

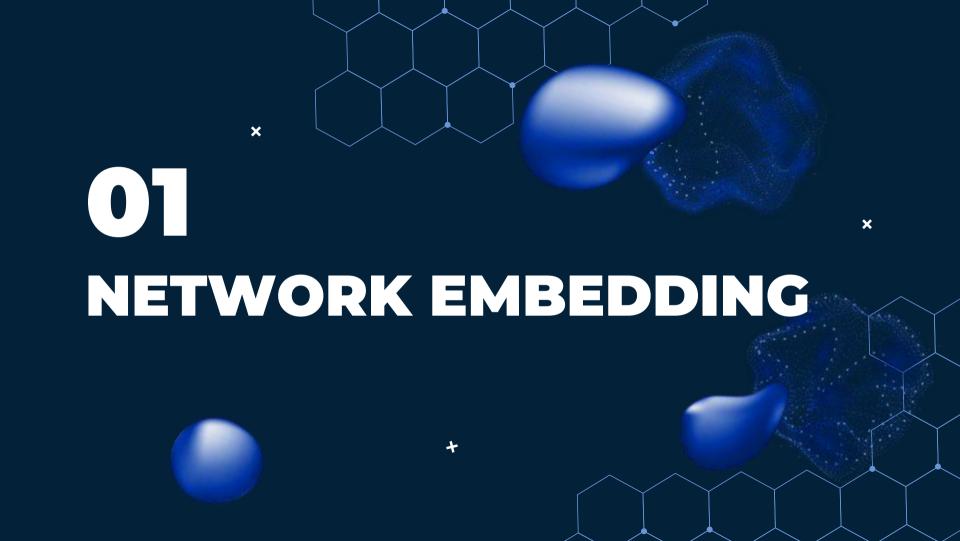
Michigan State University

# **TABLE OF CONTENTS**

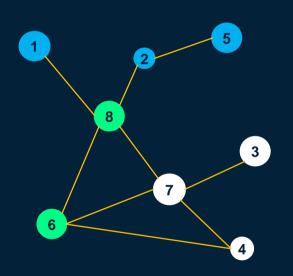
- **01** NETWORK EMBEDDING
- **02** DEEPWALK

X

- 03 NODE2VEC
- **04** EXPERIMENTS AND EVALUATION
- **05** FUTURE DIRECTIONS



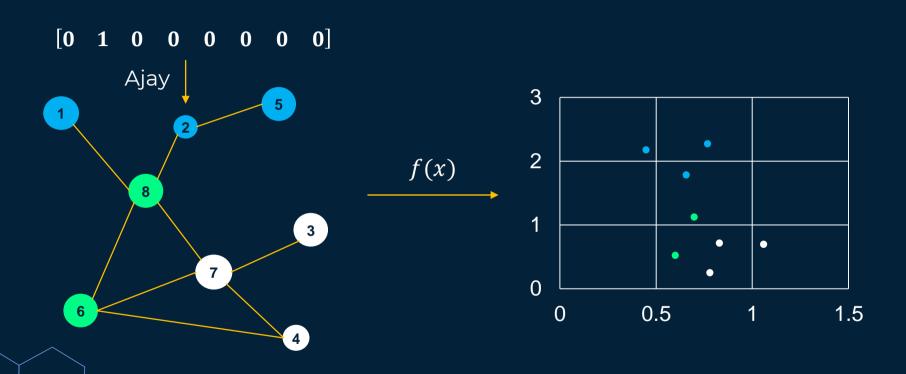
# **Understanding Graphs for Representation Learning**



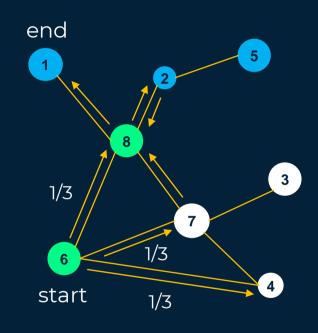
- Visualizes relationships between the data
- Consists of:
  - 1. Nodes: capture data as vectors
  - 2. Edges: connect related nodes
- Representation Learning extracts hidden features from the graph for complex analysis.

e.g. link prediction, node classification, community detection.

# **Understanding Graphs for Representation Learning**



## **Analyzing Graphs with Random Walk**



Steps = 5

- Choose starting point and steps
- Determine probabilities based on neighbor nodes
- Collect data from walks

**7** 

8

2

8

2

# SkipGrams and their relation to DeepWalk

- NLP model: words in sentence context
- Applied on the paths to maximize the probability of observing a node's neighborhood
- Nodes with similar neighborhood share similar embeddings
- The objective function of DeepWalk is the following cross entropy:

$$min_y - logP(\{v_{i-w}, ..., v_{i-1}, v_{i+1}, ..., v_{i+w}\}|y_i)$$



W is the window size which restricts the size the random walk context

#### The meaning and transformation of the SkipGram formula

Looking back at the previous formula, SkipGram removes the ordering constraint

In the end, it will be transformed into:

$$min_y - log \sum -w \le j \le w P(v_{i+j}|y_i)$$

Conditional probability  $P(v_{i+j}|y_i)$  defined using the softmax function:

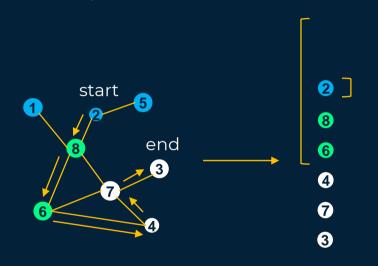
$$P(v_{i+j}|y_{j}) = \frac{\exp(y_{i+j}^{T}y_{i})}{\sum_{k=1}^{|v|} \exp(y_{k}^{T}y_{i})}$$



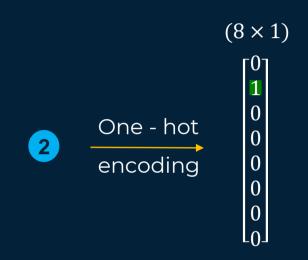


# **How DeepWalk Works**

Step 1: Random Walk



Step 2: One-hot encoding

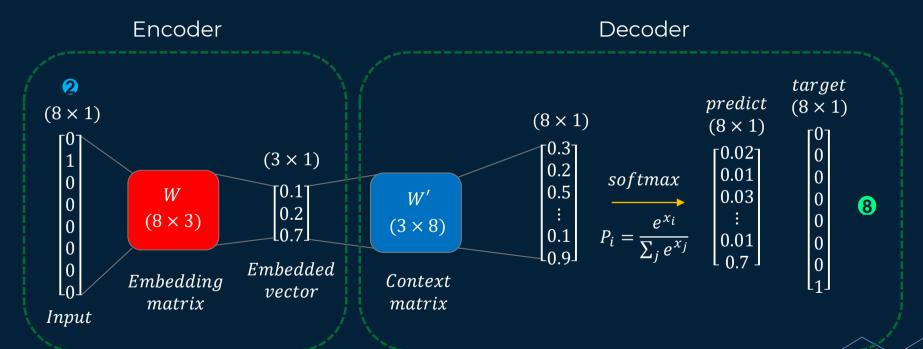


one – hot encoded vector



#### **How DeepWalk Works**

Step 3: Implement Skip-gram model







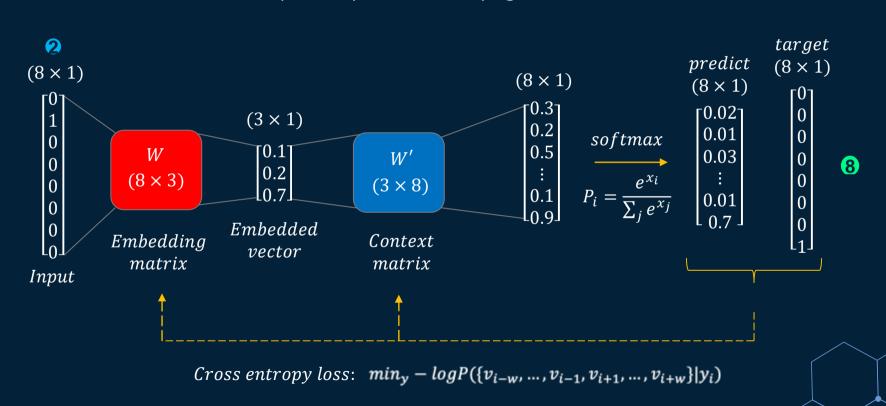






## **How DeepWalk Works**

Step 3: Implement Skip-gram model



# Why SkipGrams for DeepWalk?

#### **SkipGrams**

- Application: Texts
- Context: Relationships of words in a sentence

The quick brown fox jumps over the lazy dog

#### DeepWalk

- Application: Graphs
- Context: Relationships of nodes in Random Walks

Teacher A – Student B Student B – Student C

#### Similarities:

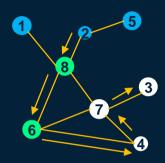
DeepWalk adapts SkipGram from natural language processing to graph data.



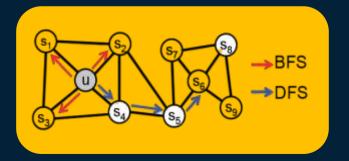
#### **How Node2Vec Works**

- Node2Vec is almost similar to DeepWalk
- Both methods use:
  - 1. Random walks
  - 2. Skipgram model
- Node2Vec have different walk algorithm to collect nodes
- Node2Vec either explores data in wide range or far range

DeepWalk



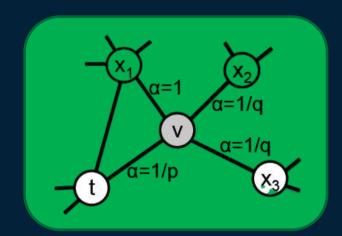
Node2Vec



#### **How Node2Vec Works**

- Current walk: t → v
- Determine probability of v → x
- Parameter  $\alpha_{pq}(t,x)$  is introduced, where:

$$\alpha_{pq}(t,x) = \begin{cases} \frac{1}{p} & \text{if } d_{tx} = 0\\ 1 & \text{if } d_{tx} = 1\\ \frac{1}{q} & \text{if } d_{tx} = 2 \end{cases}$$



- p controls likelihood of revisiting a node
- ullet q controls likelihood of walking further or locally

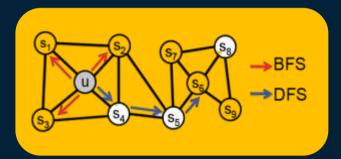


## DeepWalk & Node2Vec

# DeepWalk 1 3 7

- Explores data randomly
- Low computational cost
- Ignores important data

#### Node2Vec

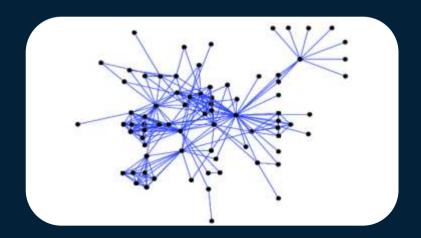


- Flexibility
- Higher computational cost
- Analyze relationships better

## Effect of p and q in Node2Vec

Step 1: Prepare dataset (Les Misérable Network)

- Contains 77 nodes, 254 edges
- Nodes: characters from Les Misérable novel
- Edges: relationships between characters



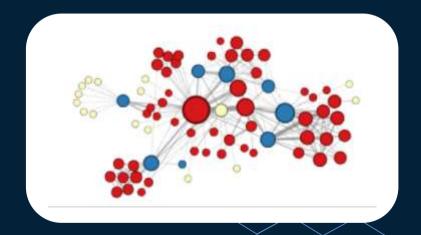


# Effect of p and q in Node2Vec

Step 2: Adjust different p and q

Set 
$$p = 1, q = 0.5$$

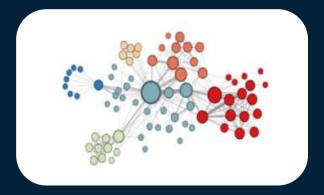
Set 
$$p = 1, q = 2$$



# Effect of p and q in Node2Vec

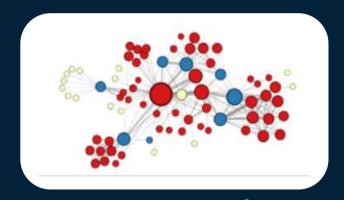
Step 3: Analyze results

Set 
$$p = 1, q = 0.5$$



 $p \gg q$ : deep exploration Homophily community

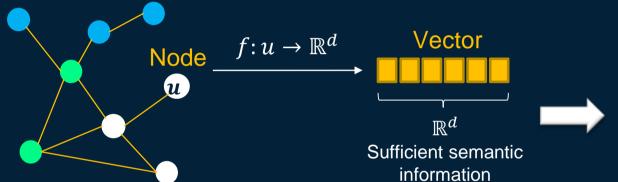
Set 
$$p = 1, q = 2$$



 $p \ll q$ : broad exploration Structural equivalence



# **Experiments**

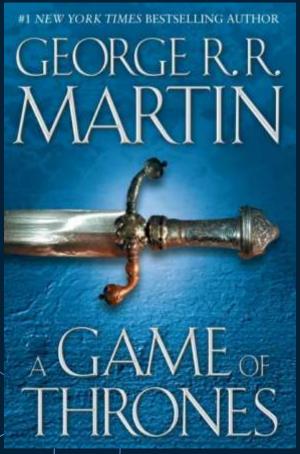


#### **Downstream Tasks**

- Node classification
- Link prediction
- Graph classification
- Anomalous node detection
- Clustering Homophily community
- structural equivalence



## **Game of Thrones**







#### From Book to Network



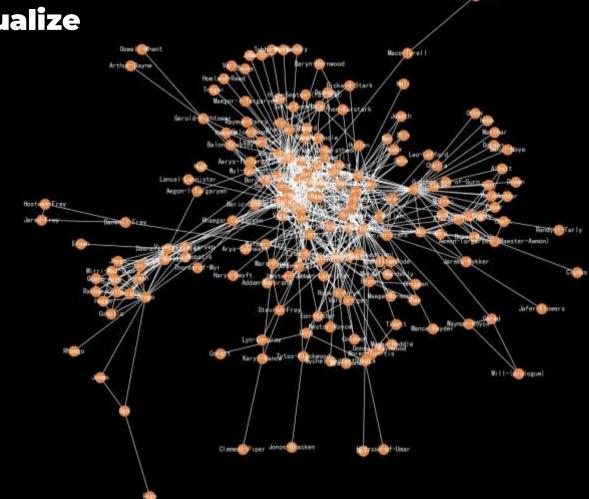
#### **Data**

Source	Target	weight
Addam-Marbrand	Jaime-Lannister	3
Addam-Marbrand	Tywin-Lannister	6
Aegon-I-Targaryen	Daenerys-Targaryen	5
Aegon-I-Targaryen	Eddard-Stark	4

Link two characters each time their names appear within 15 words.

Number of nodes: 187 Number of edges: 684 **Pre-processing-visualize** 

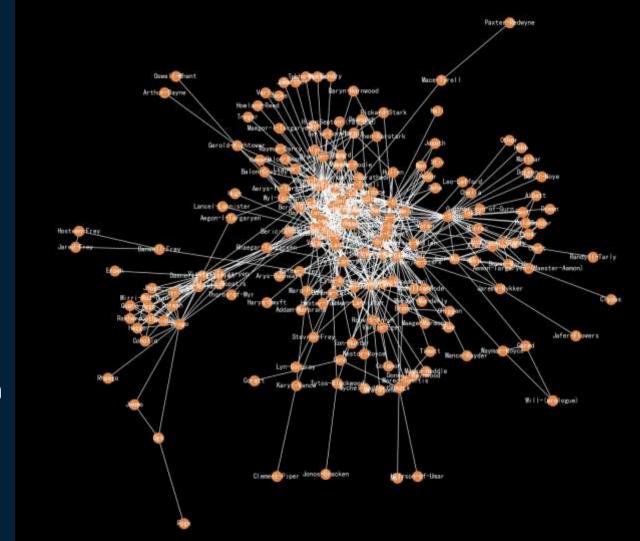
Import unweighted graph
nx.spring\_layout(G)



# **Processing-DFS**

Node2Vec (G, dimensions=32, p=5, q=0.5, walk\_length=10, num\_walks=600, workers=4)

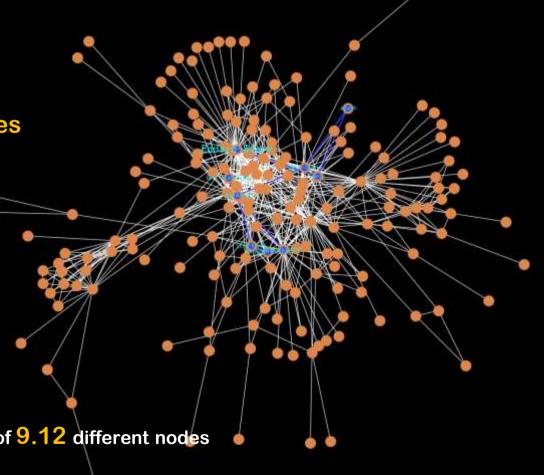
DFS depth-first search





DFS depth-first search, find homogeneous communities

array([-0.36179334, -0.11370167, ..., 0.05198546], dtype=float32)

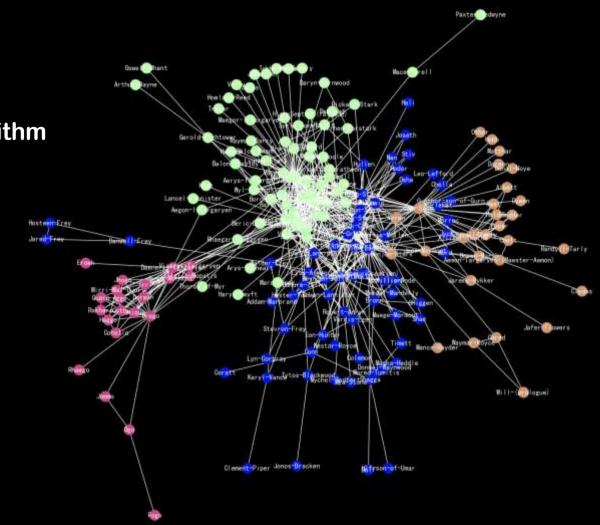


passes through an average of 9.12 different nodes

# **Processing**

Kmeans clustering algorithm

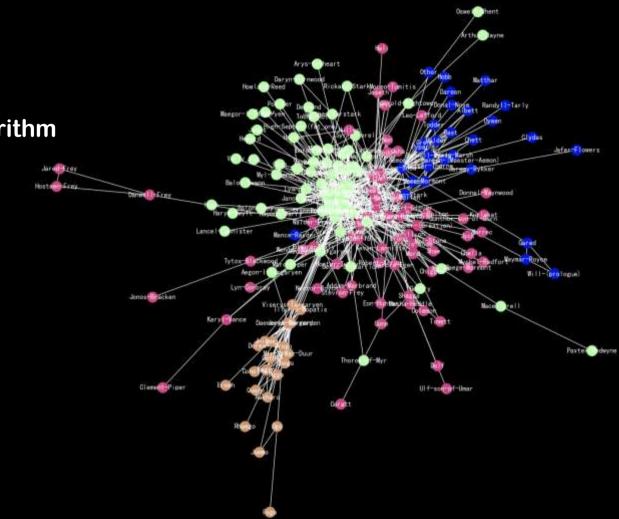
n\_clusters=4



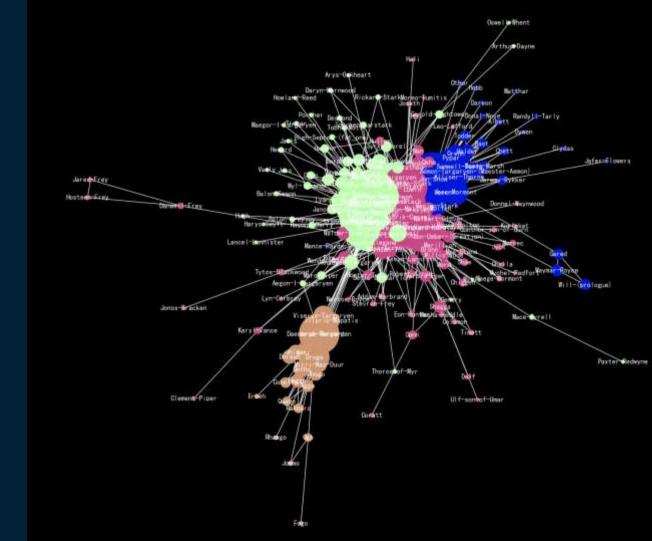
Kmeans clustering algorithm

n\_clusters=4

Consider weights between nodes

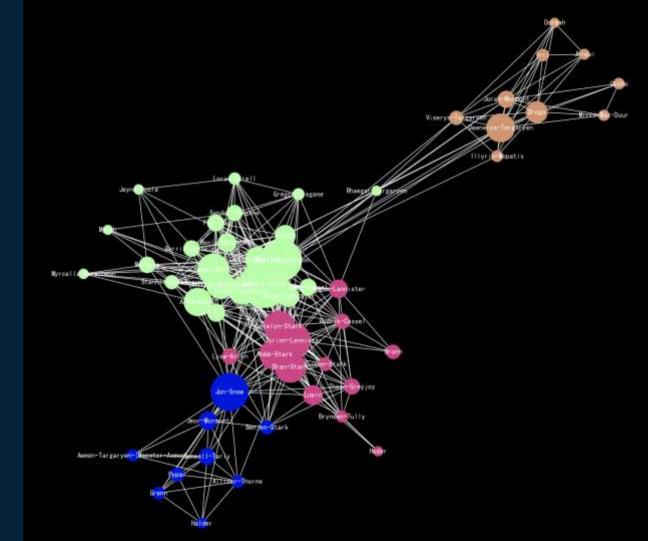


visualize the weights



visualize the weights

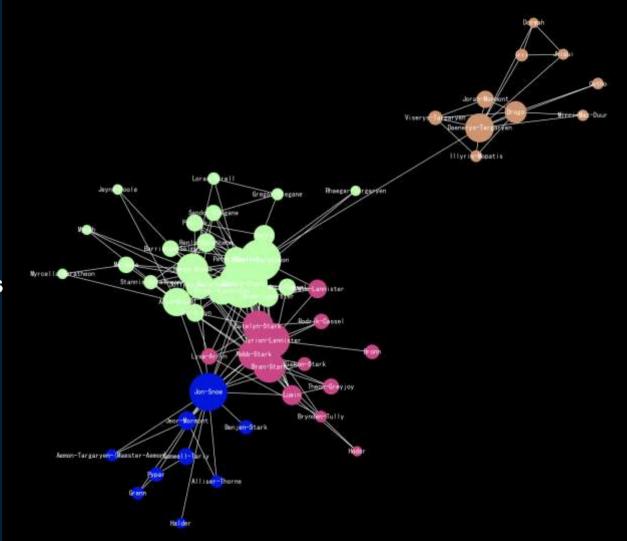
Get rid of low-weights characters

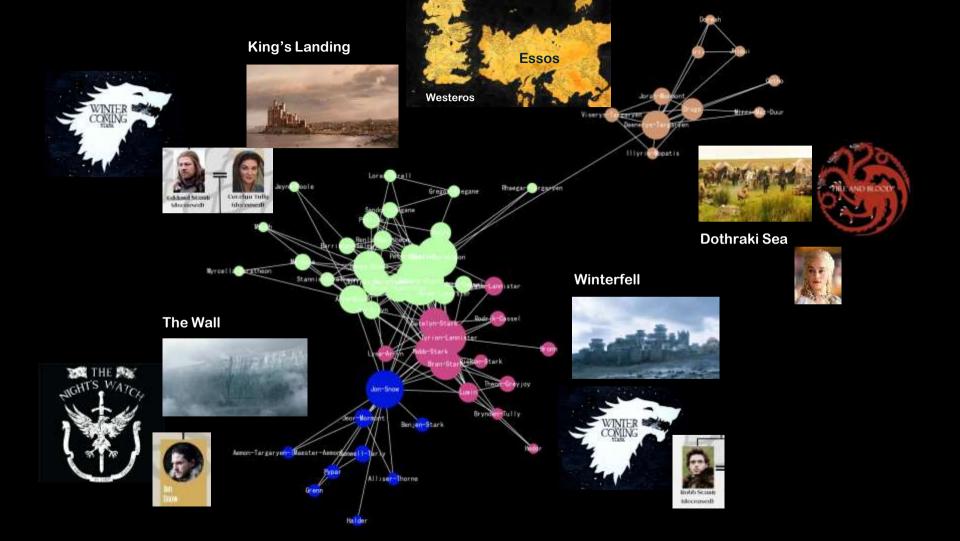


visualize the weights

Get rid of low-weights characters

Cut out some of the less frequent connections





#### Test

#### # Find similar nodes of Jon-Snow node

>>>model.wv.most\_vector('Jon-Snow')

```
<<<[( 'Dareon', 0.7889388203620911),
     ('Donal-Noye', 0.7577574849128723),# Night's Watch the blacksmith
     ('Matthar', 0.7506797313690186),
     ('Othor', 0.740899384021759),
     ('Dywen', 0.73787921667099),
     ('Rast', 0.7340330481529236),
     ('Hobb', 0.7294636368751526),
     ('Grenn', 0.7205644249916077),
     ('Albett', 0.7146326303482056),
     ('Todder', 0.706096887588501)]
```

# Same group of recruits as Jon Snow. # Same group of recruits as Jon Snow.

# Ranger of the Night's Watch



Comrades

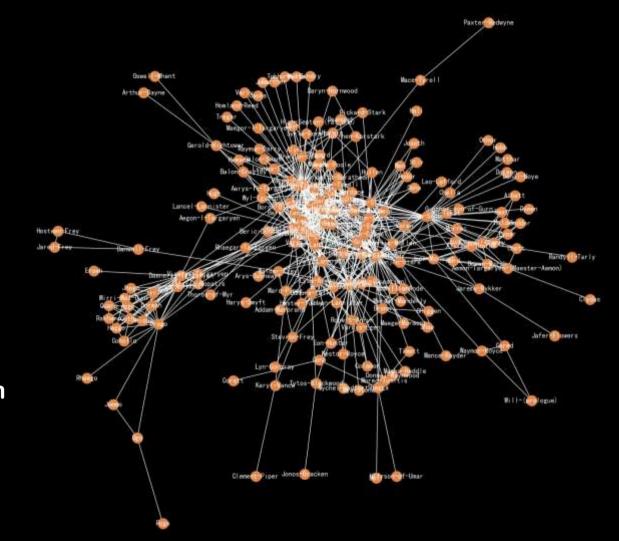


homogeneous communities

# **Processing-BFS**

Node2Vec (G, dimensions=32, p=0.1, q=100, walk\_length=10, num\_walks=600, workers=4)

BFS breadth-first search



# **Processing-BFS**

BFS breadth-first search, find structural equivalence

array([ 0.31407943, 0.24742755, ..., -0.5104667], dtype=float32)

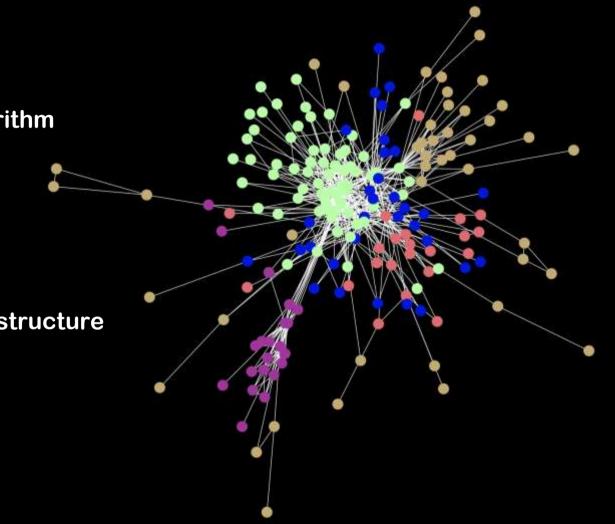
passes through an average of 4.41 different nodes

Kmeans clustering algorithm

n\_clusters=6

If we consider weights between nodes

Nodes are classified by structure



#### **Test**

#### # Find similar nodes of Jon-Snow node

>>>model.wv.most\_vector('Jon-Snow')

<<<[('Alliser-Thorne', 0.6657752394676208), # master-at-arms at Castle Black ('Bowen-Marsh', 0.6607412695884705), # First Steward at Castle Black ('Halder', 0.647807240486145), ('Grenn', 0.6452684998512268), ('Chett', 0.6423465609550476), ('Jeor-Mormont', 0.6395081877708435), # 997th Lord Commander of the Night's Watch ('Pypar', 0.6281514763832092), ('Samwell-Tarly', 0.6267001032829285), ('Dareon', 0.6214413046836853), ('Hobb', 0.6167237758636475)]

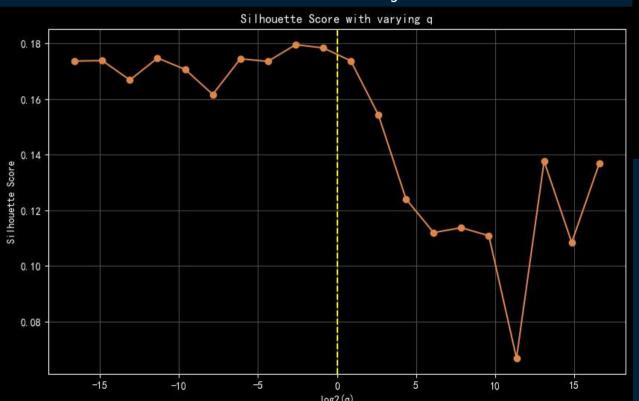
structural equivalence

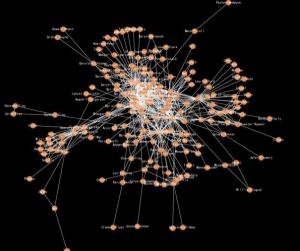


## **Evaluation**

 $p=1, q=[10^{-5}, 10^{5}]$ 

Interpretation and validation of consistency within clusters of data.





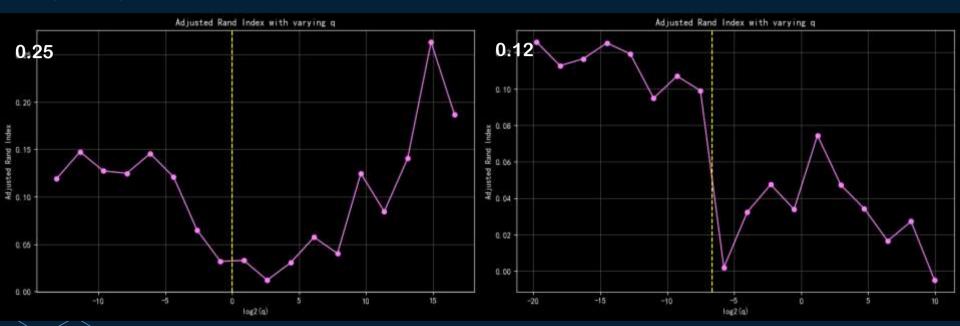
The structural differentiation is not so obvious

Lead to the results tend to be stable when q>>p

## **Evaluation-Parameter sensitivity**

$$p=1, q=[10^{-5}, 10^{5}]$$

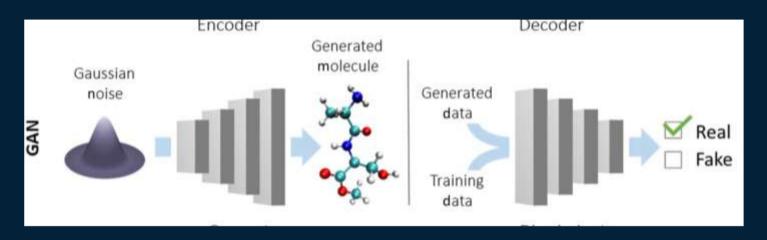
$$p=10^{-2}, q=[10^{-7}, 10^3]$$



While a low q encourages outward exploration, it is balanced by a low p which ensures that the walk does not go too far from the start node.



# Random Walk in predictions



- Their algorithms based on learning automata
- Based on Q-learning
- Based on deep learning and neural network
- Based on game theory
- Their algorithms for complex analysis.
- Chemical molecules