



# Social Data Management

## Probabilistic Graphs and Influence Algorithms

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**Graphs:** a natural way to represent data in various domains

- **transport data:** road, air links between locations
- **social networks:** relationships between humans, citation networks
- **interactions between proteins:** contacts due to biochemical processes

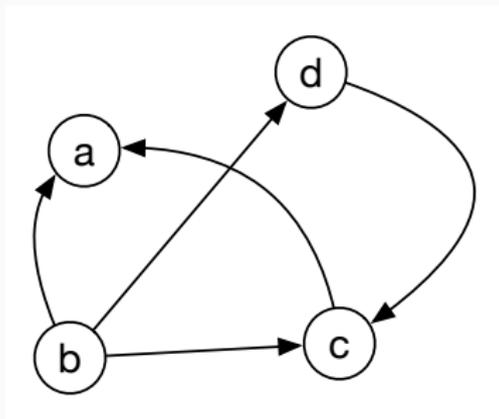
# Uncertain Graphs

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- **transport data:** road, air links between locations
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For all the above examples, the links are not exact. (*Why?*)

## (Deterministic) Graphs

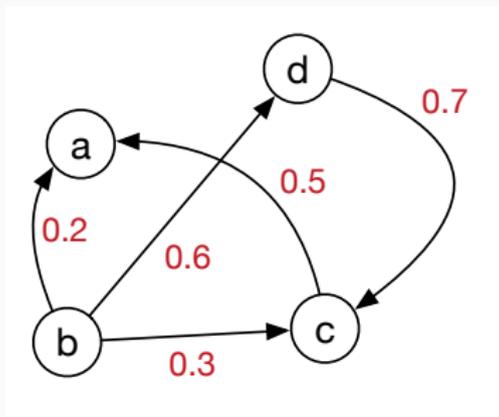


A graph  $G = (V, E)$  is formed of

- a set  $V$  of vertices (nodes)
- a set  $E \subseteq V \times V$ , of edges

# Uncertain Graphs

An **uncertain graph**  $\mathcal{G} = (V, E, p)$  is formed of



- a set  $V$  of vertices (nodes)
- a set  $E \subseteq V \times V$ , of edges
- a function  $p : E \rightarrow [0, 1]$ , representing the **probability**  $p_e$  that the edge  $e \in E$  exists or not

*What are the possible worlds and their probability for this model?*

## Uncertain Graphs: Possible Worlds

A **possible world** of  $\mathcal{G}$ , denoted  $G \sqsubseteq \mathcal{G}$  is a *deterministic* graph  $G = (V, E_G)$  where each  $e \in E_G$  is chosen from  $E$

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The probability of  $G$  is:

$$\Pr[G] = \prod_{e \in E_G} p_e \prod_{e \in E \setminus E_G} (1 - p_e)$$

*How many possible worlds are there?*

## Uncertain Graphs: Other models

Other models are possible:

- each edge is replaced by a **distribution of weights** – instead of choosing if the edge exists or not, a possible world is an instantiation of weights
- each edge has a **formula of events**, capturing **correlations**
- probabilities can be on **nodes** also – equivalent to the edge model (*Why?*)

# Queries on Uncertain Graphs

Generally, the queries we want to answer are **distance** queries:

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$$p_{s,t}(d) = \sum_{G|d_G(s,t)=d} \Pr[G]$$

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Multiple uses of distance queries:

- link prediction, social search, travel estimation

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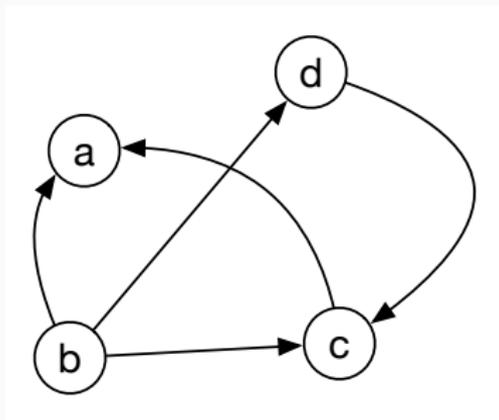
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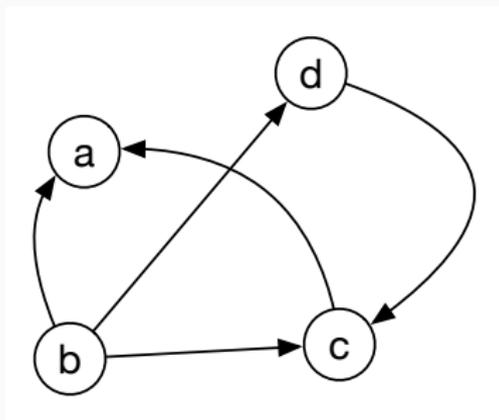
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## Queries on Uncertain Graphs



What is the distance (in hops) between  $b$  and  $a$  ?

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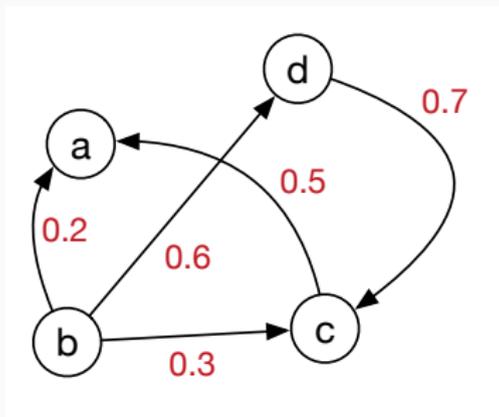


What is the distance (in hops) between  $b$  and  $a$  ?

- BFS search (or Dijkstra's algorithms) finds the edge  $b \rightarrow a$
- the cost is  $O(E)$  (linear in the size of the graph)

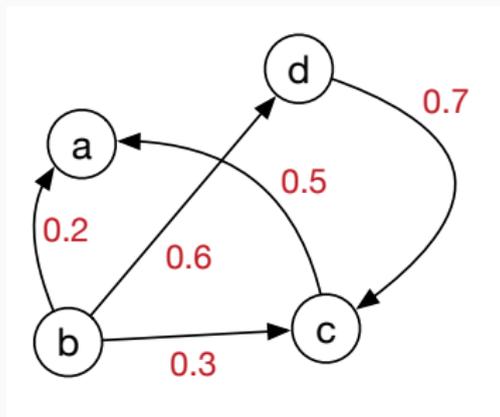
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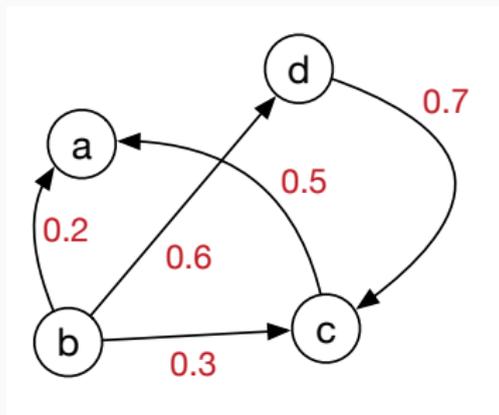


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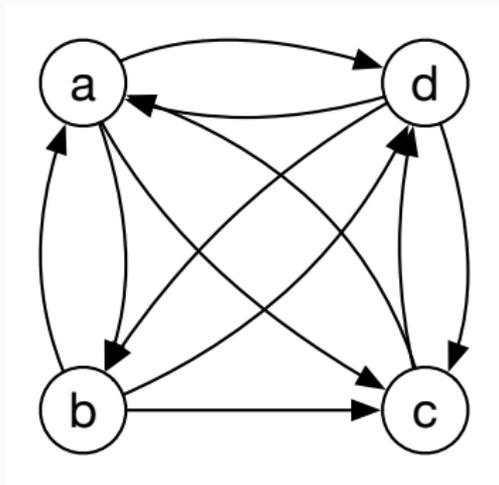
- the edge  $b \rightarrow a$  does not appear in all possible worlds:

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- there are two possible paths of distance 2 ( $b \rightarrow c \rightarrow a$ ) and 3 ( $b \rightarrow d \rightarrow c \rightarrow a$ )

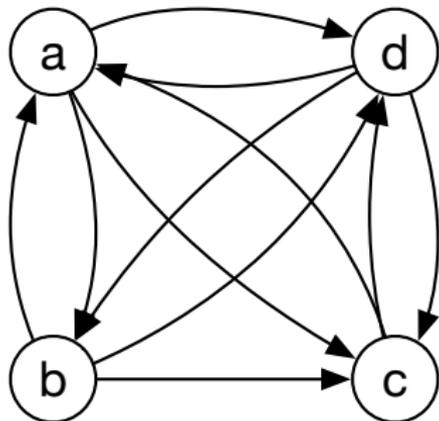
$$p_{b,a}(1) = (1 - p_{b,a}(1)) \times p(b \rightarrow c \rightarrow a)$$

## Queries on Uncertain Graphs



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## Queries on Uncertain Graphs



What is the distance (in hops) between  $b$  and  $a$  ?

- the number of paths is **exponential** in the size of the graph
- specifically, there are  $3!$  paths

## Queries on Uncertain Graphs

Distance query answering in **uncertain graphs** is at least as hard as in relational databases (*logical formulas* of paths; the number of which can be **exponential**)

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Distance query answering in **uncertain graphs** is at least as hard as in relational databases (*logical formulas* of paths; the number of which can be **exponential**)

Computing the reachability probability (i.e, computing the probability of there being a path between a source and a target) is known to be  $\#P$  hard [Valiant, SIAM J. Comp, 1979]

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1. generate sampled graphs for  $r$  rounds (is this the optimal way for an  $s, t$  distance estimation?)
2. compute the desired measure (reachability probability, distance distributions) by averaging results

# Computing Answers to Distance Queries on Probabilistic Graphs

Distance estimations in uncertain graphs can be **approximated** via Monte Carlo sampling

1. generate sampled graphs for  $r$  rounds (is this the optimal way for an  $s, t$  distance estimation?)
2. compute the desired measure (reachability probability, distance distributions) by averaging results

Same issue: *how many rounds?*

## Number of Samples: Median Distance

Median distance:

$$d_M(s, t) = \arg \max_D \left\{ \sum_{d=0}^D p_{s,t}(d) \leq \frac{1}{2} \right\}$$

## Number of Samples: Median Distance

Median distance:

$$d_M(s, t) = \arg \max_D \left\{ \sum_{d=0}^D p_{s,t}(d) \leq \frac{1}{2} \right\}$$

Let  $\mu$  be the real median, and  $\alpha$  and  $\beta$  values  $\pm \epsilon N$  away from  $\mu$ .

Then for:

$$r > \frac{c}{\epsilon^2} \log\left(\frac{2}{\delta}\right)$$

and a good choice of  $c$ :

$$\Pr(\hat{\mu} \in [\alpha, \beta]) > 1 - \delta$$

Expected reliable distance (generalization of reliability):

$$d_{\text{ER}}(s, t) = \sum_{d|d<\infty} d \cdot \frac{p_{s,t}(d)}{1 - p_{s,t}(\infty)}$$

## Number of Samples: Expected Distance

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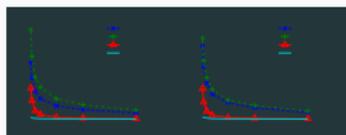
By estimating the connectivity  $\rho$ , we need to sample at least:

$$r \geq \max \left\{ \frac{3}{\epsilon^2 \rho}, \frac{(n-1)^2}{2\epsilon^2} \right\} \cdot \log \left( \frac{2}{\delta} \right)$$

for an  $(\epsilon, \delta)$  approximation.

## Number of Samples In Reality

The number of needed samples can be **surprisingly low** (but it depends on the actual probabilities)



## Sampling Graphs

Generating the entirety of the graph  $G_i$  for each round  $i < r$  is not optimal

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Generating the entirety of the graph  $G_i$  for each round  $i < r$  is not optimal

- we do not need to estimate the entire graph  $G_i$
- we can start from  $s$  and do a BFS or Dijkstra search by sampling **only the outgoing edges**
- based on the generated outgoing edges, we re-do the computation for each generated outgoing node, until we find  $t$

## Example: Median Distance $k$ -NN

$k$ -NN ( $k$  nearest neighbours) – finding the  $k$  nodes from  $s$  the “closest” by some measure

- let us consider the median distance (reminder: it is the highest distance in the distribution that has mass less or equal to 0.5)

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We only care about the top- $k$  nodes, and not their values, and we do not want to evaluate all the graph if possible

- we can evaluate a truncated distribution up to a distance  $D$

$$p_{D,s,t}(d) = \begin{cases} p_{s,t}(d) & \text{if } d < D \\ \sum_{x=D}^{\infty} p_{s,t}(x) & \text{if } d = D \\ 0 & \text{if } d > D \end{cases}$$

- for any two nodes  $t_1, t_2$ ,  $d_{D,M}(s, t_1) < d_{D,M}(s, t_2)$  implies  $d_M(s, t_1) < d_M(s, t_2)$

## Example: Median Distance $k$ -NN

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**Input:** Probabilistic graph  $\mathcal{G} = (V, E, P, W)$ , node  $s \in V$ ,  
number of samples  $r$ , number  $k$ , distance increment  $\gamma$

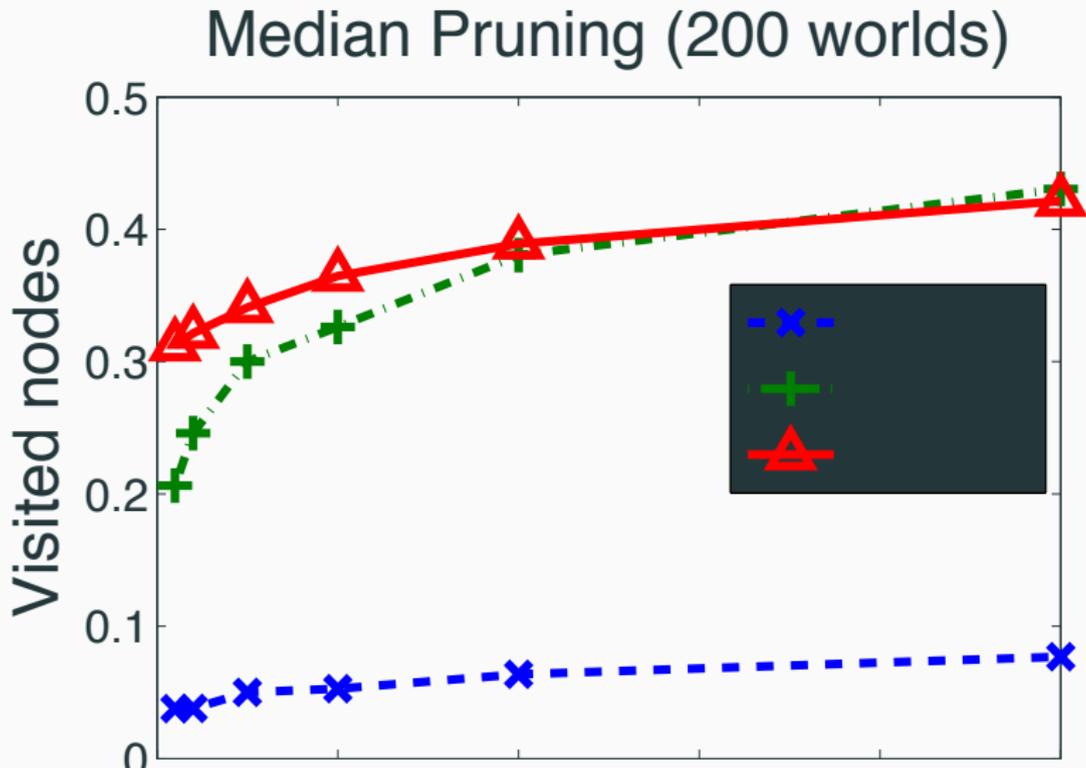
**Output:**  $T_k$ , a result set of  $k$  nodes for the  $k$ -NN query

```
1:  $T_k \leftarrow \emptyset$ ;  $D \leftarrow 0$ 
2: Initiate  $r$  executions of Dijkstra from  $s$ 
3: while  $|T_k| < k$  do
4:    $D \leftarrow D + \gamma$ 
5:   for  $i \leftarrow 1 : r$  do
6:     Continue visiting nodes in the  $i$ -th execution
       of Dijkstra until reaching distance  $D$ 
7:     For each node  $t \in V$  visited
       update the distribution  $\tilde{\mathbf{p}}_{D,s,t}$  {Create the distribu-
       tion  $\tilde{\mathbf{p}}_{D,s,t}$  if  $t$  has never been visited before}
8:   end for
9:   for all nodes  $t \notin T_k$  for which  $\tilde{\mathbf{p}}_{D,s,t}$  exists do
10:    if  $\text{median}(\tilde{\mathbf{p}}_{D,s,t}) < D$  then
11:       $T_k \leftarrow T_k \cup \{t\}$ 
12:    end if
13:  end for
14: end while
```

- start from a small distance  $D$
- decide whether there are nodes to add to the  $k$ -NN set
- increase the distance, and “re-start” each sampled graph from the new distance

## Example: Median Distance $k$ -NN

The algorithm does not need to visit all nodes



Distance Estimation in Uncertain Graphs

Influence Maximization

**Social Influence:** important problem in social network, with applications in marketing, computational advertising

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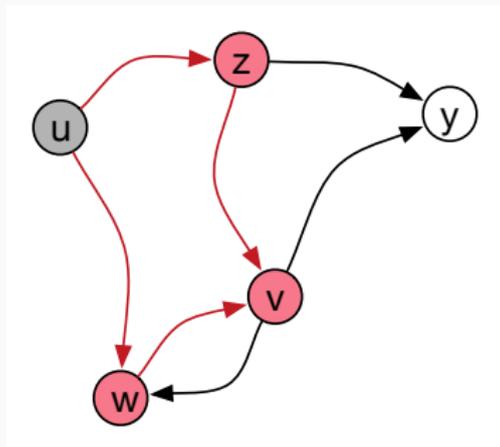
**Objective:** given a promotion budget of  $k$  social network users, maximize the expected influence spread given some influence or propagation model

Data Model: an uncertain graph  $G(V, E, p)$

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- $V$  and  $E$  are the social network
- $p$  is, on each edge, the influence probability

# Influence Spread via Cascades

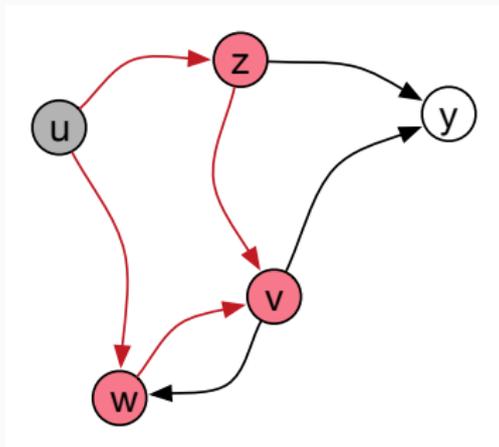


## Independent Cascade Model:

discrete time model of propagation

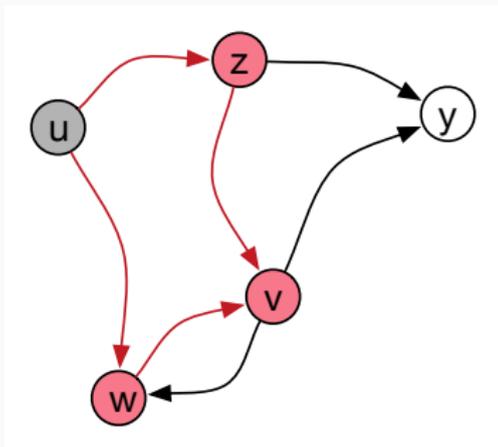
1. at time 0, activate seed  $u$
2. for a node  $i$  activated at time  $t$ :  
activate at time  $t + 1$  each  
neighbour  $v$  with probability  $p_{iv}$
3. once a node is activated, it  
cannot be activated again or  
de-activated

# Influence Spread via Cascades



We wish to compute the **expected spread** from a seed set  $S$ ,  $\sigma(S)$

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We wish to compute the **expected spread** from a seed set  $S$ ,  $\sigma(S)$

By linearity of expectation:

$$\sigma(u) = \sum_{v \in V} \Pr(u \rightarrow v)$$

- for a seed set  $S$ , more complicated
- same hardness as **reachability**

# Maximizing the Influence

Influence maximization is **computationally hard**

Two **sources of hardness**:

1. computing  $\sigma(S)$  is #P-hard (as we seen before, it is equivalent to **reachability**) – Monte Carlo with additive approximations
2. computing the selection of  $k$  seeds in  $S$  is NP-hard – maximization of a **submodular** function

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**Submodular function**: the influence spread is submodular:

$$\sigma(S \cup \{u\}) - \sigma(S) \geq \sigma(T \cup \{u\}) - \sigma(T) \quad \text{if } S \subseteq T$$

## Influence Maximization: Greedy Algorithm

We can obtain a  $(1 - \frac{1}{e})$ -approximation factor for influence maximization by using the **greedy algorithm**

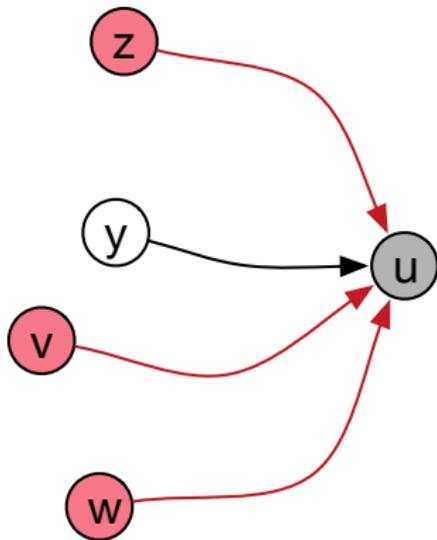
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## Steps:

1. initialize  $S = \emptyset$
2. choose the user  $u$  that maximizes  $\sigma(S \cup \{u\}) - \sigma(S)$
3.  $S = S \cup u$
4. repeat steps 2 and 3  $k$  times
5. **return**  $S$

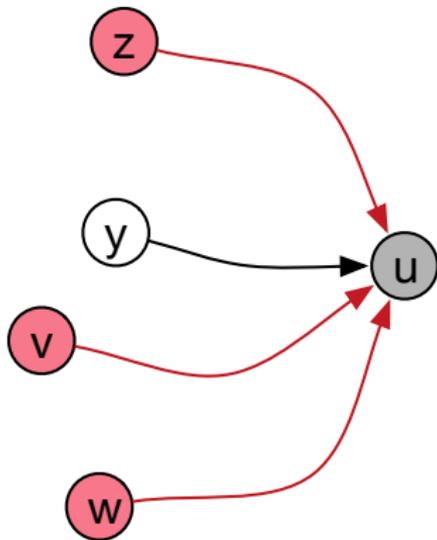
# Learning Propagation Probabilities



The probability that  $v$  is influenced by its neighbours

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Given a log of actions

$A = \{(\text{act}, u, v), \dots\}$ :

1. maximum likelihood:  $p_{vu} = \frac{A_{vu}}{A_v}$
2. Jaccard similarity:  $p_{vu} = \frac{A_{vu}}{A_{u|v}}$

# Acknowledgments

Figures in slides 16 and 20 are taken from [Potamias et al., 2010].



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