

Graph models and hypothesis testing

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Why Graph models?

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Graph models allow us to generate synthetic networks

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They allow us to capture and model properties observed in real networks

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Graph models can serve as hypotheses for mechanisms of network formation

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Graph models allow us to generate synthetic networks

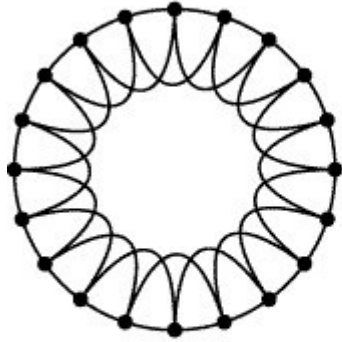
They allow us to capture and model properties observed in real networks

Graph models can serve as hypotheses for mechanisms of network formation

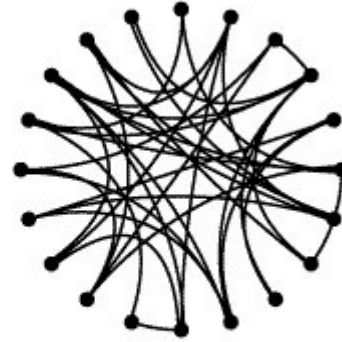
Allows us to explore how “similar” networks behave

Regular and random graphs

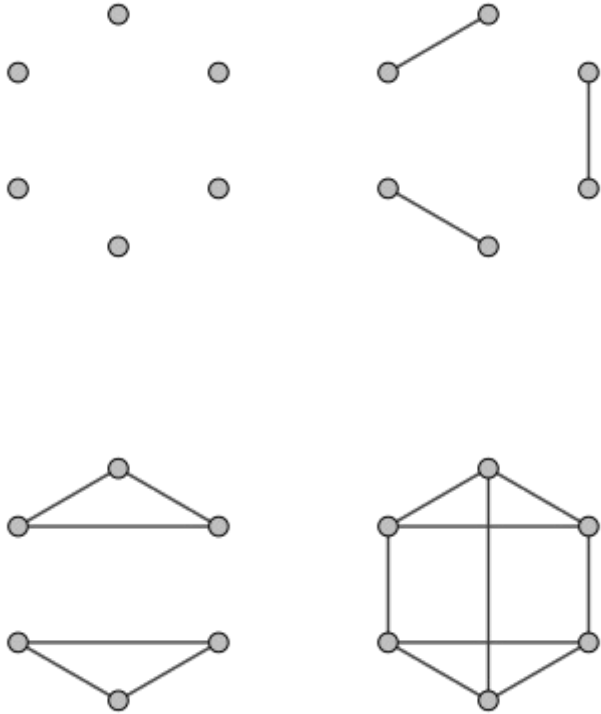
Regular



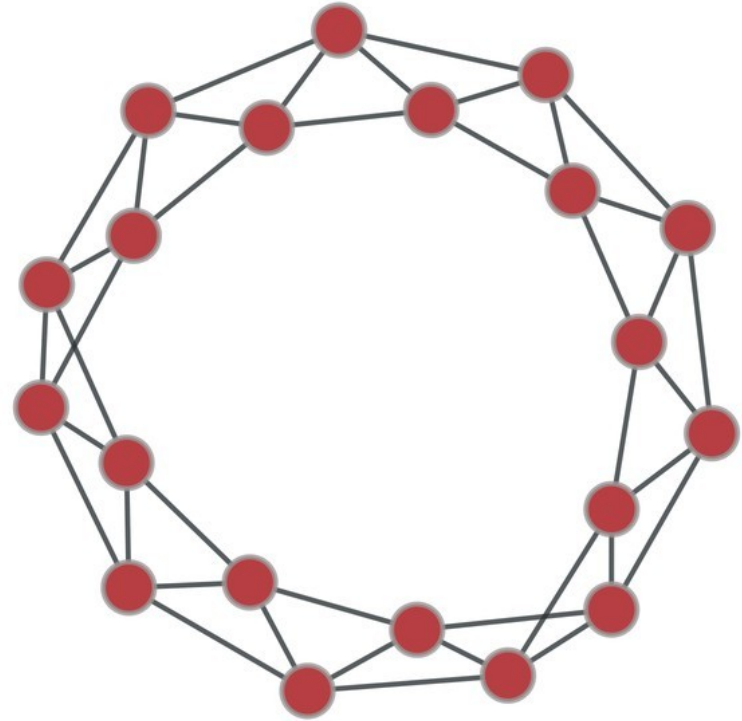
Random



Regular graphs

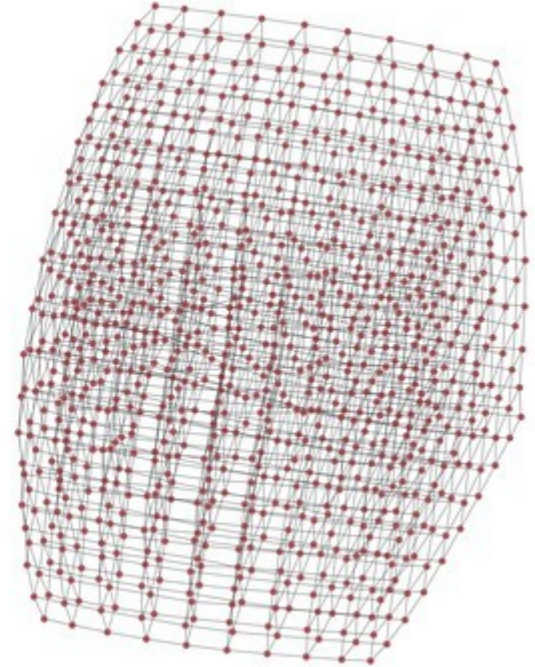
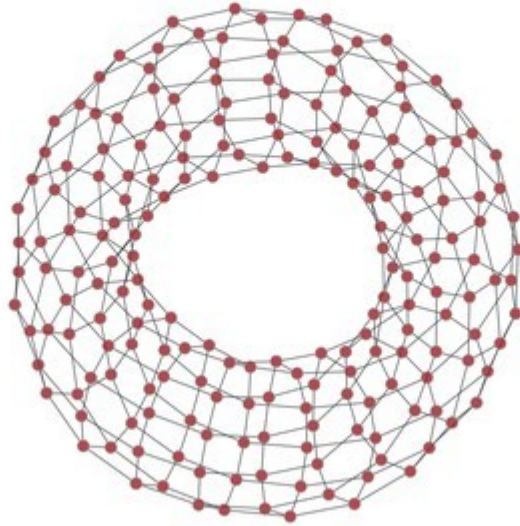
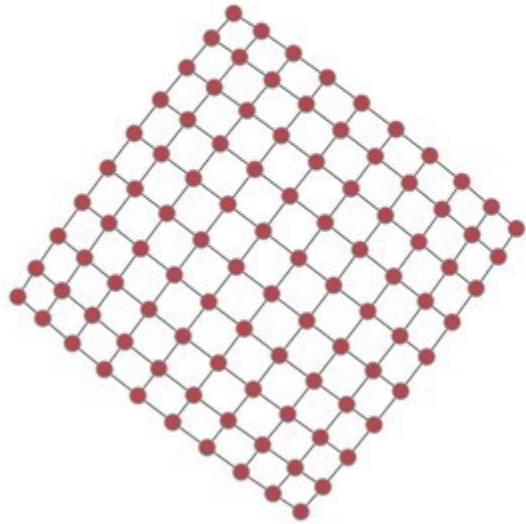


Regular graphs with 0 – 3 degree nodes



Regular Ring Lattice

Lattice graphs

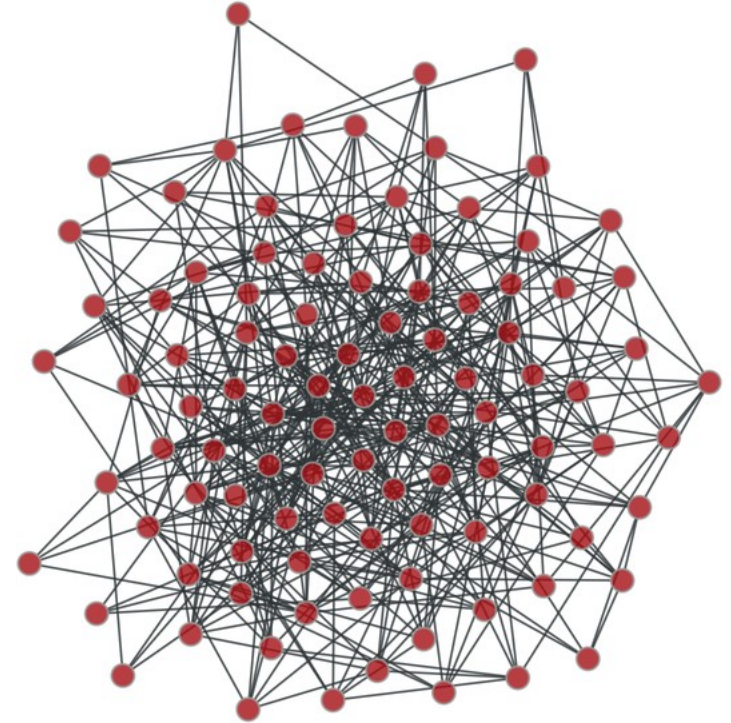


Random graphs

The Erdos-Renyi model

Specified by a number of nodes, n ,
and either:

- a number of edges, m
- a probability of connection, p



All edges are equally likely to exist

Small-world networks

It's a network after all

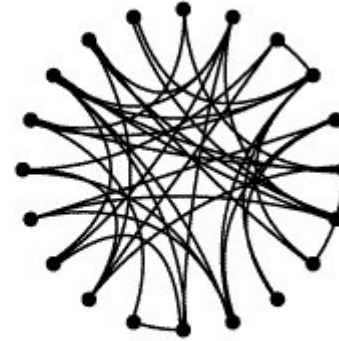
Small-world networks

It's a network after all

Regular



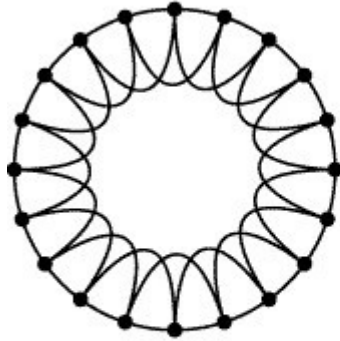
Random



Small-world networks

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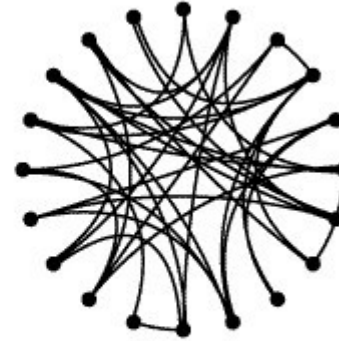
Regular



(triangles)

High **clustering coefficient**

Random

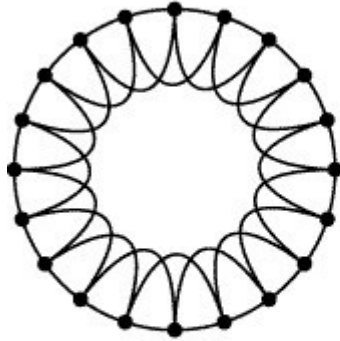


Low **clustering coefficient**

Small-world networks

It's a network after all

Regular



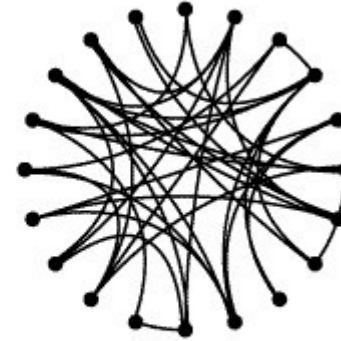
(triangles)

High **clustering coefficient**

(shortest
paths)

High mean **geodesic path**

Random



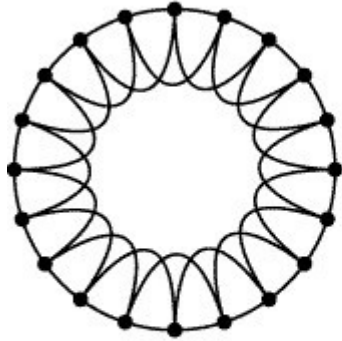
Low **clustering coefficient**

Low mean **geodesic path**

Small-world networks

It's a network after all

Regular



(triangles)

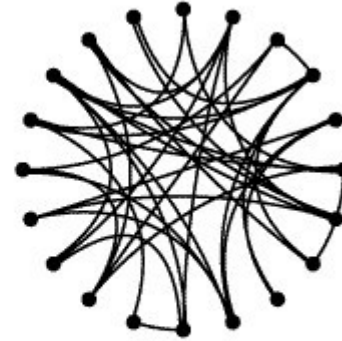
High clustering coefficient

(shortest paths)

High mean geodesic path

Real-world networks

Random

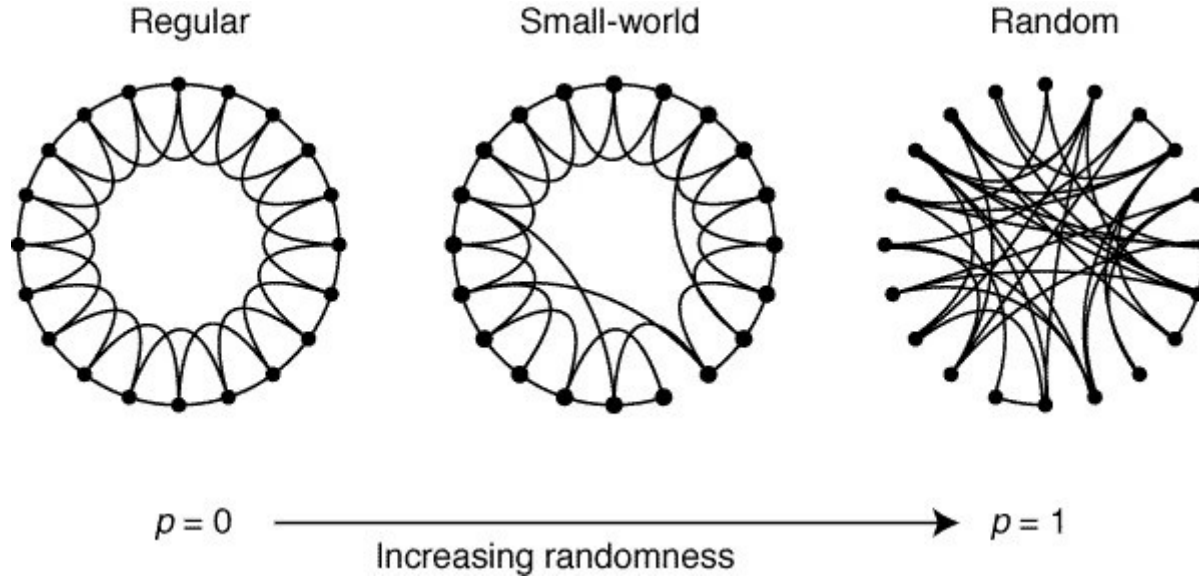


Low clustering coefficient

Low mean geodesic path

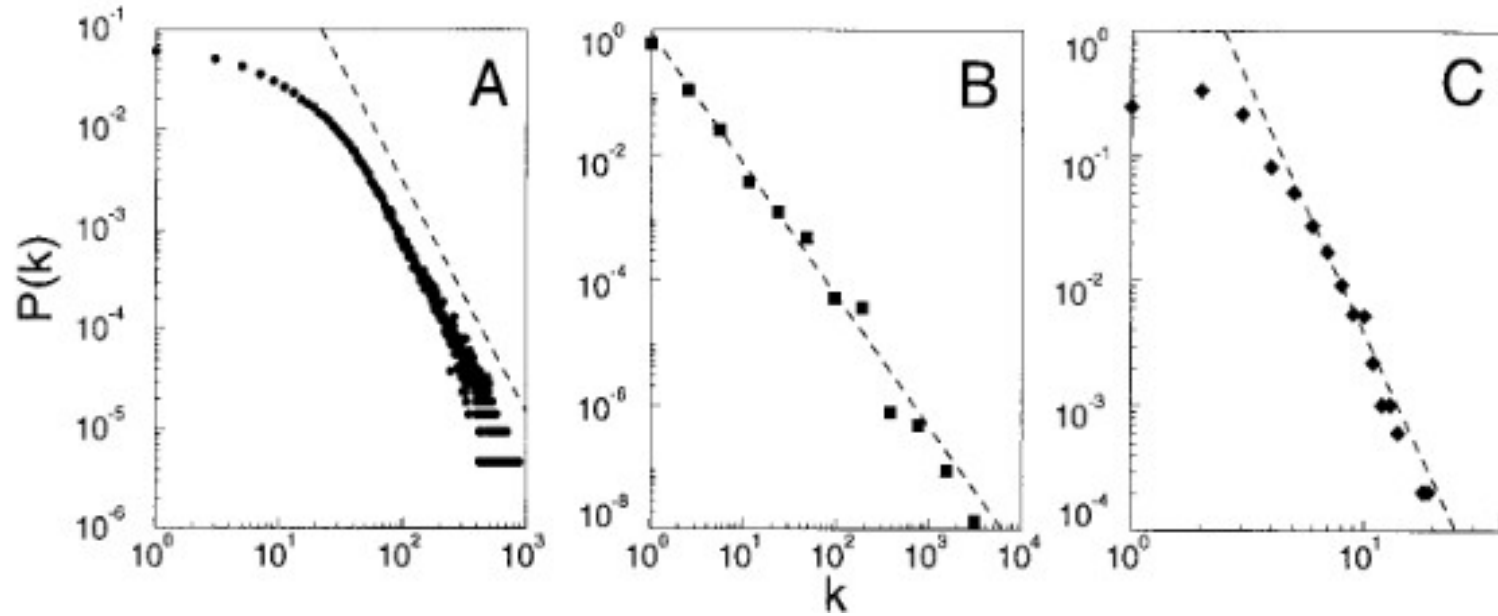
Small-world networks

It's a network after all



"Scale-free" networks

"Scale-free" networks



The Price model

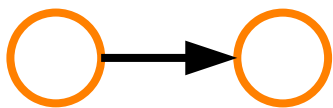
For undirected networks, this model is known as the Barabasi-Albert model

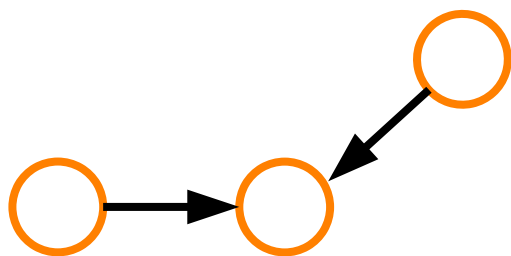
Add nodes to a network one at a time.

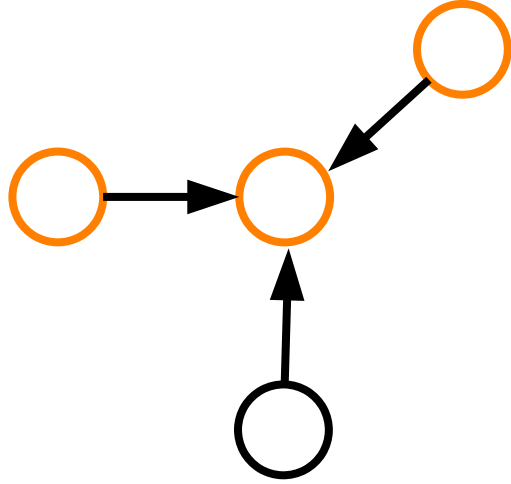
Connect to existing nodes with probability **proportional to their degree**

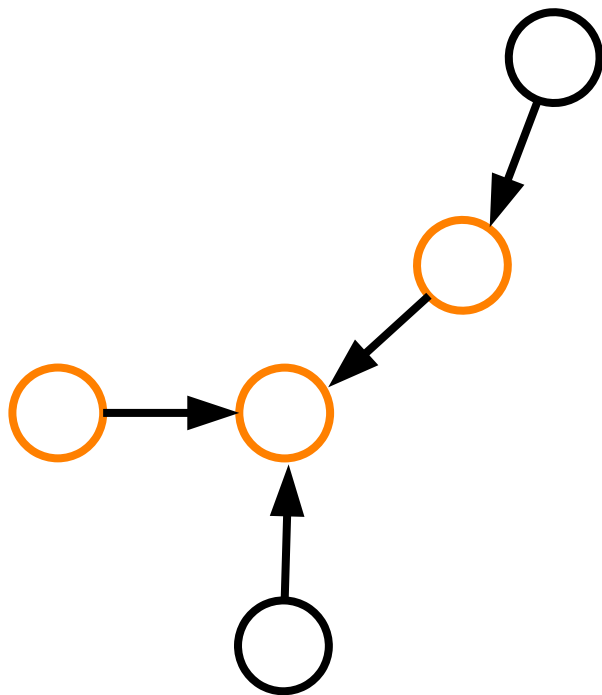
$$p_i = \frac{k_i}{\sum_j k_j},$$

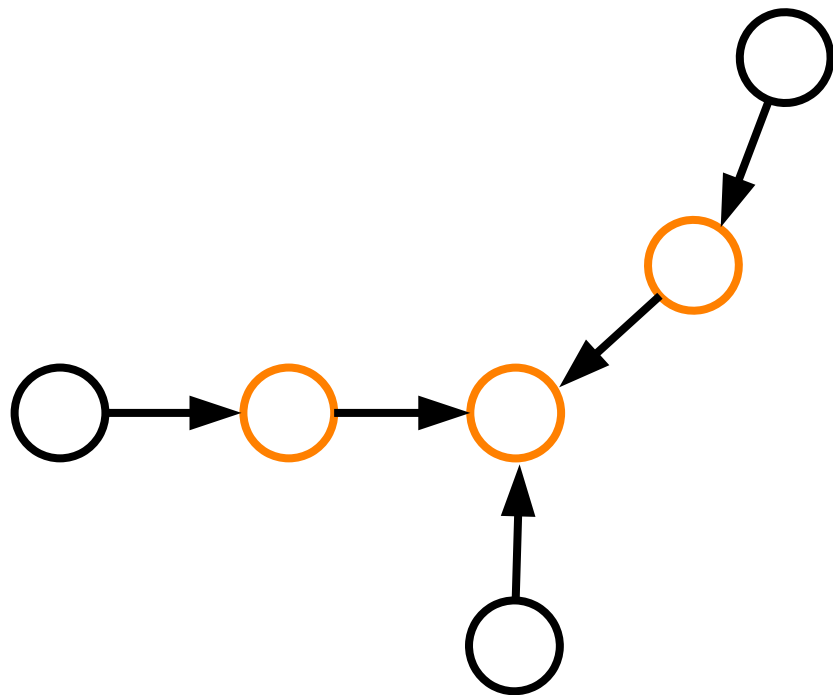


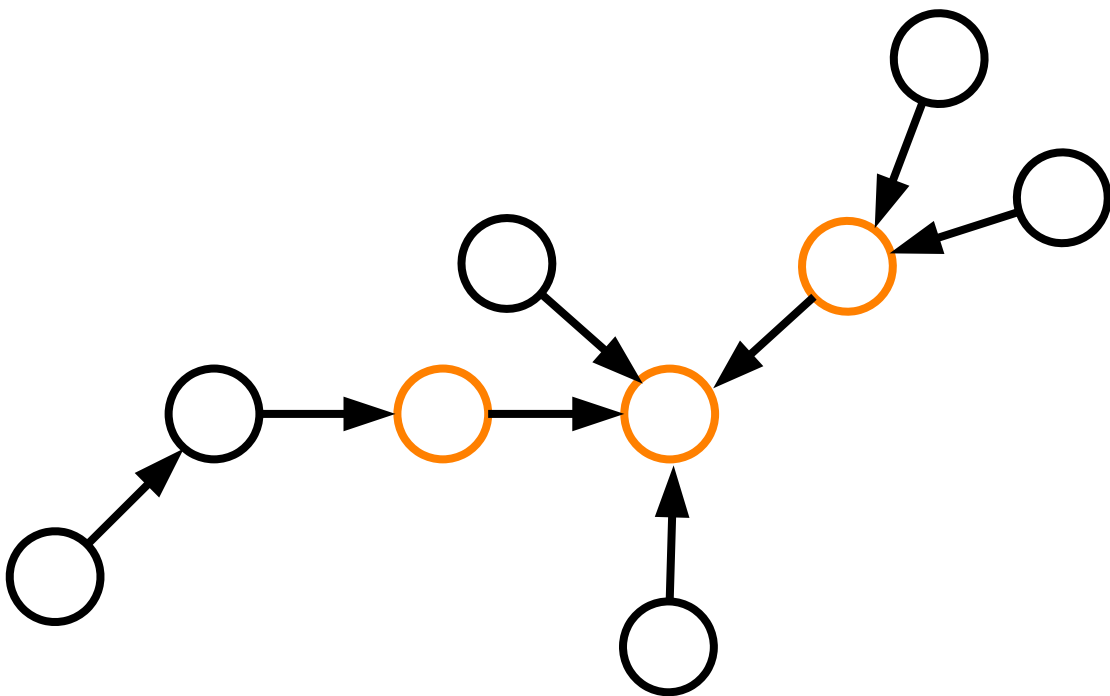


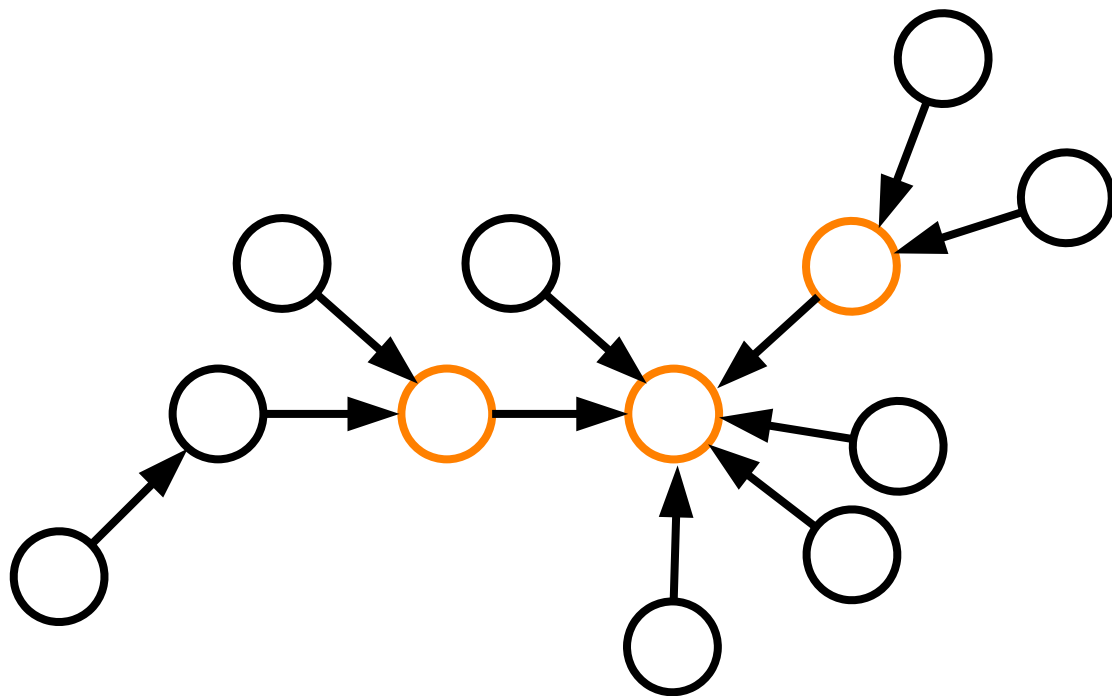


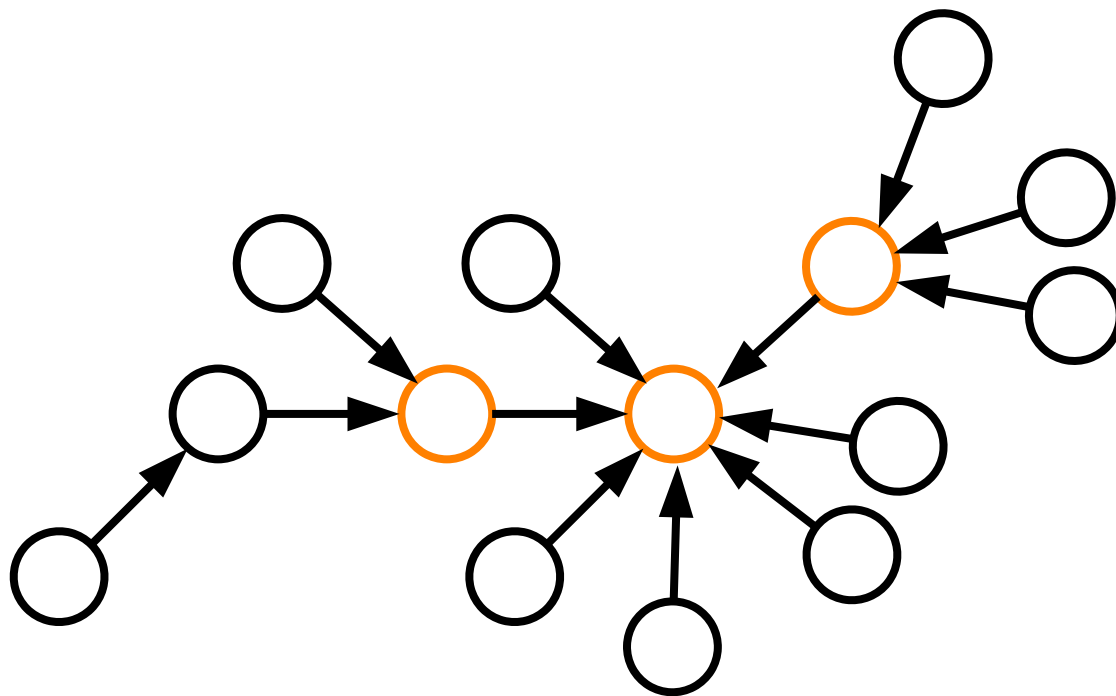




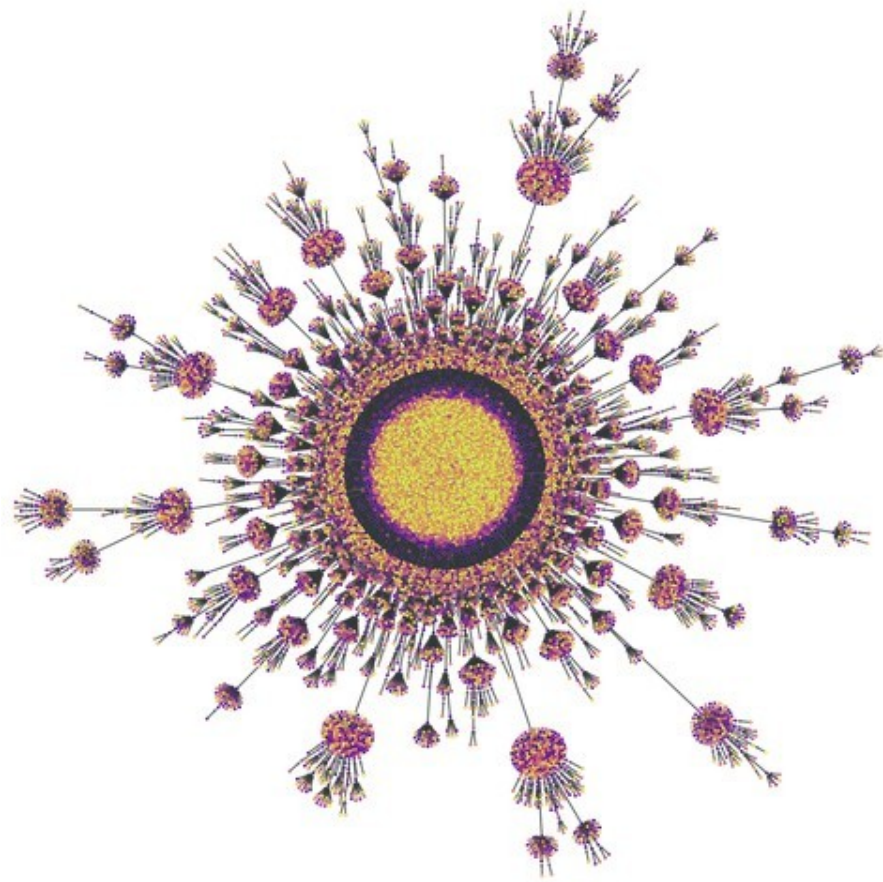
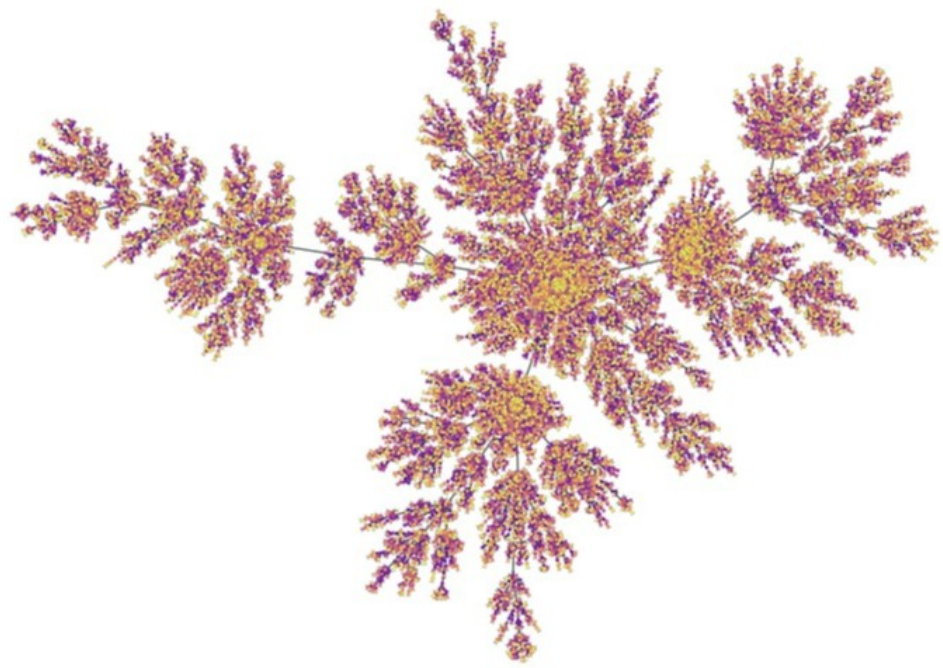








Nodes that join the network earlier have higher degree

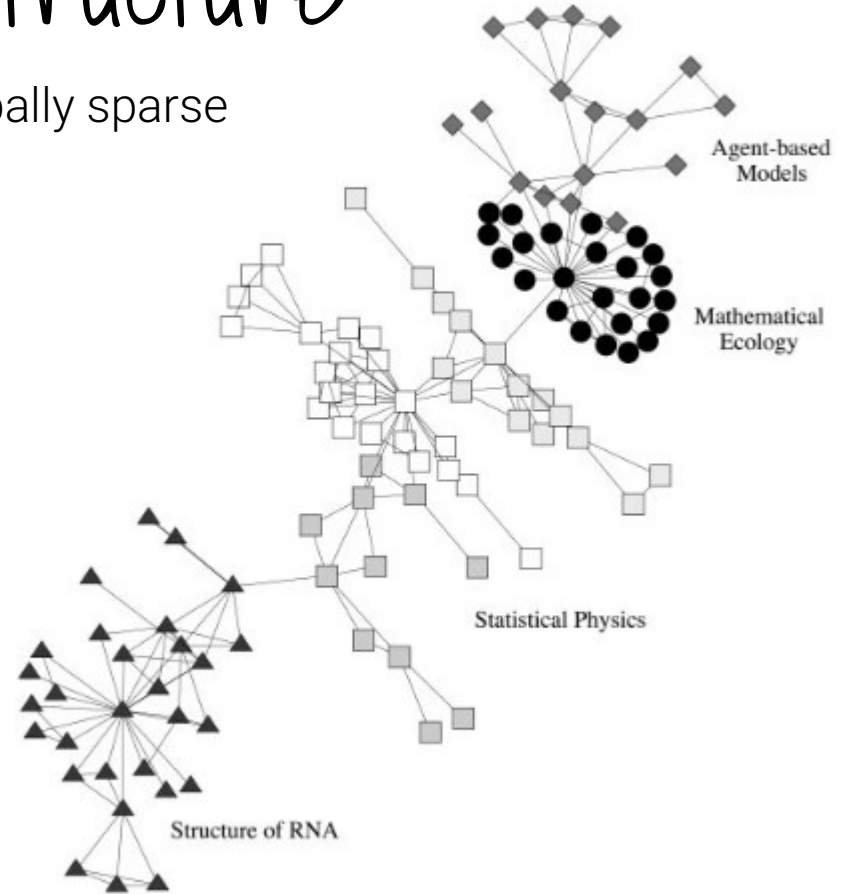
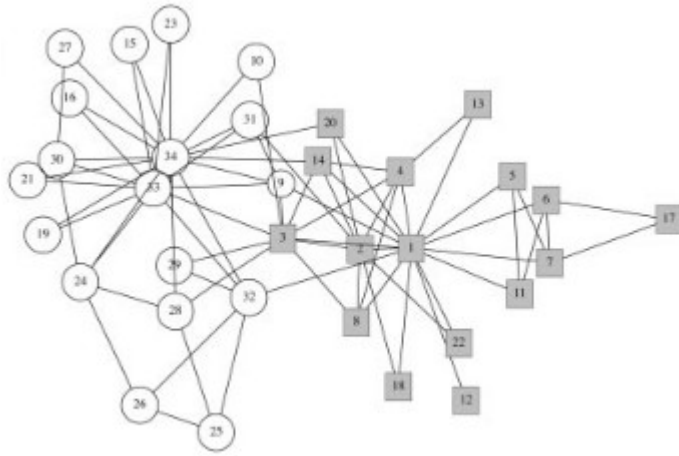


Community structure

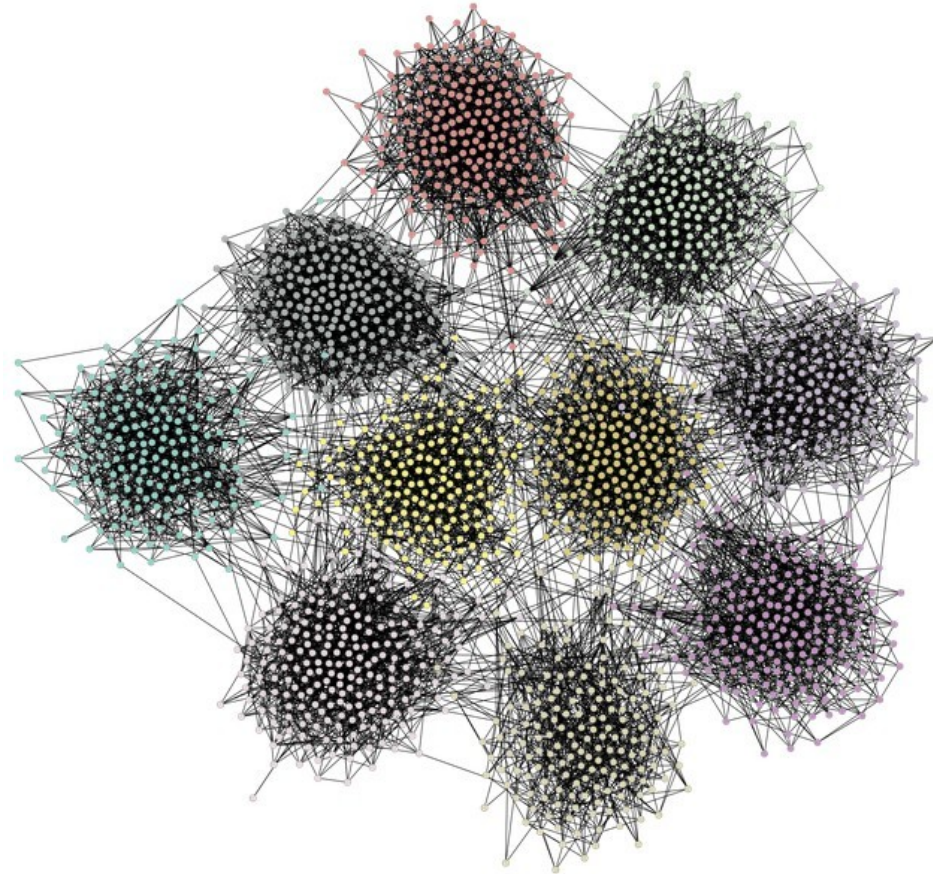
Locally dense, globally sparse

Community structure

Locally dense, globally sparse



Stochastic Block Models

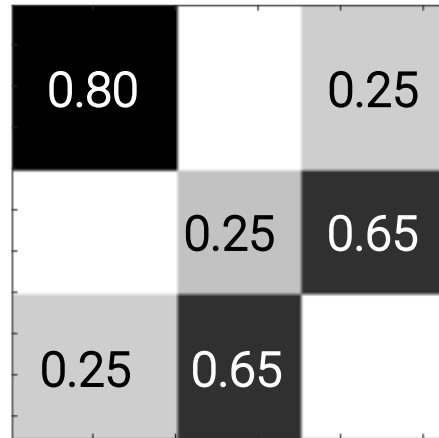


Generating a network using the SBM

Step 1 : Assign each node to a group

Generating a network using the SBM

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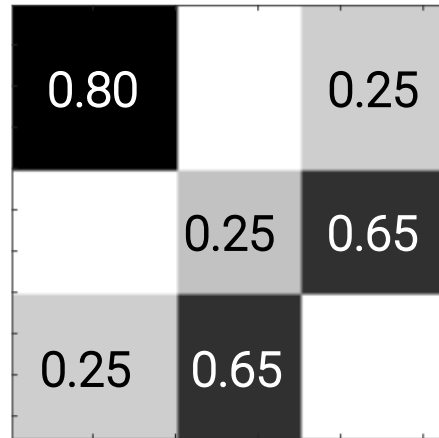


Mixing Matrix

- Step 2 : Select some connection probabilities (mixing matrix)

Generating a network using the SBM

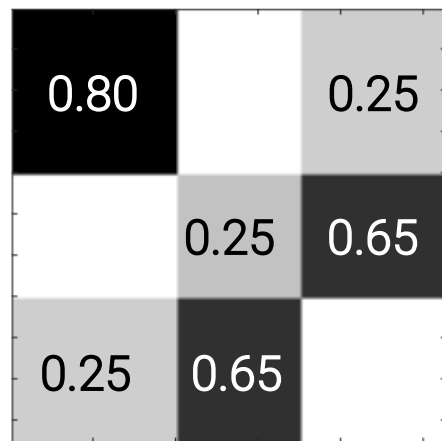
Step 1 : Assign each node to a group



Mixing Matrix

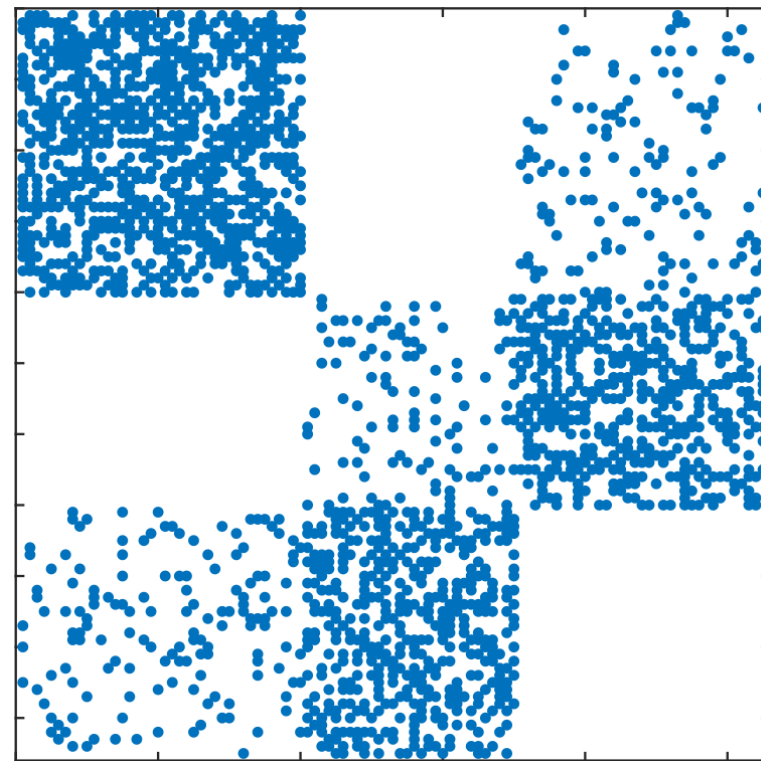

- *Step 2* : Select some connection probabilities (mixing matrix)
- *Step 3* : For each pair of nodes, add an edge with probability according to the group memberships

Generating a network using the SBM



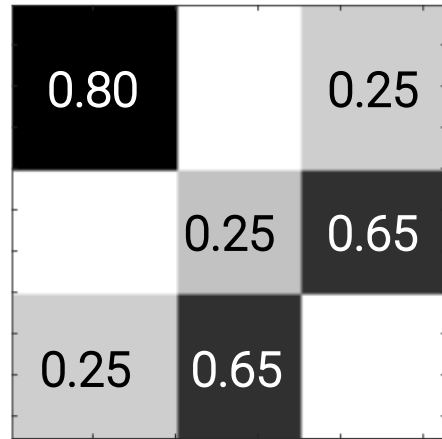
Mixing Matrix

generation



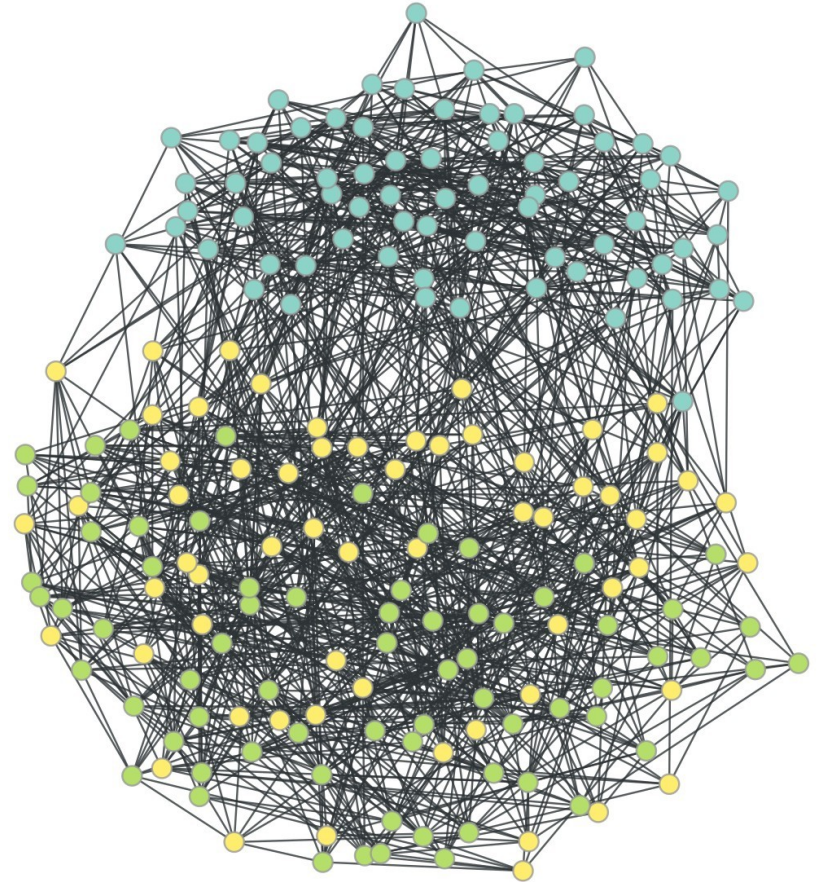
Adjacency Matrix

Generating a network using the SBM



Mixing Matrix

generation



Generating a network using the SBM

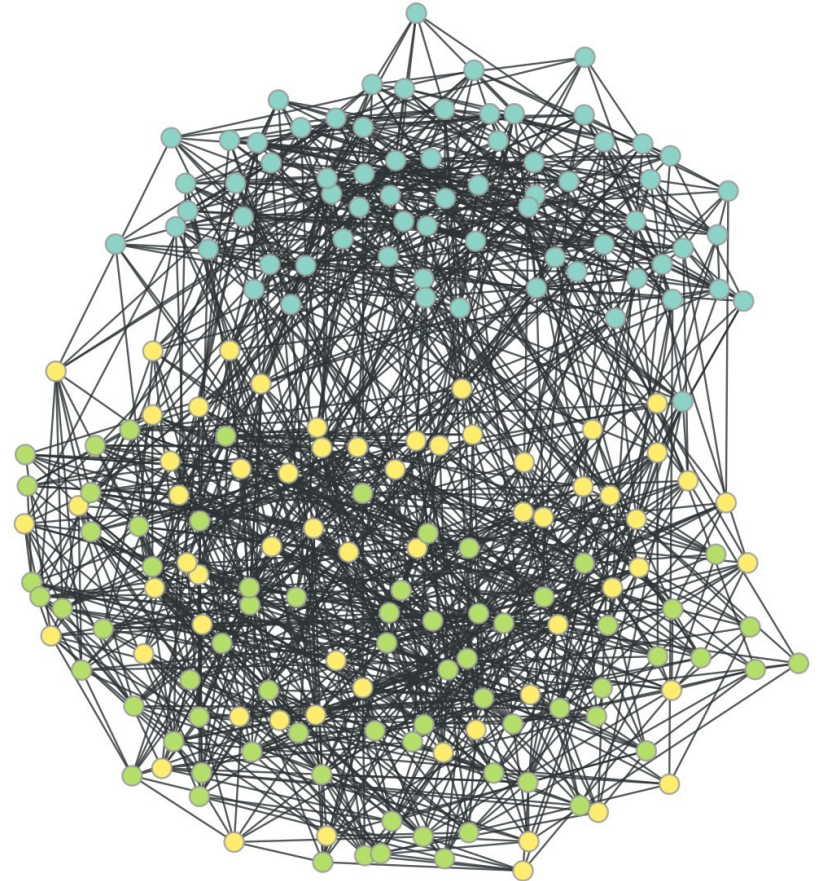
0.80		0.25
	0.25	0.65
0.25	0.65	

Mixing Matrix

generation

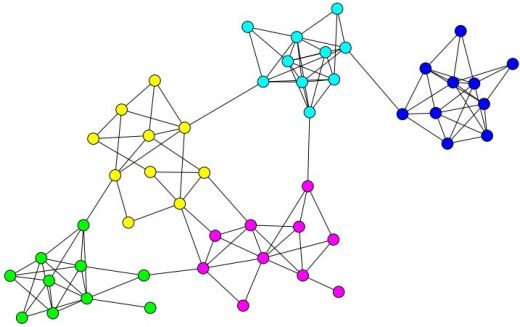
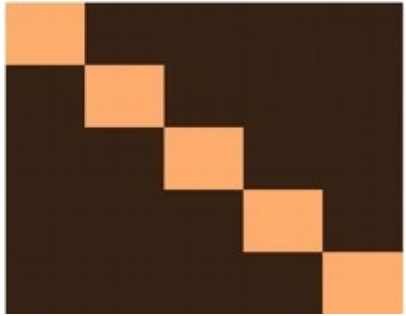


inference
(tomorrow)



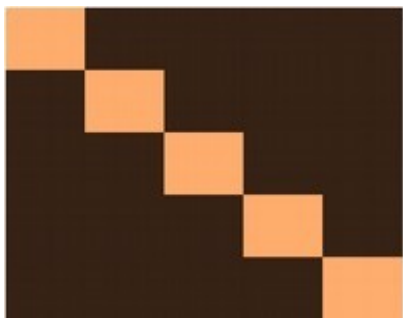
Different types of structure

assortative

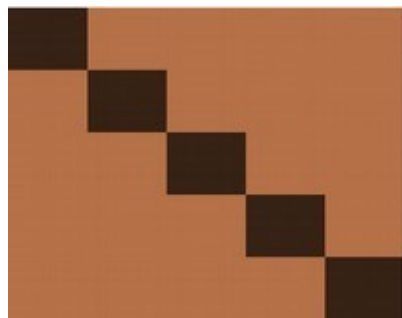


Different types of structure

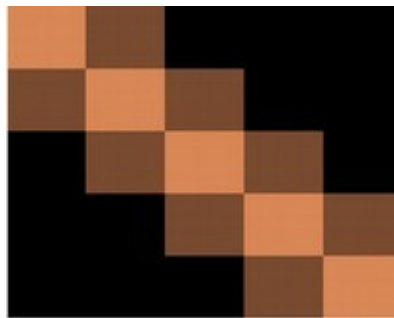
assortative



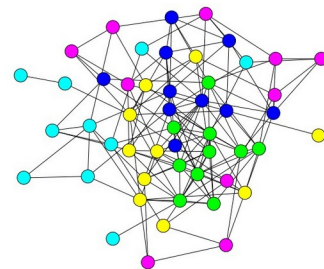
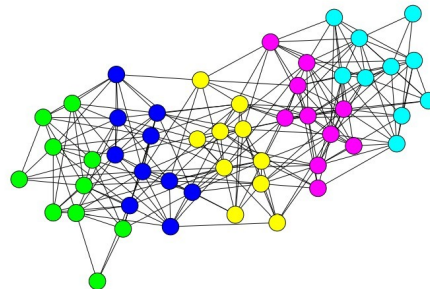
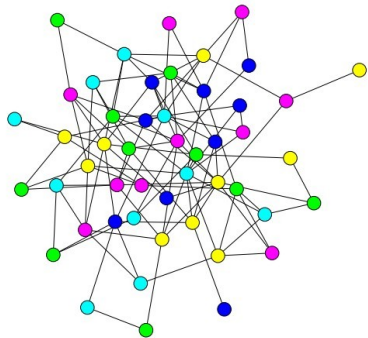
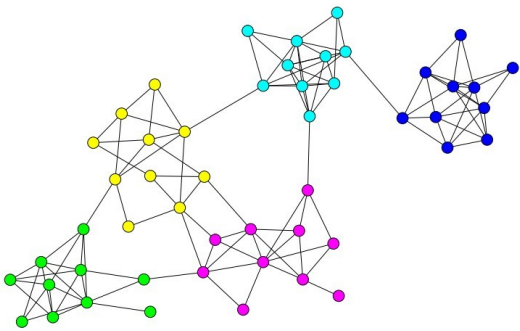
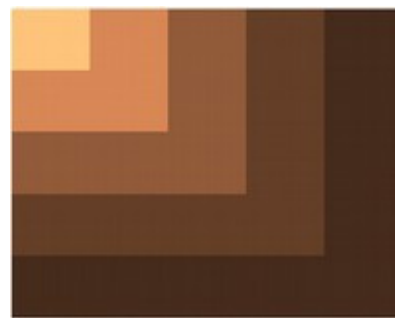
disassortative



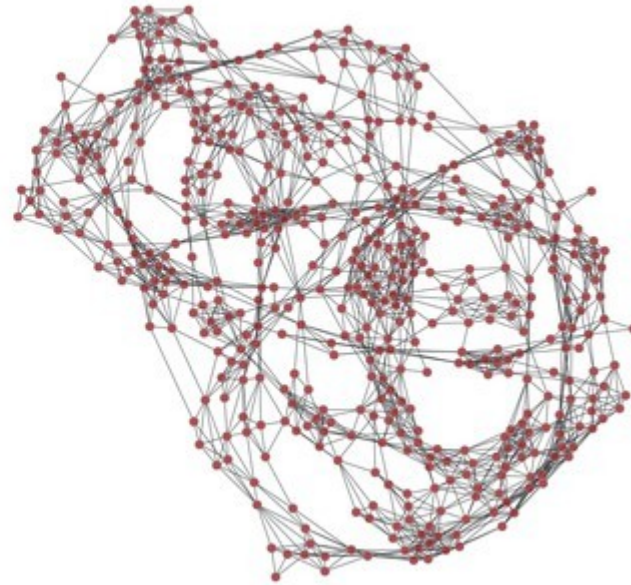
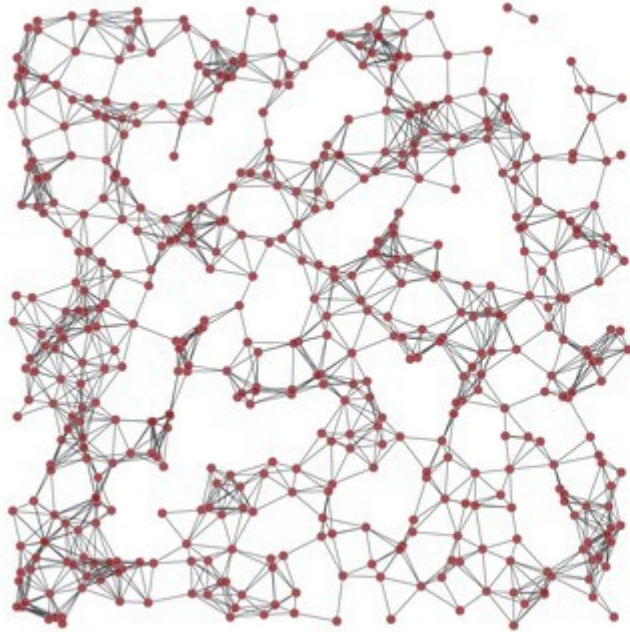
ordered



core-periphery



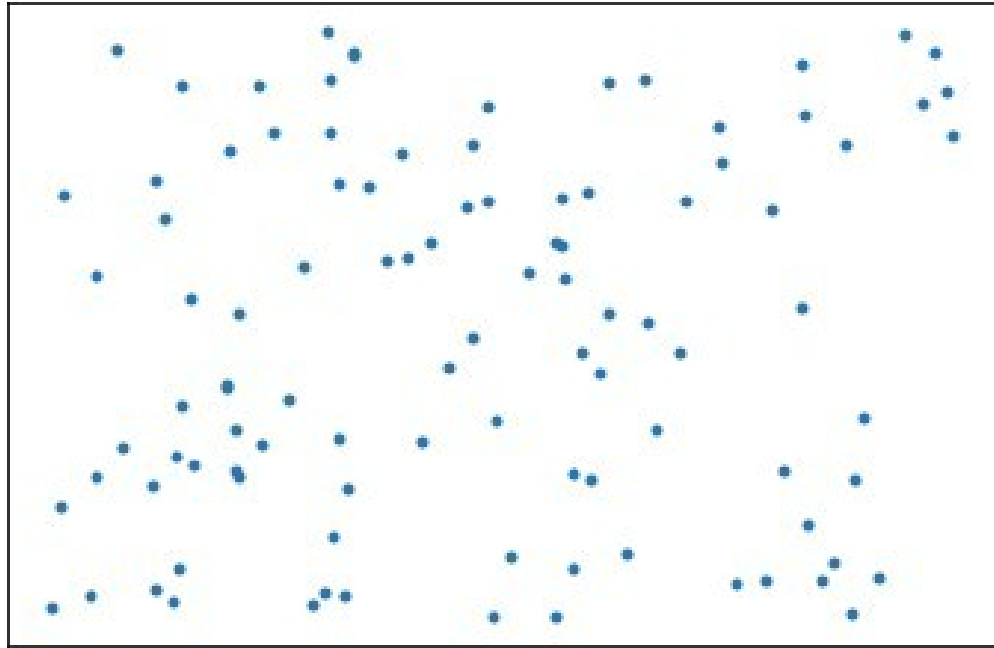
Geometric graphs



Generating a random geometric graph

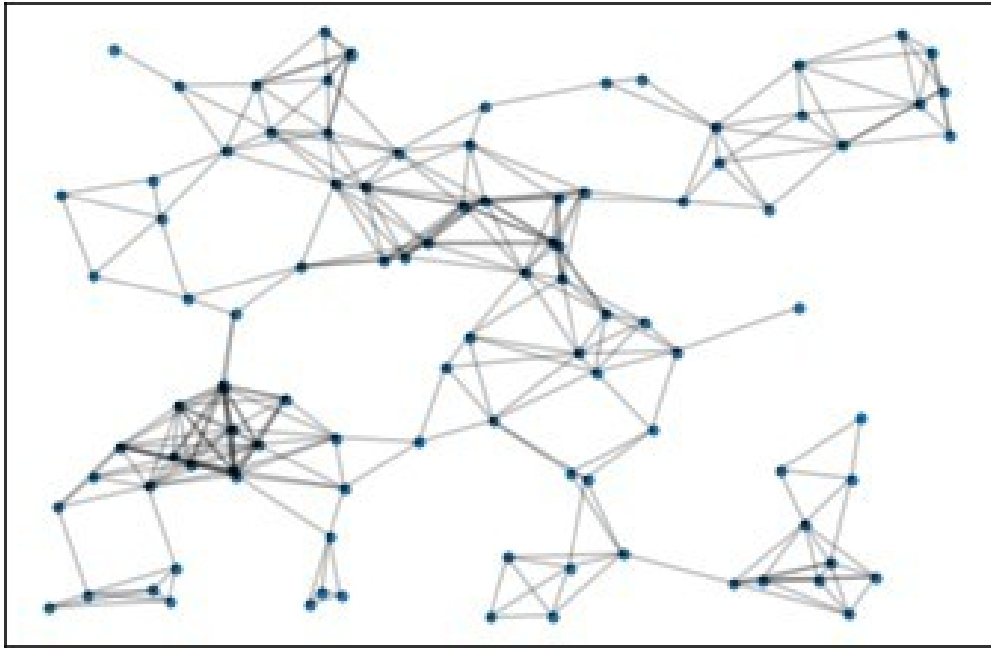


Generating a random geometric graph



Step 1 : Assign each node to a random position in a 2D space

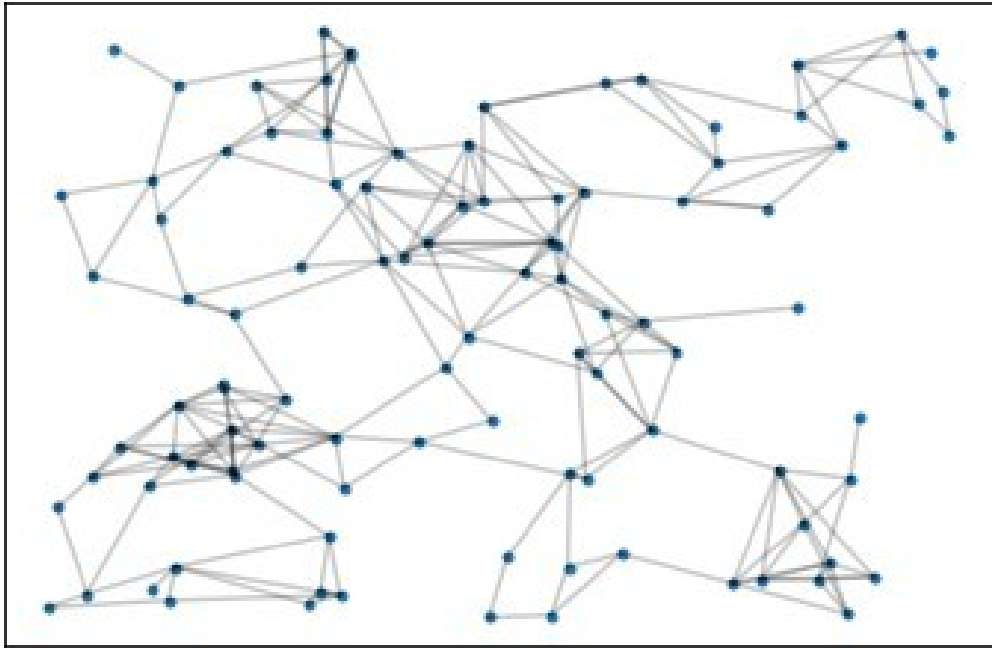
Generating a random geometric graph



Step 1 : Assign each node to a random position in a 2D space

Step 2 : Connect nodes if they are within a given radius, r

Latent space models

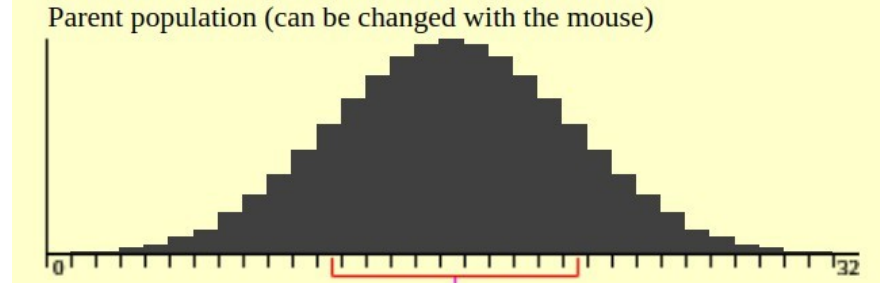


Similar to random
geometric graphs...

except edges are assigned
according to a probability as
a function of the distance

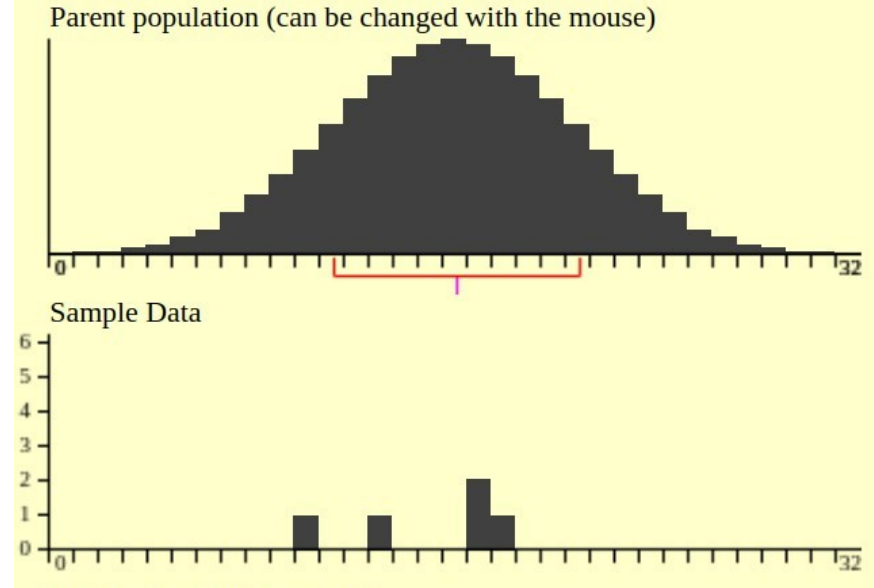
Null hypothesis testing

Population distribution



Mean = 16.00
Sd = 5.00

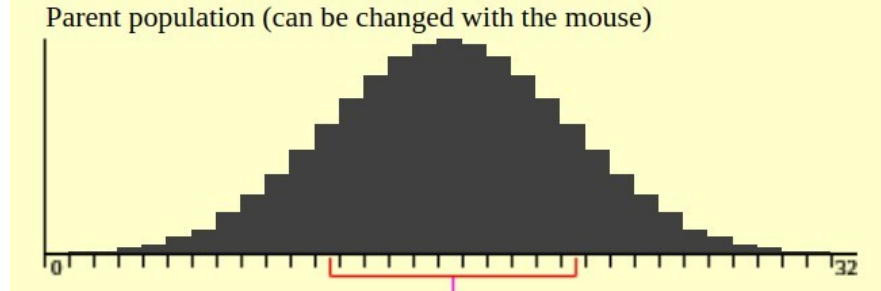
Population distribution



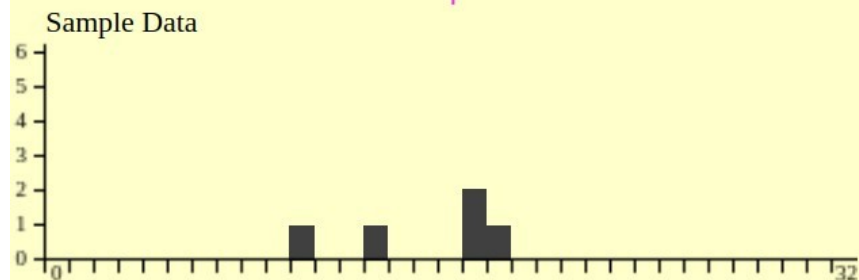
Mean = 16.00
Sd = 5.00

One sample
(size = 5)

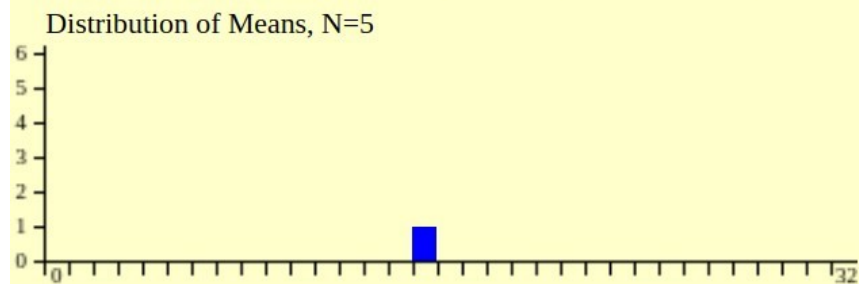
Population distribution



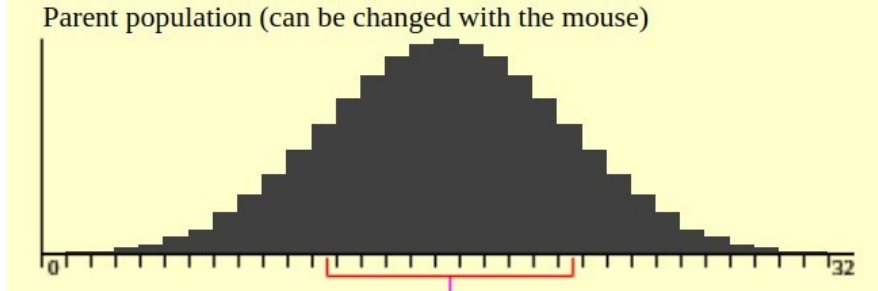
One sample
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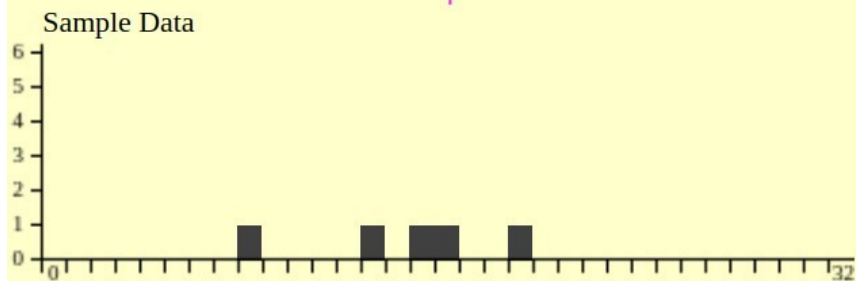
Distribution of means
(1 sample)



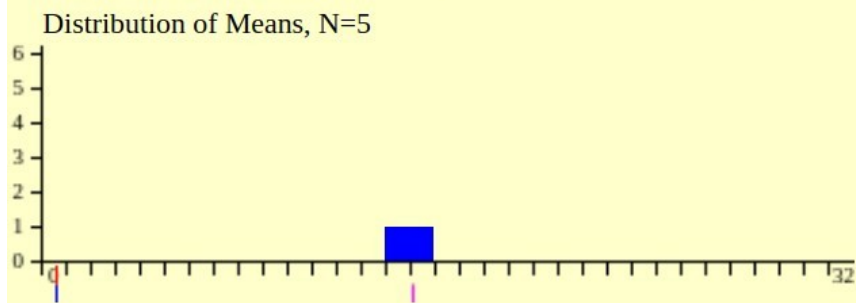
Population distribution



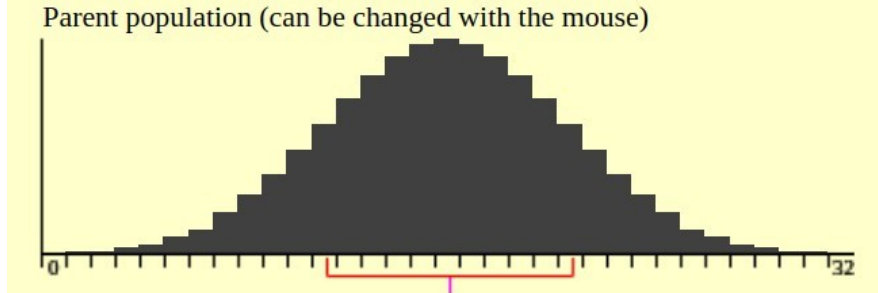
One sample
(size = 5)



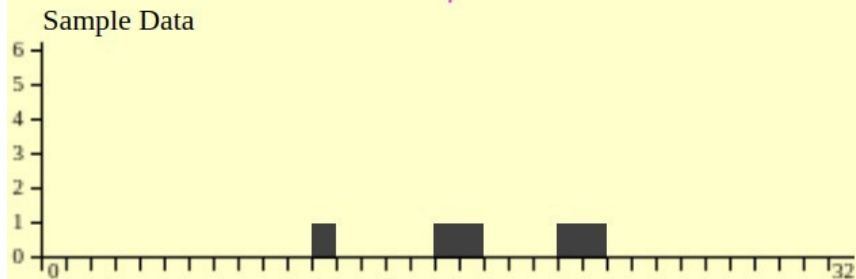
Distribution of means
(2 samples)



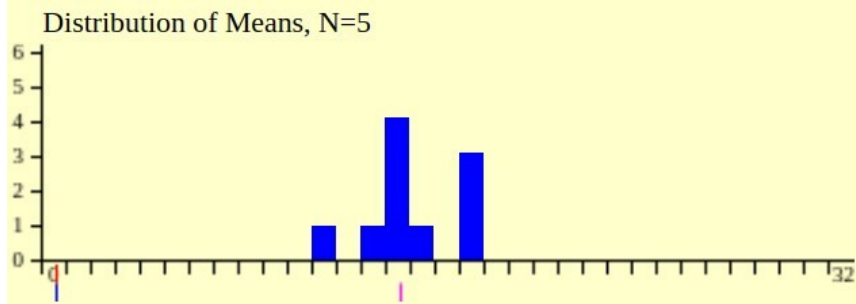
Population distribution



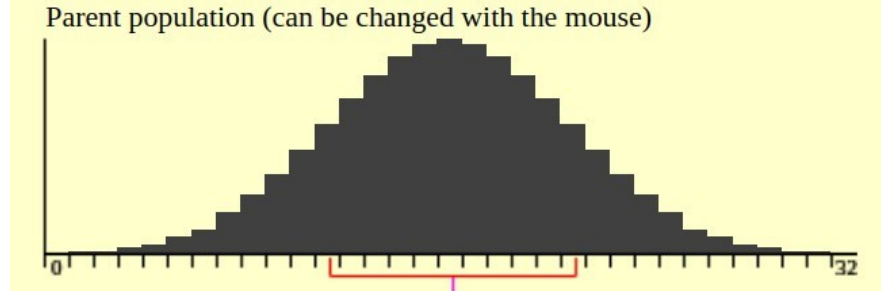
One sample
(size = 5)



Distribution of means
(10 samples)

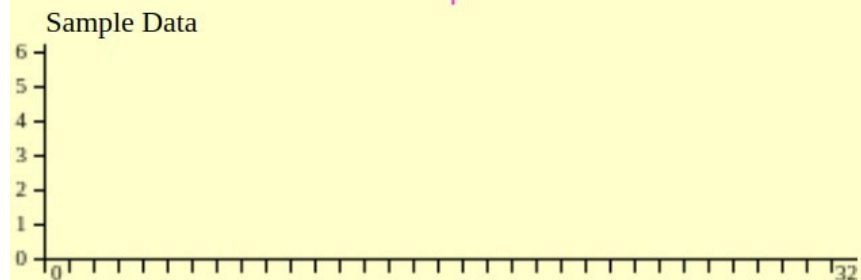


Population distribution

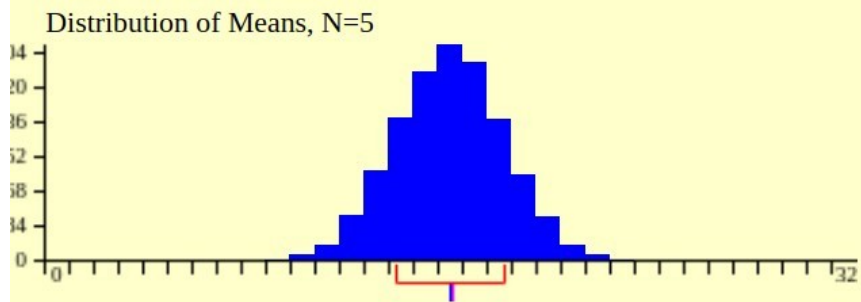


Mean = 16.00
Sd = 5.00

One sample
(size = 5)

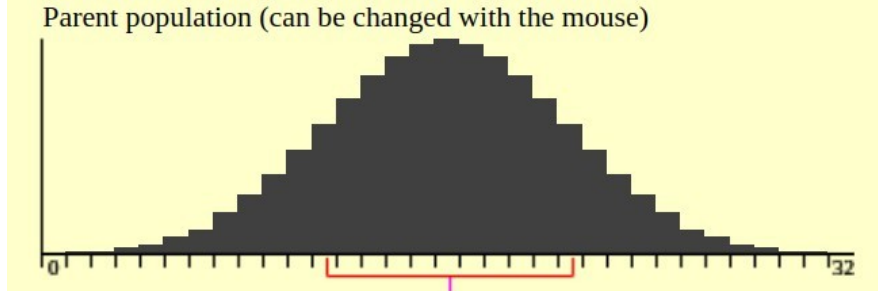


Distribution of means
(10,000 samples)



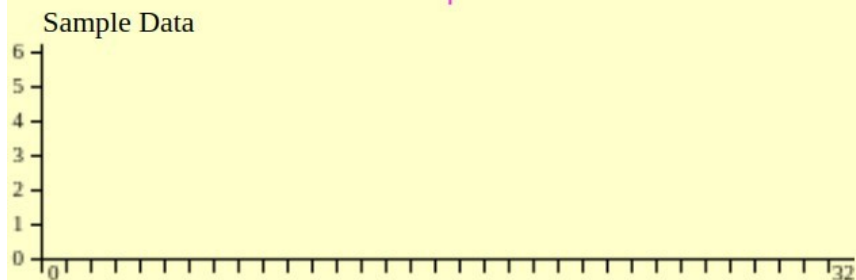
Mean = 15.99
Sd = 2.25

Population distribution

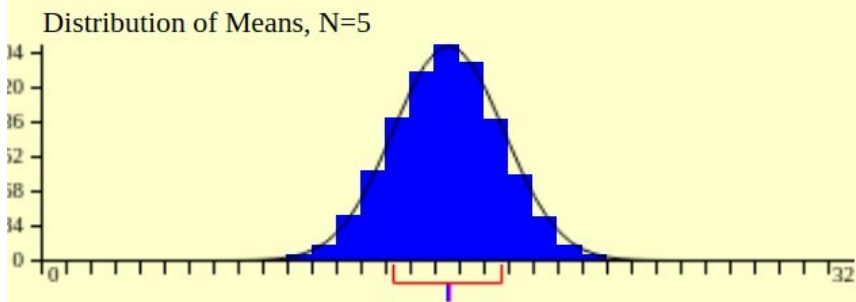


Mean = 16.00
Sd = 5.00

One sample
(size = 5)



Distribution of means
(10,000 samples)



Mean = 15.99
Sd = 2.25

Central limit theorem

The infamous P-value

P-VALUE

The probability, computed assuming that H_0 is true, that the test statistic would take a value as extreme or more extreme than that actually observed is called the ***P*-value** of the test. The smaller the *P*-value, the stronger the evidence against H_0 provided by the data.

The infamous P-value

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Definition, pg 405
Introduction to the Practice of Statistics, Fifth Edition
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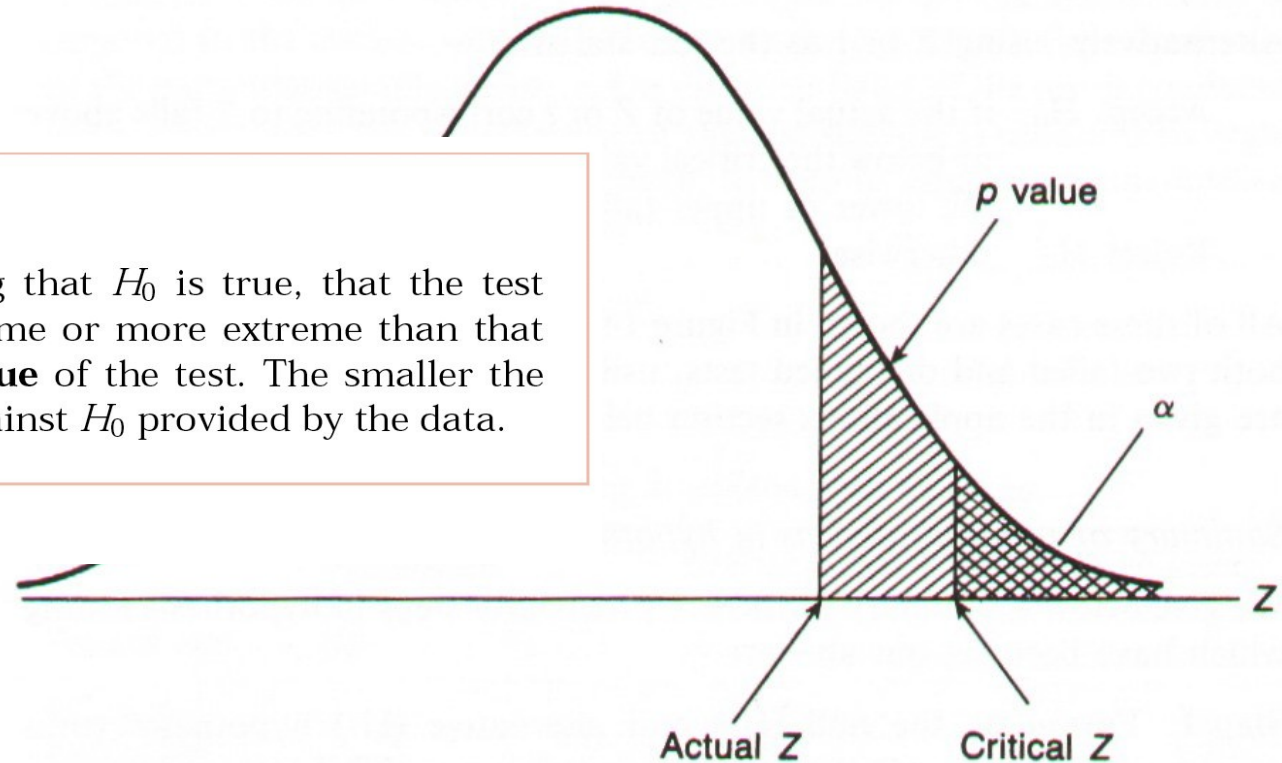


Figure 14.2 Comparison of p values and critical values of Z in a one-tailed test

The infamous P-value

P-VALUE

The probability, computed assuming that H_0 is true, that the test statistic would take a value as extreme or more extreme than that actually observed is called the **P-value** of the test. The smaller the P-value, the stronger the evidence against H_0 provided by the data.

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If $p\text{-value} < \alpha$,
then we reject the
null hypothesis

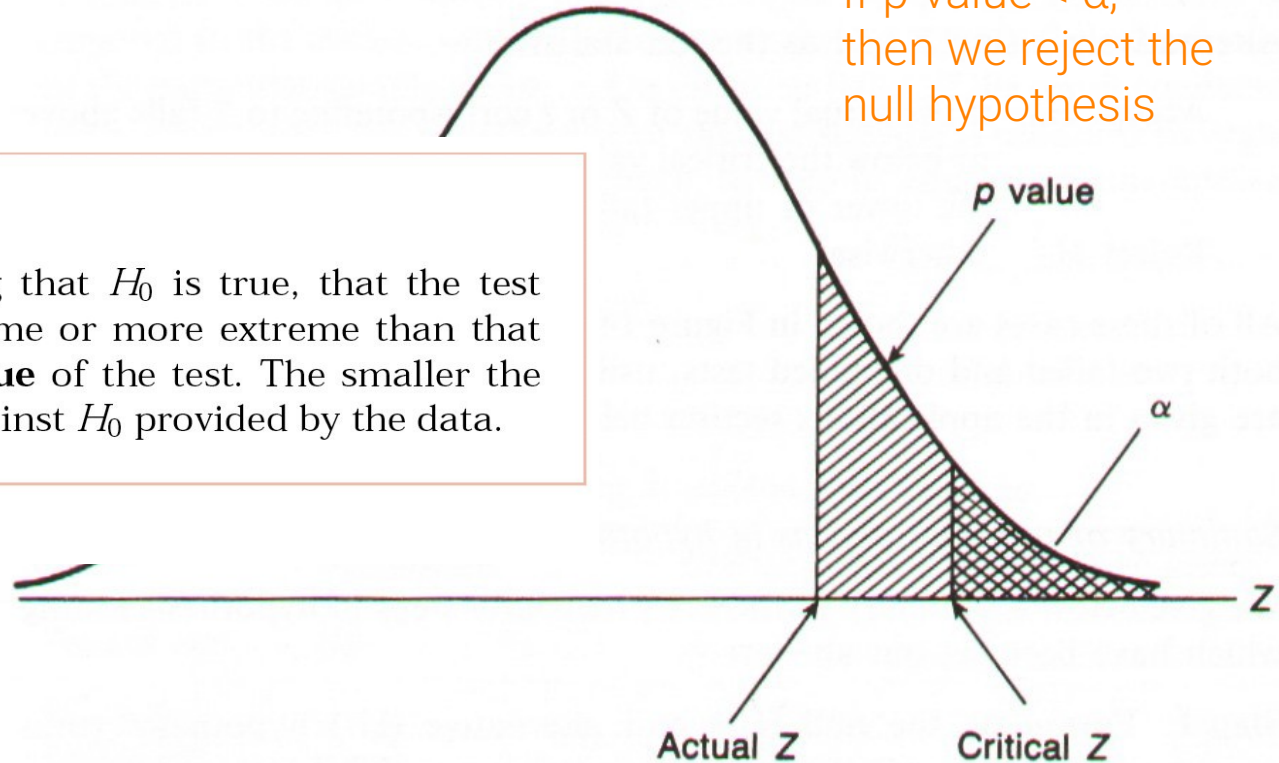
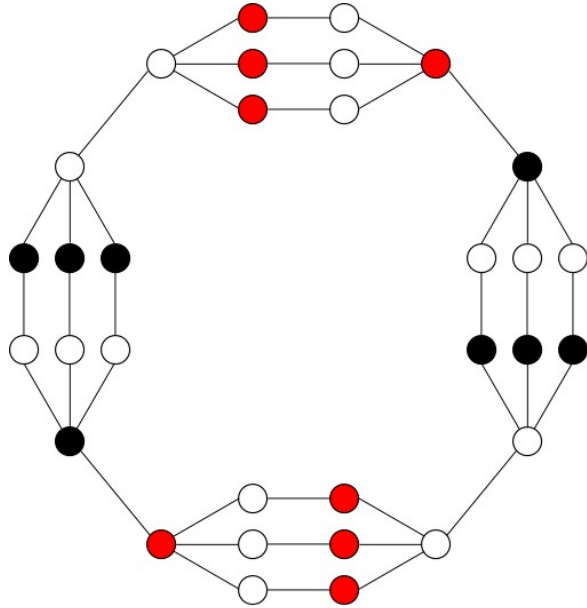


Figure 14.2 Comparison of p values and critical values of Z in a one-tailed test

For networks we can use graph models as a null model

- Does the network appear to be significantly different from a random graph?
- Can specific properties of the observed network (e.g., clustering coefficient) be explained by a particular generative process?
- Two approaches to create samples:
 - permute edges/nodes in a way that is consistent with the graph model
 - “fit” a model to an observed network and generate networks from it

Network nodes can have properties or attributes (metadata)



- Metadata (M) values
- Metadata (M) unknown

social networks *age, sex, ethnicity, race, etc.*

food webs *feeding mode, species body mass, etc.*

internet *data capacity, physical location, etc.*

protein interactions *molecular weight, association with cancer, etc.*

Blockmodel Entropy Significance Test

How well do the metadata explain the network?

Blockmodel Entropy Significance Test

How well do the metadata explain the network?

1. Divide the network G into groups according to metadata labels M .

Blockmodel Entropy Significance Test

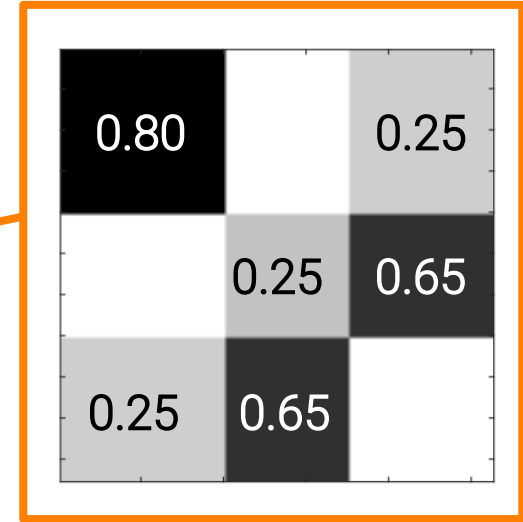
How well do the metadata explain the network?

1. Divide the network G into groups according to metadata labels M .
2. Fit the parameters of an SBM and compute the entropy $H(G, M)$

Blockmodel Entropy Significance Test

How well do the metadata explain the network?

1. Divide the network G into groups according to metadata labels M .
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Blockmodel Entropy Significance Test

How well do the metadata explain the network?

1. Divide the network G into groups according to metadata labels M .
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Test statistic



Blockmodel Entropy Significance Test

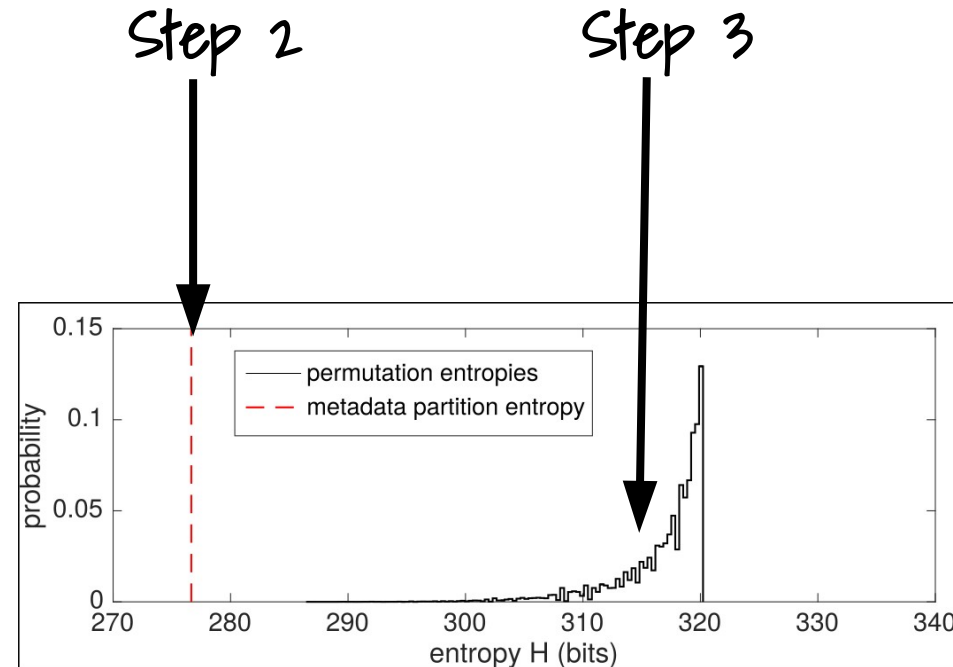
How well do the metadata explain the network?

1. Divide the network G into groups according to metadata labels M .
2. Fit the parameters of an SBM and compute the entropy $H(G, M)$
3. Compare this entropy to a distribution of entropies of networks partitioned using random permutations of the metadata labels.

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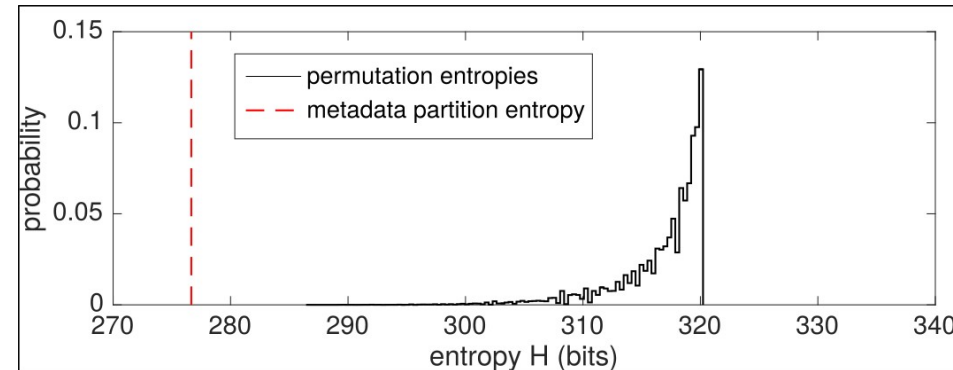
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metadata is randomly assigned
→ model gives no explanation, high H

metadata correlates with structure
→ model gives good explanation, low H



Multiple networks; multiple metadata attributes

Network	Status	Gender	Office	Practice	Law School
Friendship	$< 10^{-6}$	0.034	$< 10^{-6}$	0.033	0.134
Cowork	$< 10^{-3}$	0.094	$< 10^{-6}$	$< 10^{-6}$	0.922
Advice	$< 10^{-6}$	0.010	$< 10^{-6}$	$< 10^{-6}$	0.205

Multiple sets of metadata provide a significant explanation for multiple networks.

Summary...

We can use graph models to simulate observed properties and form hypotheses

Null hypothesis tests are a popular way to test these hypotheses

~~Note~~: we can only accept/reject the null hypothesis

(more on this tomorrow)