

# Detecting Network Effects

## Randomizing Over Randomized Experiments

Martin Saveski

(@msaveski)

MIT

# Detecting Network Effects

## Randomizing Over Randomized Experiments



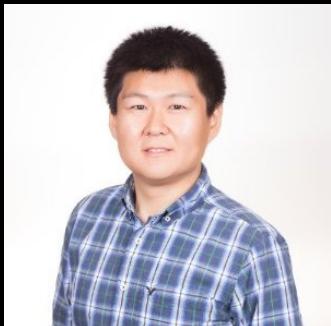
Martin Saveski  
MIT



Jean Pouget-Abadie  
Harvard



Guillaume Saint-Jacques  
MIT



Weitao Duan  
LinkedIn



Souvik Ghosh  
LinkedIn



Ya Xu  
LinkedIn

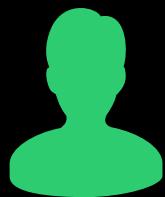


Edo Airoldi  
Harvard

**Treatment**

$$Z_i = 1$$

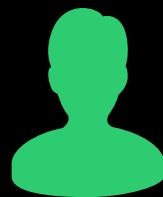
New Feed  
Ranking Algorithm



**Treatment**

$$Z_i = 1$$

New Feed  
Ranking Algorithm



**Control**

$$Z_j = 0$$

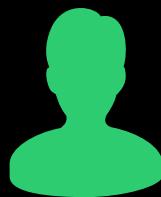
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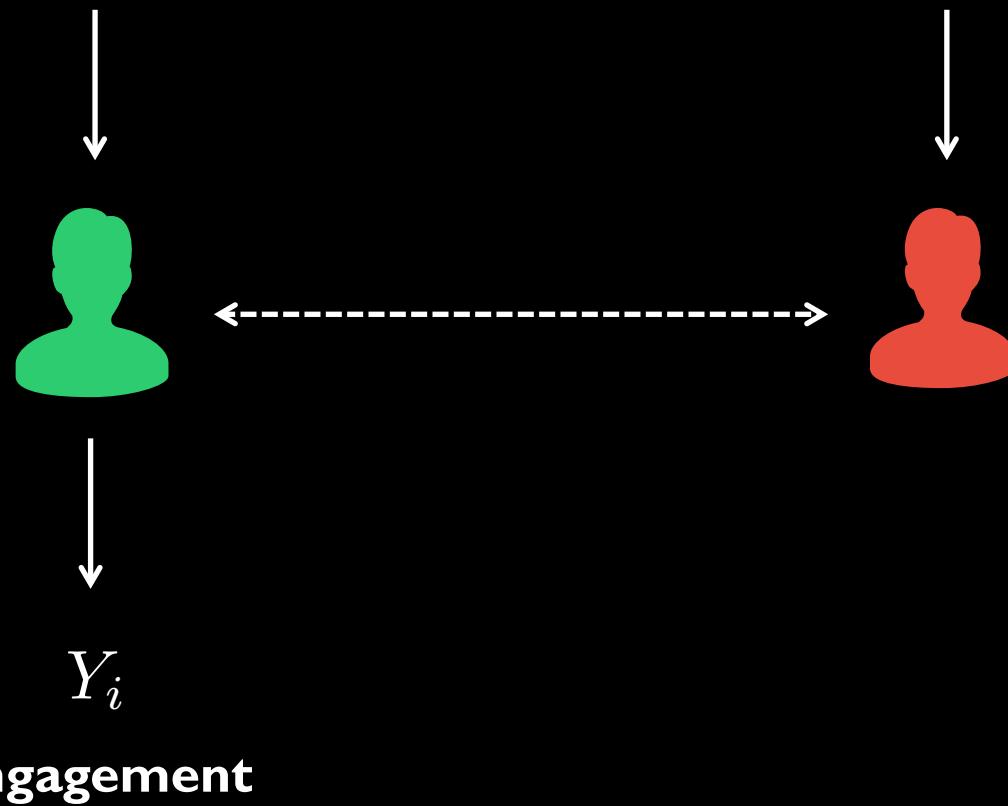
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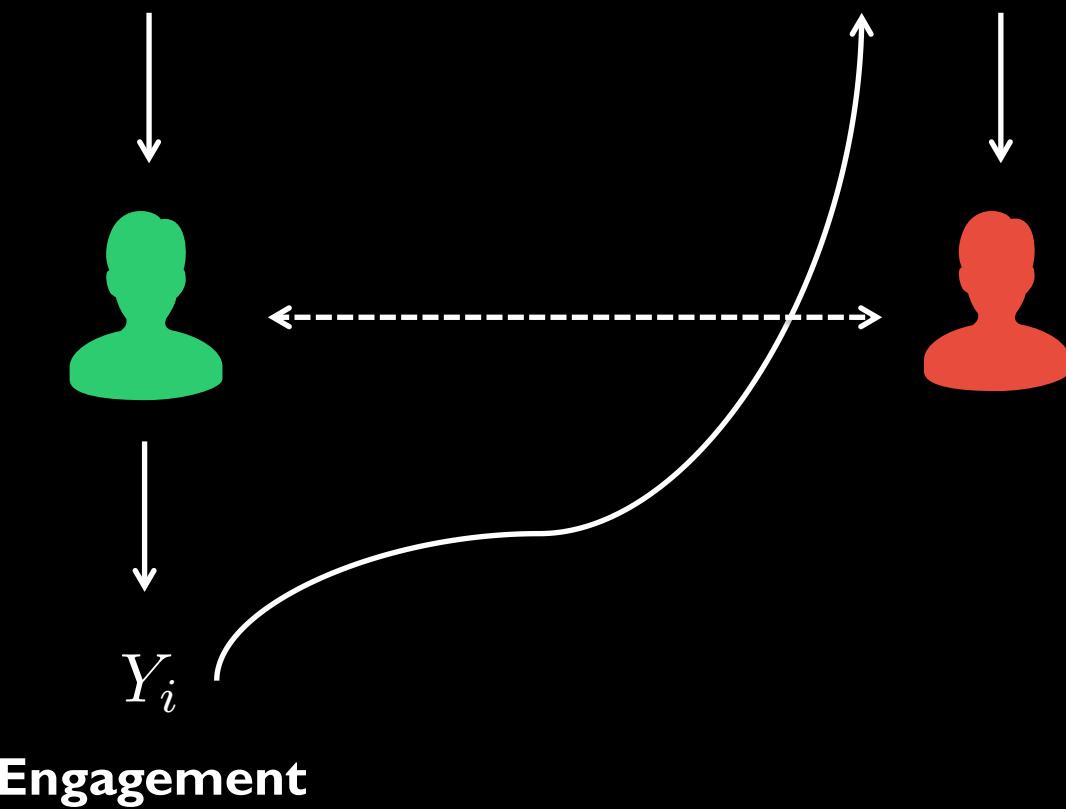
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**Treatment**

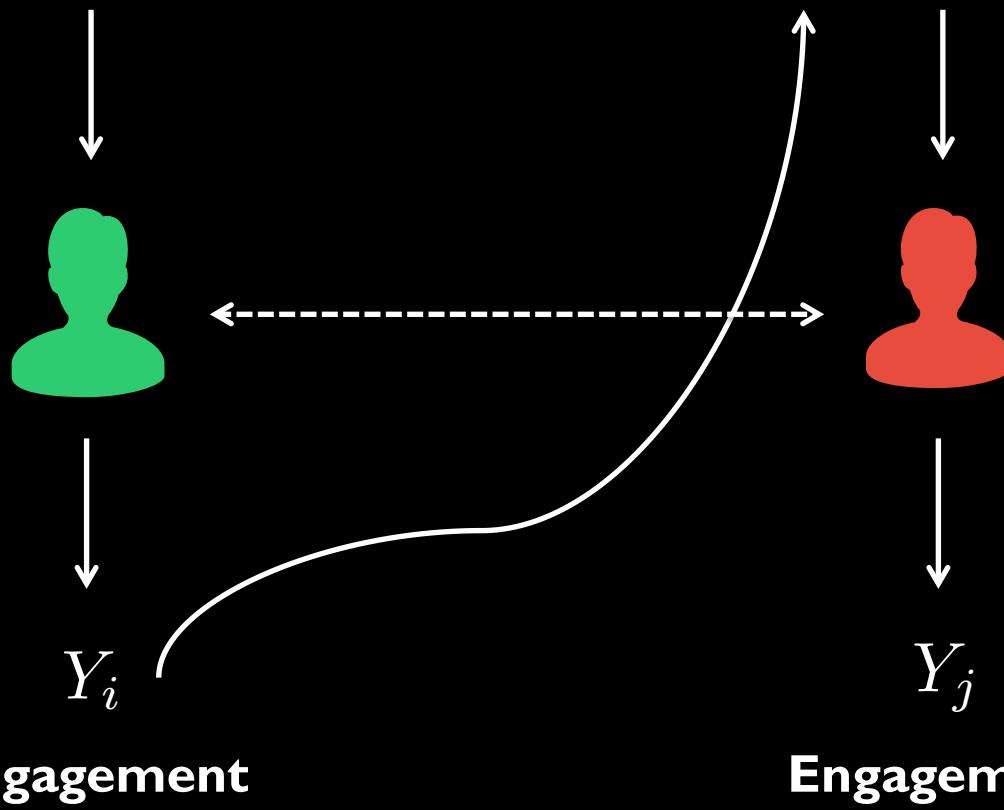
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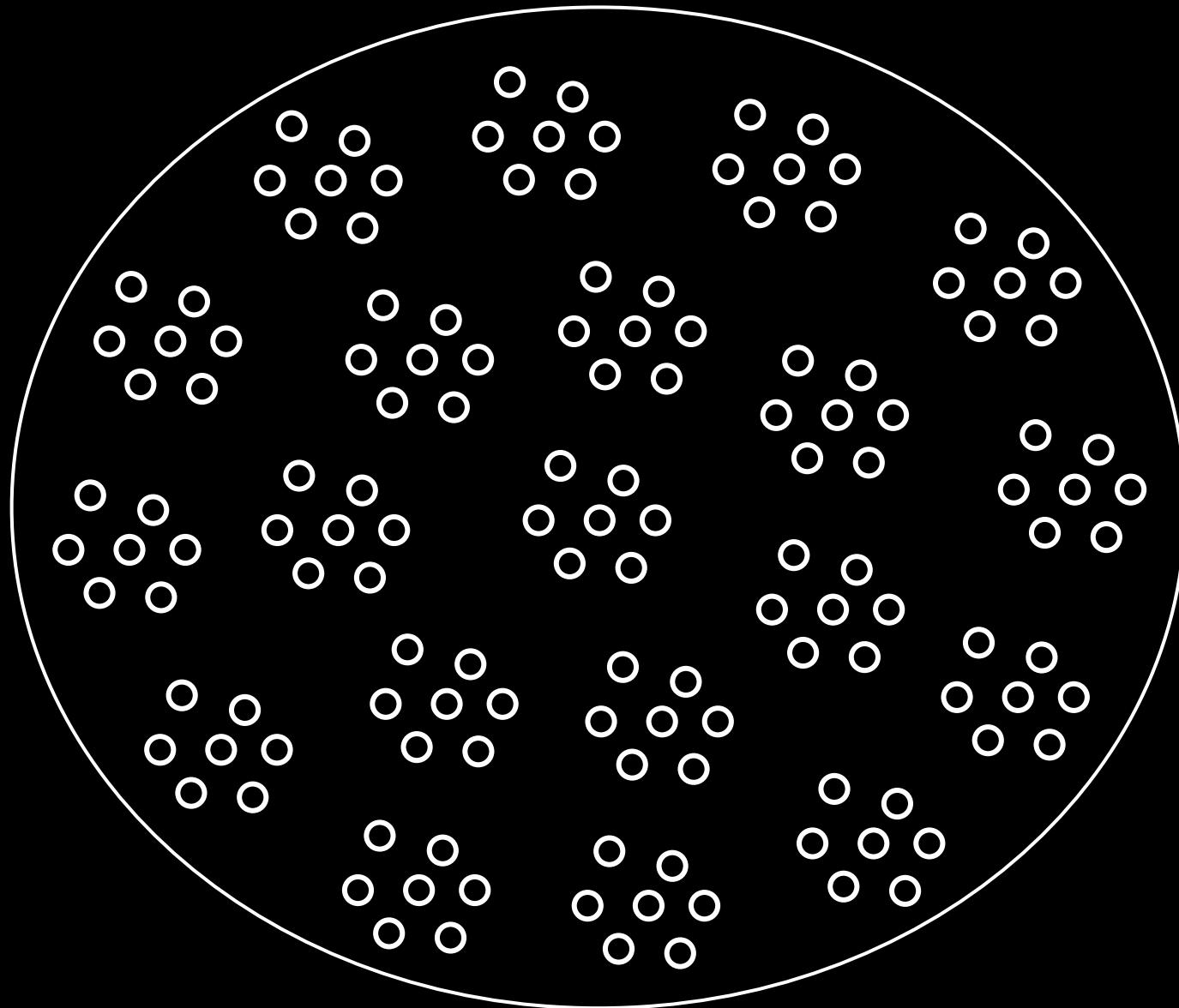
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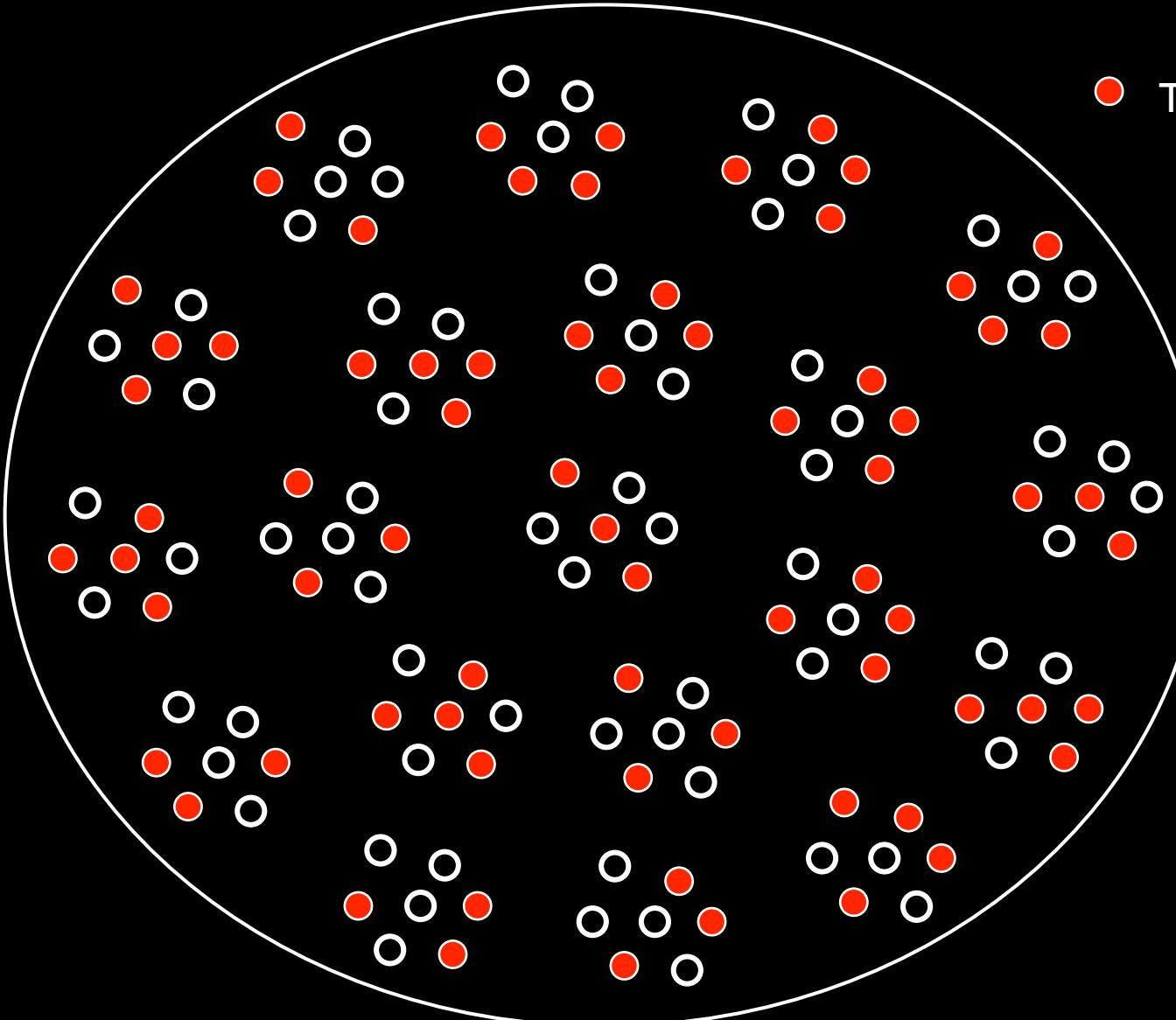
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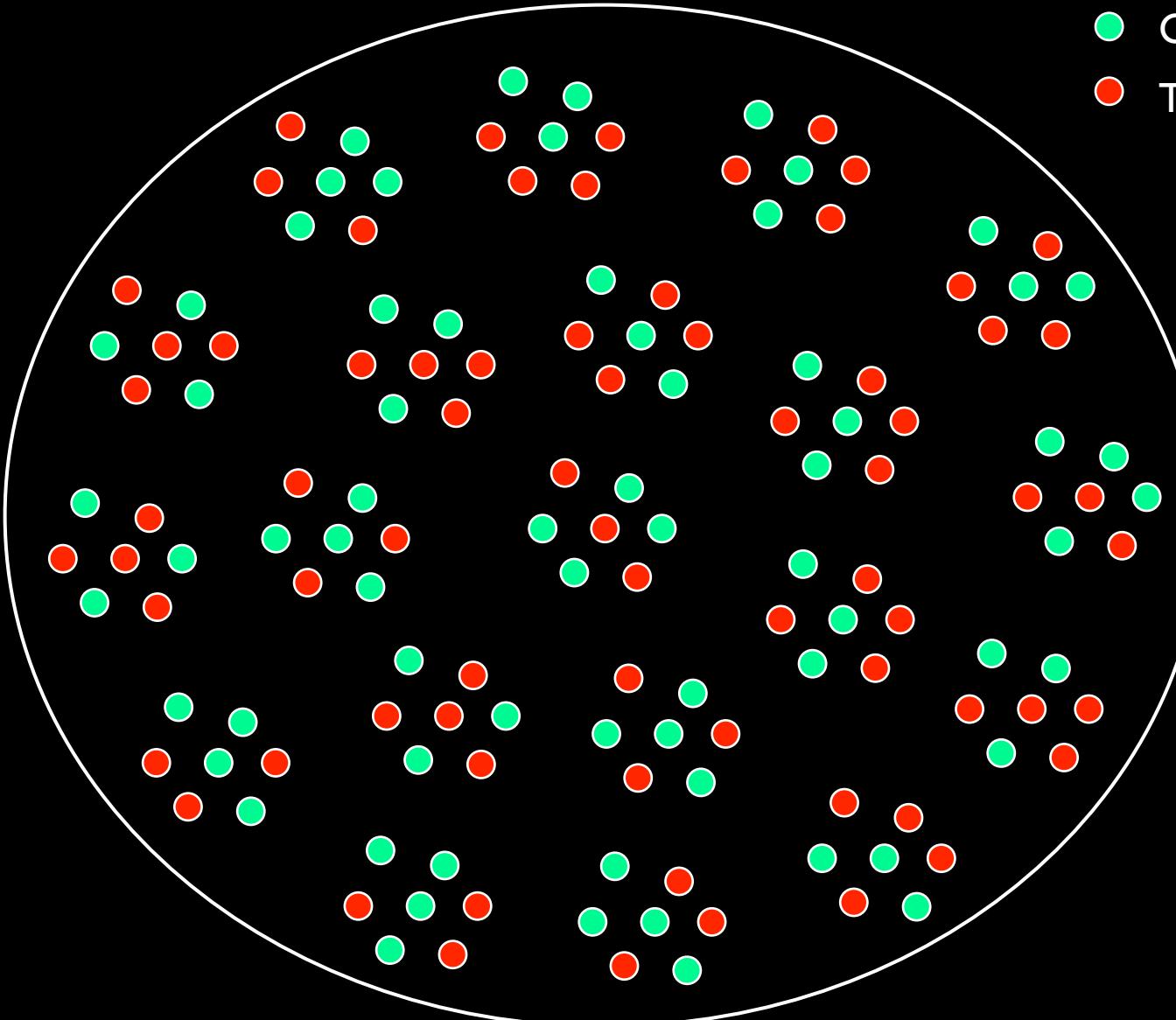


Completely-randomized Experiment



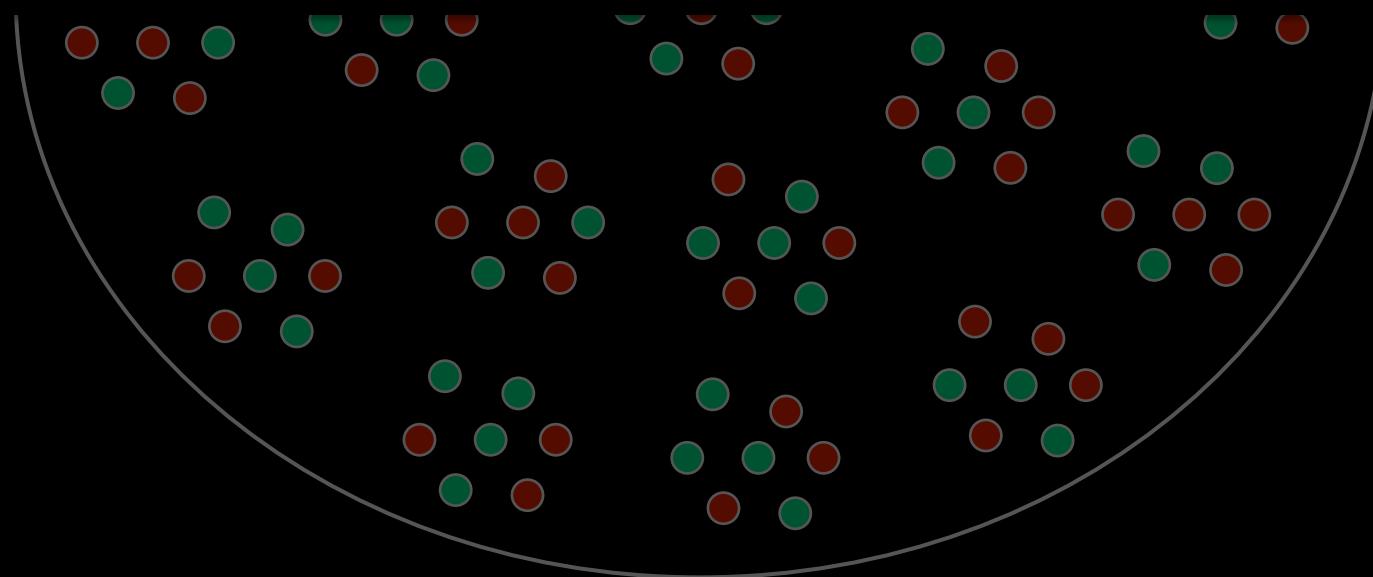
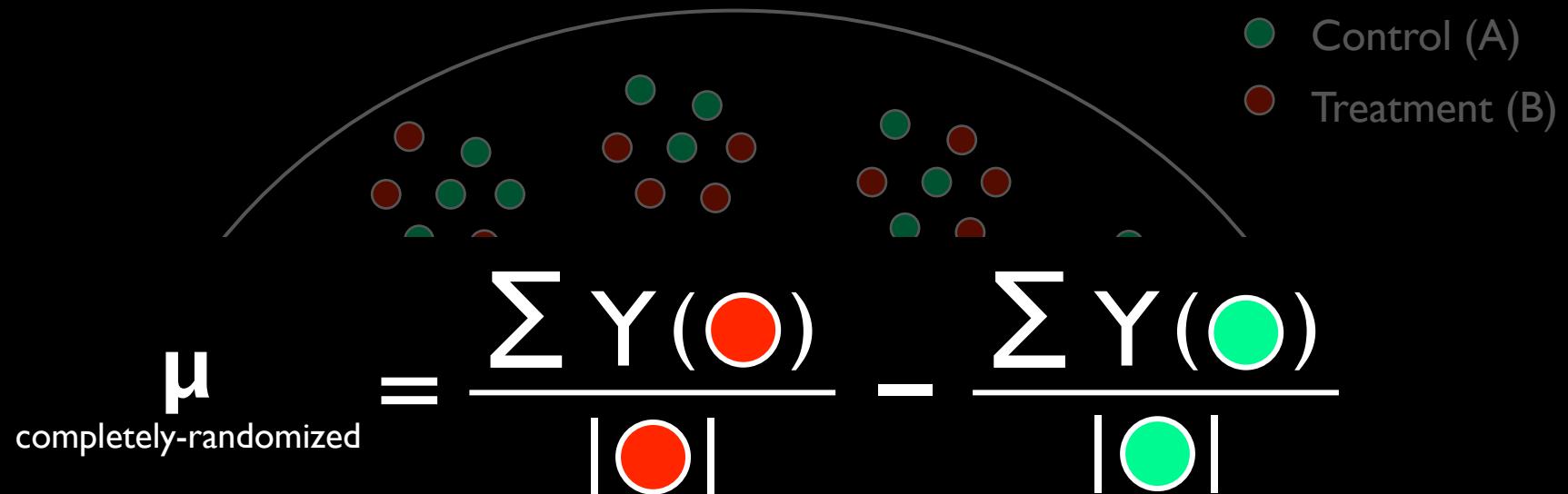
Treatment (B)

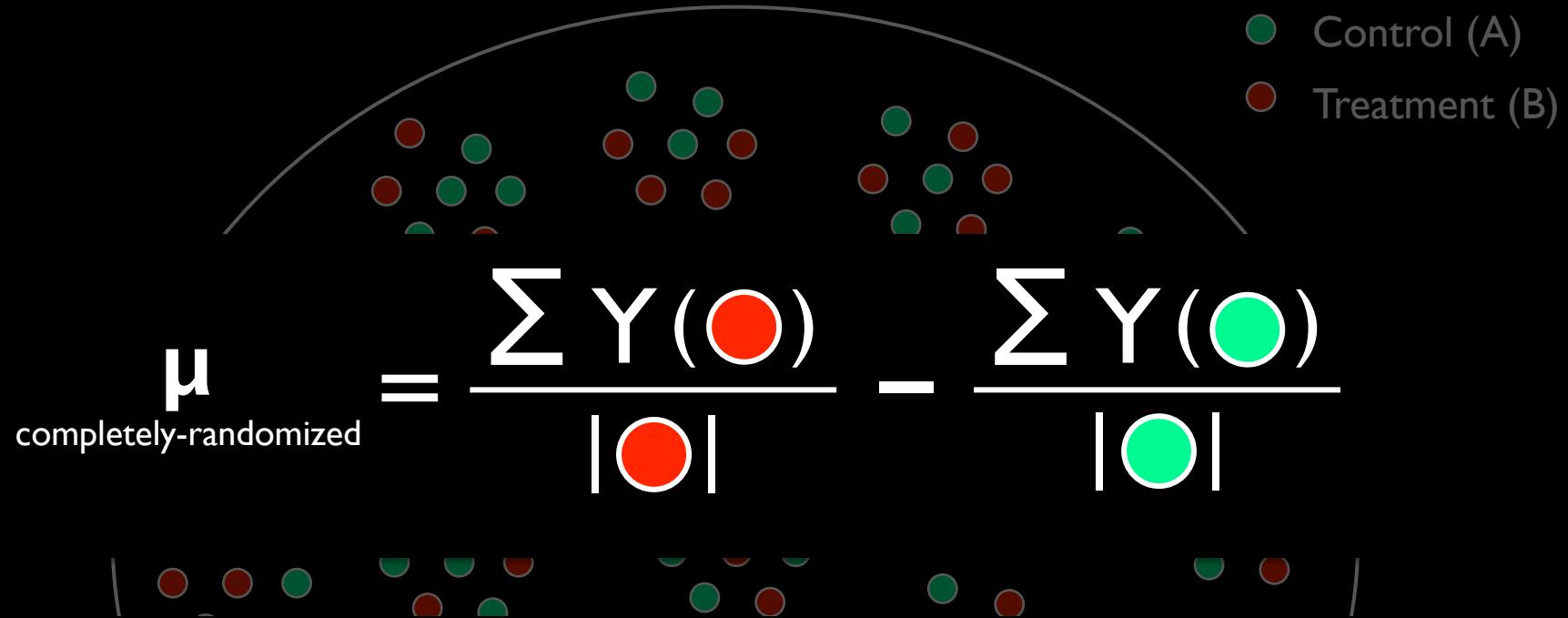
Completely-randomized Experiment



● Control (A)  
● Treatment (B)

Completely-randomized Experiment



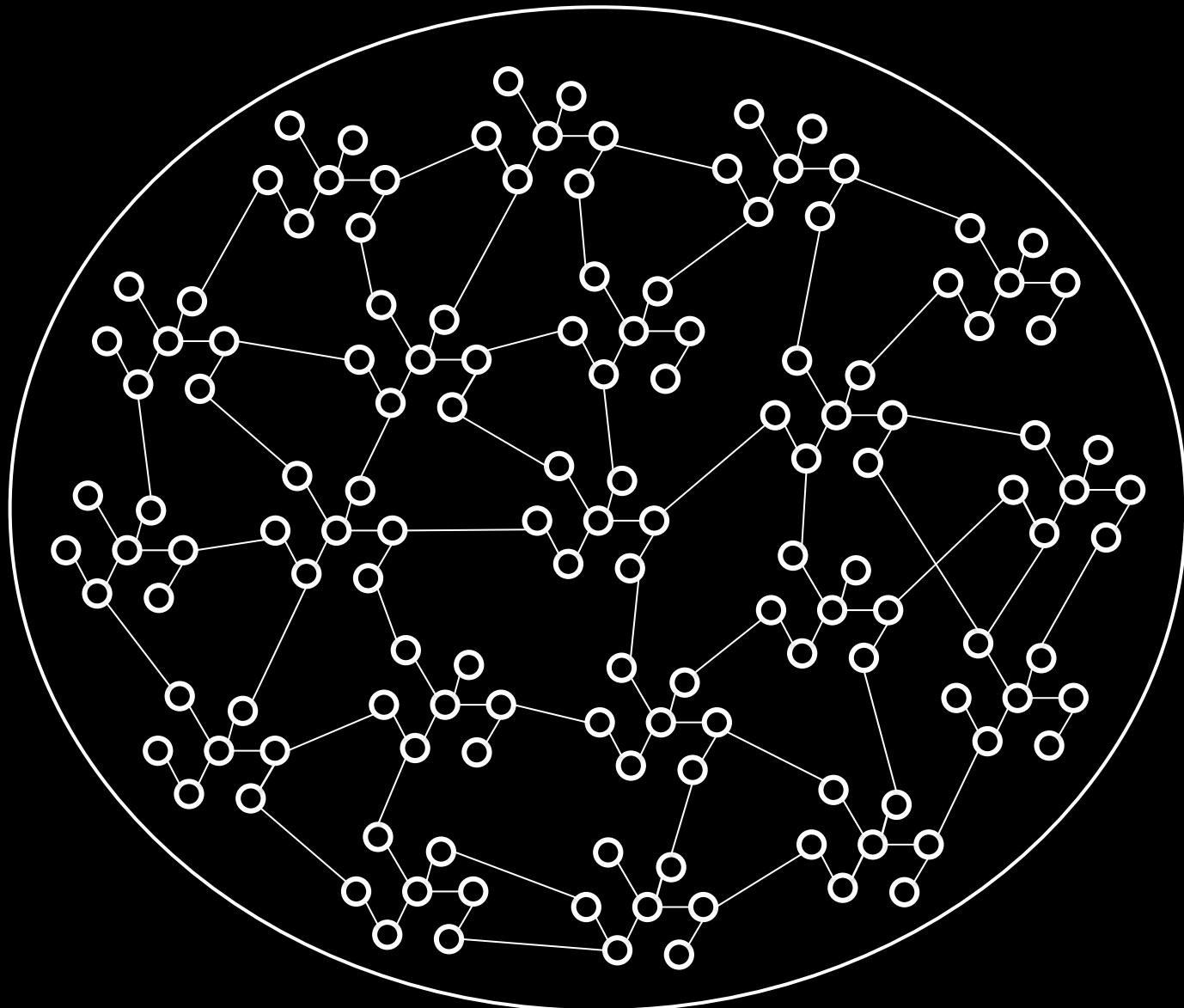


## SUTVA: Stable Unit Treatment Value Assumption

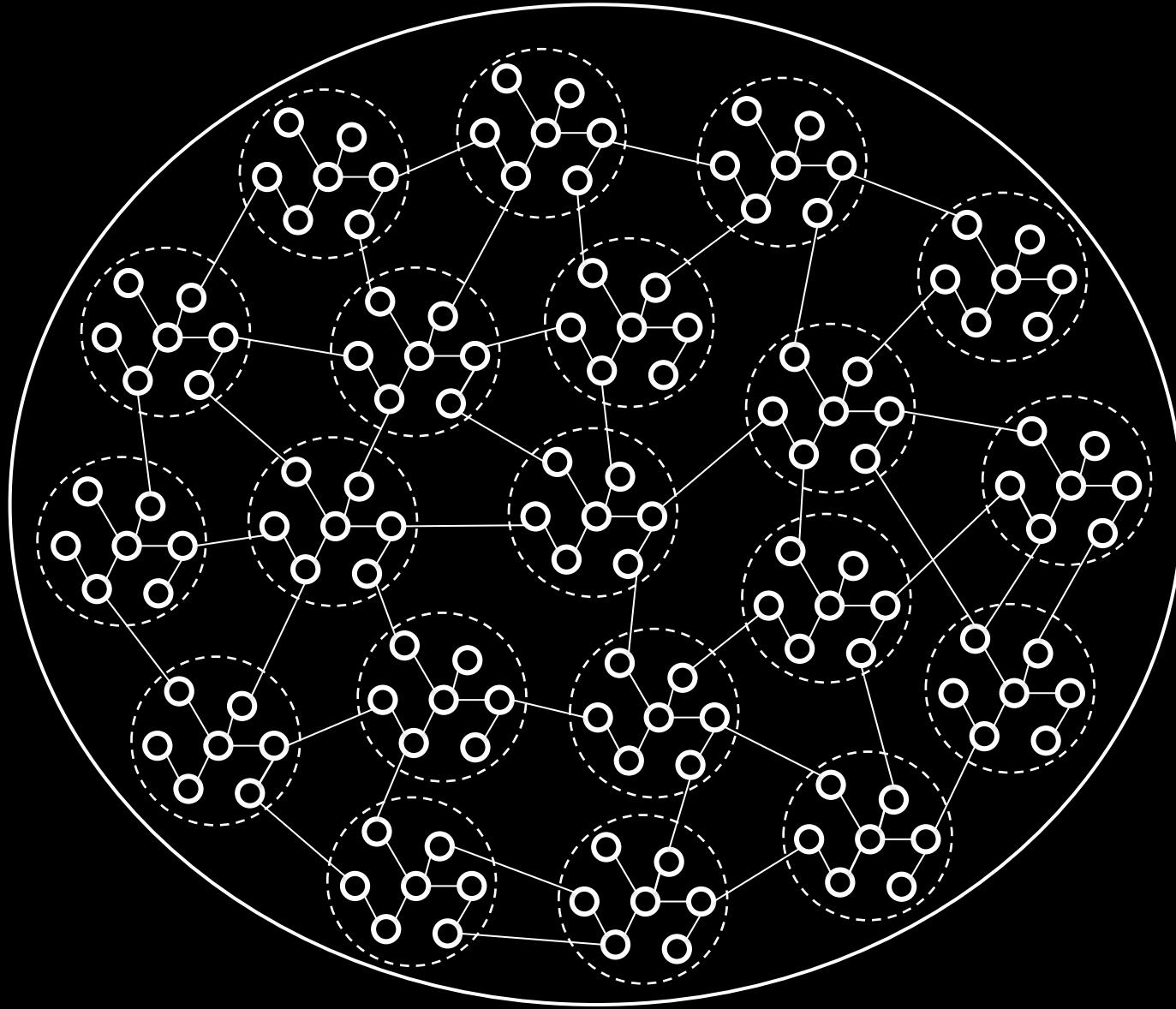
Every user's behavior is affected only by their treatment  
and NOT by the treatment of any other user

Completely-randomized Experiment

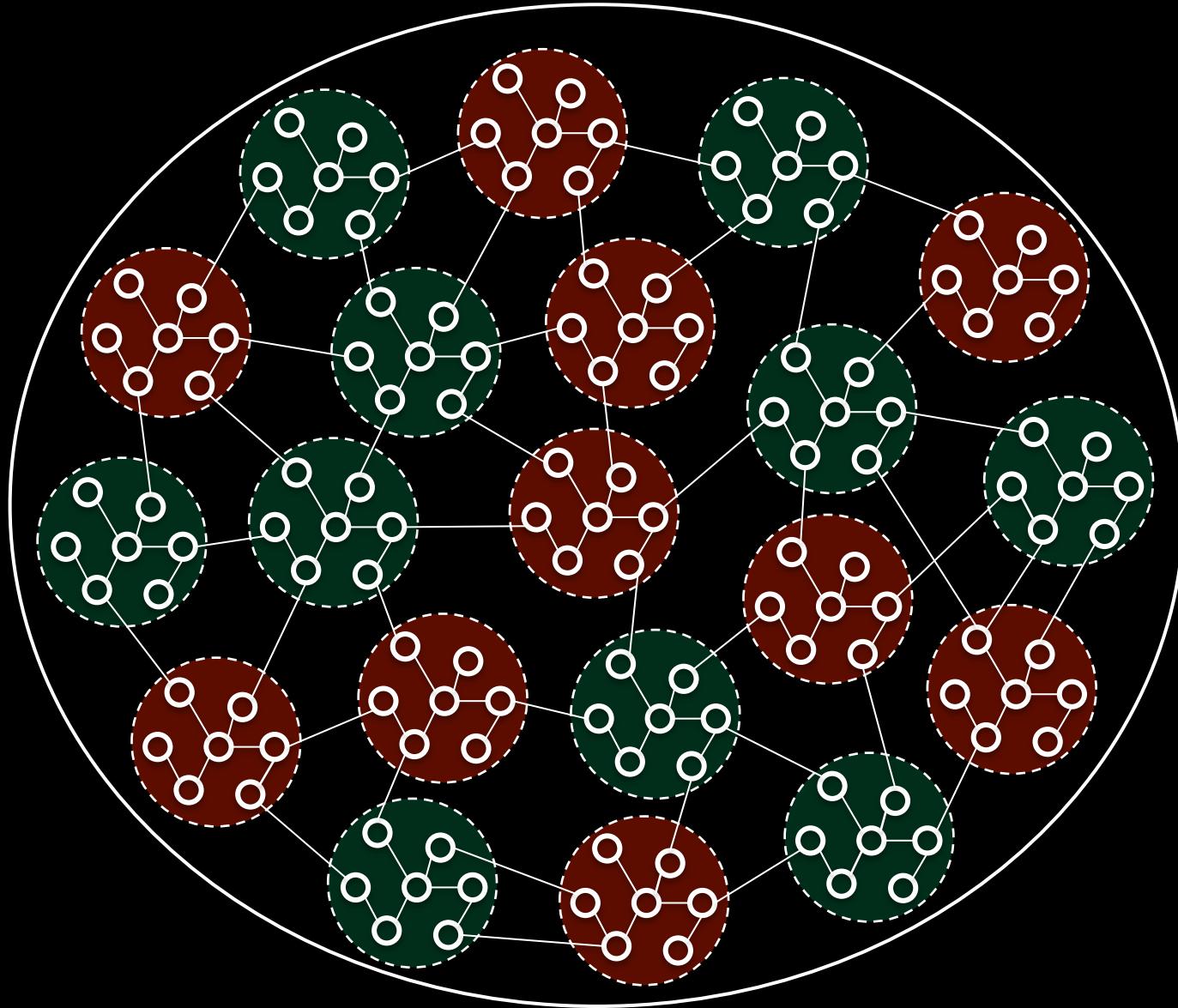
# Cluster-based Randomized Experiment



Cluster-based Randomized Experiment

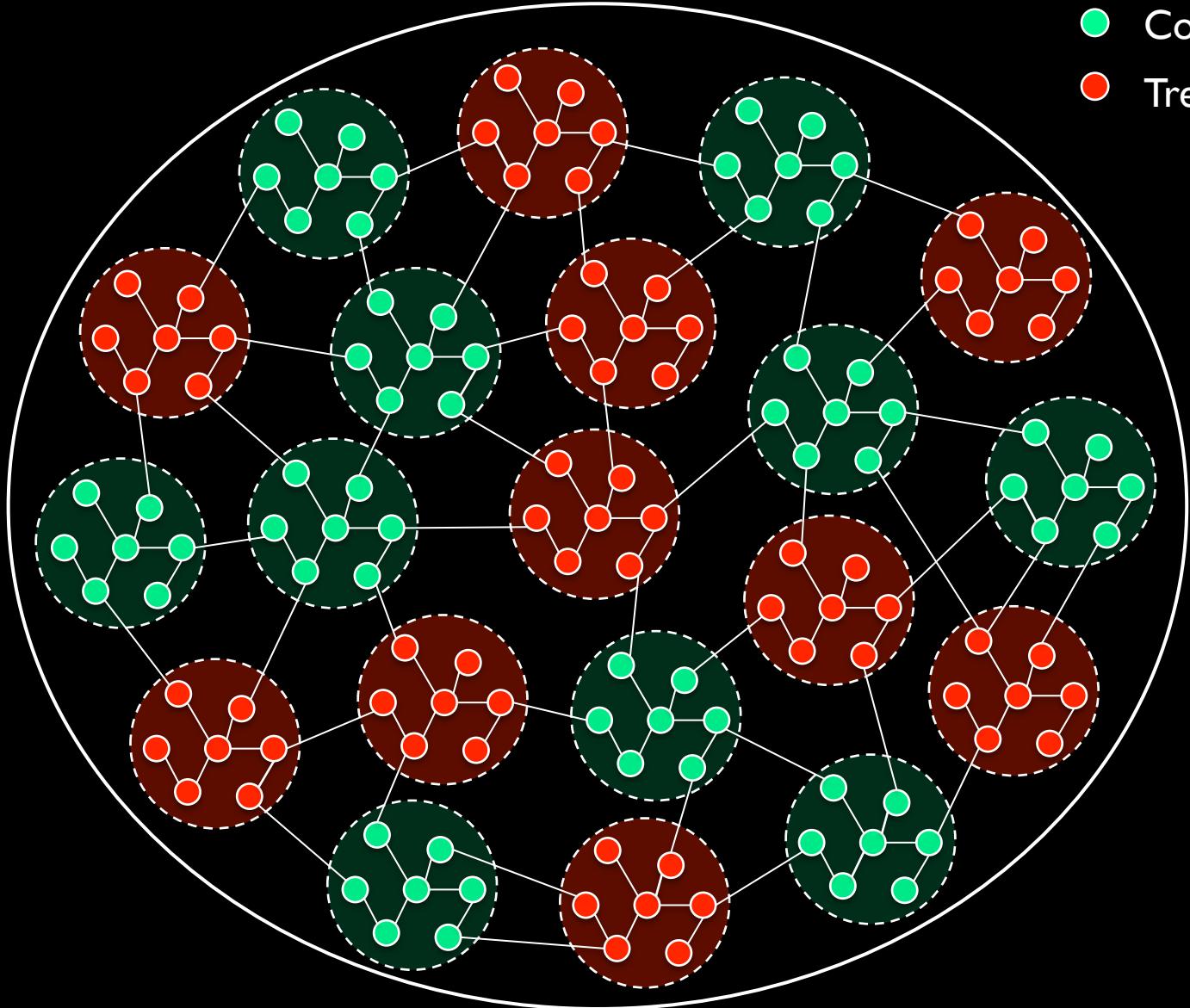


Cluster-based Randomized Experiment



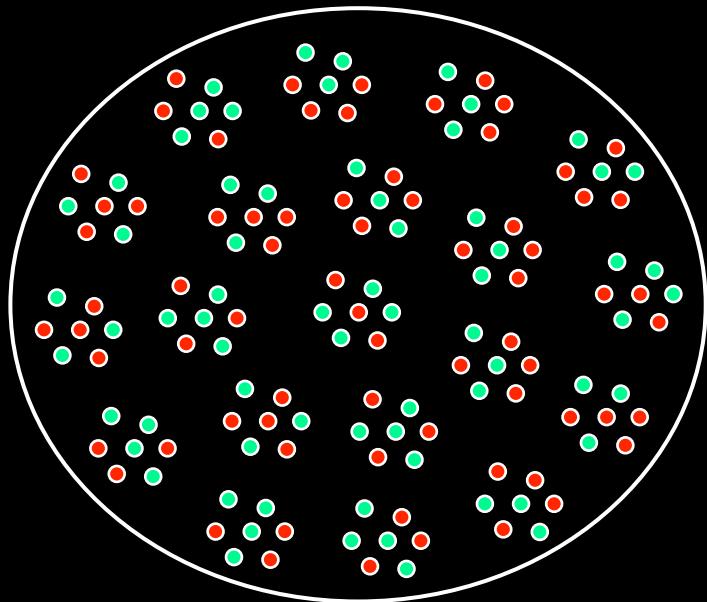
Cluster-based Randomized Experiment

- Control (A)
- Treatment (B)

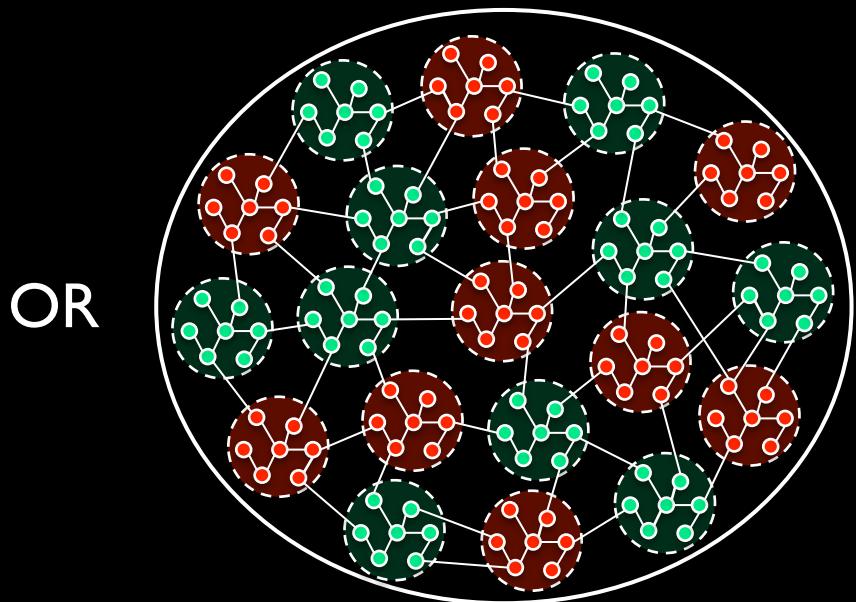


Cluster-based Randomized Experiment

Completely-randomized Experiment

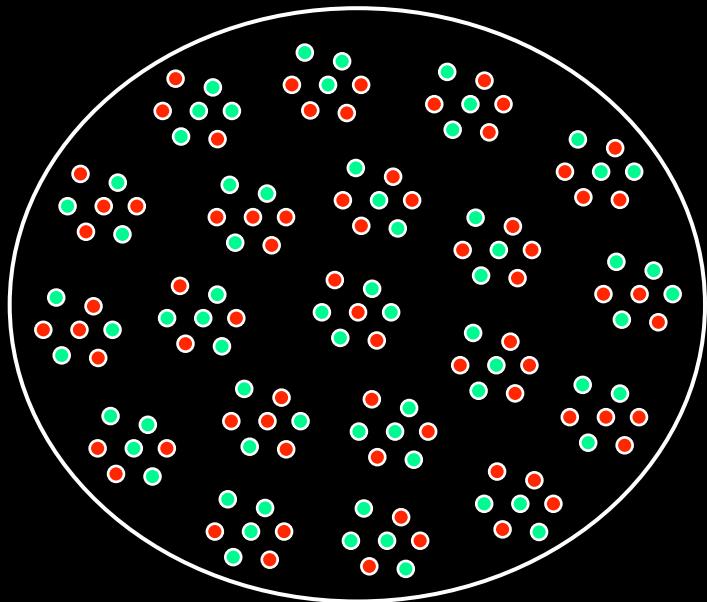


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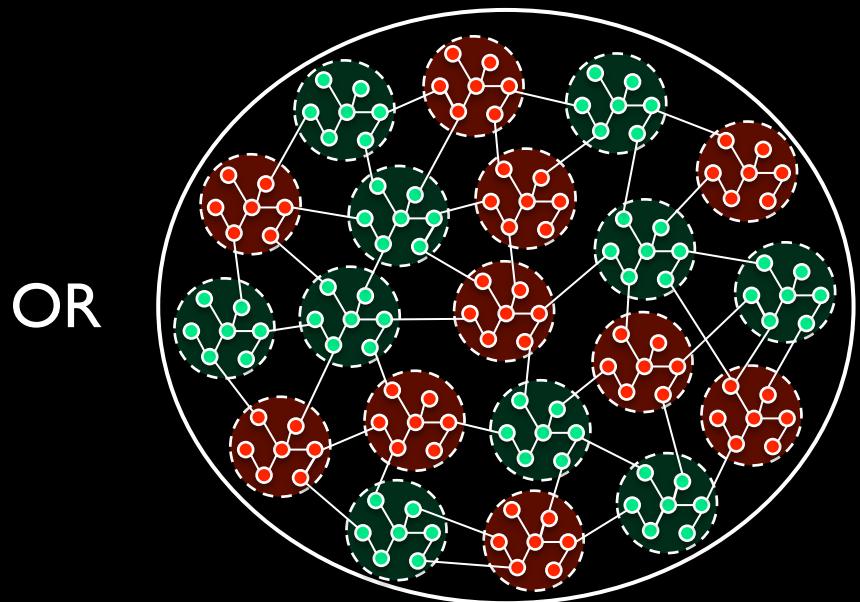


OR

Completely-randomized Experiment



Cluster-based Randomized Experiment



OR

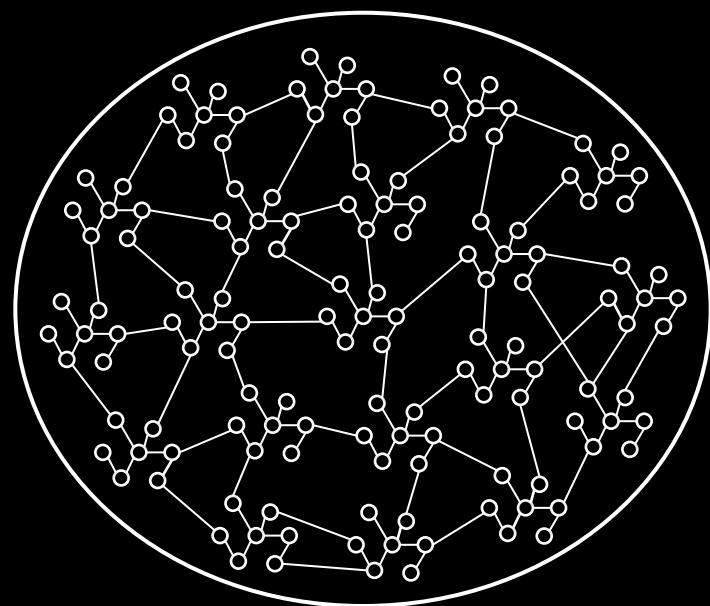
More Spillovers

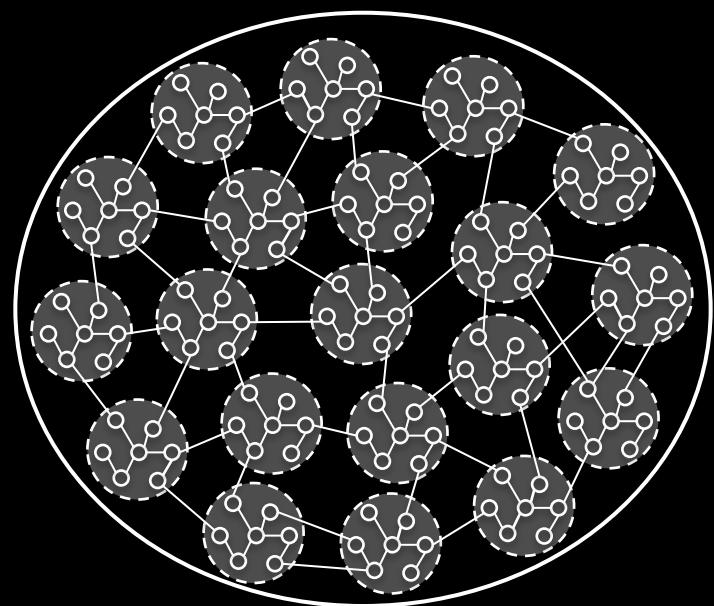
Lower Variance

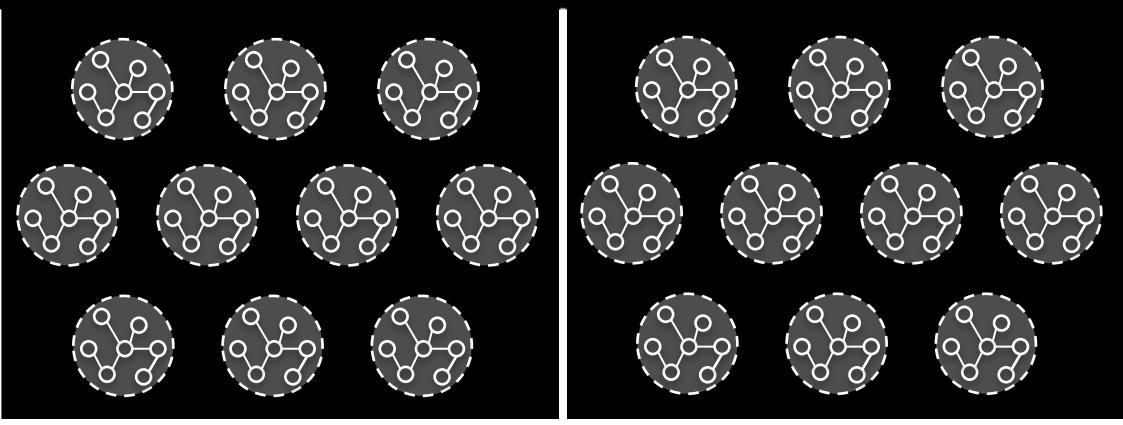
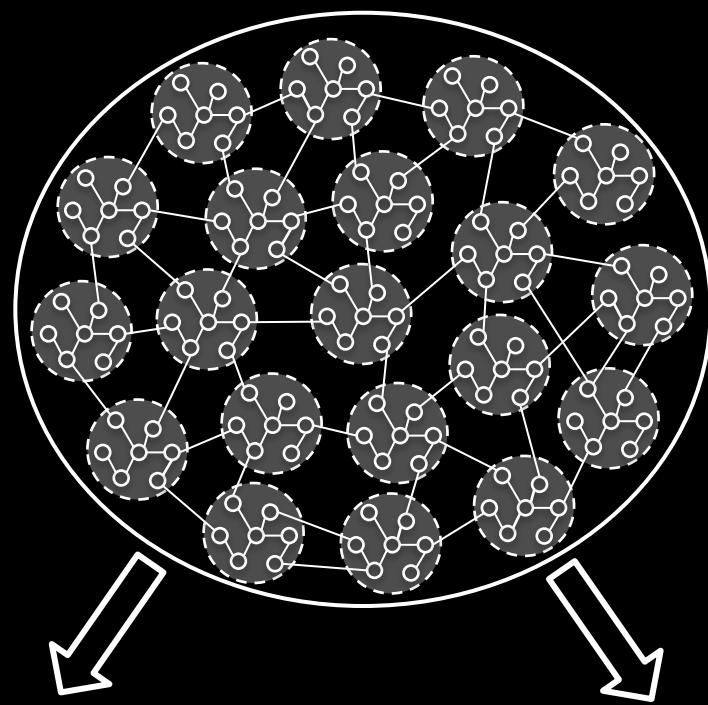
Less Spillovers

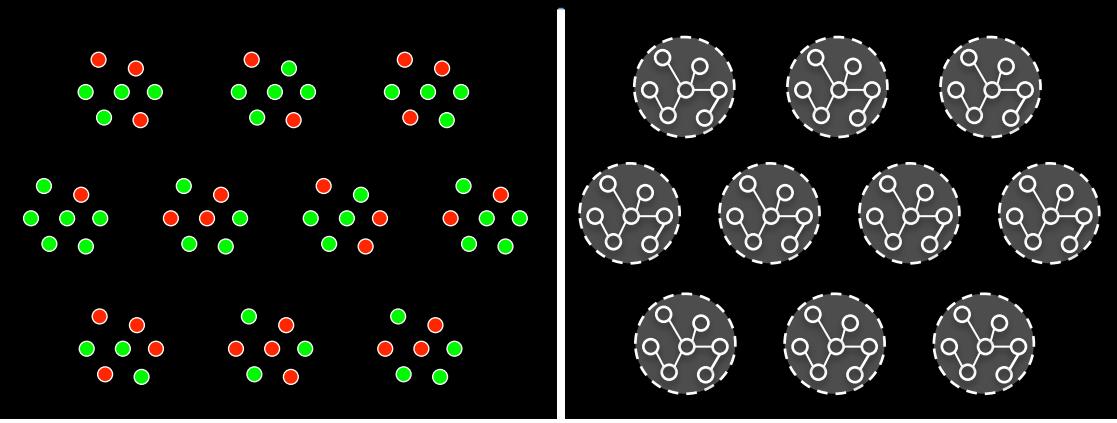
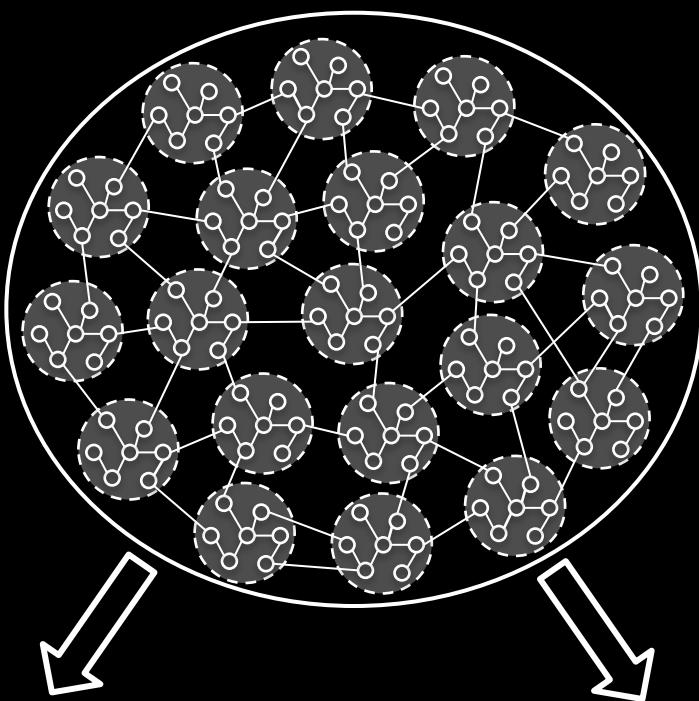
Higher Variance

# Design for Detecting Network Effects

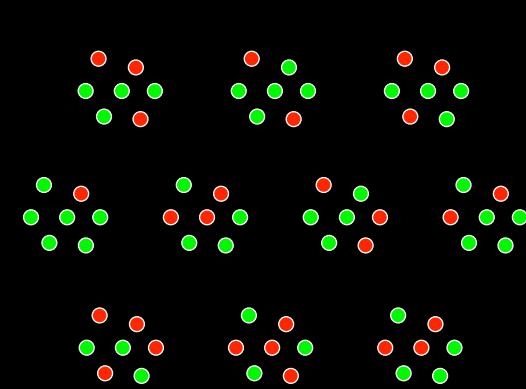
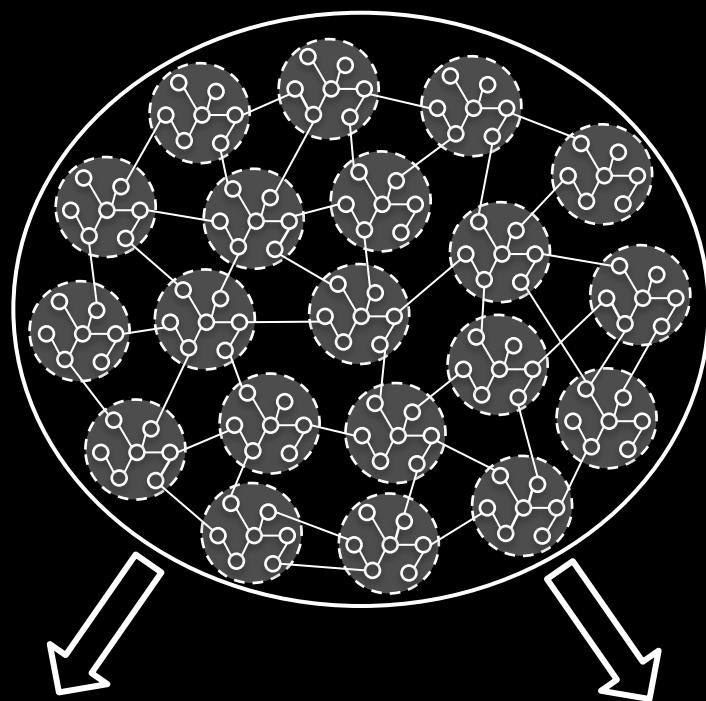




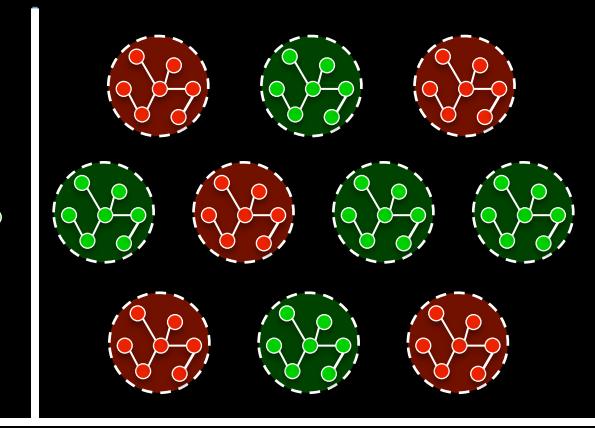




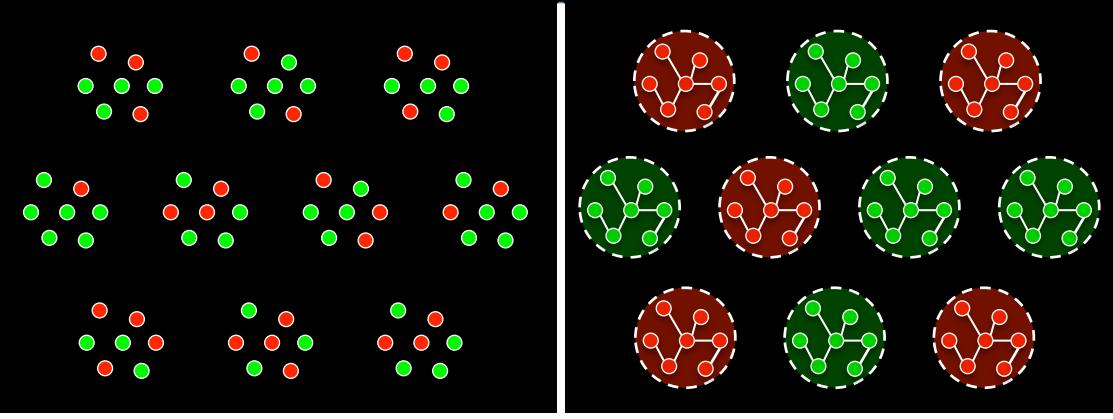
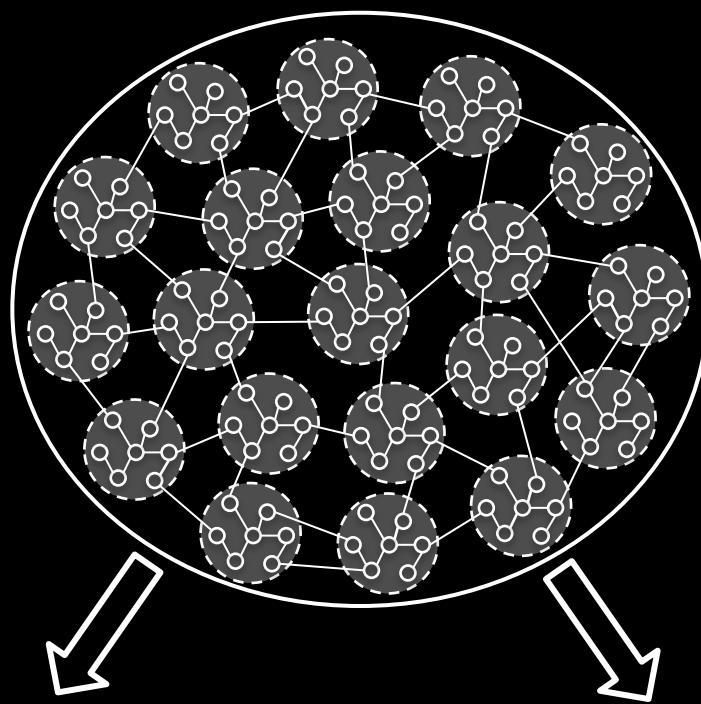
Completely Randomized  
Experiment



Completely Randomized  
Experiment



Cluster-based Randomized  
Experiment



Completely Randomized  
Experiment

Cluster-based Randomized  
Experiment

$$\mu_{\text{completely-randomized}} \stackrel{?}{=} \mu_{\text{cluster-based}}$$

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$H_0$ : SUTVA Holds

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Reject the null when:

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$$\left| \frac{\hat{\mu}_{cr} - \hat{\mu}_{cbr}}{\sqrt{\hat{\sigma}^2}} \right| \geq \frac{1}{\sqrt{\alpha}}$$

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Type I error is no greater than  $\alpha$

# Nuts and Bolts of Running Cluster-based Randomized Experiments

# Why Balanced Clustering?

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- Theoretical Motivation
  - Constants VS random variables

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- Practical Motivations
  - Variance reduction
  - Balance on pre-treatment covariates  
(homophily => large homogenous clusters)

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**Most clustering methods find skewed distributions of cluster sizes**

(Leskovec, 2009; Fortunato, 2010)

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**Restreaming Linear Deterministic Greedy**

(Nishimura & Ugander, 2013)

# Algorithms for Balanced Clustering

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=> Algorithms that enforce equal cluster sizes

## Restreaming Linear Deterministic Greedy

(Nishimura & Ugander, 2013)

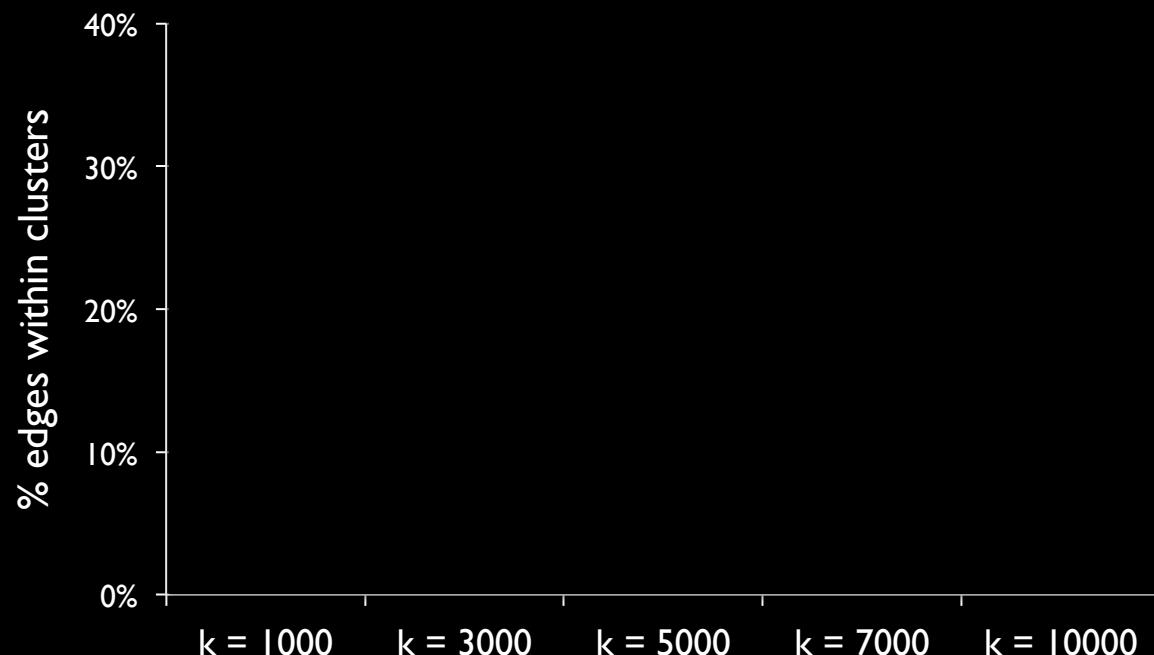
- Streaming
- Parallelizable
- Stable

# Clustering the LinkedIn Graph

- Graph: >100M nodes, >10B edges
- 350 Hadoop nodes
- 1% leniency

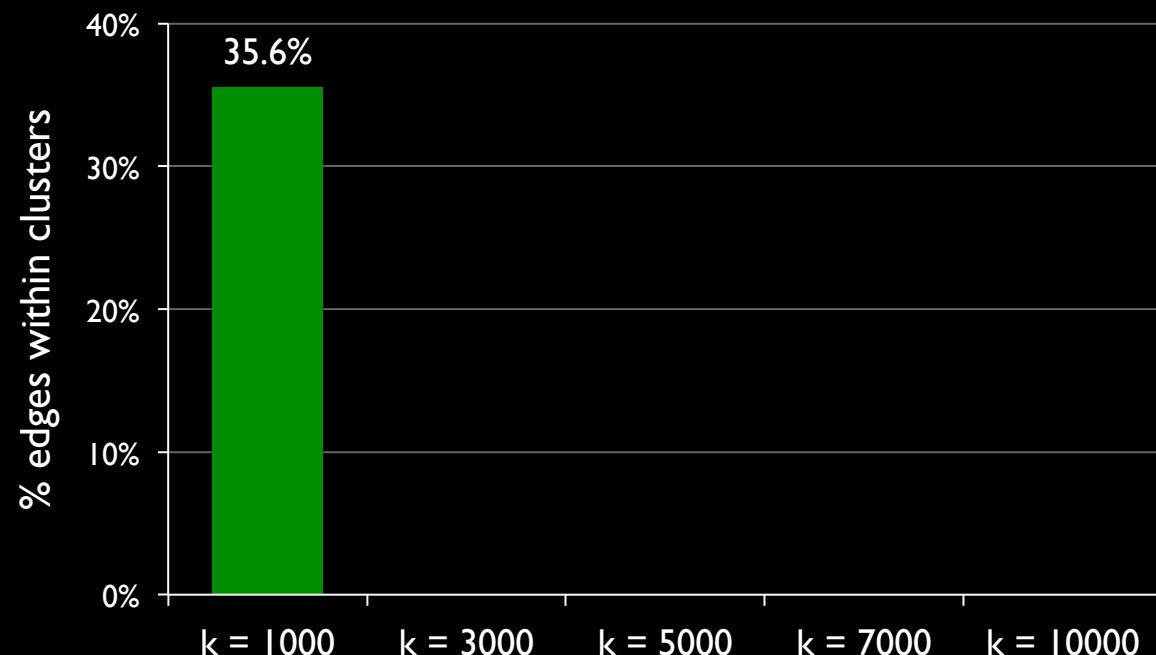
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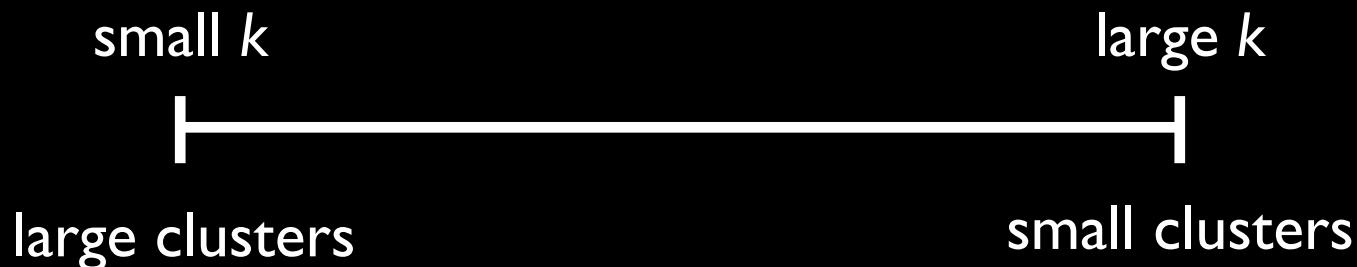
small  $k$



large  $k$



# Choosing the Number of Clusters



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Understanding the Type II error

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Assuming an interference model

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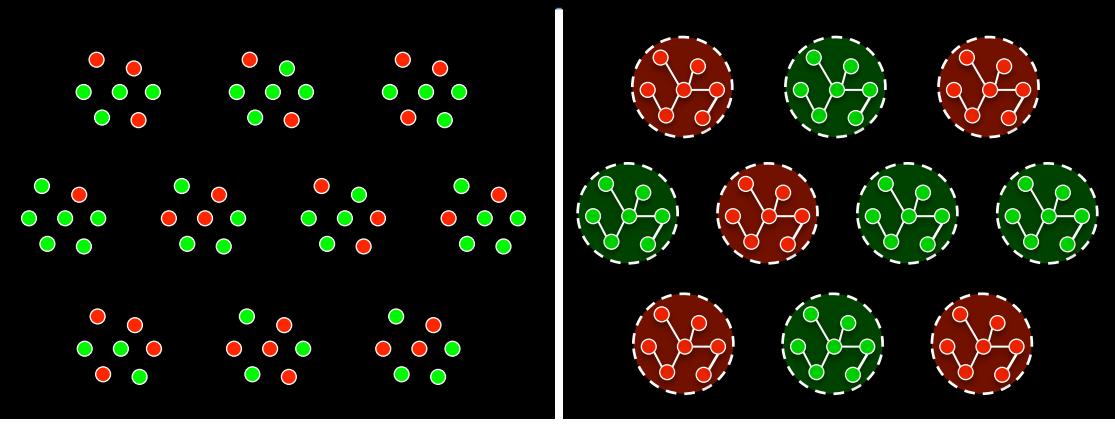
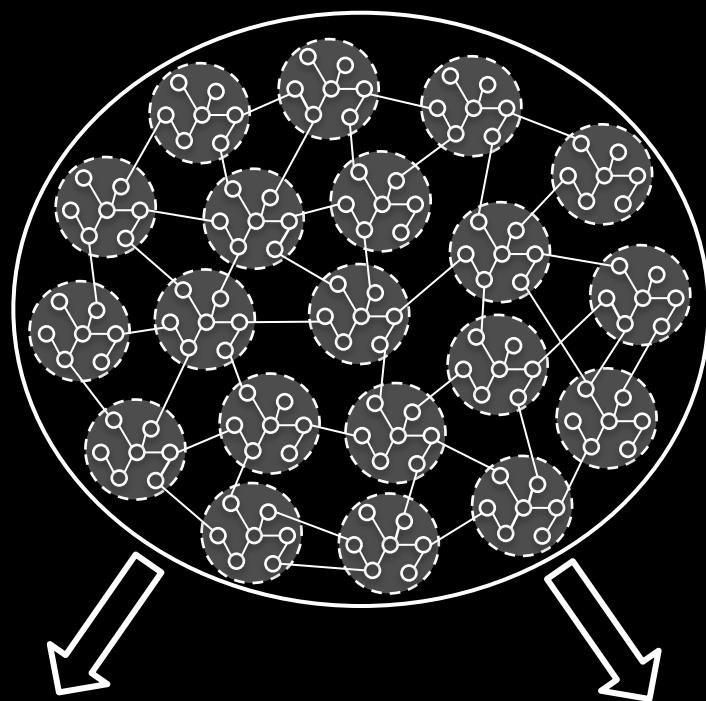
$$E [\hat{\mu}_{cbr} - \hat{\mu}_{cr}] \approx \rho \cdot \beta_2$$

$\rho$  : average fraction of a unit's neighbors contained in the cluster

Choose number of clusters  $M$  and clustering  $C$  such that

$$\max_{M,C} \frac{\rho}{\sqrt{\hat{\sigma}_C^2}}$$

# Experiments on LinkedIn



Bernoulli  
Randomized  
Experiment

$\mu_{\text{bernoulli}}$

Completely Randomized  
Experiment

Cluster-based Randomized  
Experiment

$$\mu_{\text{completely-randomized}} \stackrel{?}{=} \mu_{\text{cluster-based}}$$

# Experiment I

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- Population: 20% of all LinkedIn users [Bernoulli: 10%, Cluster-based: 10%]

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	Treatment effect	Standard Deviation
Bernoulli Randomization (BR)		
Cluster-based Randomization (CBR)		
Delta (CBR – BR)		

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Bernoulli Randomization (BR)	0.0559	0.0050
Cluster-based Randomization (CBR)		
Delta (CBR – BR)		

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	Treatment effect	Standard Deviation
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Cluster-based Randomization (CBR)	0.0771	0.0260
Delta (CBR – BR)		

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Bernoulli Randomization (BR)	0.0559	0.0050
Cluster-based Randomization (CBR)	0.0771	0.0260
Delta (CBR – BR)	-0.0211	0.0265

p-value: 0.4246

# Experiment 2

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- Population: 36% of all LinkedIn users [Bernoulli: 20%, Cluster-based: 16%]

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Bernoulli Randomization (BR)	0.2108	0.2911
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Bernoulli Randomization (BR)	0.2108	0.2911
Cluster-based Randomization (CBR)	0.5390	0.5613
Delta (CBR – BR)	-0.3281	0.5712

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Bernoulli Randomization (BR)	0.2108	0.2911
Cluster-based Randomization (CBR)	0.5390	0.5613
Delta (CBR – BR)	-0.3281	0.5712

p-value: 0.0483

**Test SUTVA null**

Test SUTVA null

reject



Test SUTVA null

reject

Use cluster-based  
experiment to estimate  
treatment effects

Test SUTVA null

reject

Use cluster-based  
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(higher variance)

## Test SUTVA null

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fail to reject

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# Papers available online

KDD'17

Arxiv

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