

Clash of the Contagions: Cooperation and Competition in Information Diffusion

Seth Myers
Jure Leskovec



Information Diffusion

Information Diffusion

- Users of a social network post and share information with their neighbors

Information Diffusion

- Users of a social network post and share information with their neighbors



Information Diffusion

- Users of a social network post and share information with their neighbors



- Users are constantly exposed to new pieces of information by their neighbors

Information Diffusion

- Users of a social network post and share information with their neighbors



- Users are constantly exposed to new pieces of information by their neighbors
- Most models assume different pieces of information spread from user to user independently

Information Diffusion

- Users of a social network post and share information with their neighbors



- Users are constantly exposed to new pieces of information by their neighbors
- Most models assume different pieces of information spread from user to user independently
- But can one piece of information promote or suppress the spread of another piece of information?

Information Diffusion - Terminology

Information Diffusion - Terminology

- We focus on single pieces of information (rumors, articles, memes, etc) called *contagions*.

Information Diffusion - Terminology

- We focus on single pieces of information (rumors, articles, memes, etc) called *contagions*.
- A user posting a contagion for their neighbors to see (“retweet”, “share”, “repost”, etc.) is called an **adoption**.

Information Diffusion - Terminology

- We focus on single pieces of information (rumors, articles, memes, etc) called *contagions*.
- A user posting a contagion for their neighbors to see (“retweet”, “share”, “repost”, etc.) is called an **adoption**.
- When a user’s neighbor adopts a contagion, the user sees the contagion and is **exposed**.

Information Diffusion - Terminology

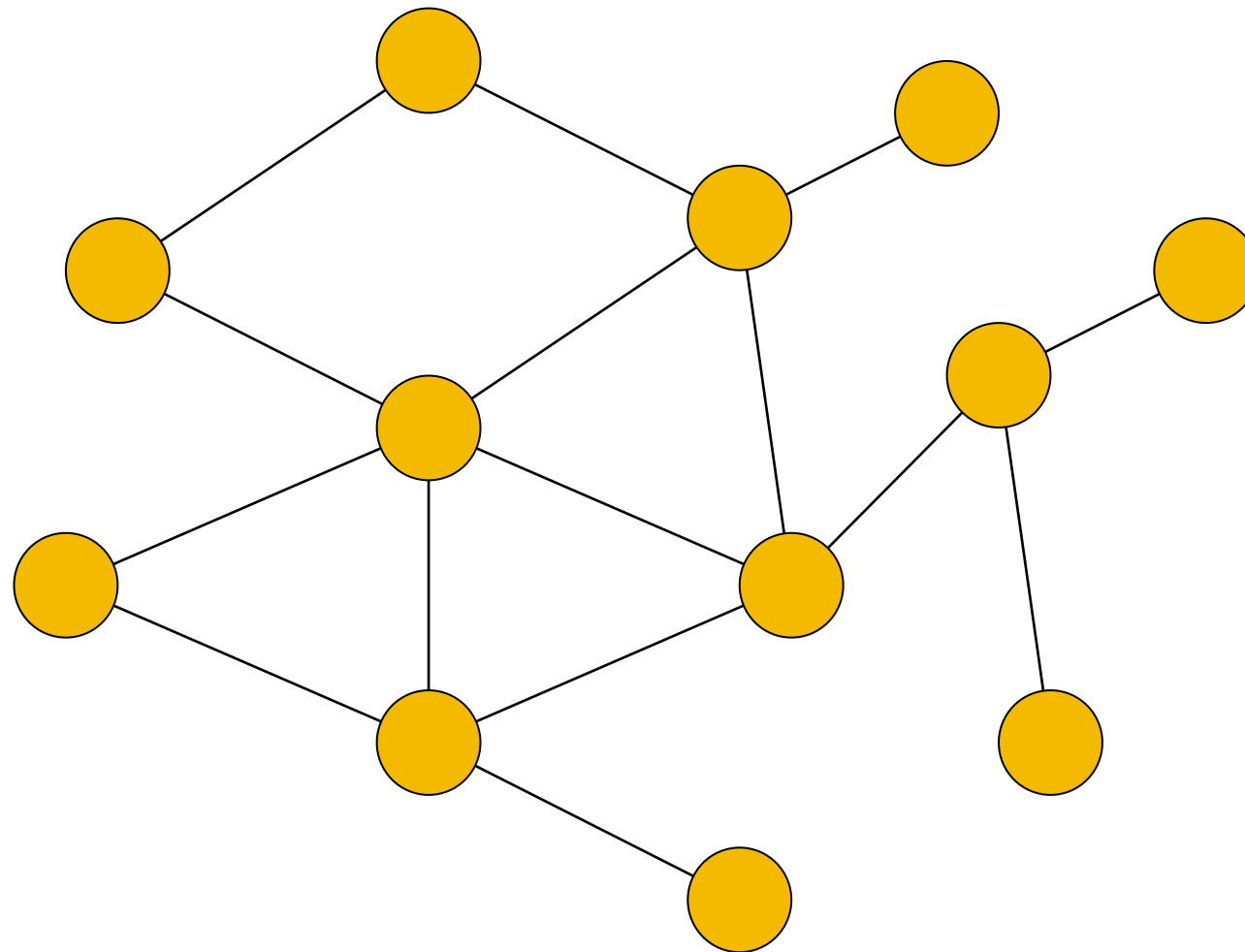
- We focus on single pieces of information (rumors, articles, memes, etc) called *contagions*.
- A user posting a contagion for their neighbors to see (“retweet”, “share”, “repost”, etc.) is called an **adoption**.
- When a user’s neighbor adopts a contagion, the user sees the contagion and is **exposed**.
- Upon **exposure** to a **contagion**, a user will **adopt** the contagion with certain probability.

Information Diffusion

Does exposure to one contagion **increase/decrease** adoption probability of another contagion?

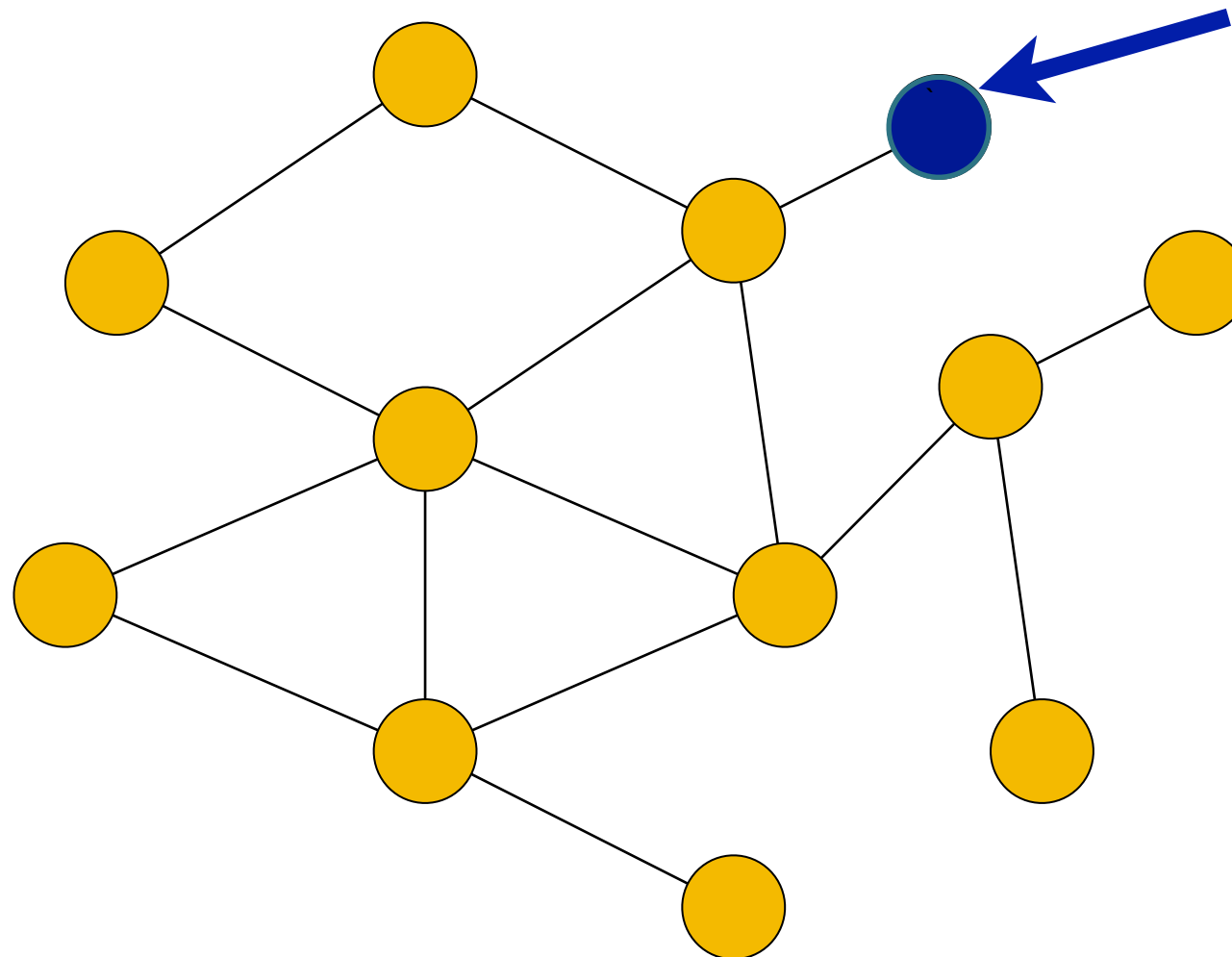
Information Diffusion

Does exposure to one contagion **increase/decrease** adoption probability of another contagion?



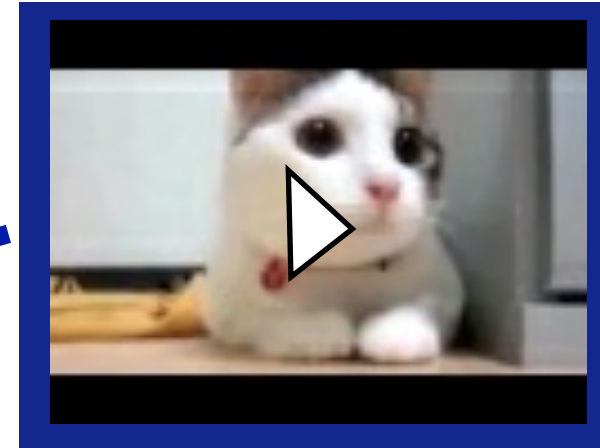
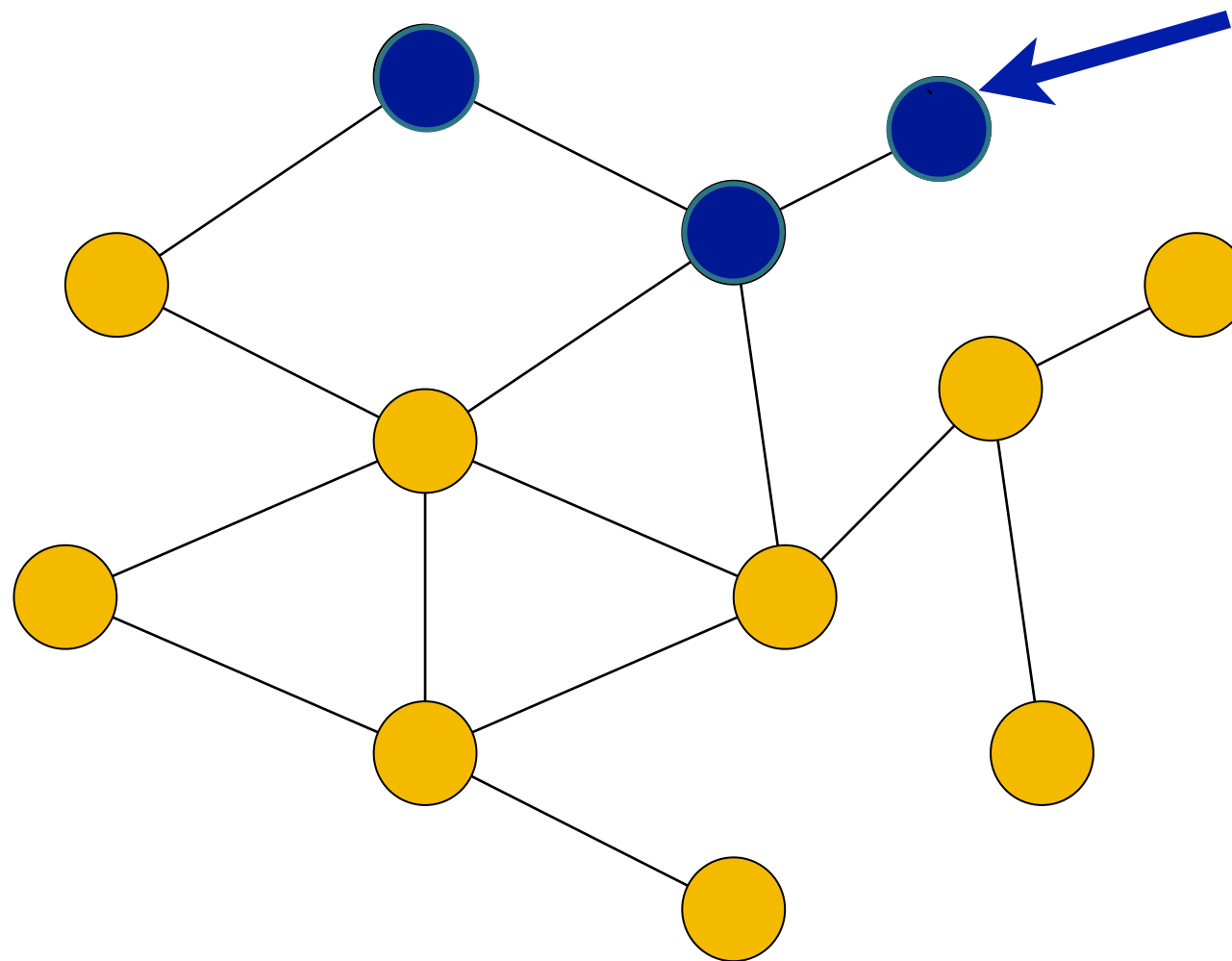
Information Diffusion

Does exposure to one contagion **increase/decrease** adoption probability of another contagion?



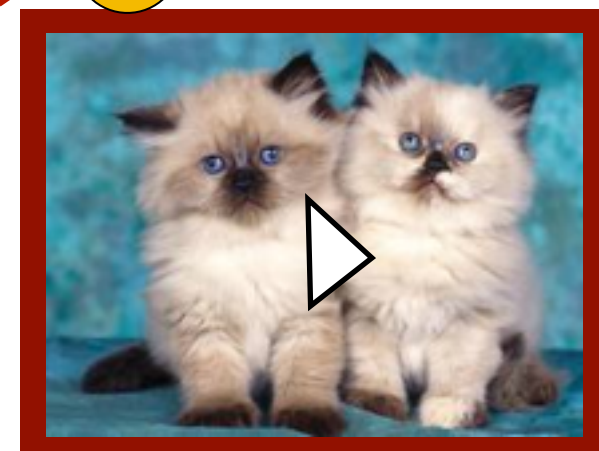
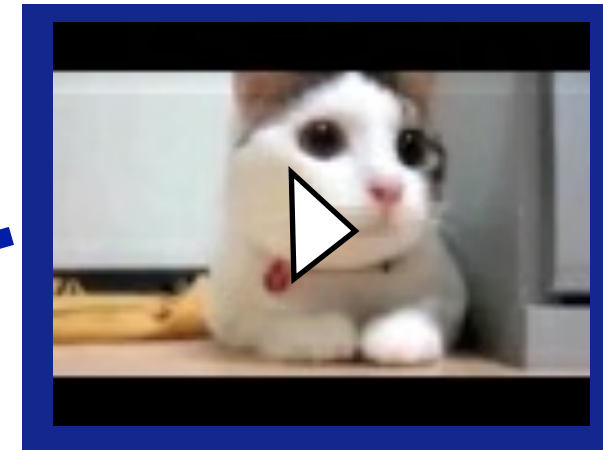
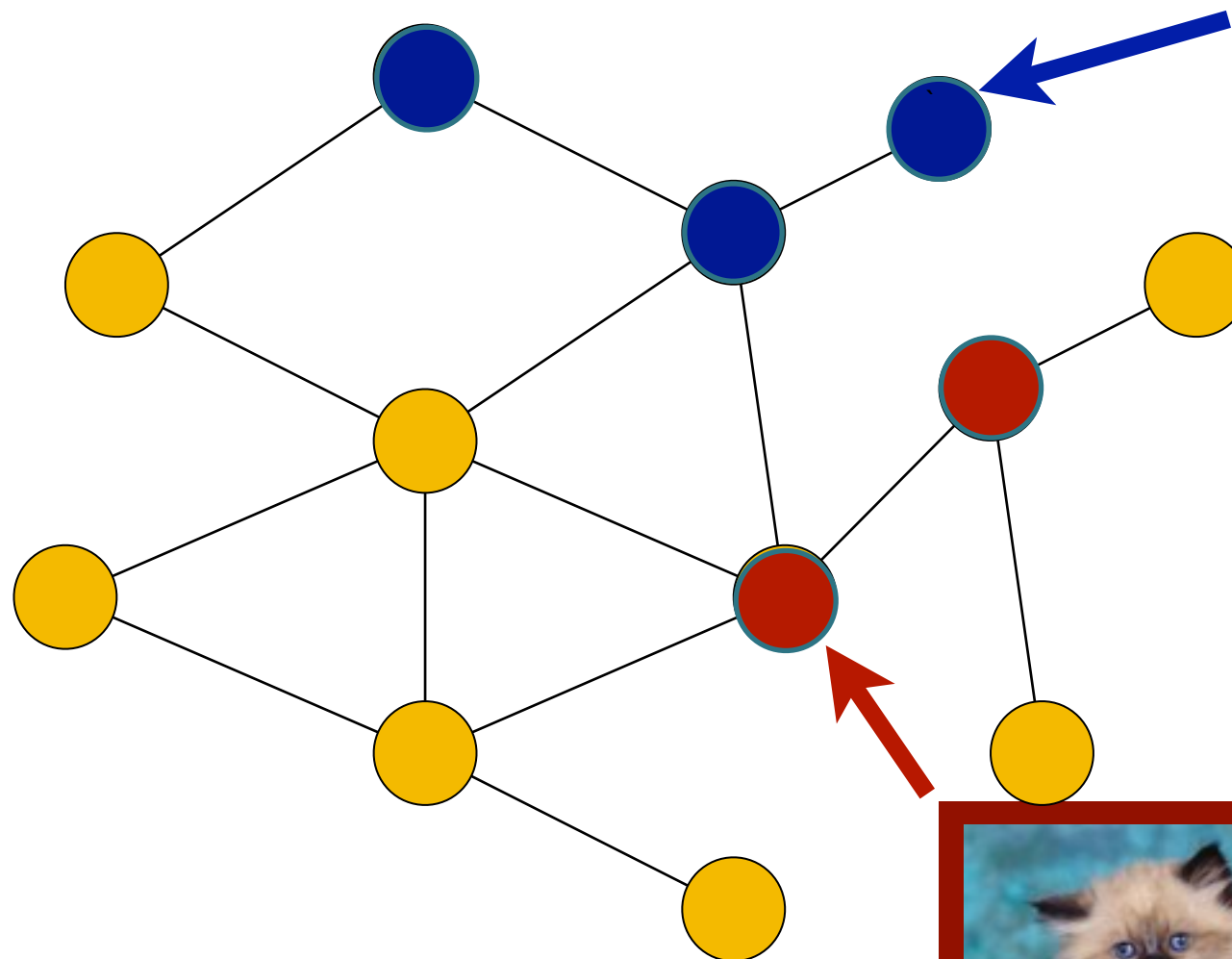
Information Diffusion

Does exposure to one contagion **increase/decrease** adoption probability of another contagion?



Information Diffusion

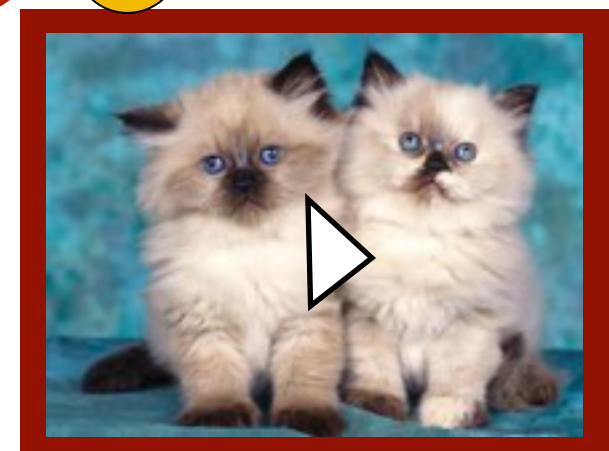
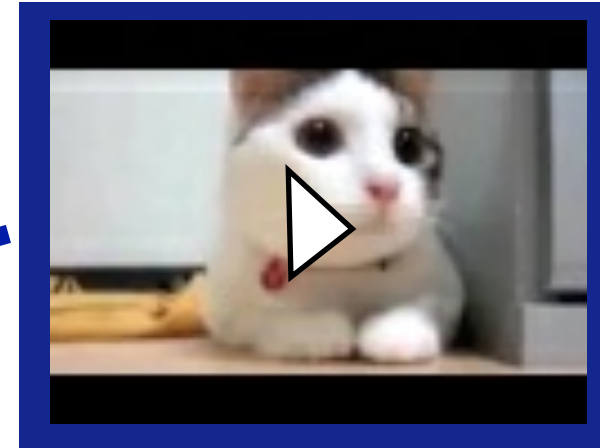
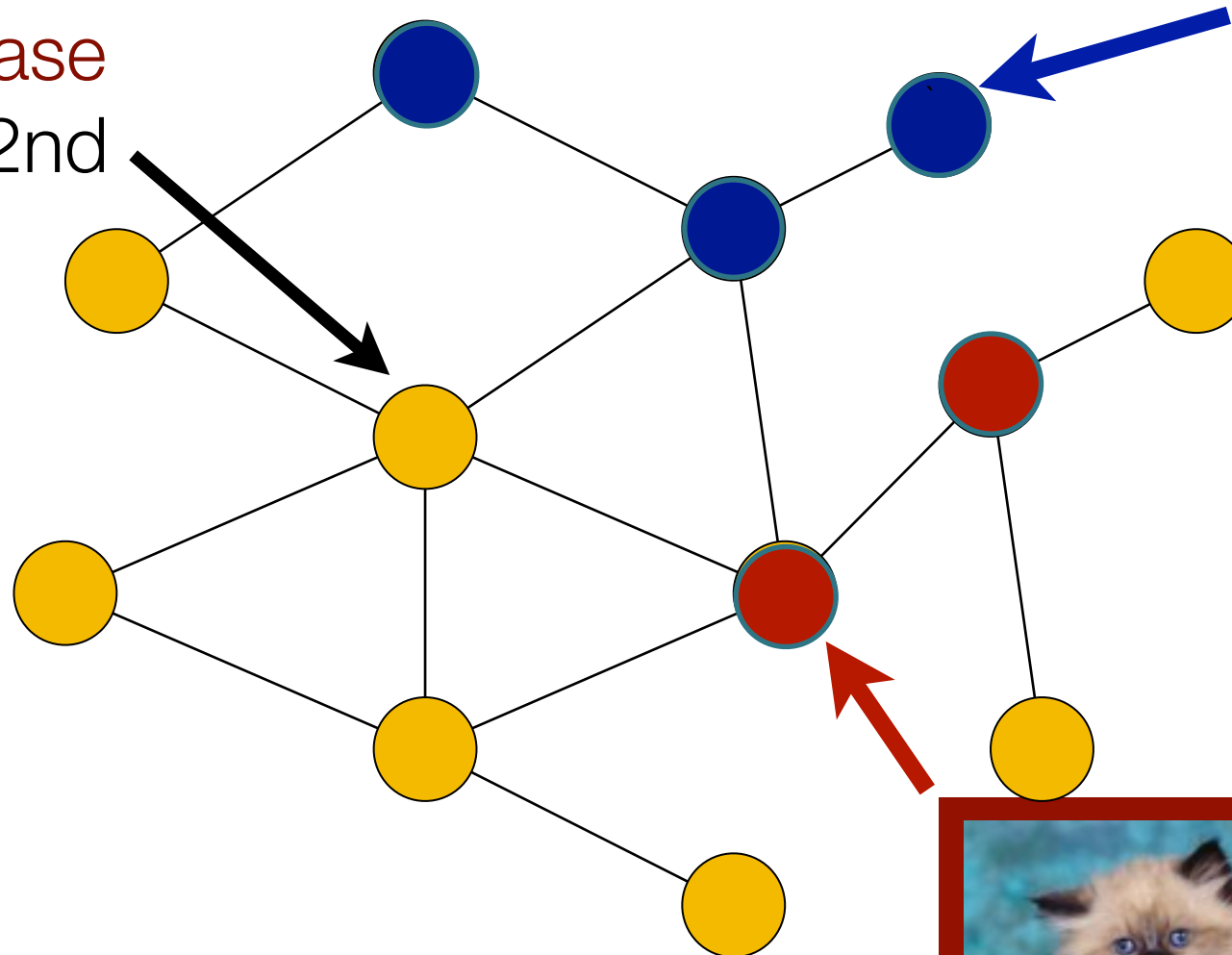
Does exposure to one contagion **increase/decrease** adoption probability of another contagion?



Information Diffusion

Does exposure to one contagion **increase/decrease** adoption probability of another contagion?

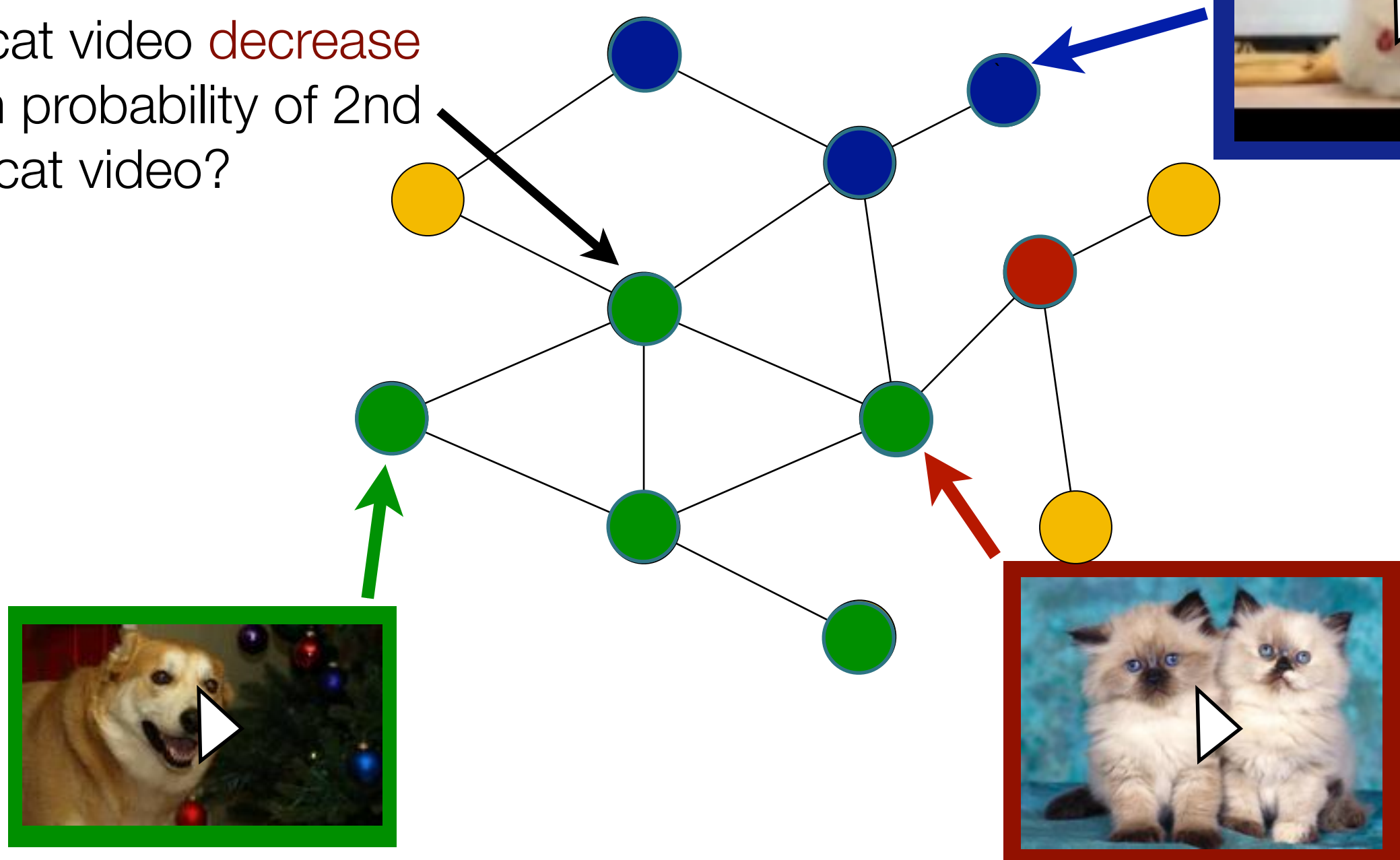
Did 1st cat video **decrease** adoption probability of 2nd cat video?



Information Diffusion

Does exposure to one contagion **increase/decrease** adoption probability of another contagion?

Did 1st cat video **decrease** adoption probability of 2nd cat video?

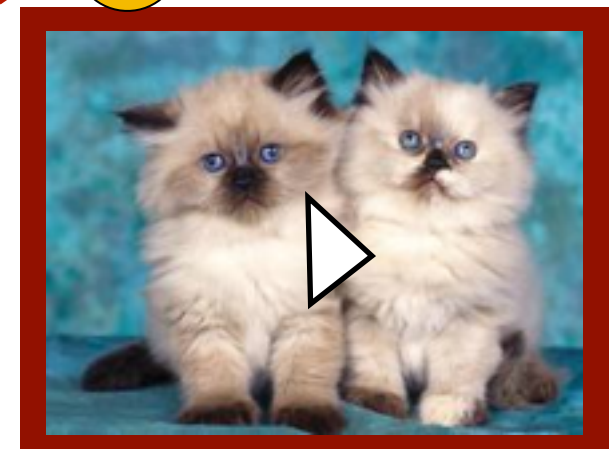
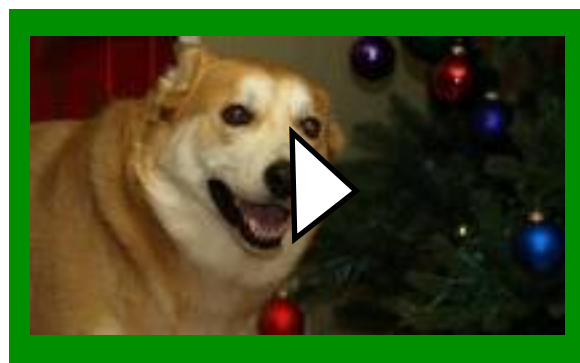
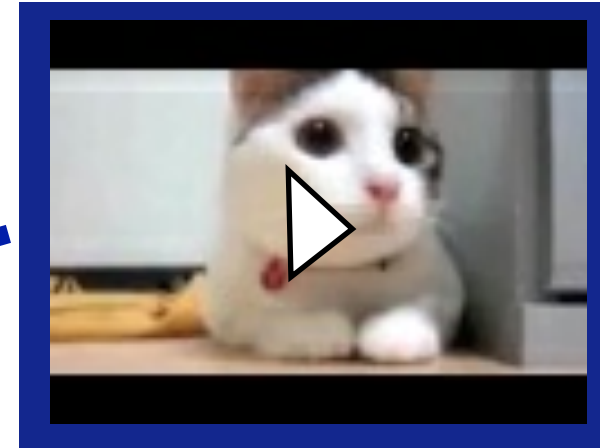
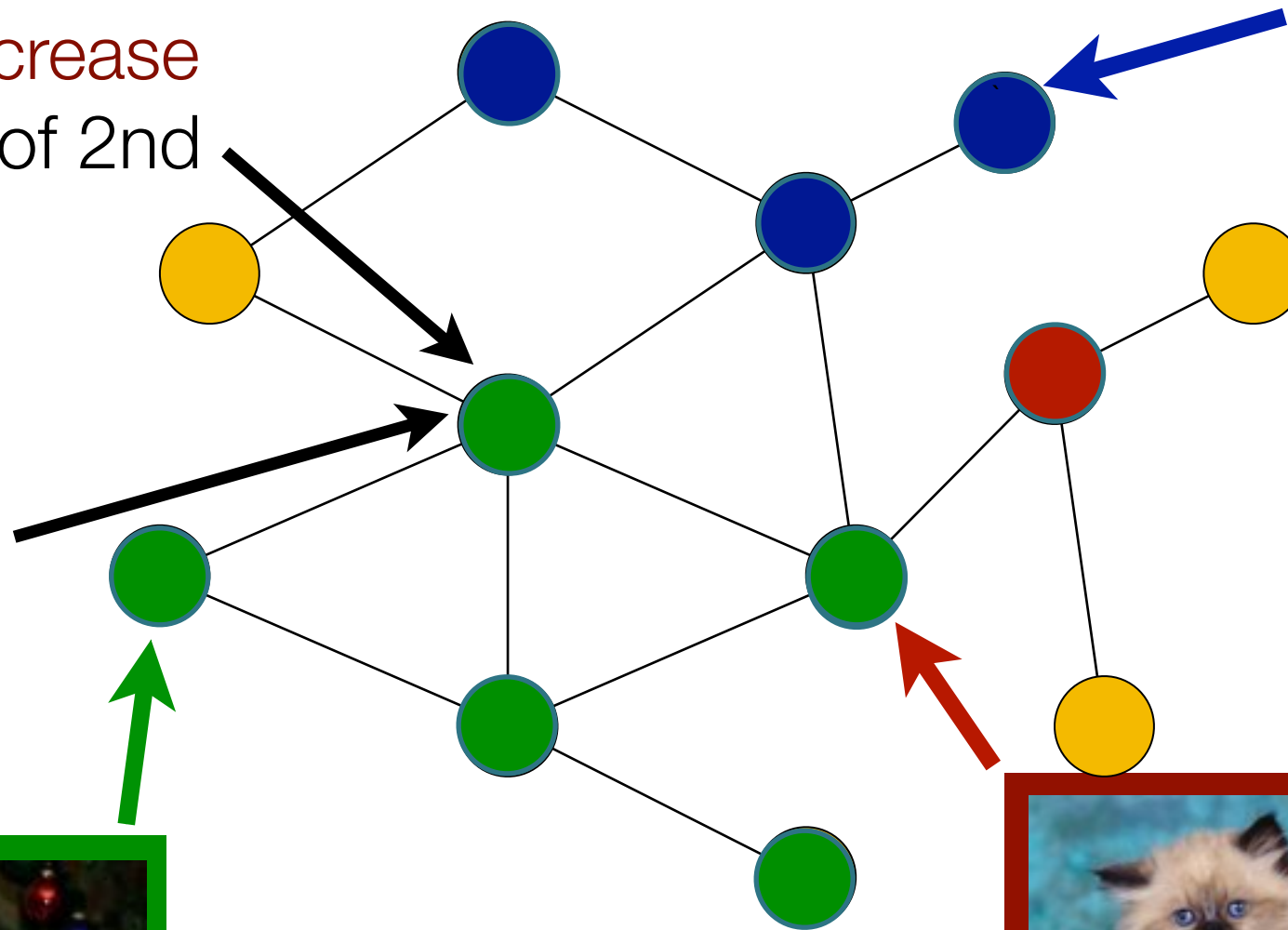


Information Diffusion

Does exposure to one contagion **increase/decrease** adoption probability of another contagion?

Did 1st cat video **decrease** adoption probability of 2nd cat video?

Did cat videos **increase** adoption probability of dog video?



Contagion Interactions

Why are interactions important?

Contagion Interactions

Why are interactions important?

- We can suppress the spread of other undesirable contagions:

Contagion Interactions

Why are interactions important?

- We can suppress the spread of other undesirable contagions:

“Fair Labor Association too easy on **Apple, Foxconn, study** says”

Contagion Interactions

Why are interactions important?

- We can suppress the spread of other undesirable contagions:

“Fair Labor Association too easy on **Apple, Foxconn, study** says”

suppressed by

Contagion Interactions

Why are interactions important?

- We can suppress the spread of other undesirable contagions:

“Fair Labor Association too easy on **Apple, Foxconn, study** says”

suppressed by

“New **iPhone 5** Sales Helps **Apple** Beat Android In The U.S.”

Contagion Interactions

Why are interactions important?

- We can suppress the spread of other undesirable contagions:

“Fair Labor Association too easy on **Apple, Foxconn, study** says”

suppressed by

“New **iPhone 5** Sales Helps **Apple** Beat Android In The U.S.”

- Contagions can promote other contagions

Contagion Interactions

Why are interactions important?

- We can suppress the spread of other undesirable contagions:

“Fair Labor Association too easy on **Apple, Foxconn, study** says”

suppressed by

“New **iPhone 5** Sales Helps **Apple** Beat Android In The U.S.”

- Contagions can promote other contagions
 - What **news stories** sequence would maximize our advertisement’s **click-through-rate**?

Contagion Interactions

Why are interactions important?

- We can suppress the spread of other undesirable contagions:

“Fair Labor Association too easy on **Apple, Foxconn, study** says”

suppressed by

“New **iPhone 5** Sales Helps **Apple** Beat Android In The U.S.”

- Contagions can promote other contagions
 - What **news stories** sequence would maximize our advertisement’s **click-through-rate**?
- In general, this leads to a more accurate diffusion model

Contagion Interactions

Why is modeling interactions difficult?

Contagion Interactions

Why is modeling interactions difficult?

- Many thousands of contagions diffusing at any time:

Contagion Interactions

Why is modeling interactions difficult?

- Many thousands of contagions diffusing at any time:
 - Observing interactions between all of them is **impossible.**

Contagion Interactions

Why is modeling interactions difficult?

- Many thousands of contagions diffusing at any time:
 - Observing interactions between all of them is **impossible.**
- The ordering of contagion exposures matters

Contagion Interactions

Why is modeling interactions difficult?

- Many thousands of contagions diffusing at any time:
 - Observing interactions between all of them is **impossible.**
- The ordering of contagion exposures matters
 - **Cat Video**, **News Article**, **Advertisement**

Contagion Interactions

Why is modeling interactions difficult?

- Many thousands of contagions diffusing at any time:
 - Observing interactions between all of them is **impossible.**
- The ordering of contagion exposures matters
 - **Cat Video**, **News Article**, **Advertisement**
vs. **News Article**, **Cat Video**, **Advertisement**

Contagion Interactions

Why is modeling interactions difficult?

- Many thousands of contagions diffusing at any time:
 - Observing interactions between all of them is **impossible.**
- The ordering of contagion exposures matters
 - **Cat Video, News Article, Advertisement**
vs. **News Article, Cat Video, Advertisement**
- The sampling of all possible interactions and exposure sequences is sparse.

Outline

1. Presentation of Model
2. Experiments of Model on real-world data
3. Insights gained from Model

Outline

1. Presentation of Model

2. Experiments of Model on real-world data

3. Insights gained from Model

Outline

1. Presentation of Model

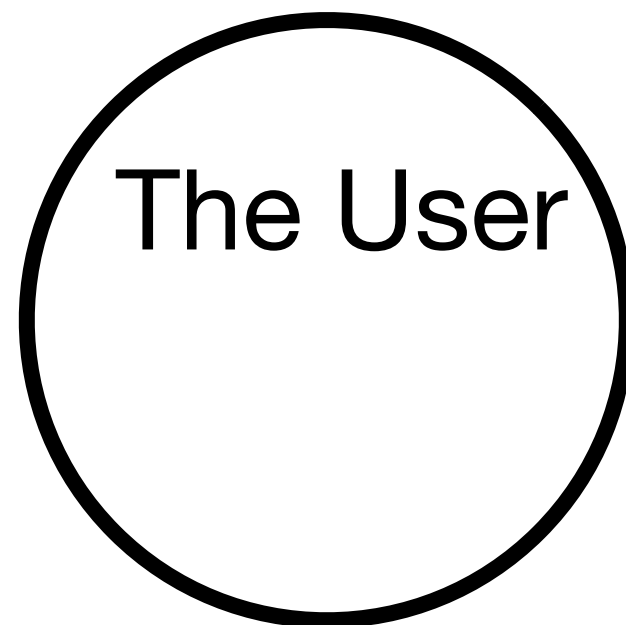
2. Experiments of Model on real-world data

3. Insights gained from Model

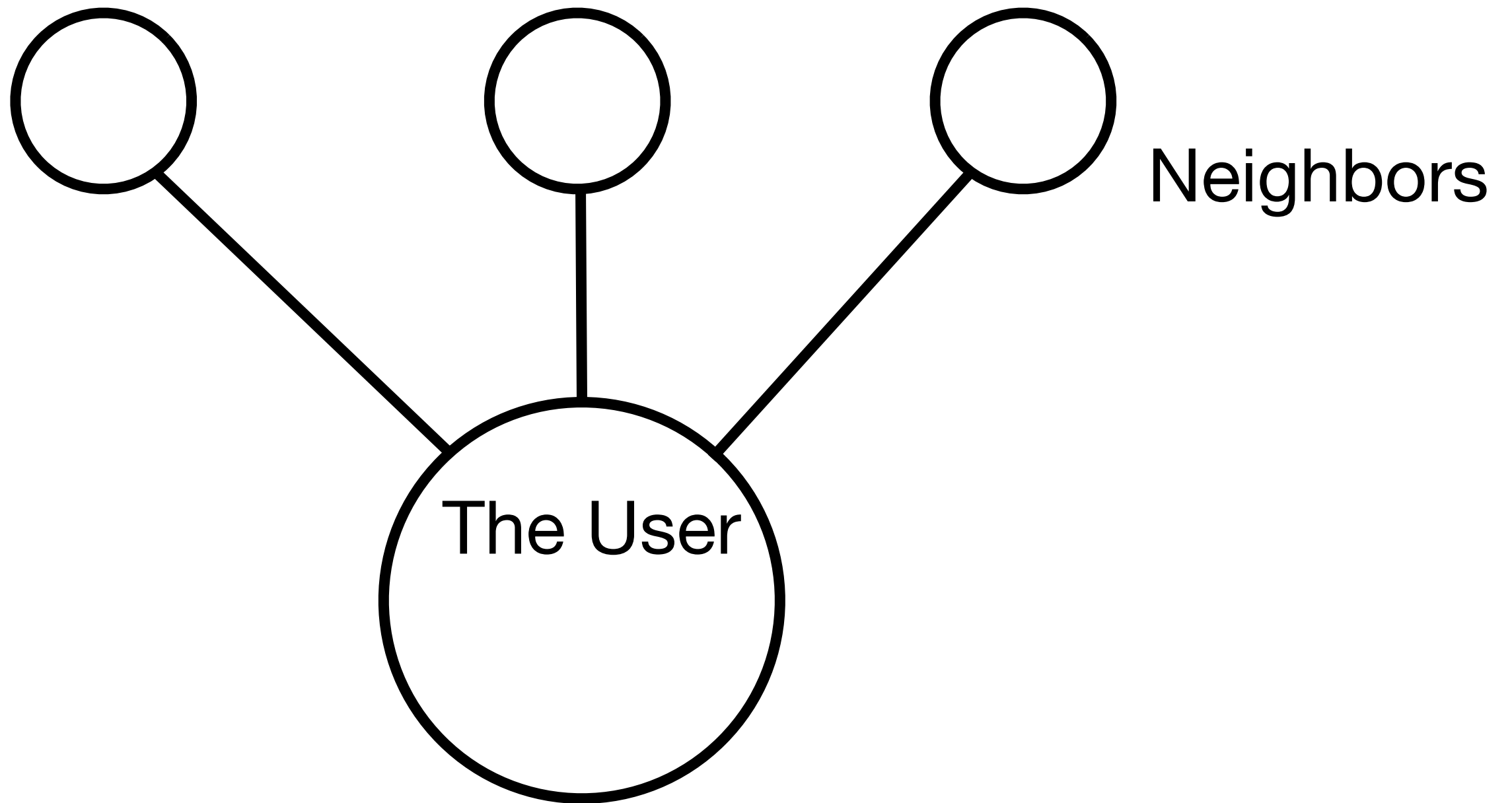
Outline

1. Presentation of Model
2. Experiments of Model on real-world data
- 3. Insights gained from Model**

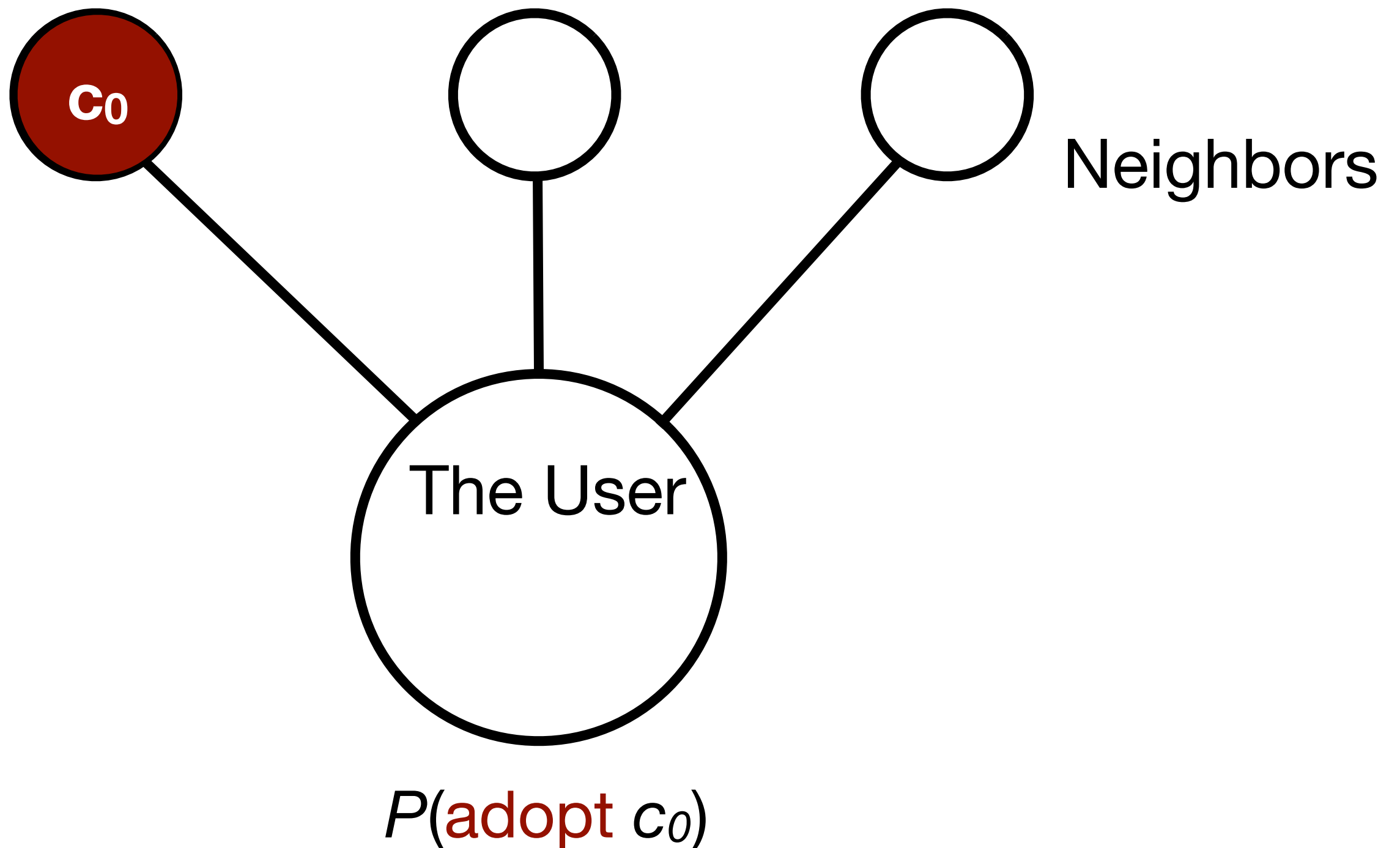
The Model



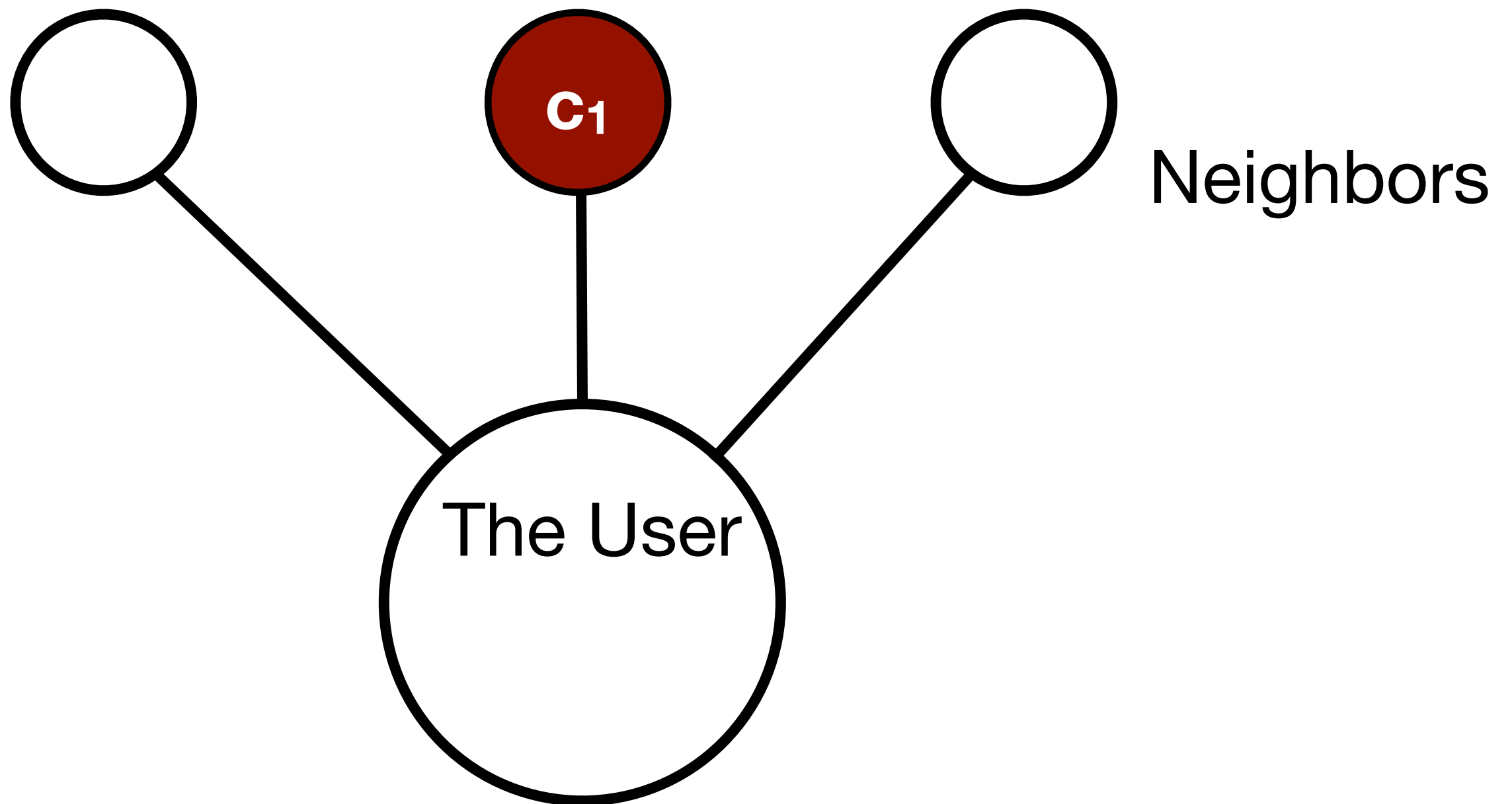
The Model



The Model

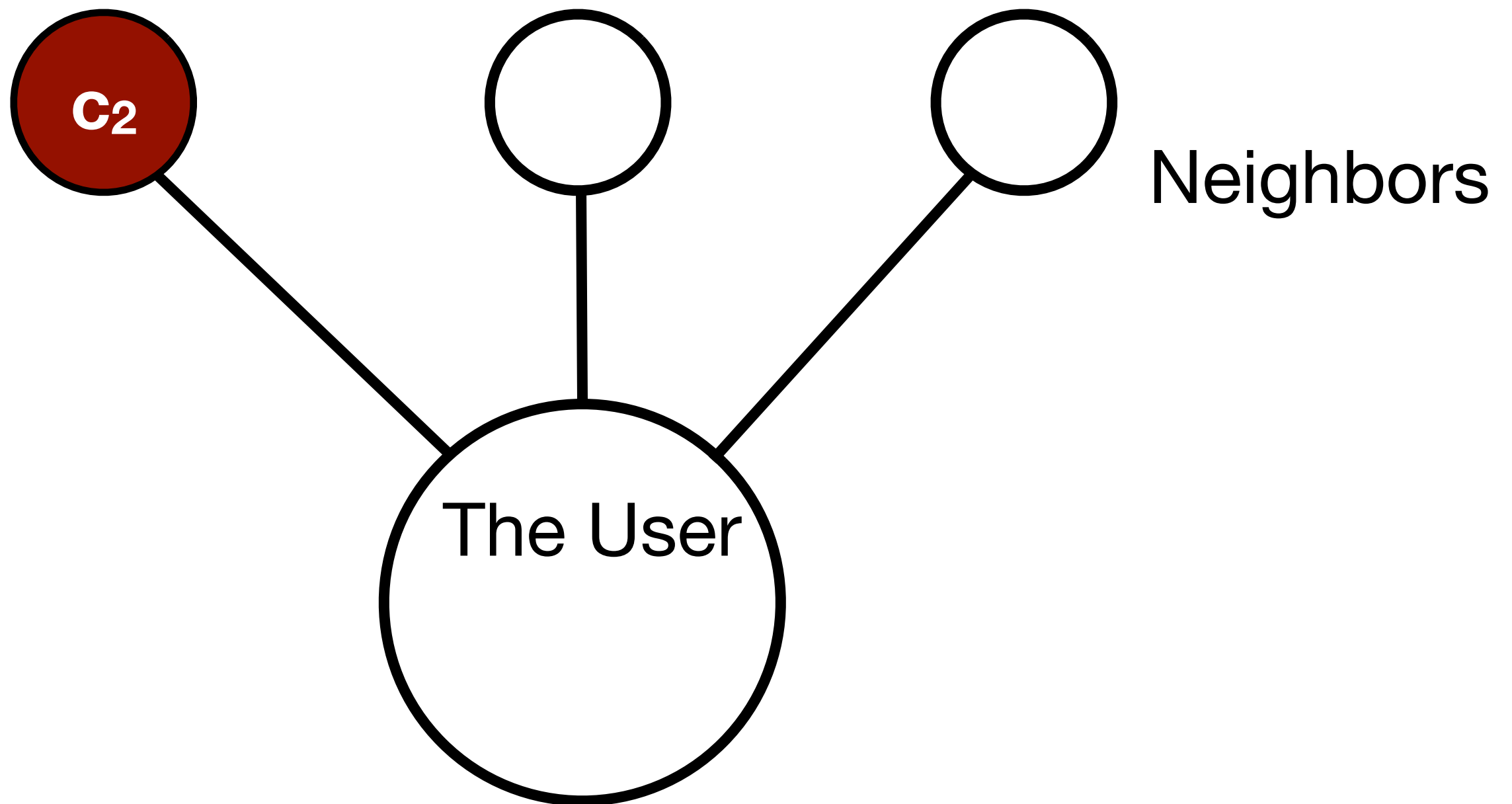


The Model



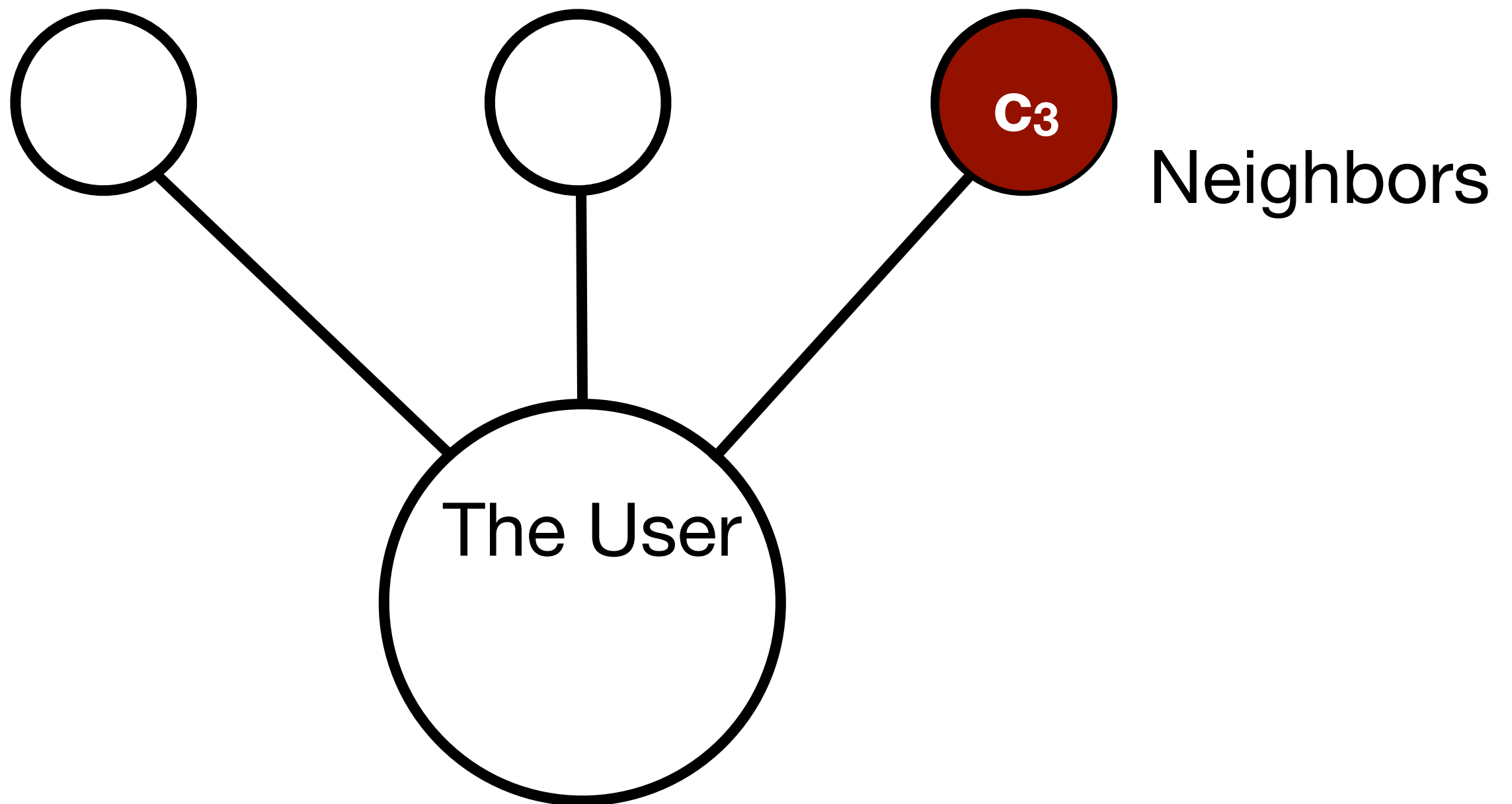
$$P(\text{adopt } c_1 \mid \text{exposed to } c_0)$$

The Model



$$P(\text{adopt } c_2 \mid \text{exposed to } c_1, c_0)$$

The Model



$$P(\text{adopt } c_3 \mid \text{exposed to } c_2, c_1, c_0)$$

The Model

The Model

- We assume K most recent exposures effect a user's adoption:

The Model

- We assume K most recent exposures effect a user's adoption:

$$P(\text{adopt } X=c_0 \mid \text{exposed } Y_1=c_1, Y_2=c_2, \dots, Y_K=c_K)$$

The Model

- We assume K most recent exposures effect a user's adoption:

$$P(\text{adopt } X=c_0 \mid \text{exposed } Y_1=c_1, Y_2=c_2, \dots, Y_K=c_K)$$

Contagion the user is viewing now.

Contagions the user previously viewed.

The Model

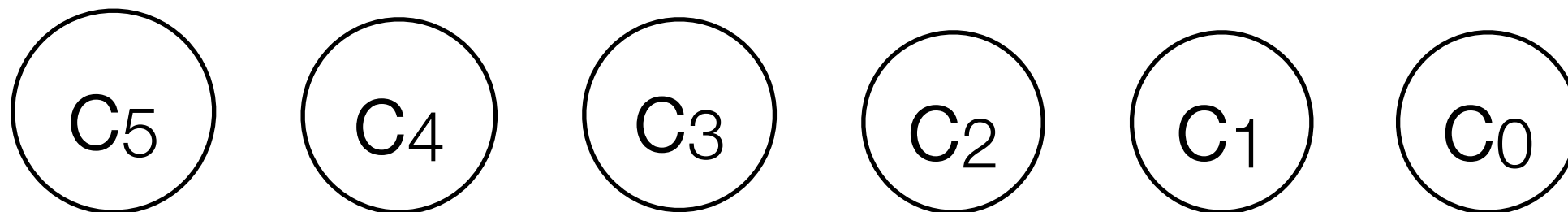
- We assume K most recent exposures effect a user's adoption:

$$P(\text{adopt } X=c_0 \mid \text{exposed } Y_1=c_1, Y_2=c_2, \dots, Y_K=c_k)$$

Contagion the user is viewing now.

Contagions the user previously viewed.

Contagions adopted by neighbors:



Time

The Model

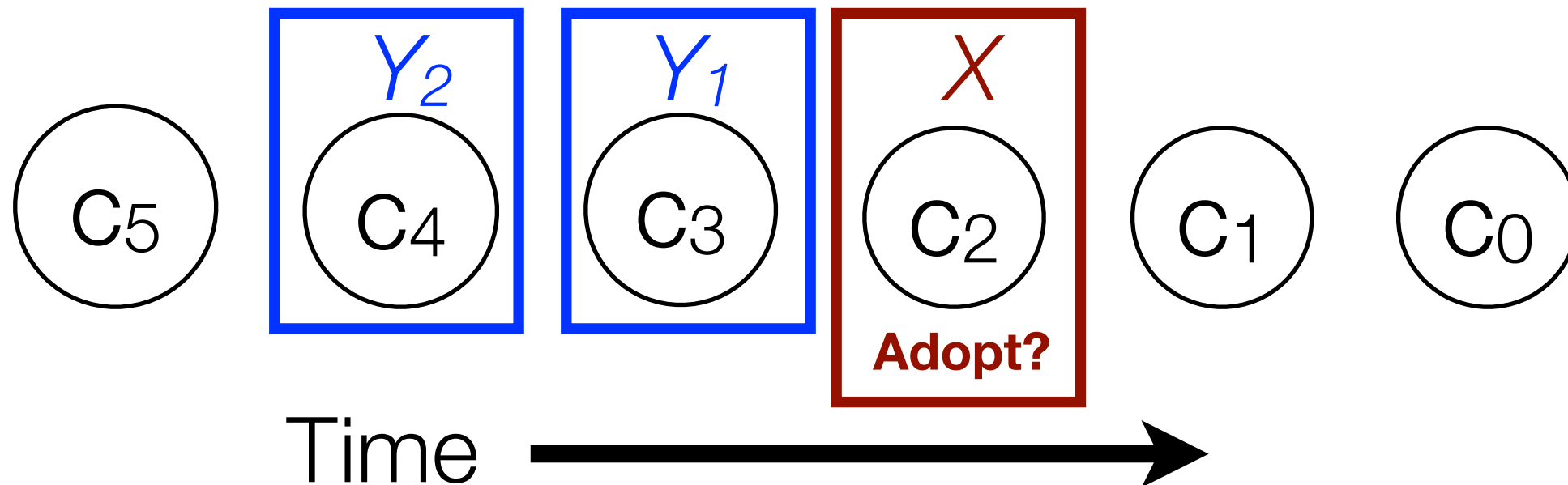
- We assume K most recent exposures effect a user's adoption:

$$P(\text{adopt } X=c_0 \mid \text{exposed } Y_1=c_1, Y_2=c_2, \dots, Y_K=c_k)$$

Contagion the user is viewing now.

Contagions the user previously viewed.

Contagions adopted by neighbors:



The Model

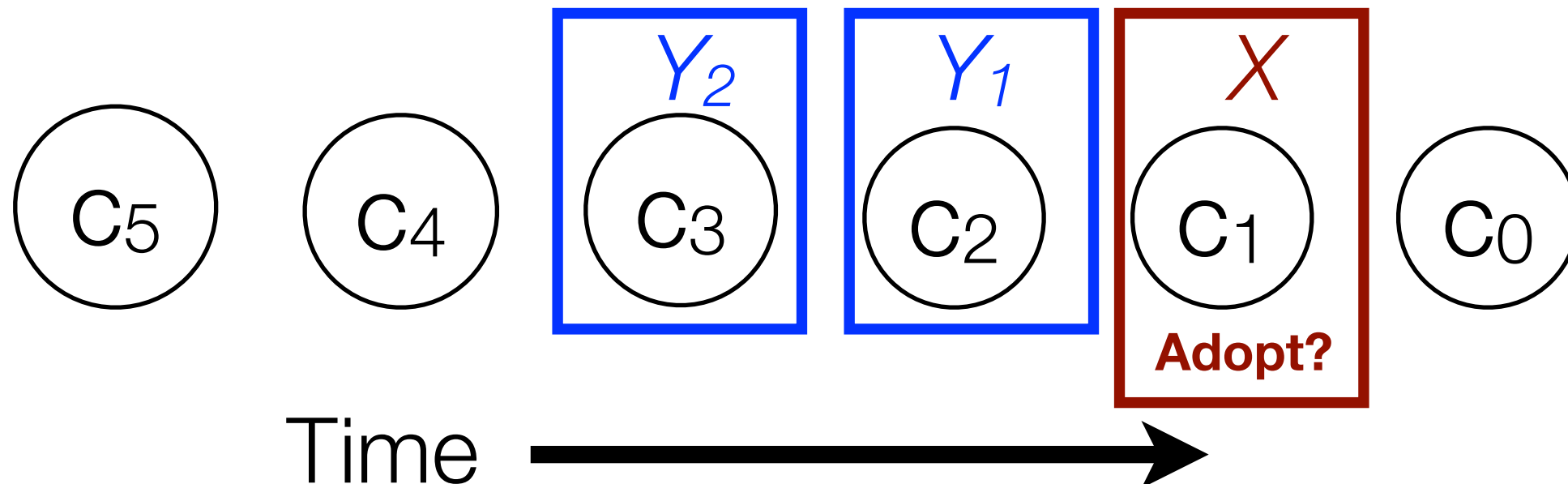
- We assume K most recent exposures effect a user's adoption:

$$P(\text{adopt } X=c_0 \mid \text{exposed } Y_1=c_1, Y_2=c_2, \dots, Y_K=c_k)$$

Contagion the user is viewing now.

Contagions the user previously viewed.

Contagions adopted by neighbors:



The Model

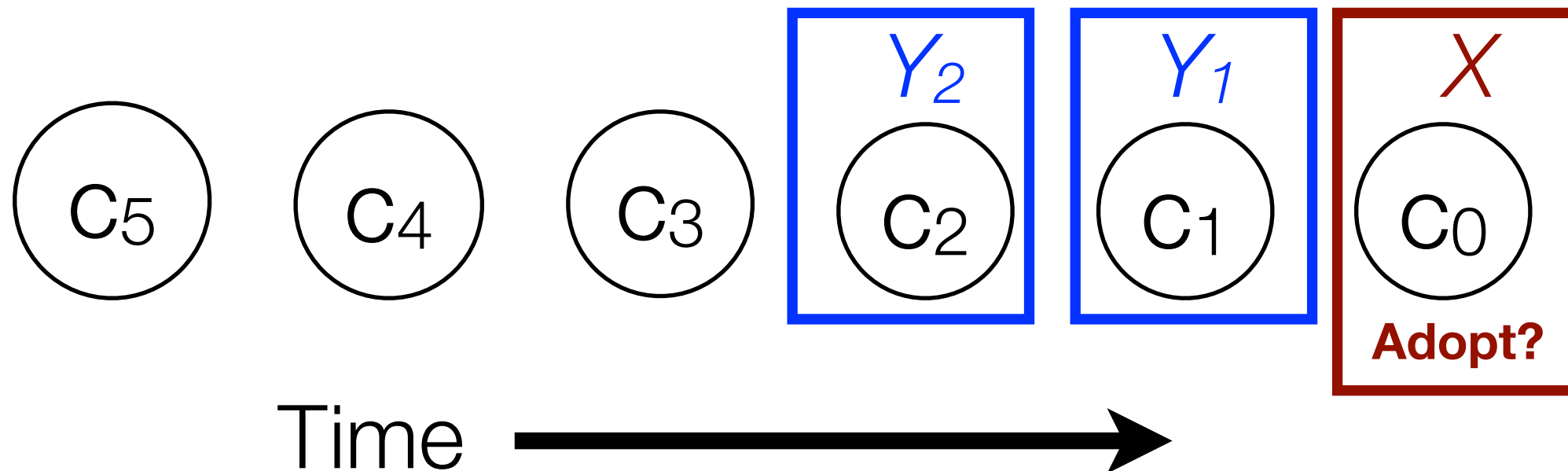
- We assume K most recent exposures effect a user's adoption:

$$P(\text{adopt } X=c_0 \mid \text{exposed } Y_1=c_1, Y_2=c_2, \dots, Y_K=c_k)$$

Contagion the user is viewing now.

Contagions the user previously viewed.

Contagions adopted by neighbors:



The Model - Simplifications

The Model - Simplifications

- Massive number of probabilities to measure:

The Model - Simplifications

- Massive number of probabilities to measure:

$$(\text{Num. Contagions})^K \approx 1.9 \times 10^{21}$$

The Model - Simplifications

- Massive number of probabilities to measure:

$$(\text{Num. Contagions})^K \approx 1.9 \times 10^{21}$$

- Simplification is necessary.

The Model - Simplifications

- Massive number of probabilities to measure:

$$(\text{Num. Contagions})^K \approx 1.9 \times 10^{21}$$

- Simplification is necessary.
- Assume Y_i is independent of Y_j . Then we apply Bayes

$$P\left(X \mid \{Y_k\}_{k=1}^K\right) = \frac{1}{P(X)^{K-1}} \prod_{k=1}^K P(X \mid Y_k)$$

The Model - Simplifications

- Massive number of probabilities to measure:

$$(\text{Num. Contagions})^K \approx 1.9 \times 10^{21}$$

- Simplification is necessary.
- Assume Y_i is independent of Y_j . Then we apply Bayes

$$P\left(X \mid \{Y_k\}_{k=1}^K\right) = \frac{1}{P(X)^{K-1}} \prod_{k=1}^K P(X \mid Y_k)$$

Easily measured empirically

Left to be modeled

The Model - Simplifications

The Model - Simplifications

- Still too many probabilities.

The Model - Simplifications

- Still too many probabilities.
- Assume:

The Model - Simplifications

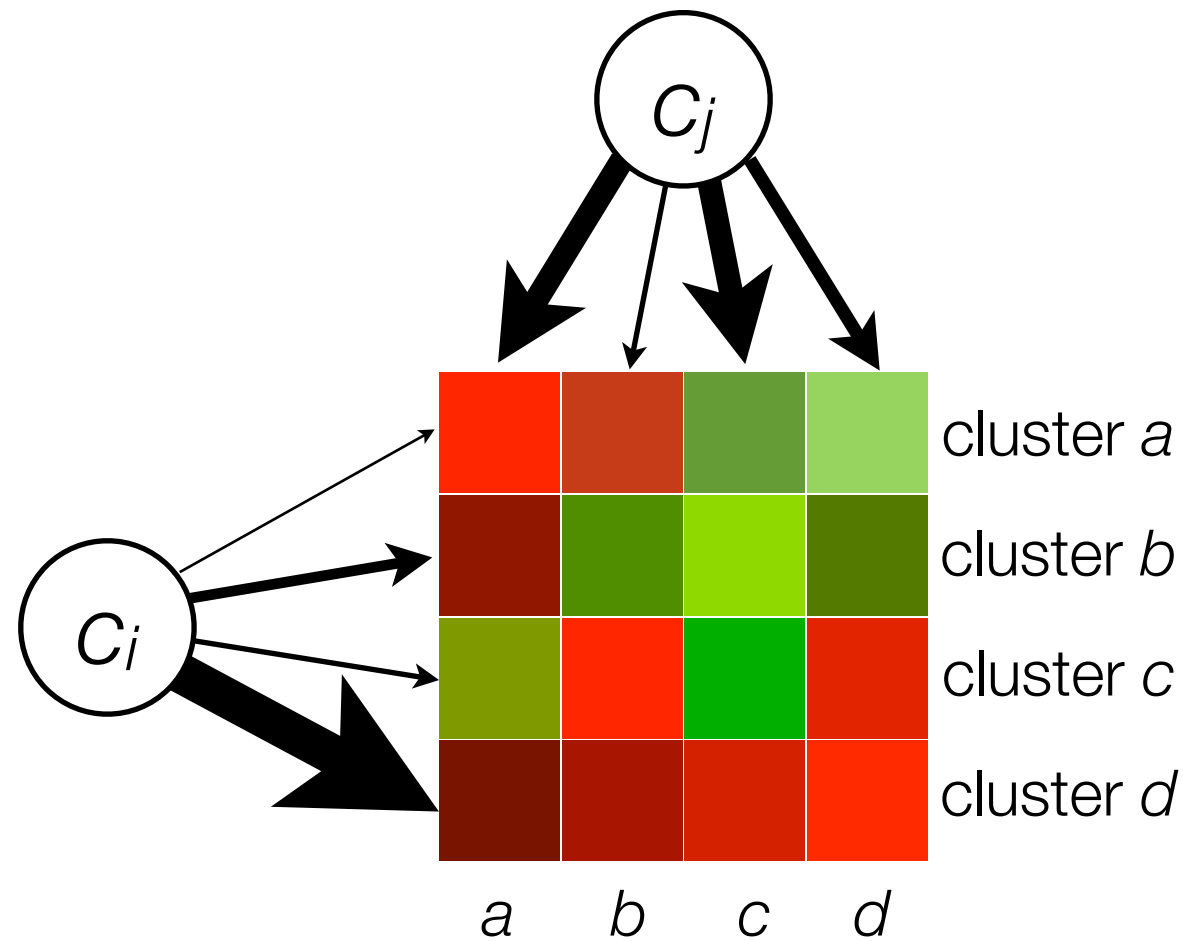
- Still too many probabilities.
- Assume: $P(X = c_i | Y_k = c_j) \approx P(X = c_i) + \textit{Interaction}(c_i, c_j)$

The Model - Simplifications

- Still too many probabilities.
- Assume: $P(X = c_i | Y_k = c_j) \approx P(X = c_i) + \textit{Interaction}(c_i, c_j)$
- Let contagion interactions come from latent interacting topics or **clusters**.

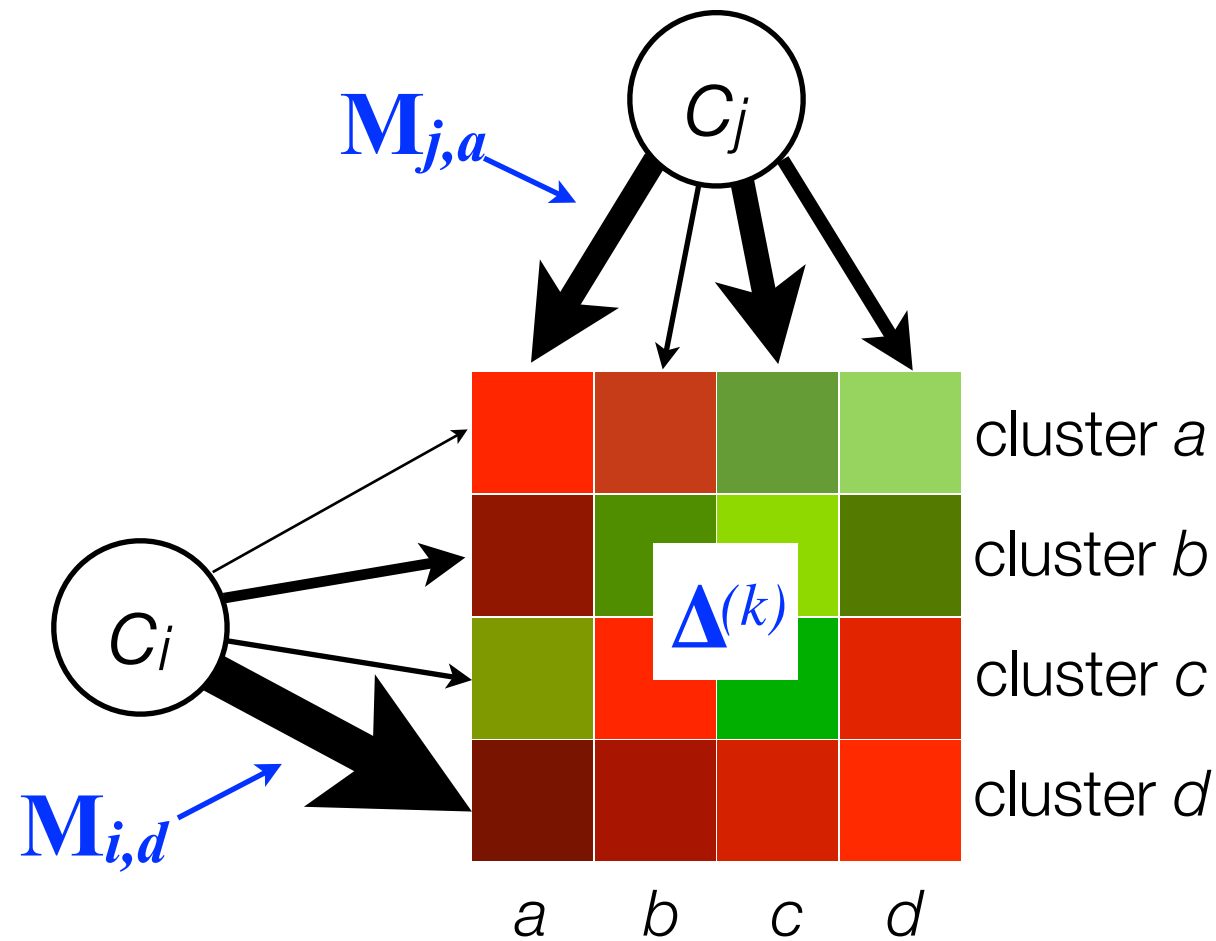
The Model - Simplifications

- Still too many probabilities.
- Assume: $P(X = c_i | Y_k = c_j) \approx P(X = c_i) + \text{Interaction}(c_i, c_j)$
- Let contagion interactions come from latent interacting topics or **clusters**.



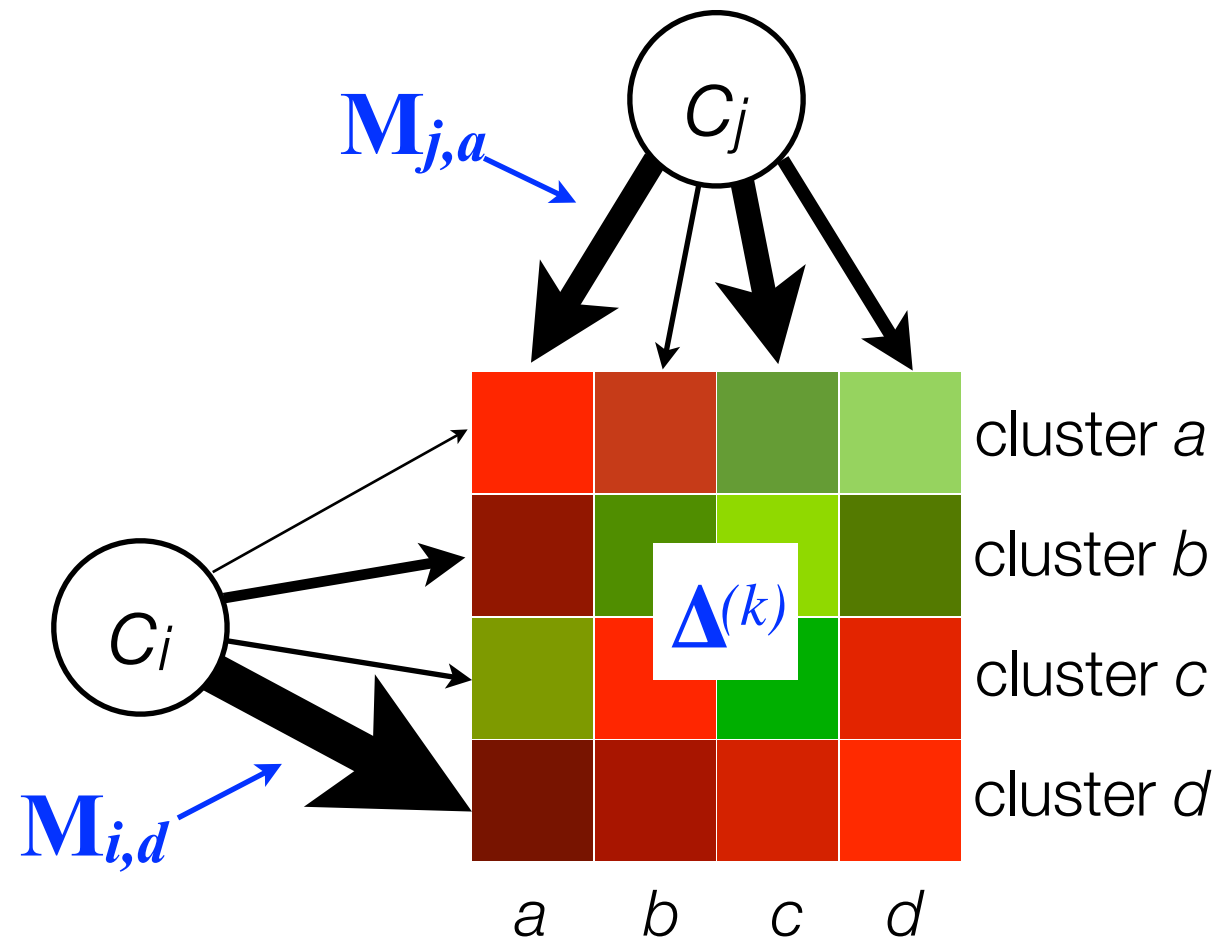
The Model - Simplifications

- Still too many probabilities.
- Assume: $P(X = c_i | Y_k = c_j) \approx P(X = c_i) + \text{Interaction}(c_i, c_j)$
- Let contagion interactions come from latent interacting topics or **clusters**.



The Model - Simplifications

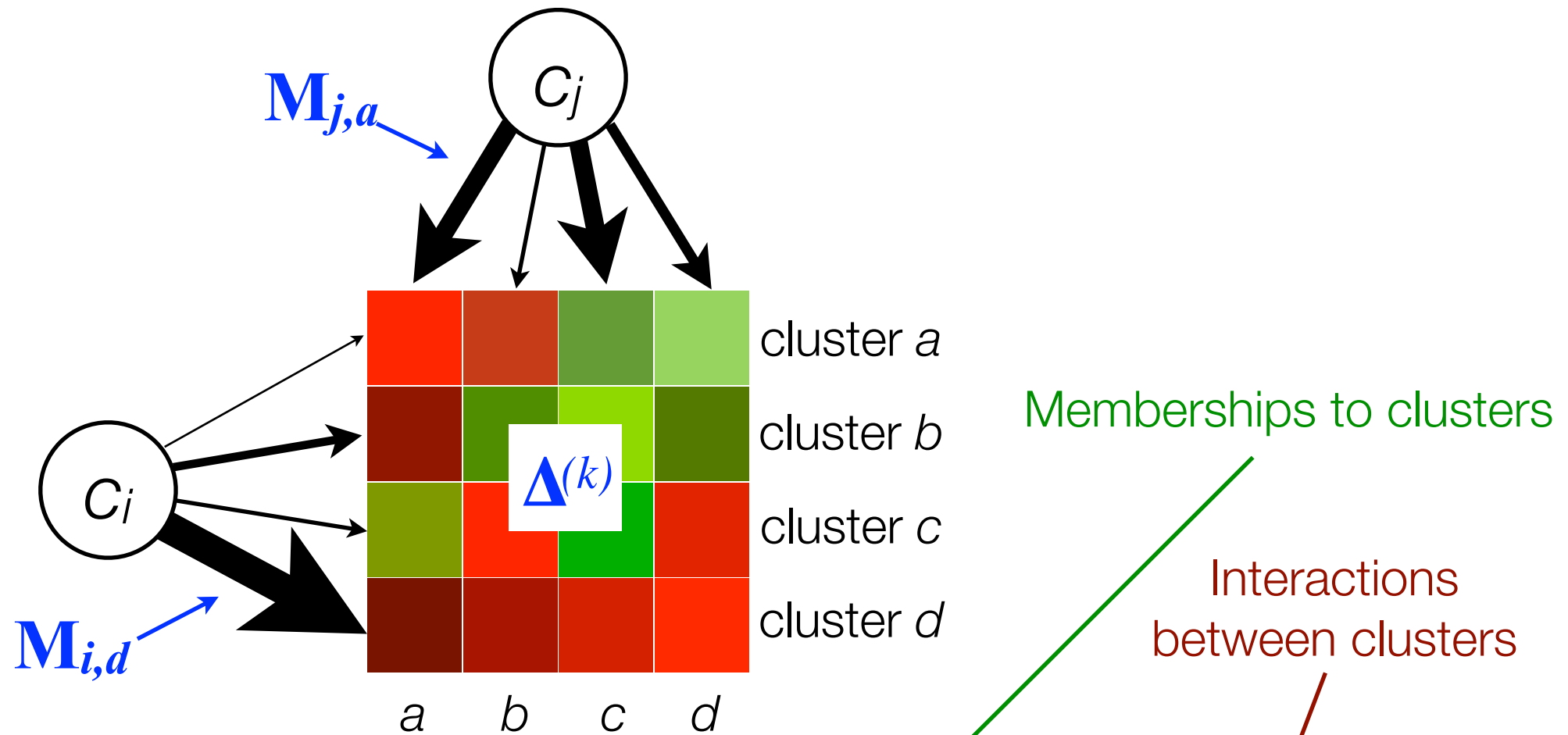
- Still too many probabilities.
- Assume: $P(X = c_i | Y_k = c_j) \approx P(X = c_i) + \text{Interaction}(c_i, c_j)$
- Let contagion interactions come from latent interacting topics or **clusters**.



$$P(X = c_i | Y_k = c_j) = P(X = c_i) + \sum_{a,b} \mathbf{M}_{i,a} \times \mathbf{M}_{ib} \times \Delta^{(k)}(a, b)$$

The Model - Simplifications

- Still too many probabilities.
- Assume: $P(X = c_i | Y_k = c_j) \approx P(X = c_i) + \text{Interaction}(c_i, c_j)$
- Let contagion interactions come from latent interacting topics or **clusters**.



$$P(X = c_i | Y_k = c_j) = P(X = c_i) + \sum_{a,b} \boxed{M_{i,a} \times M_{ib}} \times \boxed{\Delta^{(k)}(a, b)}$$

Fitting The Model

Fitting The Model

- **Given:**

Fitting The Model

- **Given:**
 - The Network - which users are connected

Fitting The Model

- **Given:**
 - The Network - which users are connected
 - Adoption times - which users adopted the contagions and when

Fitting The Model

- **Given:**
 - The Network - which users are connected
 - Adoption times - which users adopted the contagions and when
- **We measure directly:** $P(X = c_i)$ for all contagions:

Fitting The Model

- **Given:**
 - The Network - which users are connected
 - Adoption times - which users adopted the contagions and when
- **We measure directly:** $P(X = c_i)$ for all contagions:

$$P(X = c_i) = (\text{num. adoptions of } c_i) / (\text{num. exposures to } c_i)$$

Fitting The Model

- **Given:**
 - The Network - which users are connected
 - Adoption times - which users adopted the contagions and when

- **We measure directly:** $P(X = c_i)$ for all contagions:

$$P(X = c_i) = (\text{num. adoptions of } c_i) / (\text{num. exposures to } c_i)$$

- **We Infer:**

Fitting The Model

- **Given:**
 - The Network - which users are connected
 - Adoption times - which users adopted the contagions and when

- **We measure directly:** $P(X = c_i)$ for all contagions:

$$P(X = c_i) = (\text{num. adoptions of } c_i) / (\text{num. exposures to } c_i)$$

- **We Infer:**
 - The **cluster memberships** **M**.

Fitting The Model

- **Given:**
 - The Network - which users are connected
 - Adoption times - which users adopted the contagions and when

- **We measure directly:** $P(X = c_i)$ for all contagions:

$$P(X = c_i) = (\text{num. adoptions of } c_i) / (\text{num. exposures to } c_i)$$

- **We Infer:**
 - The **cluster memberships** \mathbf{M} .
 - The **cluster interactions** $\Delta^{(k)}$.

The Objective Function

The Objective Function

- We observe **exposure** sequences, and which exposures led to **adoption**.

The Objective Function

- We observe **exposure** sequences, and which exposures led to **adoption**.
- Define n_{ij}^k to be the number of times a user was exposed to c_j , then k exposures later to c_i .

The Objective Function

- We observe **exposure** sequences, and which exposures led to **adoption**.
- Define $n^{k_{ij}}$ to be the number of times a user was exposed to c_j , then k exposures later to c_i .
- Define $p^{k_{ij}}$ to be number of times this lead to adoption of c_i .

The Objective Function

- We observe **exposure** sequences, and which exposures led to **adoption**.
- Define $n^{k_{ij}}$ to be the number of times a user was exposed to c_j , then k exposures later to c_i .
- Define $p^{k_{ij}}$ to be number of times this lead to adoption of c_i .
- Then the log-likelihood is

$$\begin{aligned} \mathcal{L}(\mathbf{M}, \{\Delta\}_{k=1}^K) = & \\ & \sum_{i,j,k} p^{k_{ij}} \cdot \log \left[P(X = c_i) + \sum_{a,b} \mathbf{M}_{i,a} \cdot \Delta_{a,b}^{(k)} \cdot \mathbf{M}_{j,b} \right] \\ & + (n^{k_{ij}} - p^{k_{ij}}) \cdot \log \left[1 - P(X = c_j) - \sum_{a,b} \mathbf{M}_{i,a} \cdot \Delta_{a,b}^{(k)} \cdot \mathbf{M}_{j,b} \right] \end{aligned}$$

The Objective Function

- We observe **exposure** sequences, and which exposures led to **adoption**.
- Define $n^{k_{ij}}$ to be the number of times a user was exposed to c_j , then k exposures later to c_i .
- Define $p^{k_{ij}}$ to be number of times this lead to adoption of c_i .
- Then the log-likelihood is

$$\mathcal{L}(\mathbf{M}, \{\Delta\}_{k=1}^K) = \sum_{i,j,k} p^{k_{ij}} \cdot \log \left[P(X = c_i) + \sum_{a,b} \mathbf{M}_{i,a} \cdot \Delta_{a,b}^{(k)} \cdot \mathbf{M}_{j,b} \right] + (n^{k_{ij}} - p^{k_{ij}}) \cdot \log \left[1 - P(X = c_j) - \sum_{a,b} \mathbf{M}_{i,a} \cdot \Delta_{a,b}^{(k)} \cdot \mathbf{M}_{j,b} \right]$$

$P(X | Y_k)$ positively sampled

The Objective Function

- We observe **exposure** sequences, and which exposures led to **adoption**.
- Define $n^{k_{ij}}$ to be the number of times a user was exposed to c_j , then k exposures later to c_i .
- Define $p^{k_{ij}}$ to be number of times this lead to adoption of c_i .
- Then the log-likelihood is

$$\mathcal{L}(\mathbf{M}, \{\Delta\}_{k=1}^K) =$$

$$\sum_{i,j,k} p_{ij}^k \cdot \log \left[P(X = c_i) + \sum_{a,b} \mathbf{M}_{i,a} \cdot \Delta_{a,b}^{(k)} \cdot \mathbf{M}_{j,b} \right]$$

$P(X | Y_k)$ positively sampled

$$+ (n_{ij}^k - p_{ij}^k) \cdot \log \left[1 - P(X = c_j) - \sum_{a,b} \mathbf{M}_{i,a} \cdot \Delta_{a,b}^{(k)} \cdot \mathbf{M}_{j,b} \right]$$

$P(X | Y_k)$ negatively sampled

Optimizing the Objective Function

- We fit \mathbf{M} and $\Delta^{(k)}$ to the observed data using **stochastic gradient descent**:
- A **small subset** of the $p^{k_{ij}}$ and $n^{k_{ij}}$ values (terms in the objective function) are chosen randomly.
- The parameters are fit to this subset using gradient descent.
- After ~ 20 iterations, the $p^{k_{ij}}$ and $n^{k_{ij}}$ values are **resampled**.
- This continues until no improvement can be achieved.

The Real Dataset - Twitter

The Real Dataset - Twitter

- We iterated through every tweet sent on Twitter in January 2011, and we extracted tweeted URLs:

The Real Dataset - Twitter

- We iterated through every tweet sent on Twitter in January 2011, and we extracted tweeted URLs:
 - **18,186** high-volume URLs

The Real Dataset - Twitter

- We iterated through every tweet sent on Twitter in January 2011, and we extracted tweeted URLs:
 - **18,186** high-volume URLs
 - **1,087,033** users with **103,112,438** follower edges

The Real Dataset - Twitter

- We iterated through every tweet sent on Twitter in January 2011, and we extracted tweeted URLs:
 - **18,186** high-volume URLs
 - **1,087,033** users with **103,112,438** follower edges
 - URLs tweeted **2,664,207** times

The Real Dataset - Twitter

- We iterated through every tweet sent on Twitter in January 2011, and we extracted tweeted URLs:
 - **18,186** high-volume URLs
 - **1,087,033** users with **103,112,438** follower edges
 - URLs tweeted **2,664,207** times
 - **810,884,361** exposures to URLs.

The Real Dataset - Twitter

- We iterated through every tweet sent on Twitter in January 2011, and we extracted tweeted URLs:
 - **18,186** high-volume URLs
 - **1,087,033** users with **103,112,438** follower edges
 - URLs tweeted **2,664,207** times
 - **810,884,361** exposures to URLs.
- Each URL is a different contagion

The Real Dataset - Twitter

- We iterated through every tweet sent on Twitter in January 2011, and we extracted tweeted URLs:
 - **18,186** high-volume URLs
 - **1,087,033** users with **103,112,438** follower edges
 - URLs tweeted **2,664,207** times
 - **810,884,361** exposures to URLs.
- Each URL is a different contagion
- A user adopts a URL contagion by tweeting it.

Experiments

Experiments

- The model is **trained** on 90% of all observed exposures.

Experiments

- The model is **trained** on 90% of all observed exposures.
- The model predicts which exposures in **test set** will cause adoptions.

Experiments

- The model is **trained** on 90% of all observed exposures.
- The model predicts which exposures in **test set** will cause adoptions.
- Multiple measures of performance used

Experiments

- The model is **trained** on 90% of all observed exposures.
- The model predicts which exposures in **test set** will cause adoptions.
- Multiple measures of performance used
 - **Log-Likelihood of test set**

Experiments

- The model is **trained** on 90% of all observed exposures.
- The model predicts which exposures in **test set** will cause adoptions.
- Multiple measures of performance used
 - **Log-Likelihood of test set**
 - **maximum F_1 score**

Experiments

- The model is **trained** on 90% of all observed exposures.
- The model predicts which exposures in **test set** will cause adoptions.
- Multiple measures of performance used
 - **Log-Likelihood of test set**
 - **maximum F_1 score**
 - **Area under precision/recall curve**

Experiments - Baselines

Experiments - Baselines

- The model's performance was compared to several baseline models.

Experiments - Baselines

- The model's performance was compared to several baseline models.
 - **Prior Adoption Probability:** $P(X | Y_k) = P(X)$ for all Y_k

Experiments - Baselines

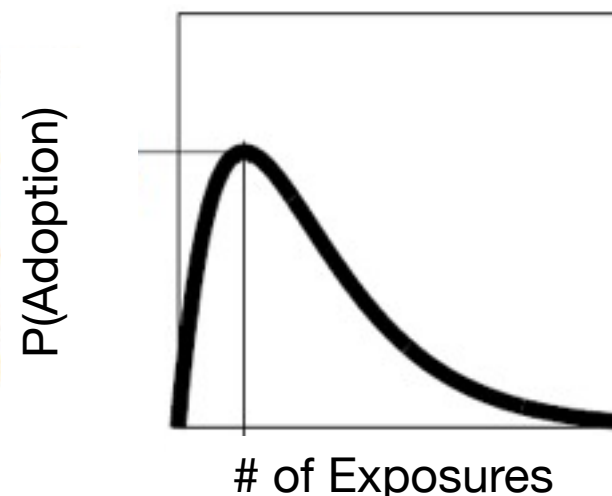
- The model's performance was compared to several baseline models.
 - **Prior Adoption Probability:** $P(X | Y_k) = P(X)$ for all Y_k
 - Independent Cascade Model [Goldenberg et al. 2001, Kempe et al 2003]

Experiments - Baselines

- The model's performance was compared to several baseline models.
 - **Prior Adoption Probability:** $P(X | Y_k) = P(X)$ for all Y_k
 - Independent Cascade Model [Goldenberg et al. 2001, Kempe et al 2003]
 - **Prior+User Bias:** $P_u(X) = P(X) + d_u$ for each user u .

Experiments - Baselines

- The model's performance was compared to several baseline models.
 - **Prior Adoption Probability:** $P(X | Y_k) = P(X)$ for all Y_k
 - Independent Cascade Model [Goldenberg et al. 2001, Kempe et al 2003]
 - **Prior+User Bias:** $P_u(X) = P(X) + d_u$ for each user u .
 - **Exposure Curve** [Romero et al 2011, Myers et al 2012]: Adoption probability of X as a function of exposure count.



Experiments - Results

	Log-Like.	Area under PR	max F₁
Prior Adoption Probability	-335,550.39	0.0157	0.0157
Prior+User Bias	-338,821.54	0.0123	0.0112
Exposure Curve	-338,367.86	0.0250	0.0181
Our Model	-299,884.86	0.1238	0.0465

Experiments - Results

	Log-Like.	Area under PR	max F₁
Prior Adoption Probability	-335,550.39	0.0157	0.0157
Prior+User Bias	-338,821.54	0.0123	0.0112
Exposure Curve	-338,367.86	0.0250	0.0181
Our Model	-299,884.86	0.1238	0.0465

11% Improvement 400% Improvement! 168% Improvement!

Experiments - Results

	Log-Like.	Area under PR	max F₁
Prior Adoption Probability	-335,550.39	0.0157	0.0157
Prior+User Bias	-338,821.54	0.0123	0.0112
Exposure Curve	-338,367.86	0.0250	0.0181
Our Model	-299,884.86	0.1238	0.0465

11% Improvement 400% Improvement! 168% Improvement!

Including a **user bias** parameter offered no improvement in performance.

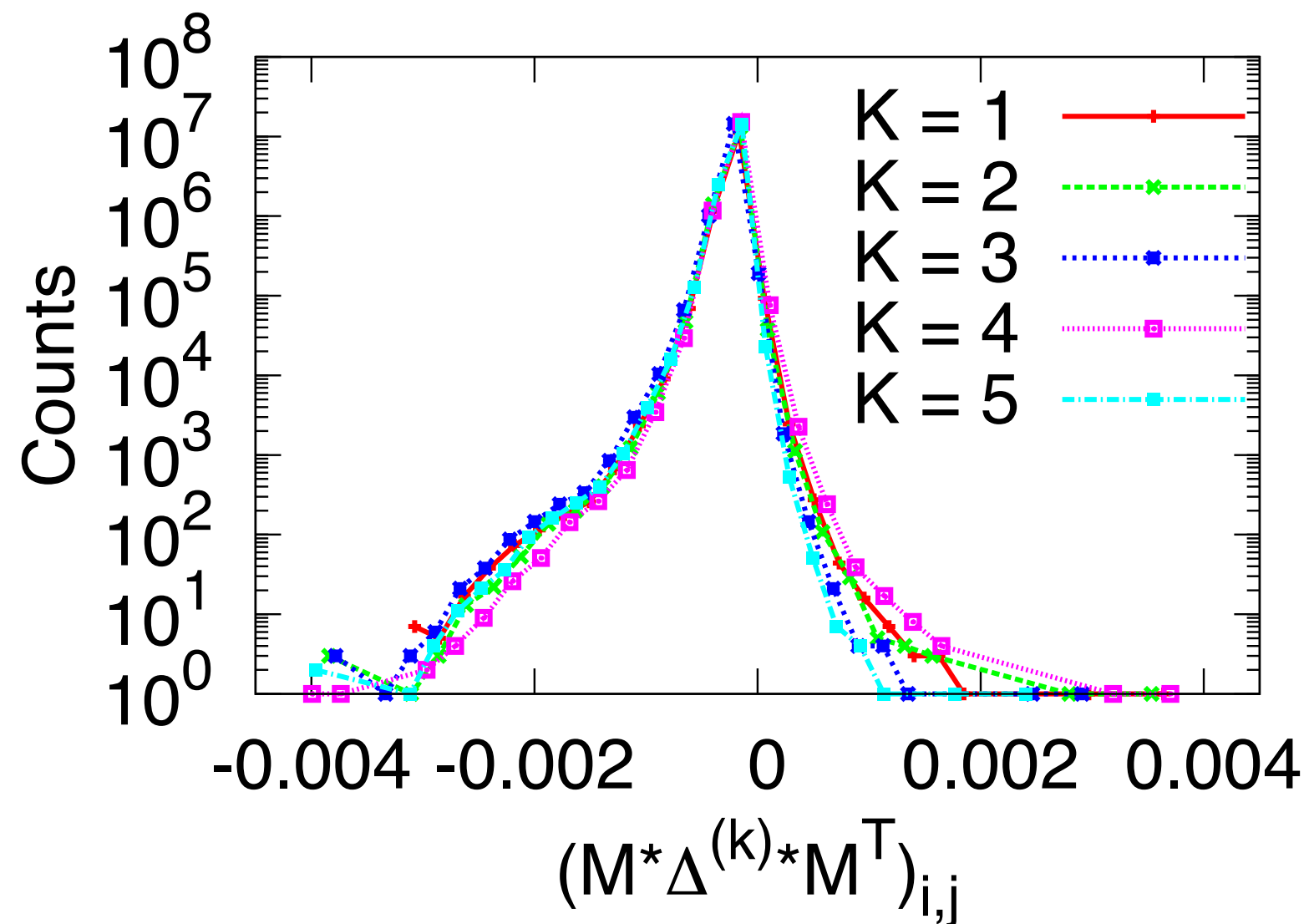
Insights from the Model

Insights from the Model

- Most interactions are slightly negative (suppressive):

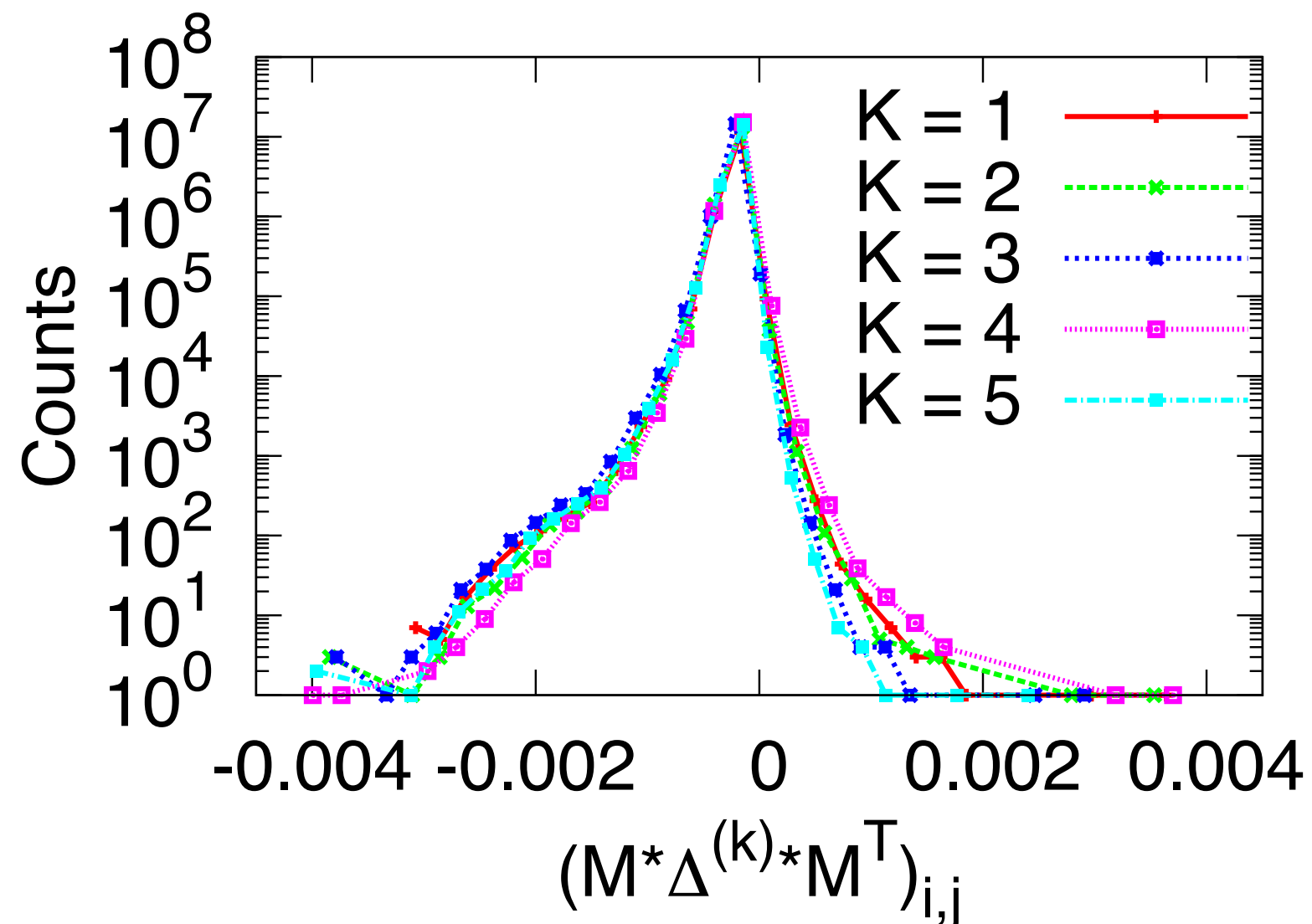
Insights from the Model

- Most interactions are slightly negative (suppressive):



Insights from the Model

- Most interactions are slightly negative (suppressive):



- In all, interactions between other contagions change adoption probability by **71% on average!**

Insights from the Model

Insights from the Model

- Contagions with **higher prior adoption probabilities** interact strongly with lower adoption probability contagions

Insights from the Model

- Contagions with **higher prior adoption probabilities** interact strongly with lower adoption probability contagions

$$P(X = c_i | Y_k = c_j)$$

Insights from the Model

- Contagions with **higher prior adoption probabilities** interact strongly with lower adoption probability contagions

$$P(X = \boxed{c_i} \mid Y_k = \boxed{c_j})$$

Lower prior adoption prob. → $\boxed{c_i}$ $\boxed{c_j}$ ← *Higher prior adoption prob.*

Insights from the Model

- Contagions with **higher prior adoption probabilities** interact strongly with lower adoption probability contagions

$$P(X = \boxed{c_i} \mid Y_k = \boxed{c_j})$$

Lower prior adoption prob. → $\boxed{c_i}$ $\boxed{c_j}$ ← *Higher prior adoption prob.*

- If highly related in subject matter, the interaction is positive.

Insights from the Model

- Contagions with **higher prior adoption probabilities** interact strongly with lower adoption probability contagions

$$P(X = \boxed{c_i} \mid Y_k = \boxed{c_j})$$

Lower prior adoption prob. → $\boxed{c_i}$ $\boxed{c_j}$ ← *Higher prior adoption prob.*

- If highly related in subject matter, the interaction is positive.
- If they are unrelated, the interaction is negative.

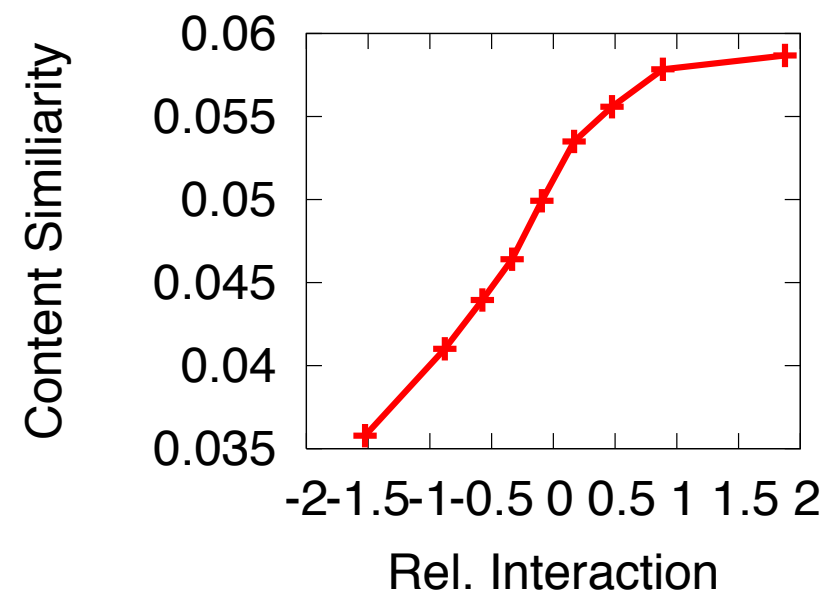
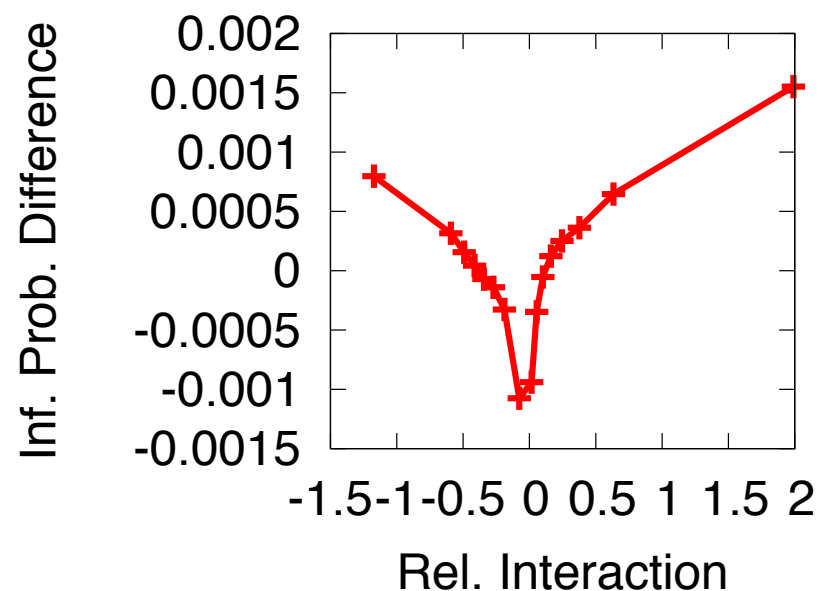
Insights from the Model

- Contagions with **higher prior adoption probabilities** interact strongly with lower adoption probability contagions

$$P(X=c_i | Y_k=c_j)$$

Lower prior adoption prob. → c_i c_j ← *Higher prior adoption prob.*

- If highly related in subject matter, the interaction is positive.
- If they are unrelated, the interaction is negative.



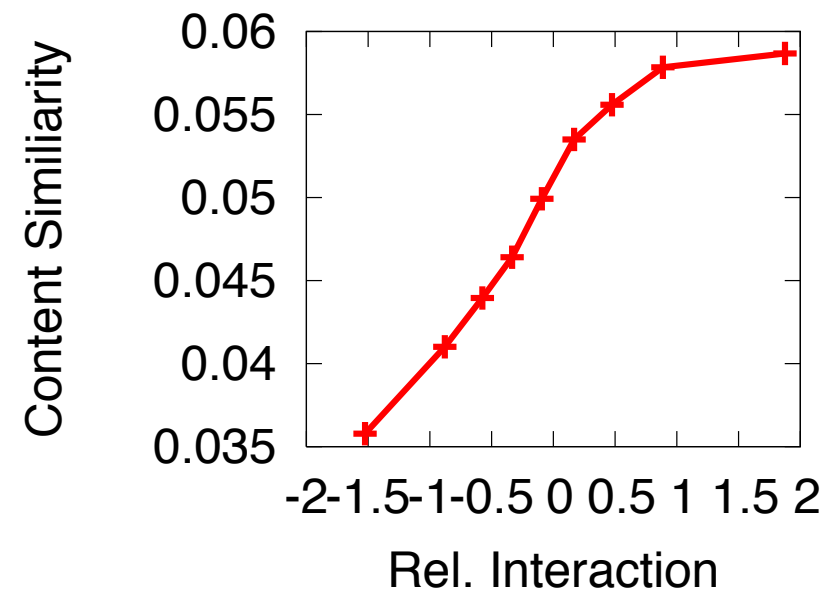
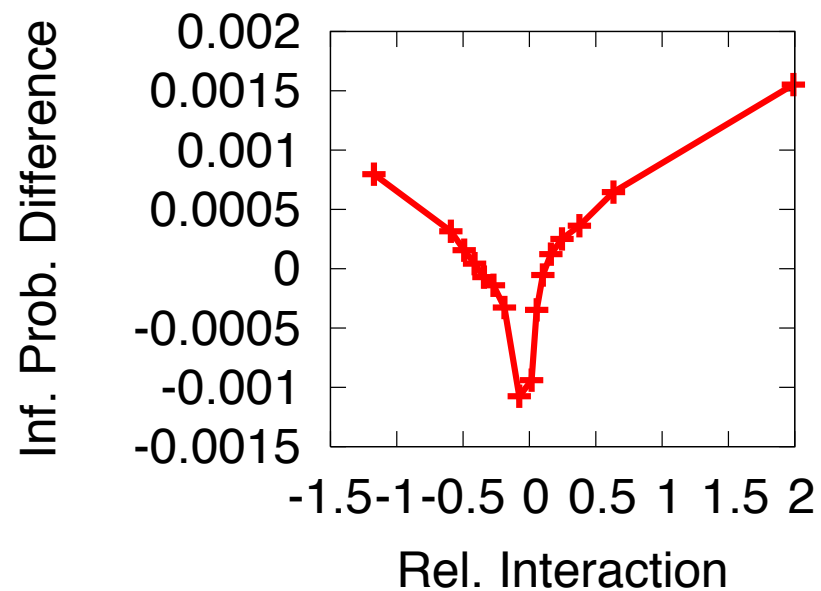
Insights from the Model

- Contagions with **higher prior adoption probabilities** interact strongly with lower adoption probability contagions

$$P(X=c_i | Y_k=c_j)$$

Lower prior adoption prob. → c_i c_j ← *Higher prior adoption prob.*

- If highly related in subject matter, the interaction is positive.
- If they are unrelated, the interaction is negative.



- This is evidence of an underlying process of interactions...

The Interaction Process - An example

“Paint continues to dry without incident.”

Golf.



The Interaction Process - An example

“Paint continues to dry without incident.”

Boring...

Golf.

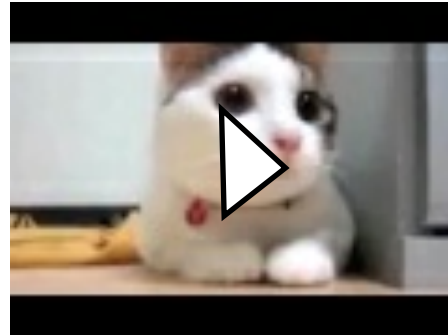


The Interaction Process - An example

“Paint continues to dry without incident.”

Golf.

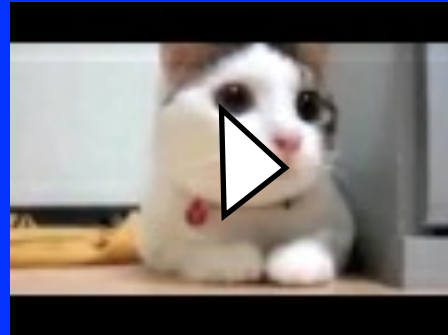
[yawn]



The Interaction Process - An example

“Paint continues to dry without incident.”

Golf.



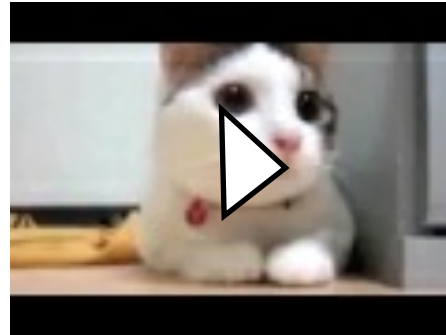
I love cat videos!



The Interaction Process - An example

“Paint continues to dry without incident.”

Golf.



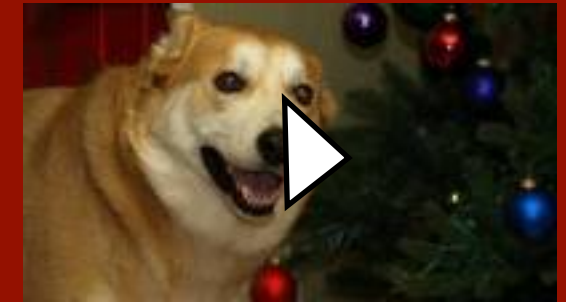
I love cat videos!



The Interaction Process - An example

“Paint continues to dry without incident.”

Golf.



I love cat videos!

The Interaction Process - An example

“Paint continues to dry without incident.”

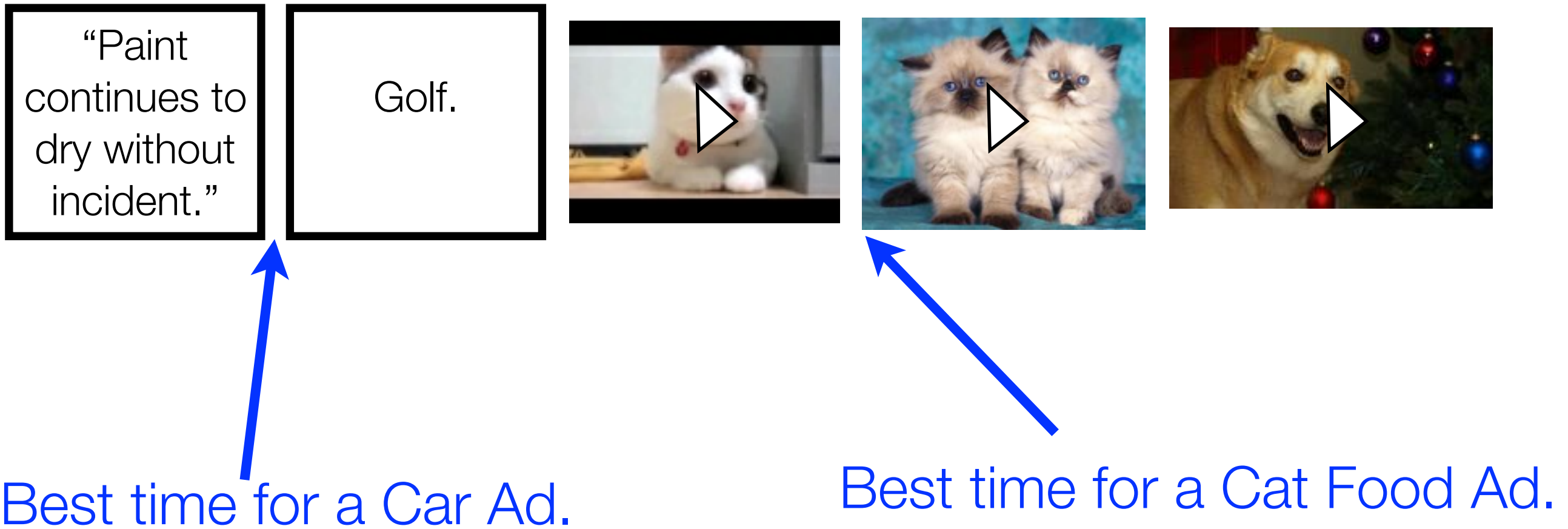
Golf.



Best time for a Cat Food Ad.



The Interaction Process - An example



Conclusion

Conclusion

- We presented a novel information diffusion model to account for interactions between diffusing contagions

Conclusion

- We presented a novel information diffusion model to account for interactions between diffusing contagions
- We developed a **scalable** algorithm to fit the model to observable data.

Conclusion

- We presented a novel information diffusion model to account for interactions between diffusing contagions
- We developed a **scalable** algorithm to fit the model to observable data.
- The model **outperforms** several baselines.

Conclusion

- We presented a novel information diffusion model to account for interactions between diffusing contagions
- We developed a **scalable** algorithm to fit the model to observable data.
- The model **outperforms** several baselines.
- Our model provides **insight** into the process of interactions between spreading contagions.