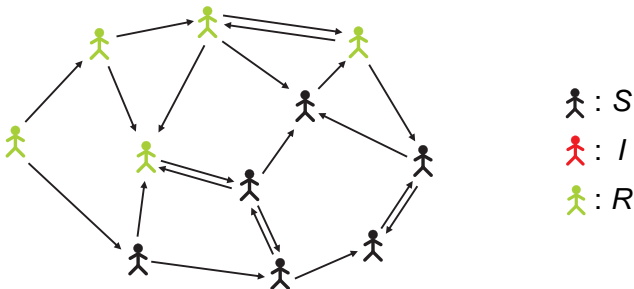
Every node has an associated recovery time and every edge an associated infection rate.



Final state of the disease spreading.

## SIR model (example simulation)

Every node has an associated recovery time and every edge an associated infection rate.

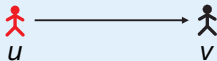


What is the probability that some fixed individual gets infected?

# Mapping SIR models to reliability problems

The **final state** of a disease spreading can be **expressed by a classical reliability model** on the social network with edge intactness probabilities defined as follows. [Grassberger, 1983]

$q(u, v) = 1 - p(u, v)$ : Probability that  $u$  infects  $v$  in the situation where  $u$  just switched to state  $I$  and  $v$  is in state  $S$ .



## Correspondence between the models

The set of all vertices reachable from the source  $s$  of the disease in the reliability model (after edge failures) corresponds to the part of the population who got infected.

## s-t reliability models in SIR spreadings

Considering the spread of a disease starting at node  $s$ , the  $s$ - $t$  reliability corresponds to the probability that node  $t$  gets infected.

We are interested in estimating the probability of the **rare event** that the disease spreads over a long distance from  $s$  to  $t$ .

Very small  $s$ - $t$  reliabilities have to be estimated.

Standard Monte-Carlo approach is not efficient for this problem as for getting an “accurate” estimate, the number of samples  $N$  must satisfy:

$$N \propto \frac{1}{\text{REL}_{s,t}(G)}$$

## s-t reliability models in SIR spreadings

Considering the spread of a disease starting at node  $s$ , the  $s$ - $t$  reliability corresponds to the probability that node  $t$  gets infected.

We are interested in estimating the probability of the **rare event** that the disease spreads over a long distance from  $s$  to  $t$ .

**Very small  $s$ - $t$  reliabilities have to be estimated.**

Standard Monte-Carlo approach is not efficient for this problem as for getting an “accurate” estimate, the number of samples  $N$  must satisfy:

$$N \propto \frac{1}{\text{REL}_{s,t}(G)}$$

## s-t reliability models in SIR spreadings

Considering the spread of a disease starting at node  $s$ , the  $s$ - $t$  reliability corresponds to the probability that node  $t$  gets infected.

We are interested in estimating the probability of the **rare event** that the disease spreads over a long distance from  $s$  to  $t$ .

Very small  $s$ - $t$  reliabilities have to be estimated.

Standard Monte-Carlo approach is not efficient for this problem as for getting an “accurate” estimate, the number of samples  $N$  must satisfy:

$$N \propto \frac{1}{\text{REL}_{s,t}(G)}$$

# Outline

## 1 Introduction

- The  $s$ - $t$  reliability problem
- Motivation: spreading processes on networks
- Complexity results

## 2 New algorithm for $s$ - $t$ reliability estimation in DAGs

- The method
- Worst-case bounds on the running time
- Computational results

## 3 Conclusions

# Exact computation of $\text{REL}_{s,t}(G)$

Even on planar acyclic graphs with uniform edge failure probability (i.e.  $p(e) = \bar{p} \forall e \in E$ ) the problem is **#P-complete**.

## Network classes where computation of $\text{REL}_{s,t}(G)$ is easy

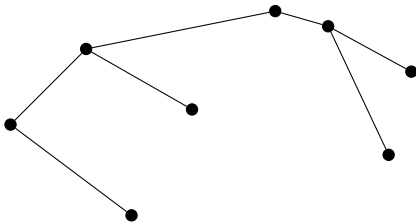
- Trees
- Series-parallel graphs
- Other graphs with bounded tree-width

# Exact computation of $\text{REL}_{s,t}(G)$

Even on planar acyclic graphs with uniform edge failure probability (i.e.  $p(e) = \bar{p} \forall e \in E$ ) the problem is **#P-complete**.

## Network classes where computation of $\text{REL}_{s,t}(G)$ is easy

- **Trees**
- Series-parallel graphs
- Other graphs with bounded tree-width

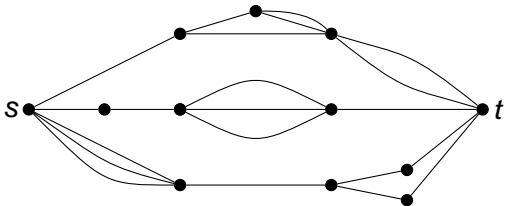


# Exact computation of $\text{REL}_{s,t}(G)$

Even on planar acyclic graphs with uniform edge failure probability (i.e.  $p(e) = \bar{p} \forall e \in E$ ) the problem is **#P-complete**.

## Network classes where computation of $\text{REL}_{s,t}(G)$ is easy

- Trees
- **Series-parallel graphs**
- Other graphs with bounded tree-width

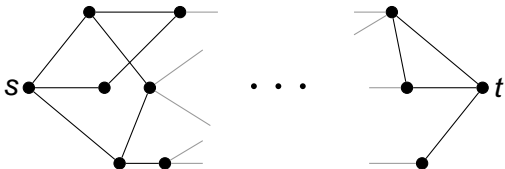


# Exact computation of $\text{REL}_{s,t}(G)$

Even on planar acyclic graphs with uniform edge failure probability (i.e.  $p(e) = \bar{p} \forall e \in E$ ) the problem is **#P-complete**.

## Network classes where computation of $\text{REL}_{s,t}(G)$ is easy

- Trees
- Series-parallel graphs
- Other graphs with bounded tree-width



# Approximating $REL_{s,t}(G)$

## Definition

An  $\epsilon - \delta$  approximation is an algorithm returning with probability at least  $1 - \delta$  a (relative)  $\epsilon$ -approximation.

## Open question

Are there  $\epsilon - \delta$  approximations with running time polynomial in the input size and  $1/\epsilon$  (i.e. FPRAS)?

Even practically useful algorithms for non-trivial large instances are missing.

# Approximating $REL_{s,t}(G)$

## Definition

An  $\epsilon - \delta$  approximation is an algorithm returning with probability at least  $1 - \delta$  a (relative)  $\epsilon$ -approximation.

## Open question

Are there  $\epsilon - \delta$  approximations with running time polynomial in the input size and  $1/\epsilon$  (i.e. FPRAS)?

Even practically useful algorithms for non-trivial large instances are missing.

# Outline

## 1 Introduction

- The  $s$ - $t$  reliability problem
- Motivation: spreading processes on networks
- Complexity results

## 2 New algorithm for $s$ - $t$ reliability estimation in DAGs

- The method
- Worst-case bounds on the running time
- Computational results

## 3 Conclusions

# General remarks

## Goal

Construction of an algorithm allowing to estimate small  $s$ - $t$  reliabilities in large-scale acyclic networks.

## Main idea

Creation of a Monte-Carlo algorithm taking samples over a different space than  $2^E$ , giving more influence on intact states (**importance sampling**).

## Terminology and Notations

We represent the state of a network after edge failures by the set of intact (not failed) edges.

### Notations

- $w(z)$  (for  $z \subseteq E$ ): probability that state  $z$  occurs, i.e.

$$w(z) = \prod_{e \in z} q(e) \prod_{e \in E \setminus z} p(e)$$

- $\mathcal{P} \subseteq 2^E$ : set of all paths from  $s$  to  $t$ .
- $\mathcal{P}(z) \subseteq \mathcal{P}$  (for  $z \subseteq E$ ): set of  $s - t$  paths contained in state  $z$ .
- $A$ : set of all intact states, i.e.  $A = \{a \subseteq E \mid \mathcal{P}(a) \neq \emptyset\}$

We thus have  $w(A) := \sum_{a \in A} w(a) = \text{REL}_{s,t}(G)$ .

## Terminology and Notations

We represent the state of a network after edge failures by the set of intact (not failed) edges.

### Notations

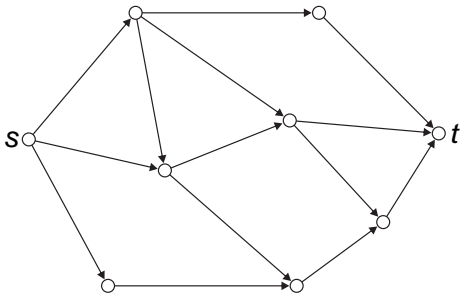
- $w(z)$  (for  $z \subseteq E$ ): probability that state  $z$  occurs, i.e.

$$w(z) = \prod_{e \in z} q(e) \prod_{e \in E \setminus z} p(e)$$

- $\mathcal{P} \subseteq 2^E$ : set of all paths from  $s$  to  $t$ .
- $\mathcal{P}(z) \subseteq \mathcal{P}$  (for  $z \subseteq E$ ): set of  $s - t$  paths contained in state  $z$ .
- $A$ : set of all intact states, i.e.  $A = \{a \subseteq E \mid \mathcal{P}(a) \neq \emptyset\}$

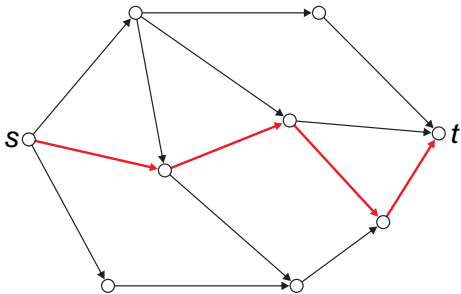
We thus have  $w(A) := \sum_{a \in A} w(a) = \text{REL}_{s,t}(G)$ .

# Sampling intact states



- 1 sample  $s$ - $t$  path
- 2 sample remaining edges

# Sampling intact states

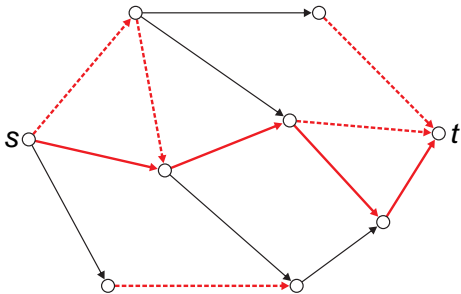


- 1 sample  $s$ - $t$  path
- 2 sample remaining edges

A path  $\gamma \in \mathcal{P}$  is sampled with probability proportional to

$$\prod_{e \in \gamma} q(\gamma).$$

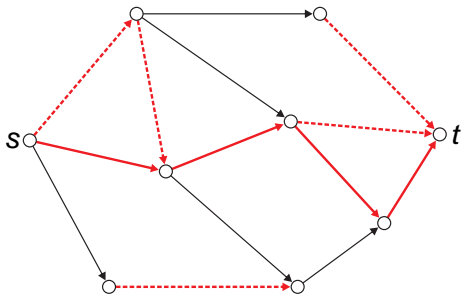
# Sampling intact states



- 1 sample  $s$ - $t$  path
- 2 sample remaining edges

Every edge  $e \in E \setminus \gamma$  is intact with probability equal to  $q(e)$ .

# Sampling intact states



- 1 sample  $s$ - $t$  path
- 2 sample remaining edges

$$\begin{aligned} \{ \text{---} \rightarrow, \text{- - -} \rightarrow \} & : a \\ \{ \text{---} \rightarrow \} & : \gamma \end{aligned}$$

Sample space:  $\Omega = \{(a, \gamma) \in A \times \mathcal{P} \mid \gamma \in \mathcal{P}(a)\}$

Each element  $\omega = (\gamma, a) \in \Omega$  is sampled with probability  $w(\omega) \propto w(a)$  (i.e.,  $w(\omega) = w(a) / \sum_{(a', \gamma') \in \Omega} w(a')$ ).

## Remarks on $\Omega$

Every element  $a \in A$  has  $|\mathcal{P}(a)|$  corresponding elements in  $\Omega$  (**overcounting**).

$w(\Omega) = \sum_{\omega \in \Omega} w(\omega)$ : mean number of intact paths in a state after edge failures (easy to calculate in acyclic graphs).

$w(\Omega)/w(A)$ : mean number of intact paths in an intact state.

$w(\Omega)$  is often a reasonable estimate for low  $s$ - $t$  reliabilities.

### Approach

We will estimate  $\text{REL}_{s,t}(G) = w(A)$  by estimating  $w(A)/w(\Omega)$  with a Monte-Carlo method and multiplying this estimate by  $w(\Omega)$ .

## Remarks on $\Omega$

Every element  $a \in A$  has  $|\mathcal{P}(a)|$  corresponding elements in  $\Omega$  (**overcounting**).

$w(\Omega) = \sum_{\omega \in \Omega} w(\omega)$ : mean number of intact paths in a state after edge failures (easy to calculate in acyclic graphs).

$w(\Omega)/w(A)$ : mean number of intact paths in an intact state.

$w(\Omega)$  is often a reasonable estimate for low  $s$ - $t$  reliabilities.

### Approach

We will estimate  $\text{REL}_{s,t}(G) = w(A)$  by estimating  $w(A)/w(\Omega)$  with a Monte-Carlo method and multiplying this estimate by  $w(\Omega)$ .

## Introducing influence values on $\Omega$ for estimating $w(A)/w(\Omega)$

Influence values will compensate the overcounting.

We associate to each  $\omega = (\gamma, \mathbf{a}) \in \Omega$  an influence value  $m(\omega) = 1/|\mathcal{P}(\mathbf{a})|$ .

The mean influence value of  $\Omega$  is  $w(A)/w(\Omega)$ .

# Outline

## 1 Introduction

- The  $s$ - $t$  reliability problem
- Motivation: spreading processes on networks
- Complexity results

## 2 New algorithm for $s$ - $t$ reliability estimation in DAGs

- The method
- Worst-case bounds on the running time
- Computational results

## 3 Conclusions

# Number of samples to draw

By the theory of zero-one estimators we have:

## Theorem

*The proposed algorithm is an  $\epsilon - \delta$  approximation when the number of samples  $N$  satisfies*

$$N \geq 4(e - 2) \ln \left( \frac{2}{\delta} \right) \frac{1}{\epsilon^2} \frac{w(\Omega)}{\text{REL}_{s,t}(\mathbf{G})} .$$

Finding upper bounds for  $\frac{w(\Omega)}{\text{REL}_{s,t}(\mathbf{G})}$  allows to eliminate the unknown value  $\text{REL}_{s,t}(\mathbf{G})$  in the above theorem.

## Upper bounds for $\frac{w(\Omega)}{\text{REL}_{s,t}(G)}$

An algebraic argument due to Karp and Luby gives the following bound.

$$\frac{w(\Omega)}{\text{REL}_{s,t}(G)} \leq \prod_{e \in E} (1 + q(e))$$

In the case of uniform edge failure probability  $\bar{p} = 1 - \bar{q}$ , the above bound can be sharpened to

$$\frac{w(\Omega)}{\text{REL}_{s,t}(G)} \leq \min \left\{ \left( \frac{2}{2 - \bar{q}} \right)^{m - l_{\min}}, 2^{1 + \frac{\mu}{m} [(\bar{q}m + \ln(2))(\bar{q}m + \ln(2) + l_{\max})]} \right\}$$

where  $l_{\min}$  resp.  $l_{\max}$  are the lengths of an  $s - t$  path in  $G$  with minimal resp. maximal cardinality and  $\mu$  is the maximal edge-to-vertex ratio over all subgraphs of  $G$ .

# Outline

## 1 Introduction

- The  $s$ - $t$  reliability problem
- Motivation: spreading processes on networks
- Complexity results

## 2 New algorithm for $s$ - $t$ reliability estimation in DAGs

- The method
- Worst-case bounds on the running time
- Computational results

## 3 Conclusions

## Instances

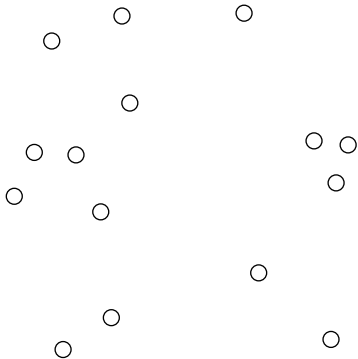
For numerical tests two types of random generators were used for the creation of directed acyclic networks:

- DEL (Delaunay) generator
- TC (Topological construction) generator

## DEL (Delaunay) instances

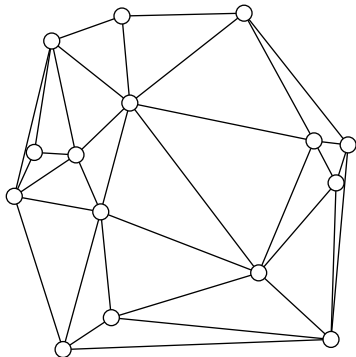
- 1  $N$  vertices are chosen in  $[0, 1]^2$ .
- 2 Construction of Delaunay triangulation.
- 3 Two vertices with maximal Euclidian distance are declared as  $s$  and  $t$ .
- 4 All edges are directed “from  $s$  to  $t$ ”.
- 5 All arcs get uniform intactness probability  $\bar{q}$ .

## DEL (Delaunay) instances



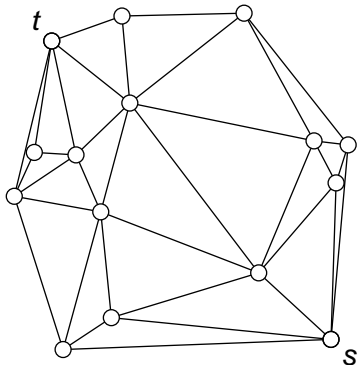
- 1**  $N$  vertices are chosen in  $[0, 1]^2$ .
- 2** Construction of Delaunay triangulation.
- 3** Two vertices with maximal Euclidian distance are declared as  $s$  and  $t$ .
- 4** All edges are directed “from  $s$  to  $t$ ”.
- 5** All arcs get uniform intactness probability  $\bar{q}$ .

## DEL (Delaunay) instances



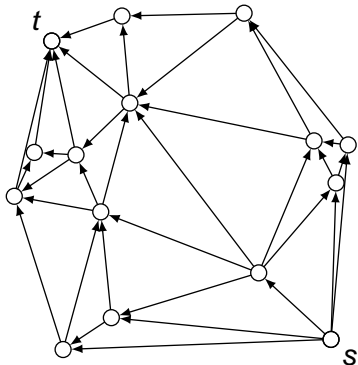
- 1**  $N$  vertices are chosen in  $[0, 1]^2$ .
- 2** Construction of Delaunay triangulation.
- 3** Two vertices with maximal Euclidian distance are declared as  $s$  and  $t$ .
- 4** All edges are directed “from  $s$  to  $t$ ”.
- 5** All arcs get uniform intactness probability  $\bar{q}$ .

## DEL (Delaunay) instances



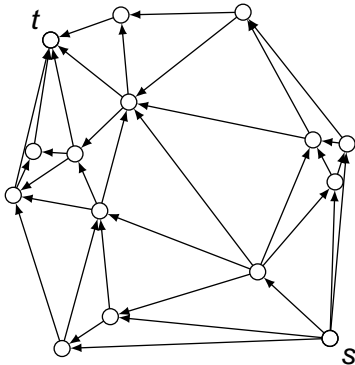
- 1  $N$  vertices are chosen in  $[0, 1]^2$ .
- 2 Construction of Delaunay triangulation.
- 3 Two vertices with maximal Euclidian distance are declared as  $s$  and  $t$ .
- 4 All edges are directed “from  $s$  to  $t$ ”.
- 5 All arcs get uniform intactness probability  $\bar{q}$ .

## DEL (Delaunay) instances



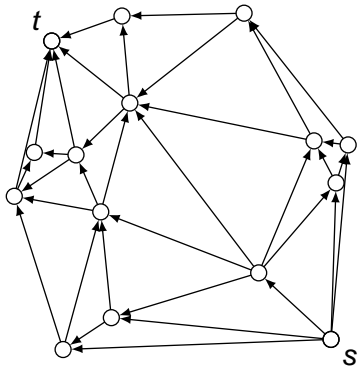
- 1  $N$  vertices are chosen in  $[0, 1]^2$ .
- 2 Construction of Delaunay triangulation.
- 3 Two vertices with maximal Euclidian distance are declared as  $s$  and  $t$ .
- 4 All edges are directed “from  $s$  to  $t$ ”.
- 5 All arcs get uniform intactness probability  $\bar{q}$ .

# DEL (Delaunay) instances



- 1  $N$  vertices are chosen in  $[0, 1]^2$ .
- 2 Construction of Delaunay triangulation.
- 3 Two vertices with maximal Euclidian distance are declared as  $s$  and  $t$ .
- 4 All edges are directed “from  $s$  to  $t$ ”.
- 5 All arcs get uniform intactness probability  $\bar{q}$ .

## DEL (Delaunay) instances



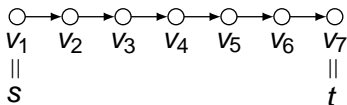
Parameters:  $N, \bar{q}$

- 1  $N$  vertices are chosen in  $[0, 1]^2$ .
- 2 Construction of Delaunay triangulation.
- 3 Two vertices with maximal Euclidian distance are declared as  $s$  and  $t$ .
- 4 All edges are directed “from  $s$  to  $t$ ”.
- 5 All arcs get uniform intactness probability  $\bar{q}$ .

## TC (Topological construction) instances

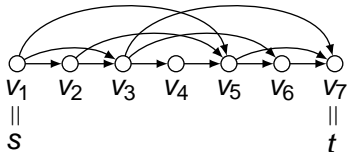
- 1 Start with a path over  $N$  vertices:  
 $s = v_1 \rightarrow v_2 \rightarrow \dots \rightarrow v_n = t$ .
- 2 Every other possible arc is inserted with probability  $\lambda$ .
- 3 Edge  $(v_i, v_j)$  gets intactness probability uniformly distributed in  $[0, 1/(j-i)^{1-\alpha}]$ .

# TC (Topological construction) instances



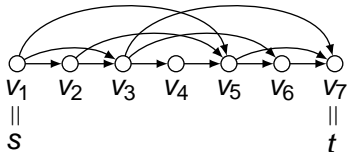
- 1 Start with a path over  $N$  vertices:  
 $S = v_1 \rightarrow v_2 \rightarrow \dots \rightarrow v_n = t$ .
- 2 Every other possible arc is inserted with probability  $\lambda$ .
- 3 Edge  $(v_i, v_j)$  gets intactness probability uniformly distributed in  $[0, 1/(j-i)^{1-\alpha}]$ .

# TC (Topological construction) instances



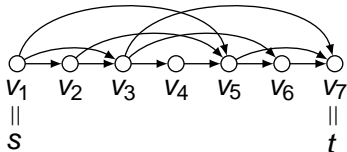
- 1 Start with a path over  $N$  vertices:  
 $s = v_1 \rightarrow v_2 \rightarrow \dots \rightarrow v_n = t$ .
- 2 Every other possible arc is inserted with probability  $\lambda$ .
- 3 Edge  $(v_i, v_j)$  gets intactness probability uniformly distributed in  $[0, 1/(j-i)^{1-\alpha}]$ .

# TC (Topological construction) instances



- 1 Start with a path over  $N$  vertices:  
 $s = v_1 \rightarrow v_2 \rightarrow \dots \rightarrow v_n = t$ .
- 2 Every other possible arc is inserted with probability  $\lambda$ .
- 3 Edge  $(v_i, v_j)$  gets intactness probability uniformly distributed in  $[0, 1/(j-i)^{1-\alpha}]$ .

## TC (Topological construction) instances

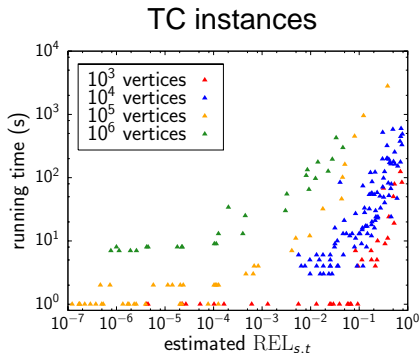
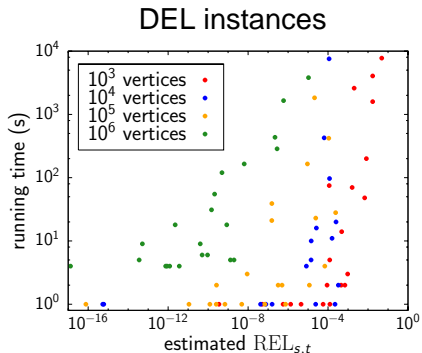


- 1 Start with a path over  $N$  vertices:  
 $s = v_1 \rightarrow v_2 \rightarrow \dots \rightarrow v_n = t$ .
- 2 Every other possible arc is inserted with probability  $\lambda$ .
- 3 Edge  $(v_i, v_j)$  gets intactness probability uniformly distributed in  $[0, 1/(j-i)^{1-\alpha}]$ .

Parameters:  $N, \lambda, \alpha$

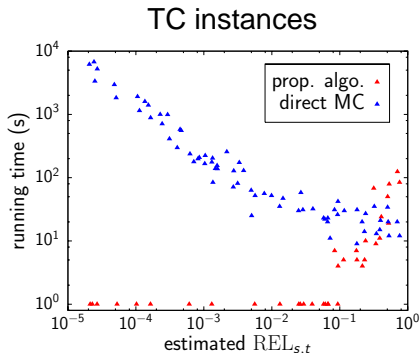
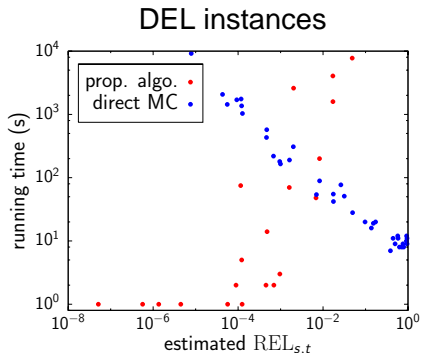
$\lambda$  was chosen such that the expected degree of the vertices is equal to ten.

# Running time of proposed algorithm



The dependence of the running time on  $\epsilon$  and  $\delta$  was essentially as predicted, i.e.  $\propto \ln(1/\delta)/\epsilon^2 \rightarrow$  we fixed  $\epsilon = 0.1$ ,  $\delta = 0.001$ .

# Comparison with direct Monte-Carlo



Instances with 1000 vertices were used for these comparisons.

## Conclusions

- A **new algorithm** for estimating small  $s - t$  reliabilities in large acyclic networks was presented.
- Computational results show the successful application of the proposed algorithm on two types of **large-scale instances** and its **advantage compared to the direct Monte-Carlo approach**.
- For uniform edge failure probabilities, a **worst-case bound on the number of samples to be drawn** was given which sharpens a previous result by Karp and Luby.