

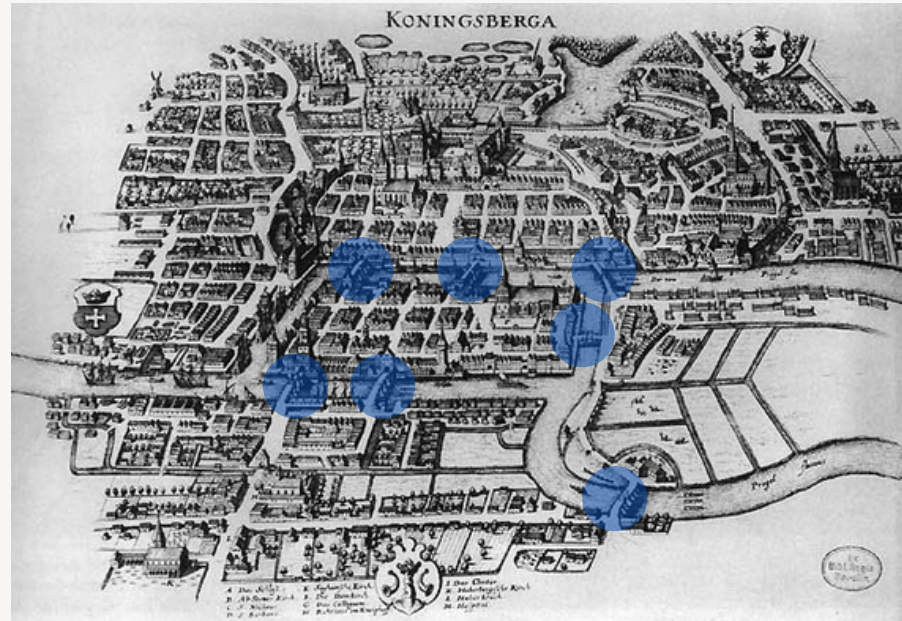
Análise de Redes Biológicas com a Linguagem Python

Prof. Dr. Gilderlanio Santana de Araújo
Arthur Ribeiro dos Santos

PPGBM0102 – Tópicos Avançados III – T01

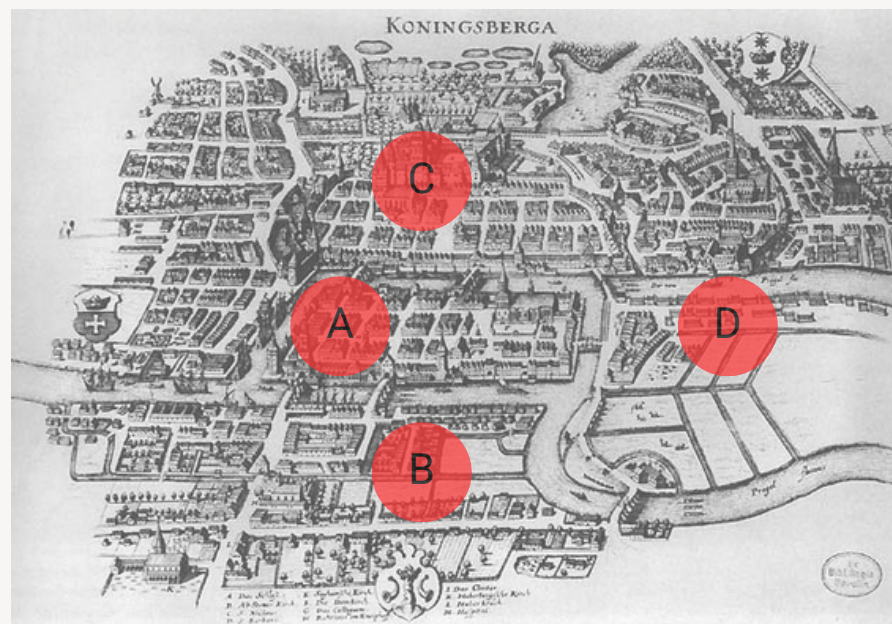
Sete Pontes de Königsberg

- Desafio contemporâneo
- É possível atravessar todas pontes da cidade sem repetições?



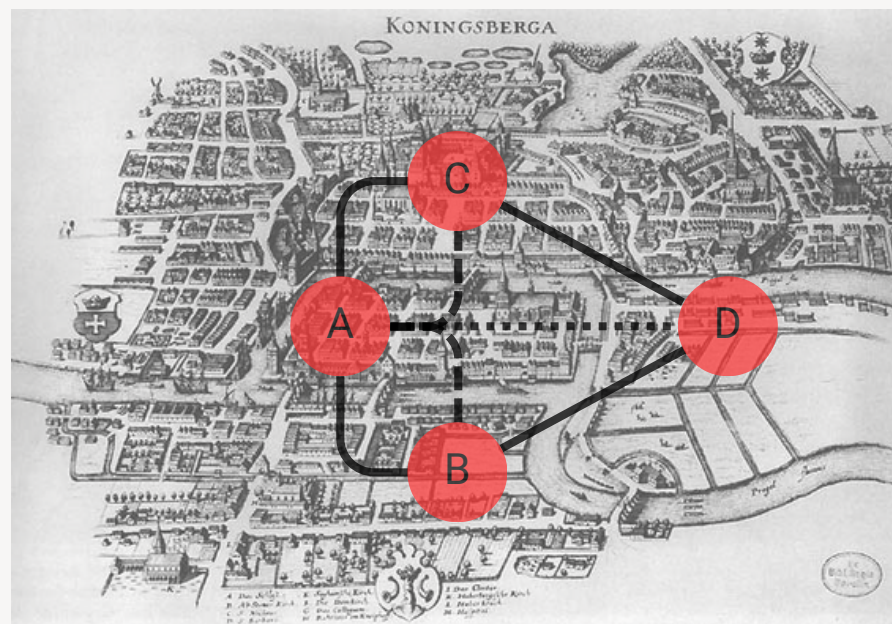
Sete Pontes de Königsberg

- Resolução do problema por Leonhard Euler (1736):
 - Comprovou matematicamente a solução negativa do problema
 - Simplificação do problema em pontos e linhas



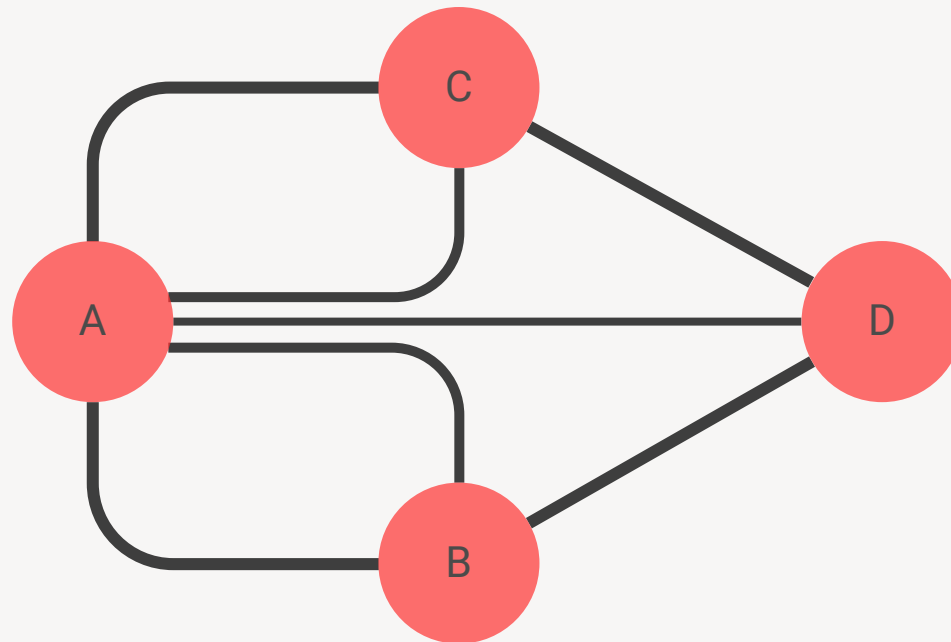
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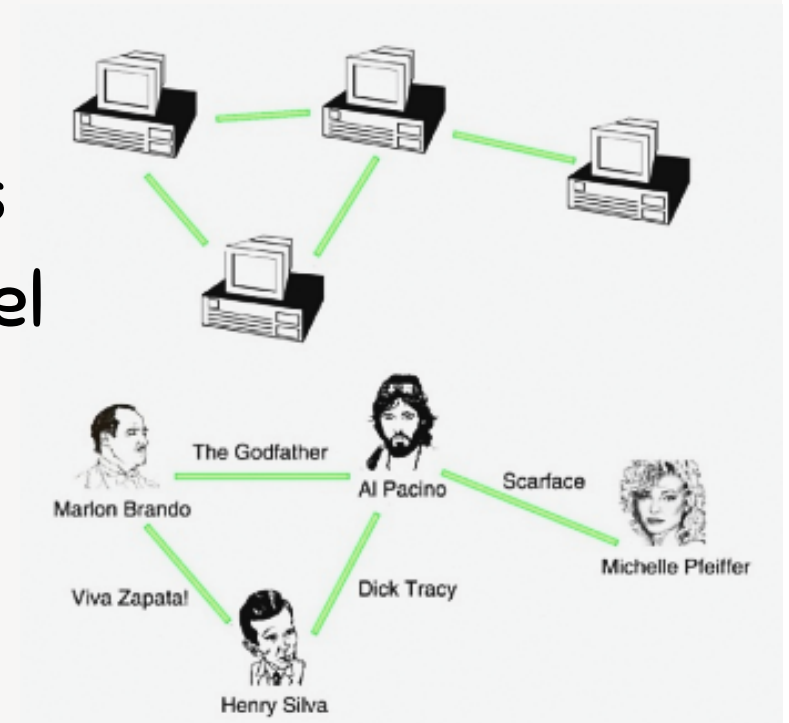
Sete Pontes de Königsberg

- Resolução do problema por Leonhard Euler (1736):
 - Comprovou matematicamente a solução negativa do problema
 - Simplificação do problema em pontos e linhas



≡ Motivações

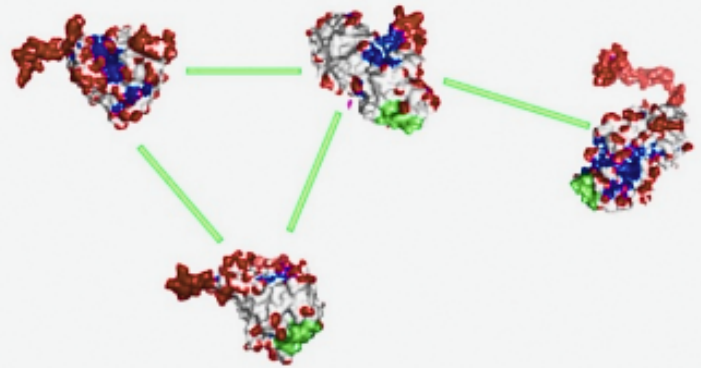
- Maior integração de dados
- Possibilitam análise a nível sistêmico



<http://networksciencebook.com>

≡ Motivações



- Grande volume de dados biológicos públicos
- Transformam problemas biológicos como questões de redes



<http://networksciencebook.com>

≡ Exemplos de redes biológicas

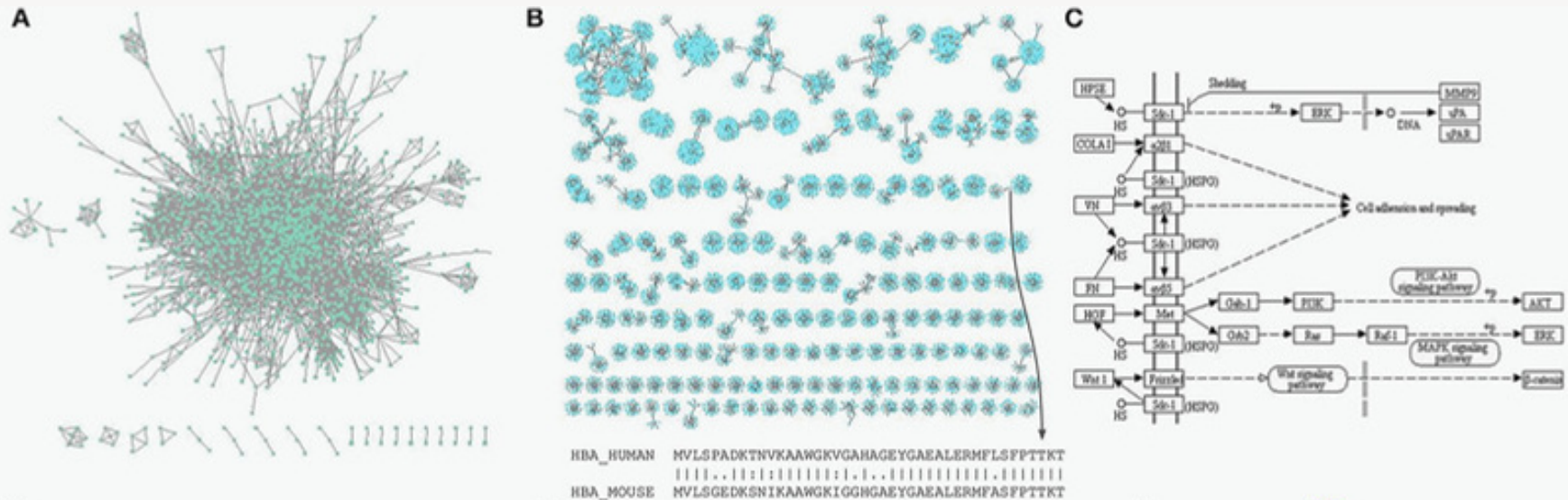
A Guide to Conquer the Biological Network Era Using Graph Theory

 Mikaela Koutrouli^{1*}  Evangelos Karatzas^{1,2*}  David Paez-Espino³  Georgios A. Pavlopoulos^{1*}

¹ Institute for Fundamental Biomedical Research, BSRC "Alexander Fleming", Vari, Greece

² Department of Informatics and Telecommunications, University of Athens, Athens, Greece

³ Lawrence Berkeley National Laboratory, Department of Energy, Joint Genome Institute, Walnut Creek, CA, United States







Rede proteína-proteína

Vias metabólicas do KEGG

Rede de similaridade de sequência

≡ Exemplos de redes biológicas

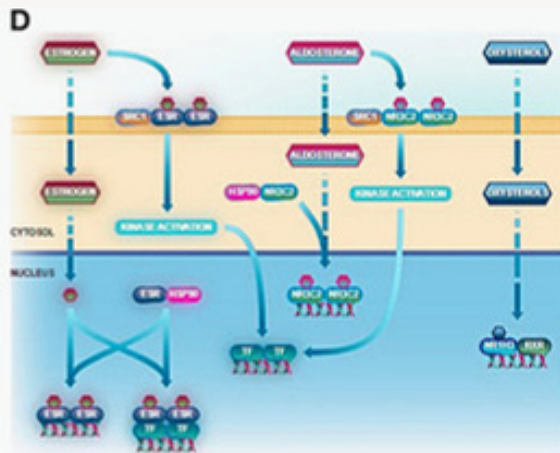
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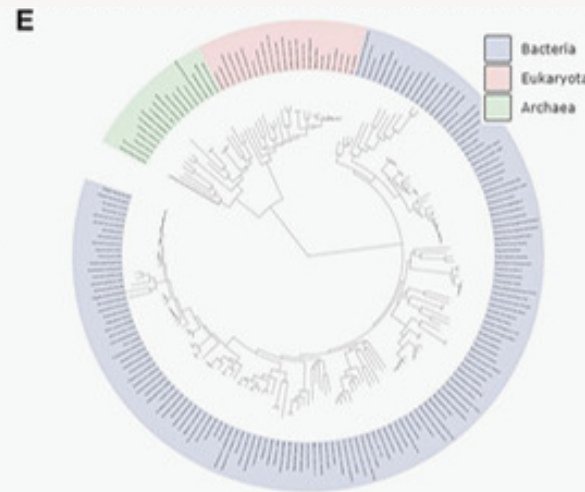
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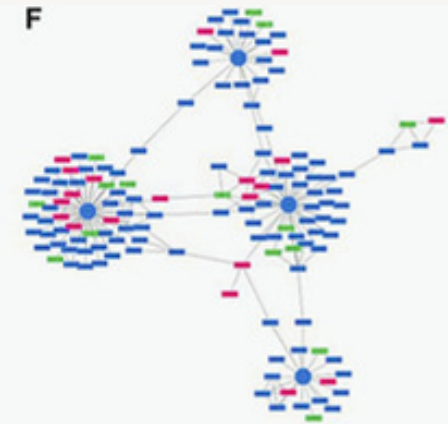
³ Lawrence Berkeley National Laboratory, Department of Energy, Joint Genome Institute, Walnut Creek, CA, United States



Rede transdutora de sinais
(REACTOME)



Árvore filogenética



Rede de expressão gênica

≡ Exemplos de redes biológicas

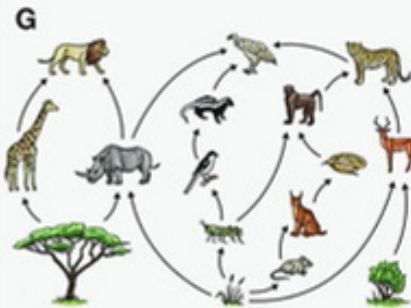
A Guide to Conquer the Biological Network Era Using Graph Theory

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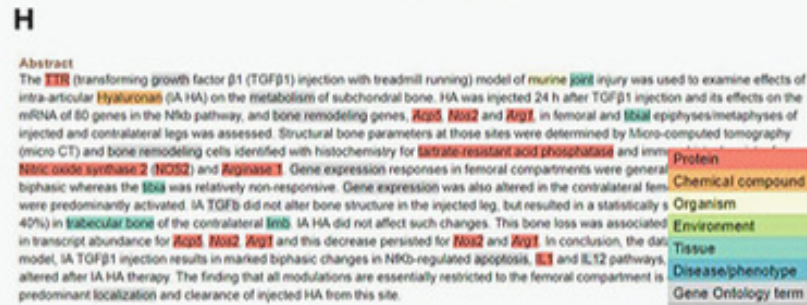
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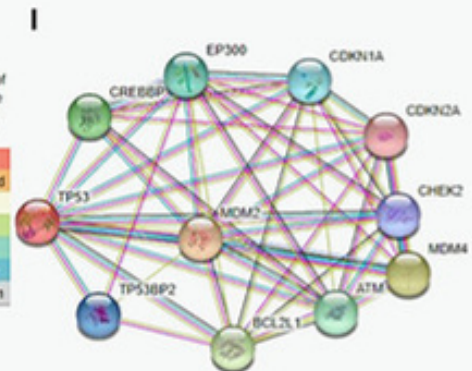
³ Lawrence Berkeley National Laboratory, Department of Energy, Joint Genome Institute, Walnut Creek, CA, United States



Cadeia alimentar



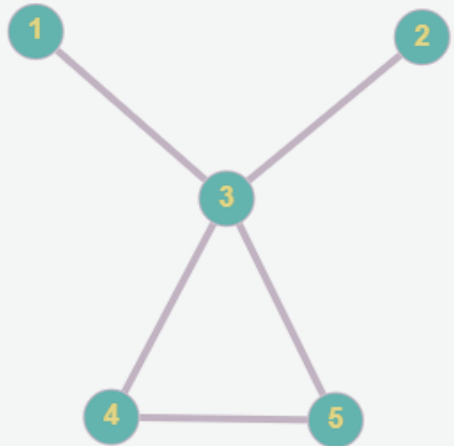
Co-ocorrências



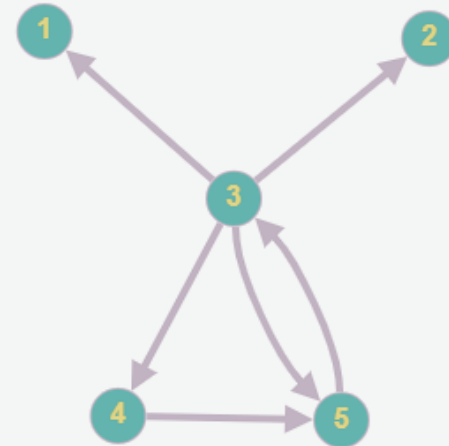
Rede PPI de múltiplas arestas

≡ Tipos de grafos

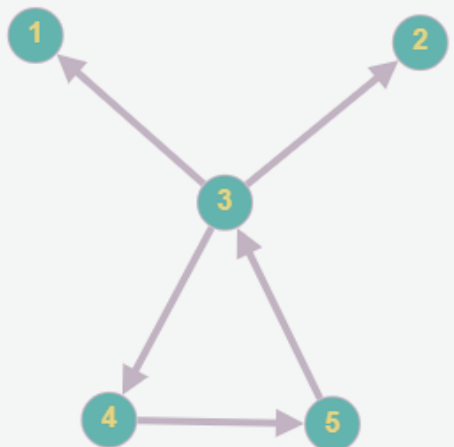
Não direcionado



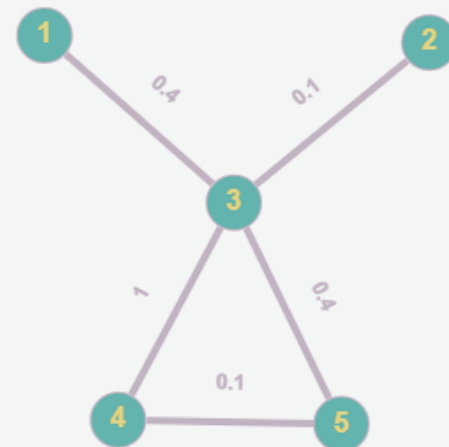
Misto



Direcionado

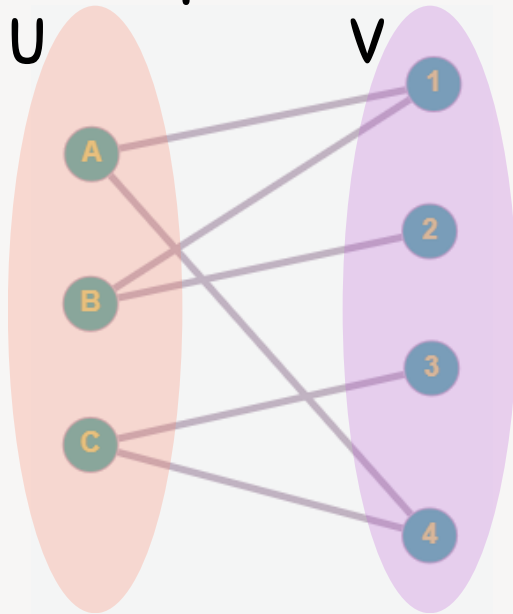


Com peso

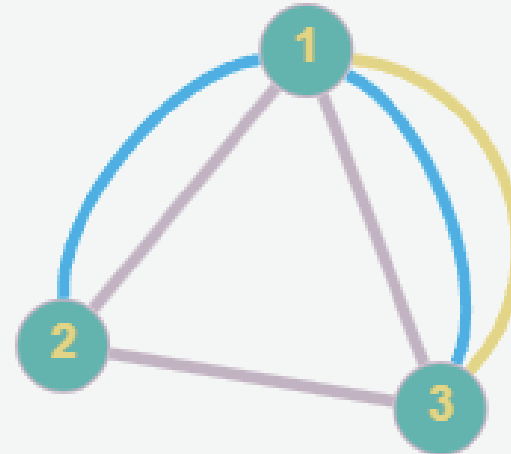


≡ Tipos de grafos

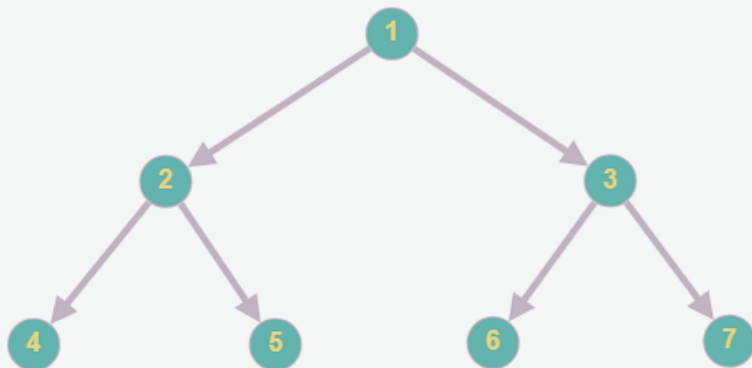
Bipartido



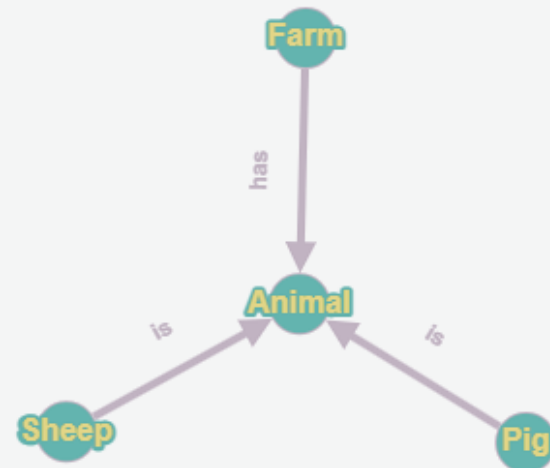
Multi-aresta



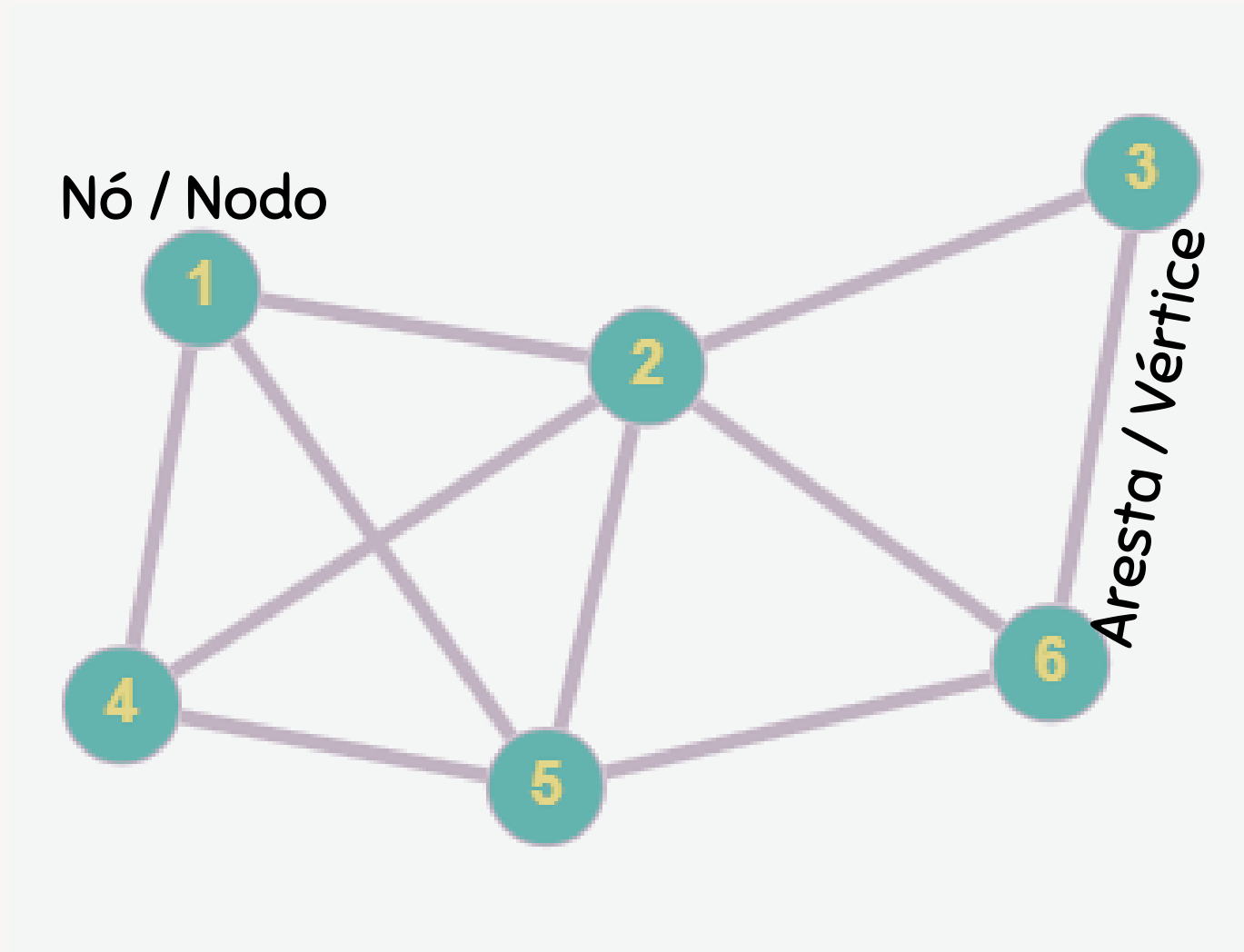
Árvore



Semântico

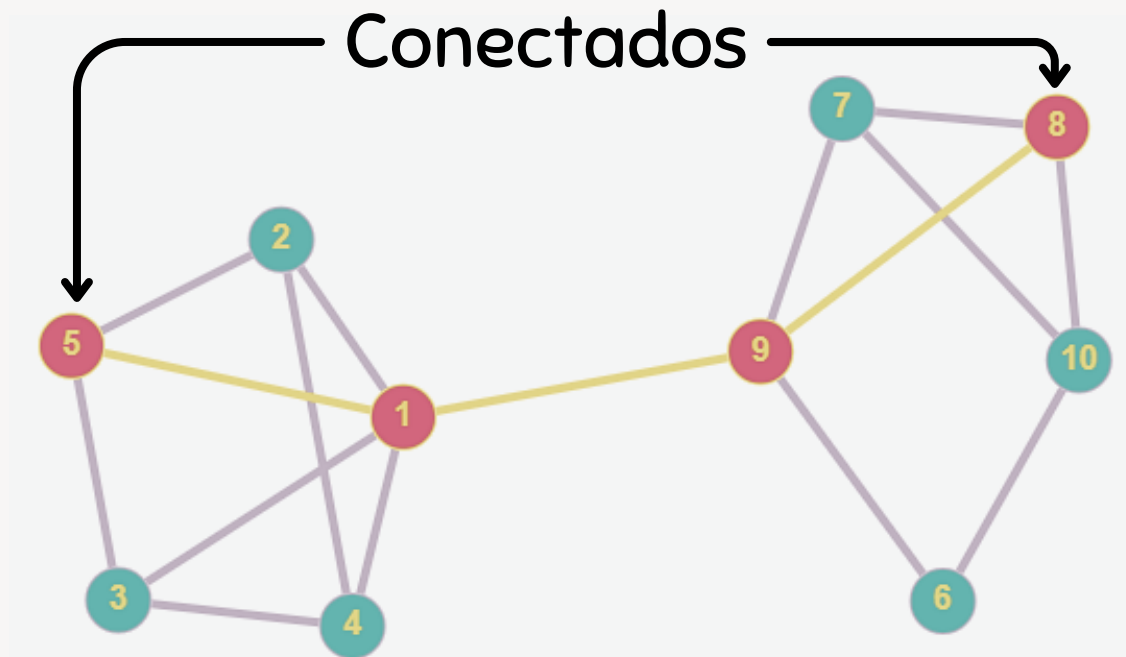


≡ Estrutura básica de grafos



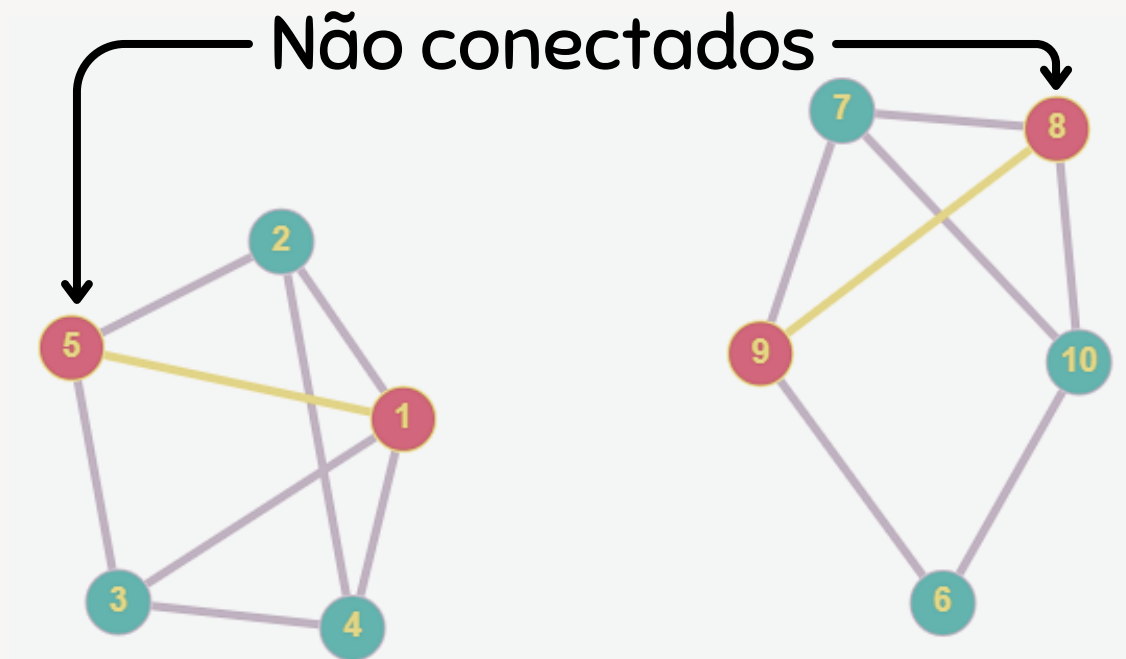
≡ Conectividade

- Conceito relacionado a ligação entre elementos de uma rede
- Mede a robustez de um sistema

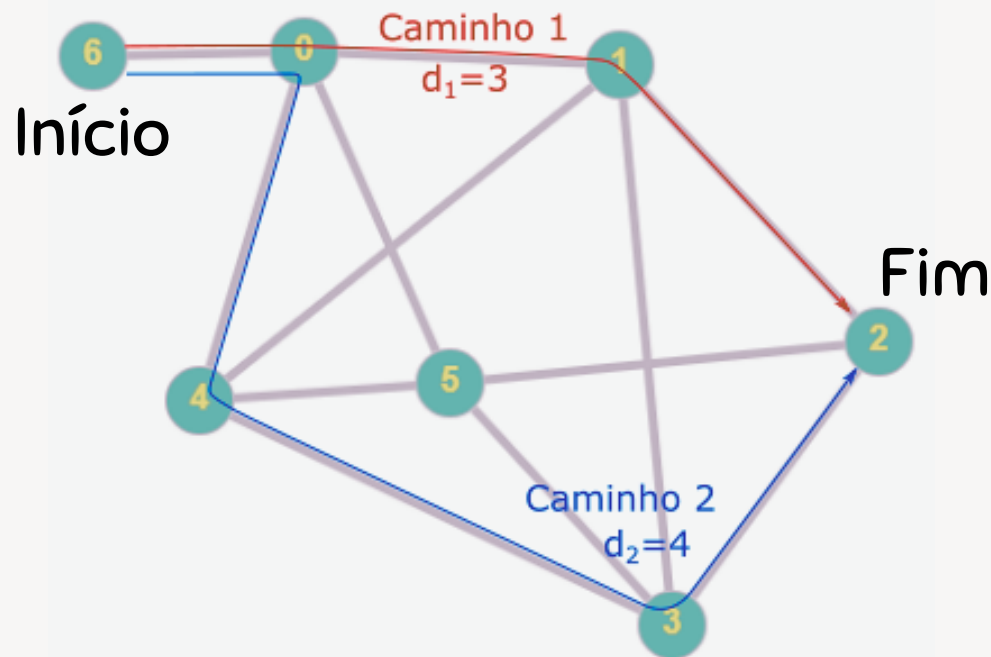


≡ Conectividade

- Conceito relacionado a ligação entre elementos de uma rede
- Mede a robustez de um sistema

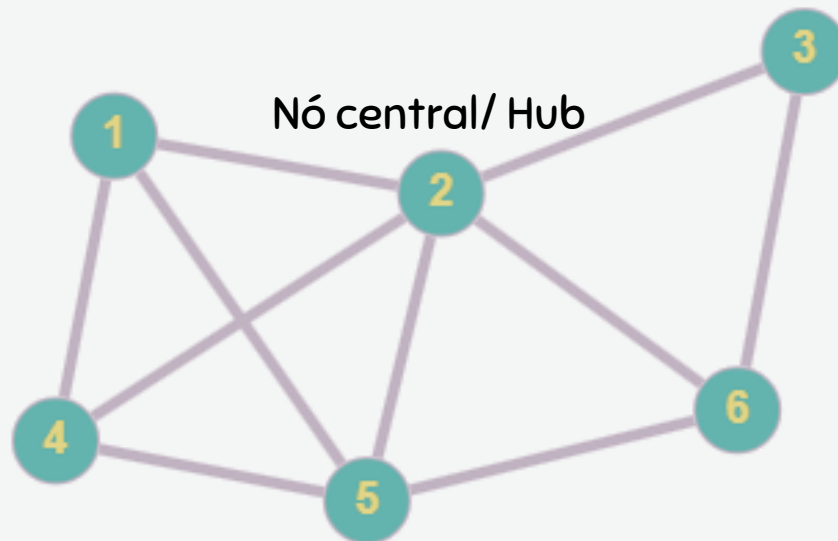


- Fortemente ligado a teoria dos problemas de fluxo
 - Medições de distância e caminho



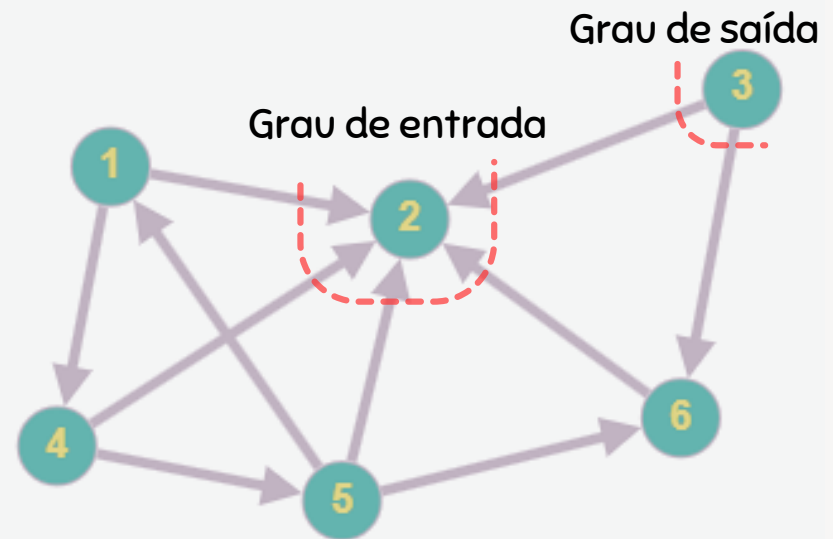
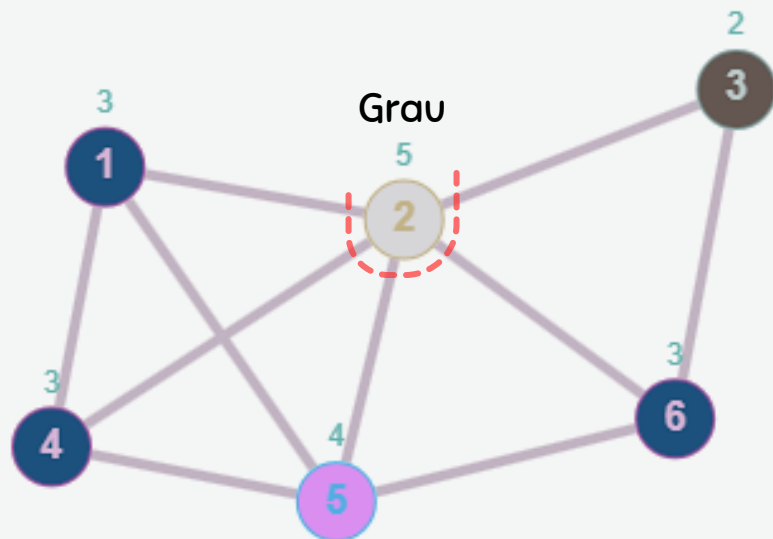
≡ Centralidade

- Conceito que mede a influência das conexões de uma rede
- Permite a estimação de importância de um elemento dentro de um sistema



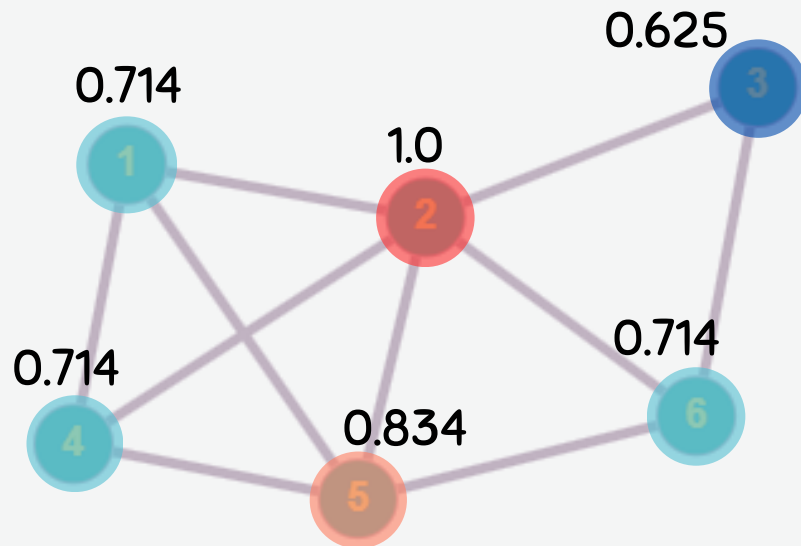
≡ Centralidade de grau

- Mede o número de ligações associados a cada elemento de uma rede
 - Aumento da chance de um elemento receber informações do sistema



≡ Centralidade de proximidade

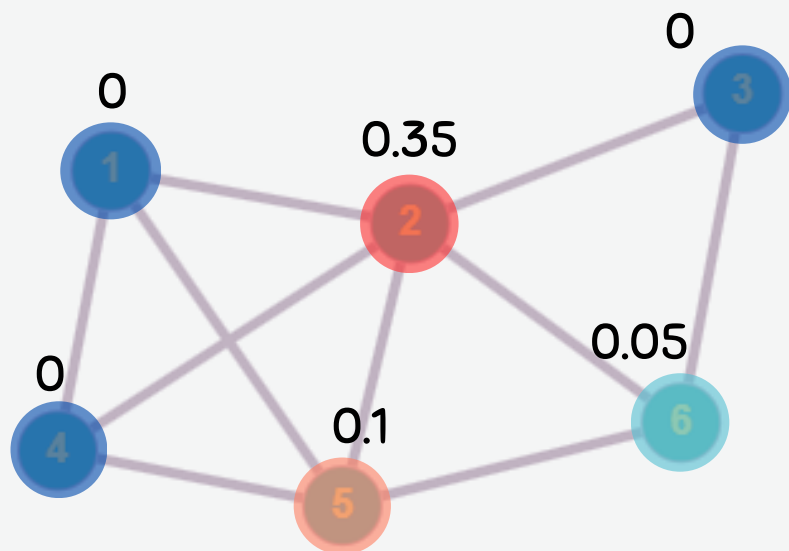
- Mede a menor distância entre todos pares de elementos de uma rede
 - Identifica o elemento com maior facilidade de transmitir informações



$$C(u) = \frac{n - 1}{\sum_{v=1}^{n-1} d(v, u)}$$

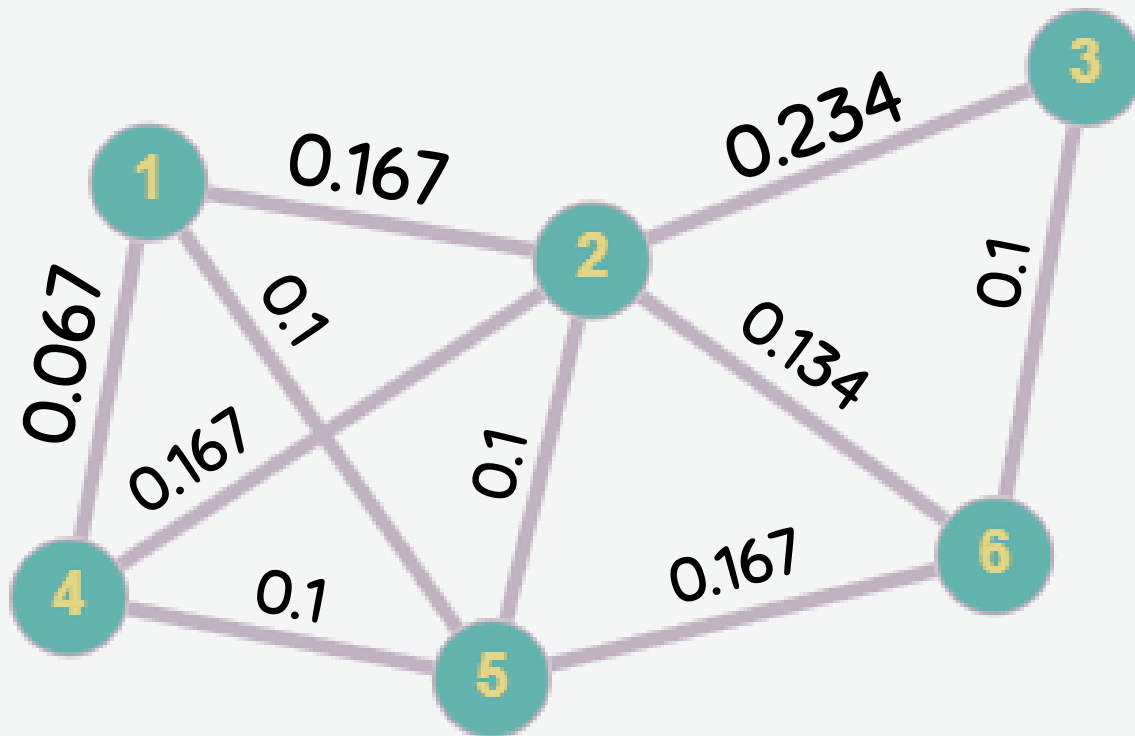
≡ Centralidade de intermediação

- Mede o número de menores caminhos que atravessam um elemento/conexão
 - Identifica as estruturas principais envolvidas na transmissão de informação



$$c_B(v) = \sum_{s,t \in V} \frac{\sigma(s, t|v)}{\sigma(s, t)}$$

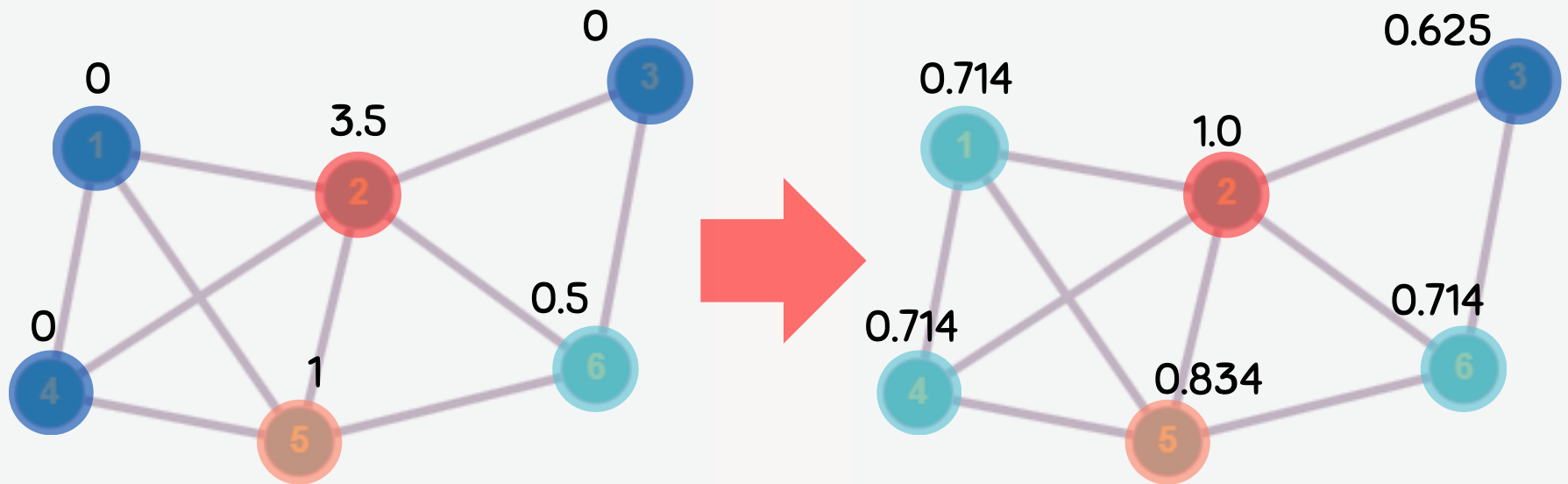
≡ Centralidade de intermediação



$$c_B(e) = \sum_{s,t \in V} \frac{\sigma(s,t|e)}{\sigma(s,t)}$$

≡ Centralidade de intermediação

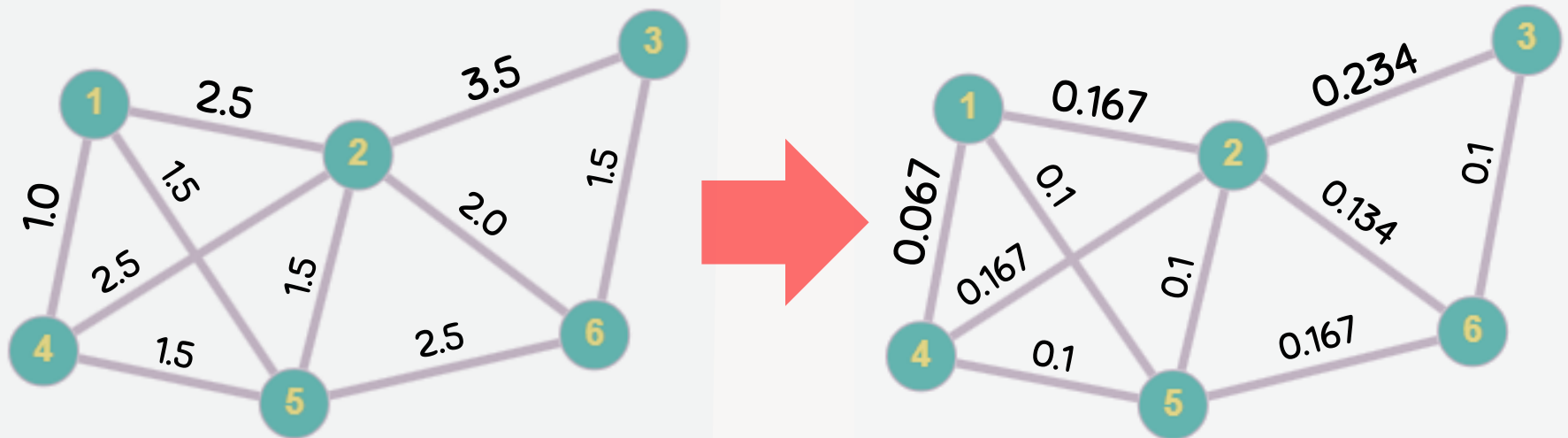
Coeficientes de intermediação não-normalizados



$$G_n = G * 2 / ((N-1)(N-2))$$

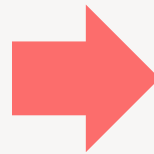
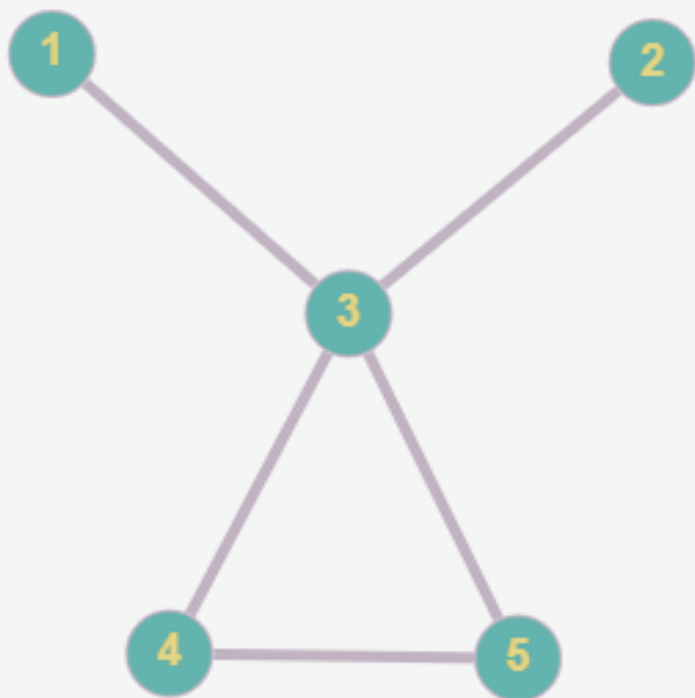
≡ Centralidade de intermediação

Coeficientes de intermediação não-normalizados



$$G_n = G * 2 / (N * (N - 1))$$

≡ Matriz e lista de adjacência



i\j	1	2	3	4	5
1	0	1	0	1	1
2	1	0	1	1	1
3	0	1	0	0	0
4	1	1	0	0	1
5	1	1	0	1	0

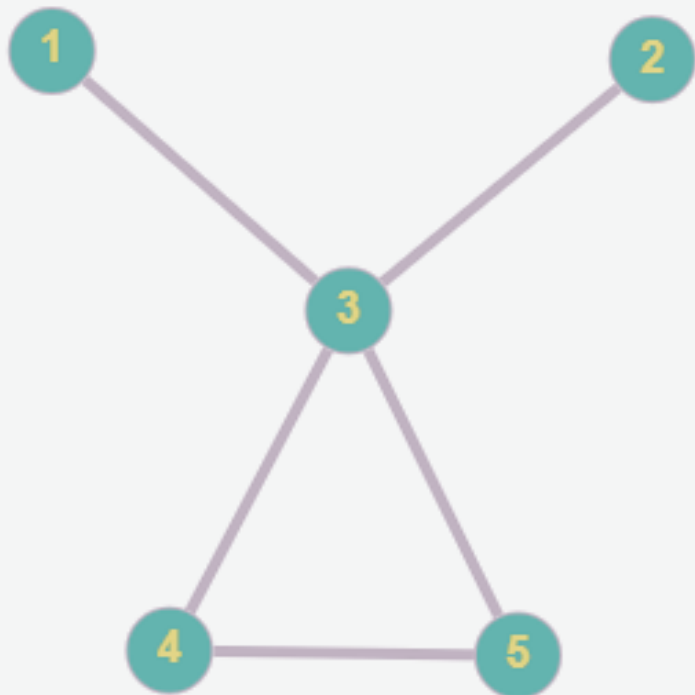
≡ Matriz e lista de adjacência

i \ j	1	2	3	4	5
1	0	1	0	1	1
2	1	0	1	1	1
3	0	1	0	0	0
4	1	1	0	0	1
5	1	1	0	1	0



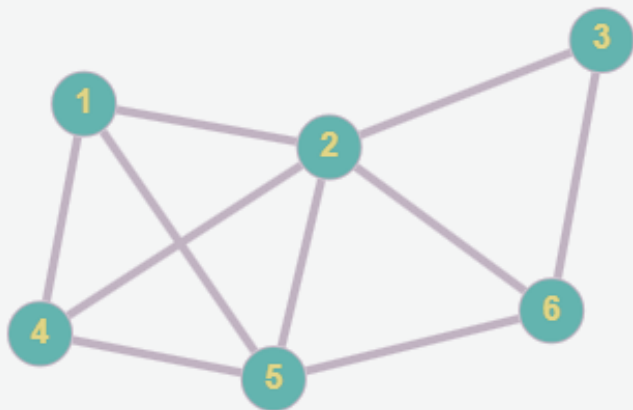
Origem	Destino	Valor
1	2	1
1	3	0
1	4	1
1	5	1
2	3	1
2	4	1
2	5	1
3	4	0
3	5	0
4	5	1

≡ Matriz e lista de adjacência

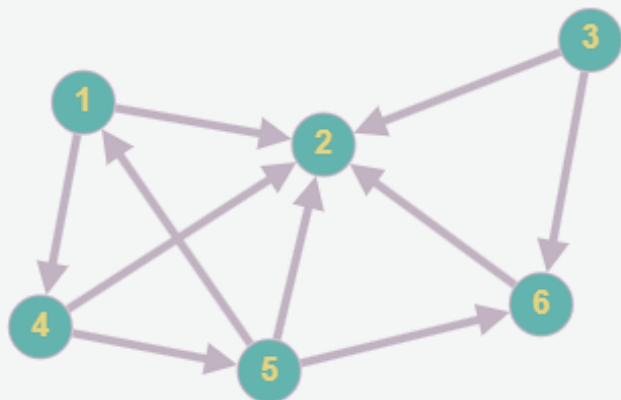


Origem	Destino	Valor
1	2	1
1	3	0
1	4	1
1	5	1
2	3	1
2	4	1
2	5	1
3	4	0
3	5	0
4	5	1

≡ Matriz e lista de adjacência



i\j	1	2	3	4	5	6
1	0	1	0	1	1	0
2	1	0	1	1	1	1
3	0	1	0	0	0	1
4	1	1	0	0	1	0
5	1	1	0	1	0	1
6	0	1	1	0	1	0



i\j	1	2	3	4	5	6
1	0	1	0	1	0	0
2	0	0	0	0	0	0
3	0	1	0	0	0	1
4	0	1	0	0	1	0
5	1	1	0	1	0	1
6	0	1	0	0	0	0

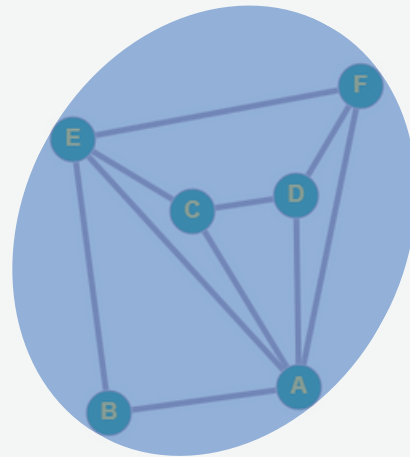
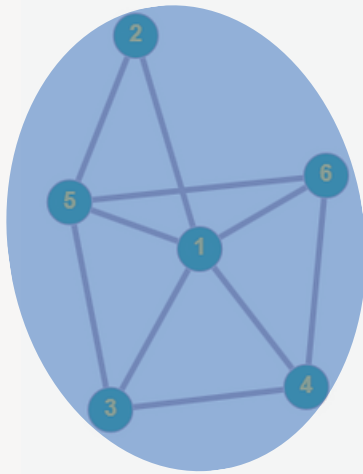
≡ Matriz e lista de adjacência

i\j	1	2	3	4	5	6
1	0	1	0	1	0	0
2	0	0	0	0	0	0
3	0	1	0	0	0	1
4	0	1	0	0	1	0
5	1	1	0	1	0	1
6	0	1	0	0	0	0



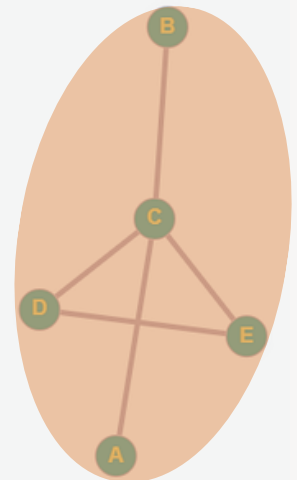
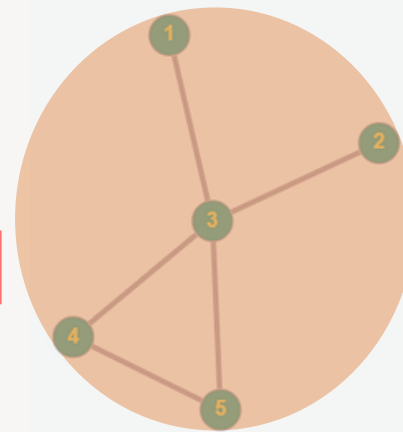
Origem	Destino	Valor
1	2	1
1	3	0
...		
2	1	0
2	3	0
...		
3	1	0
3	2	1
...		

≡ Isomorfismo entre grafos



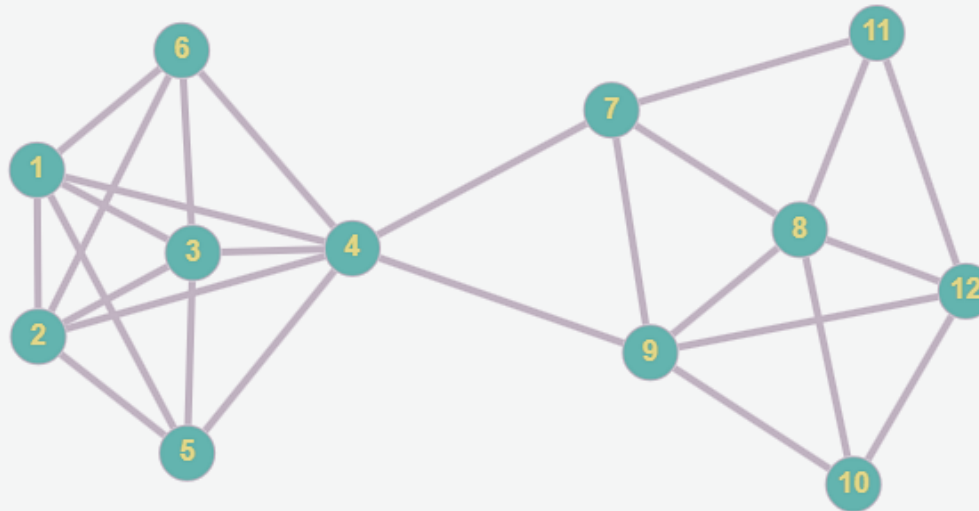
0	1	1	1	1	1
1	0	0	0	1	0
1	0	0	1	1	0
1	0	1	0	0	1
1	1	1	0	0	1
1	0	0	1	1	0

0	0	1	0	0
0	0	1	0	0
1	1	0	1	1
0	0	1	0	1
0	0	1	1	0



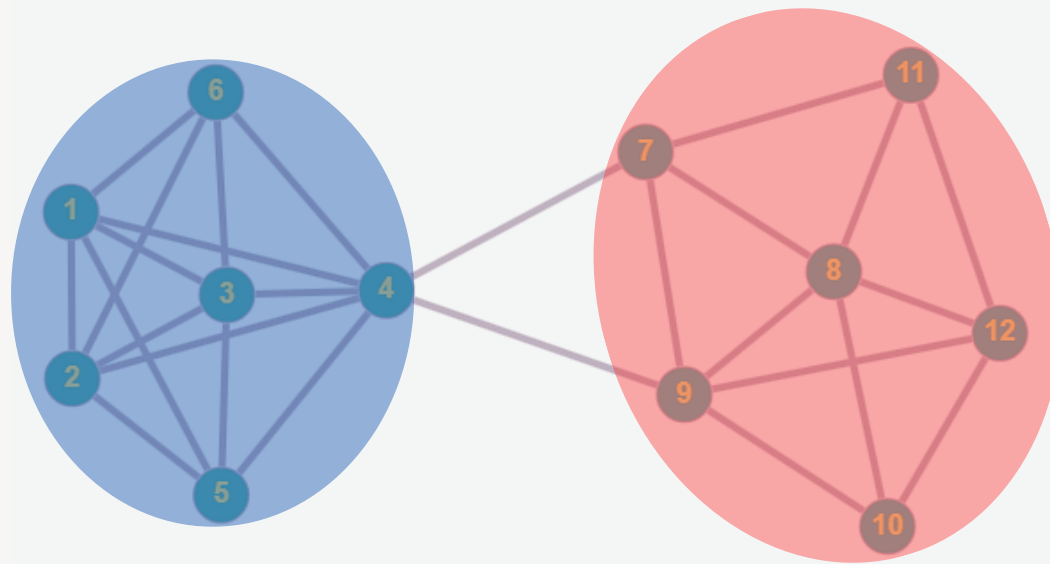
≡ Partição de grafos

- Problema de teoria de grafos
- Capacidade de isolar um grafo em componentes menores
 - Compartilhamento de características

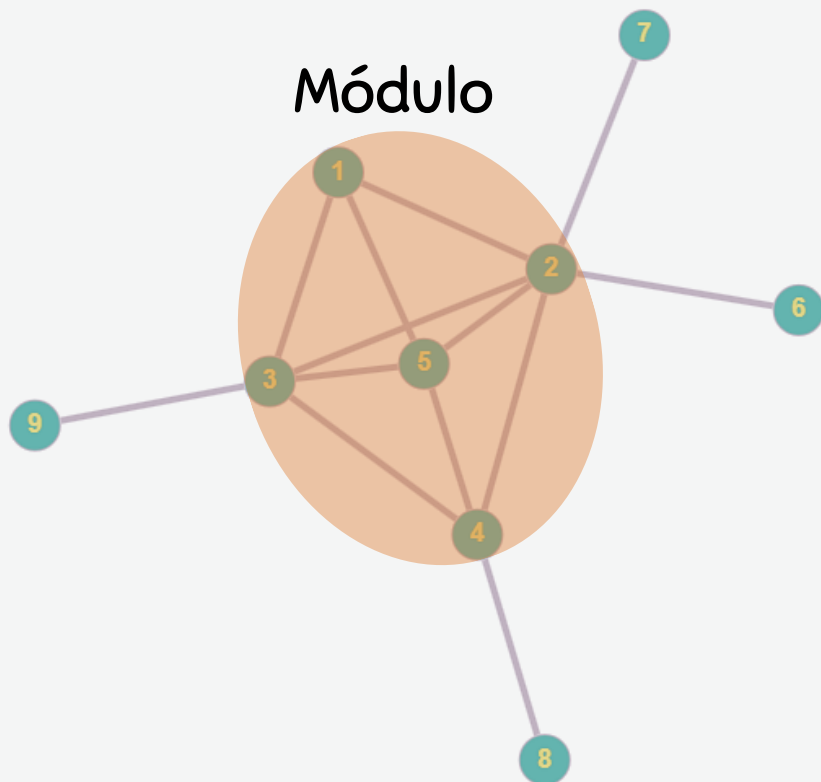


≡ Partição de grafos

- Problema de teoria de grafos
- Capacidade de isolar um grafo em componentes menores
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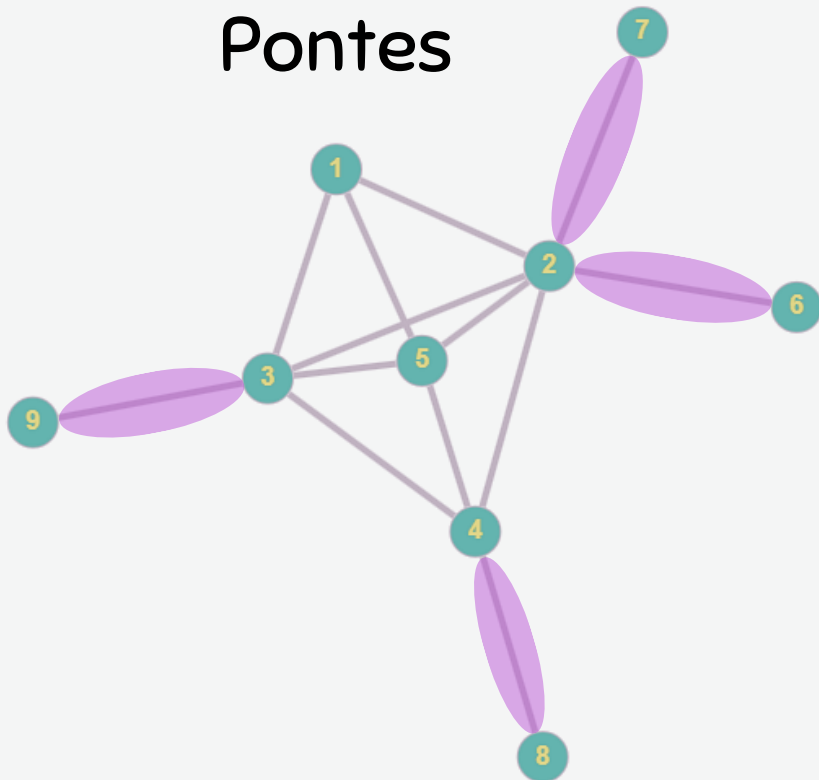
≡ Partição de grafos



i\j	1	2	3	4	5	6	7	8	9
1	0	1	1	0	1	0	0	0	0
2	1	0	1	1	1	1	1	0	0
3	1	1	0	1	1	0	0	0	1
4	0	1	1	0	1	0	0	1	0
5	1	1	1	1	0	0	0	0	0
6	0	1	0	0	0	0	0	0	0
7	0	1	0	0	0	0	0	0	0
8	0	0	0	1	0	0	0	0	0
9	0	0	1	0	0	0	0	0	0

≡ Partição de grafos

Pontes



i\j	1	2	3	4	5	6	7	8	9
1	0	1	1	0	1	0	0	0	0
2	1	0	1	1	1	1	1	0	0
3	1	1	0	1	1	0	0	0	1
4	0	1	1	0	1	0	0	1	0
5	1	1	1	1	0	0	0	0	0
6	0	1	0	0	0	0	0	0	0
7	0	1	0	0	0	0	0	0	0
8	0	0	0	1	0	0	0	0	0
9	0	0	1	0	0	0	0	0	0



NetworkX

Network Analysis in Python

NetworkX is a Python package for the creation, manipulation, and study of the structure, dynamics, and functions of complex networks.



- Algoritmos para análise de métricas de grafos
- Métodos de visualização e layouts para redes



NetworkX

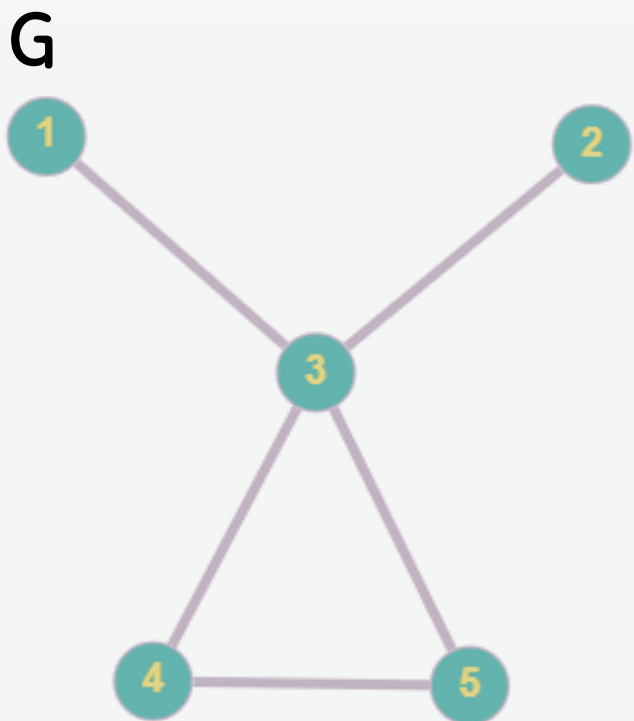
Network Analysis in Python

NetworkX is a Python package for the creation, manipulation, and study of the structure, dynamics, and functions of complex networks.



```
$ pip install networkx[default]
```

≡ Manipulação de grafos no NetworkX



```
>>> import networkx as nx
```

```
>>> G = nx.Graph()
```

```
>>> G.add_edge(1, 3)
```

```
>>> G.add_edge(2, 3)
```

```
...
```


≡ Manipulação de grafos no NetworkX

```
>>> G.edges()
```

```
EdgeView([(1, 3), (3, 2), (3, 4), (3, 5), (4, 5)])
```

```
>>> G.nodes()
```

```
NodeView((1, 3, 2, 4, 5))
```

```
>>> G.degree()
```

```
DegreeView({1: 1, 3: 4, 2: 1, 4: 2, 5: 2})
```

≡ Manipulação de grafos no NetworkX

```
>>> nx.degree_centrality(G)
```

```
EdgeView([(1, 3), (3, 2), (3, 4), (3, 5), (4, 5)])
```

```
>>> nx.closeness_centrality(G)
```

```
{1: 0.5714285714285714,  
 3: 1.0,  
 2: 0.5714285714285714,  
 4: 0.6666666666666666,  
 5: 0.6666666666666666}
```

≡ Manipulação de grafos no NetworkX

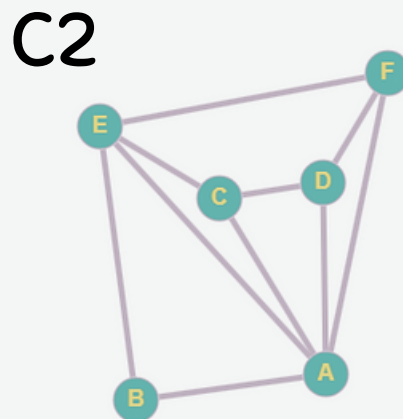
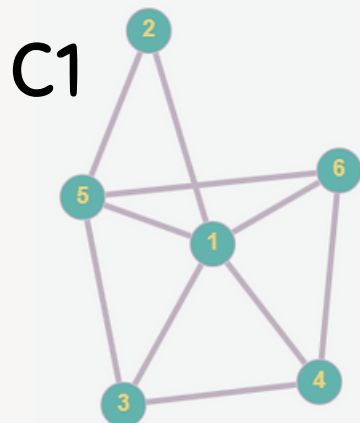
```
>>> nx.betweenness_centrality(G)
```

```
{1: 0.0, 3: 0.8333333333333333, 2: 0.0, 4: 0.0, 5: 0.0}
```

```
>>> nx.edge_betweenness_centrality(G)
```

```
{(1, 3): 0.4,  
 (3, 2): 0.4,  
 (3, 4): 0.30000000000000004,  
 (3, 5): 0.30000000000000004,  
 (4, 5): 0.1}
```

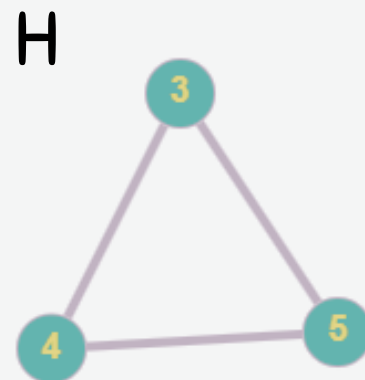
≡ Manipulação de grafos no NetworkX



```
>>> nx.is_isomorphic(C1, C2)
```

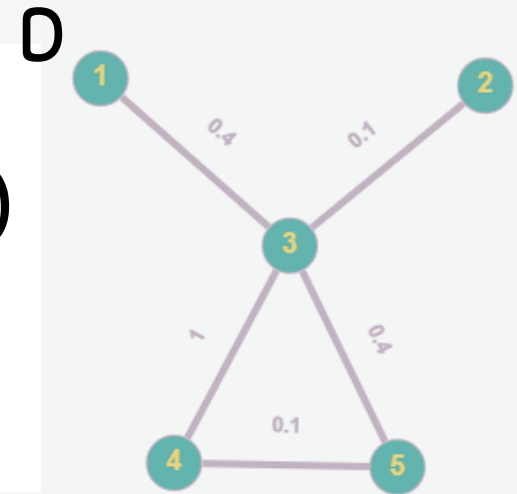
True

```
>>> H = G.subgraph([3, 4, 5])
```

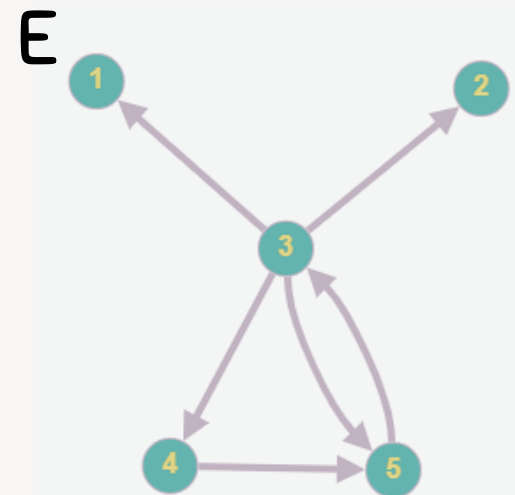


≡ Manipulação de grafos no NetworkX

```
>>> D = nx.Graph()
>>> D.add_edge(3, 5, weight = 0.4)
>>> D.add_edge(4, 5, weight = 0.1)
...
```



```
>>> E = nx.DiGraph()
>>> E.add_edge(3, 5)
>>> E.add_edge(5, 3)
...
```

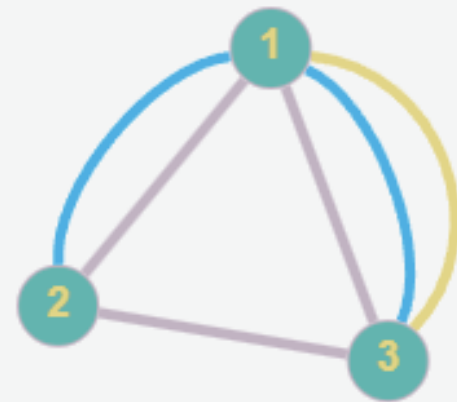


≡ Manipulação de grafos no NetworkX

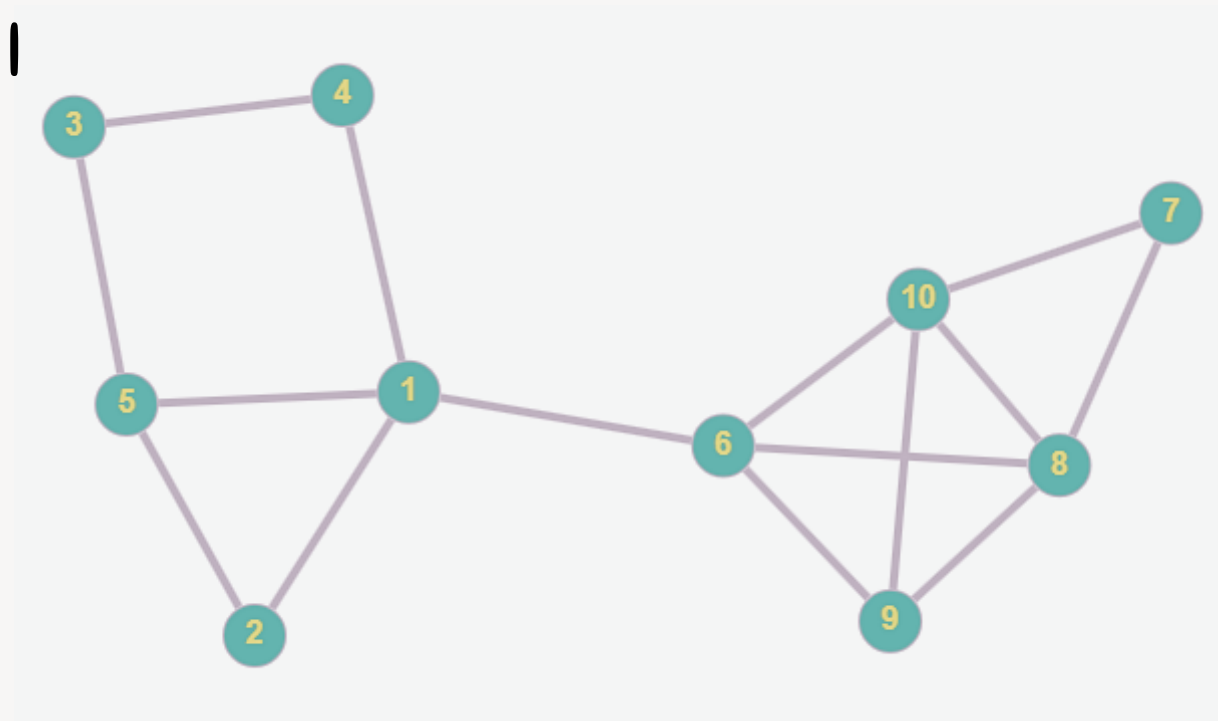
```
>>> F = nx.MultiGraph()  
>>> F.add_edge(1, 2, color = "gray")  
>>> F.add_edge(1, 2, color = "blue")  
...
```

- Armazenamento de atributos:
 - Dicionário

F



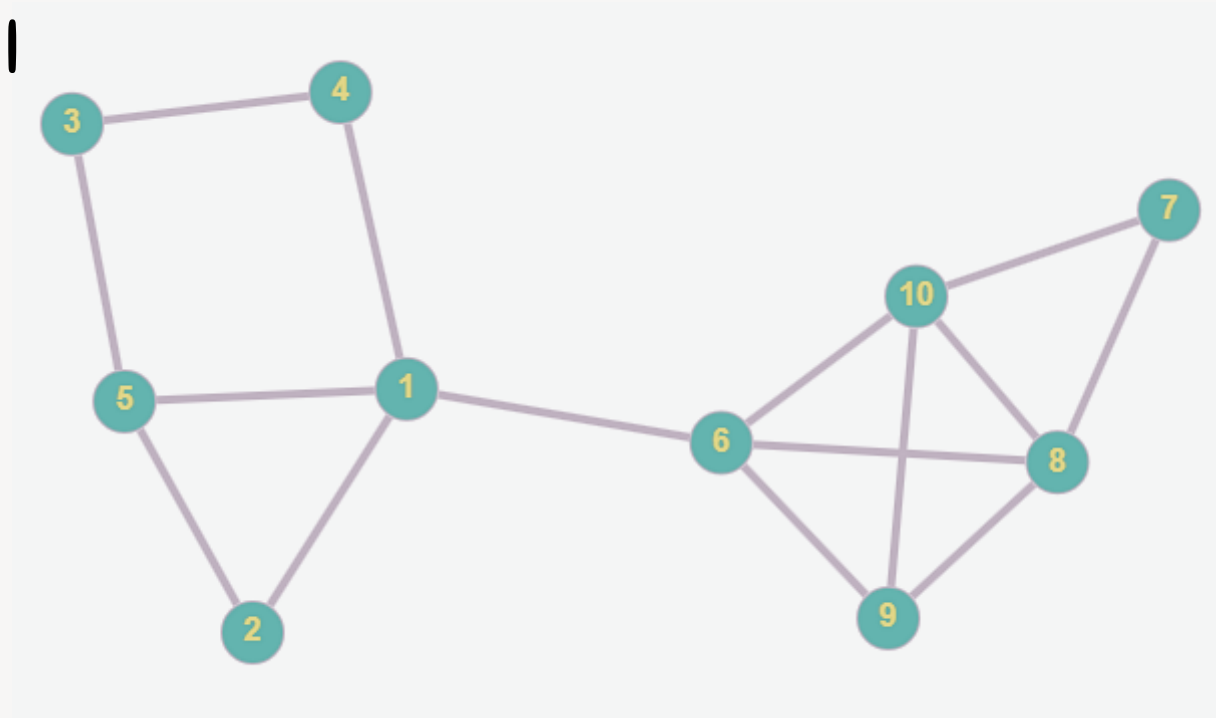
≡ Manipulação de grafos no NetworkX



```
>>> nx.has_bridges(I)  
True
```

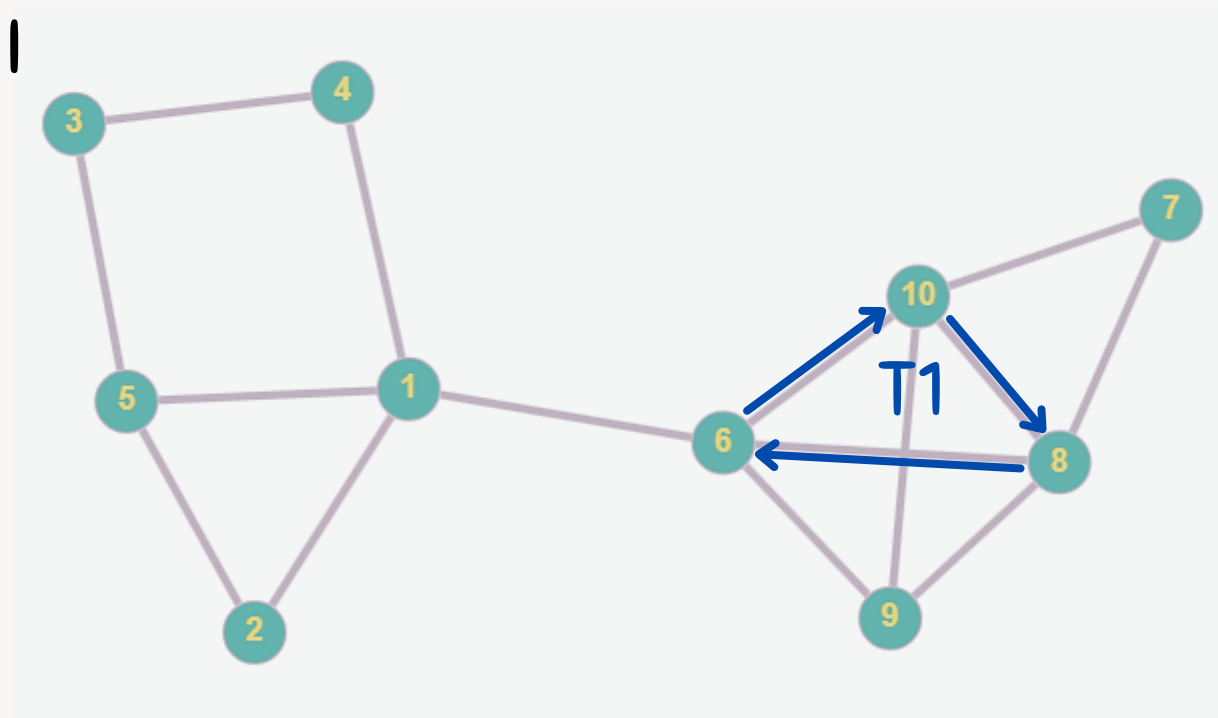
```
>>> list(nx.bridges(I))  
[(1, 6)]
```

Coeficiente de agrupamento



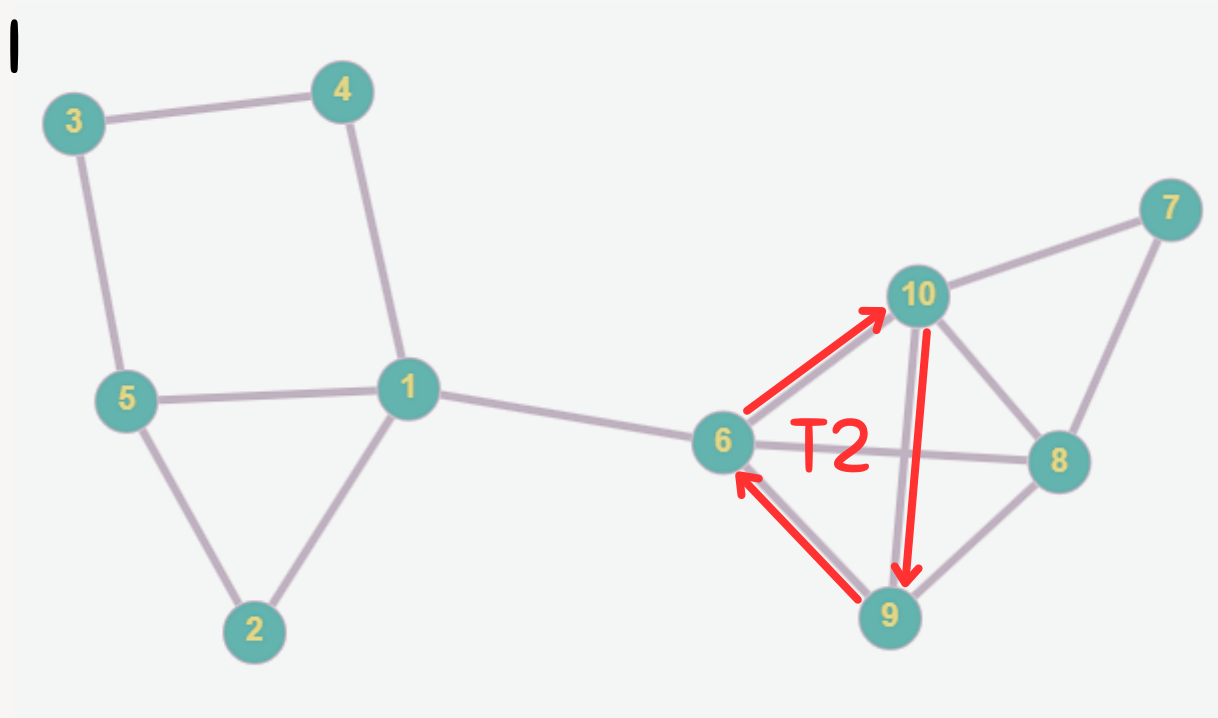
$$c_u = \frac{2T(u)}{\deg(u)(\deg(u) - 1)}$$

Coeficiente de agrupamento



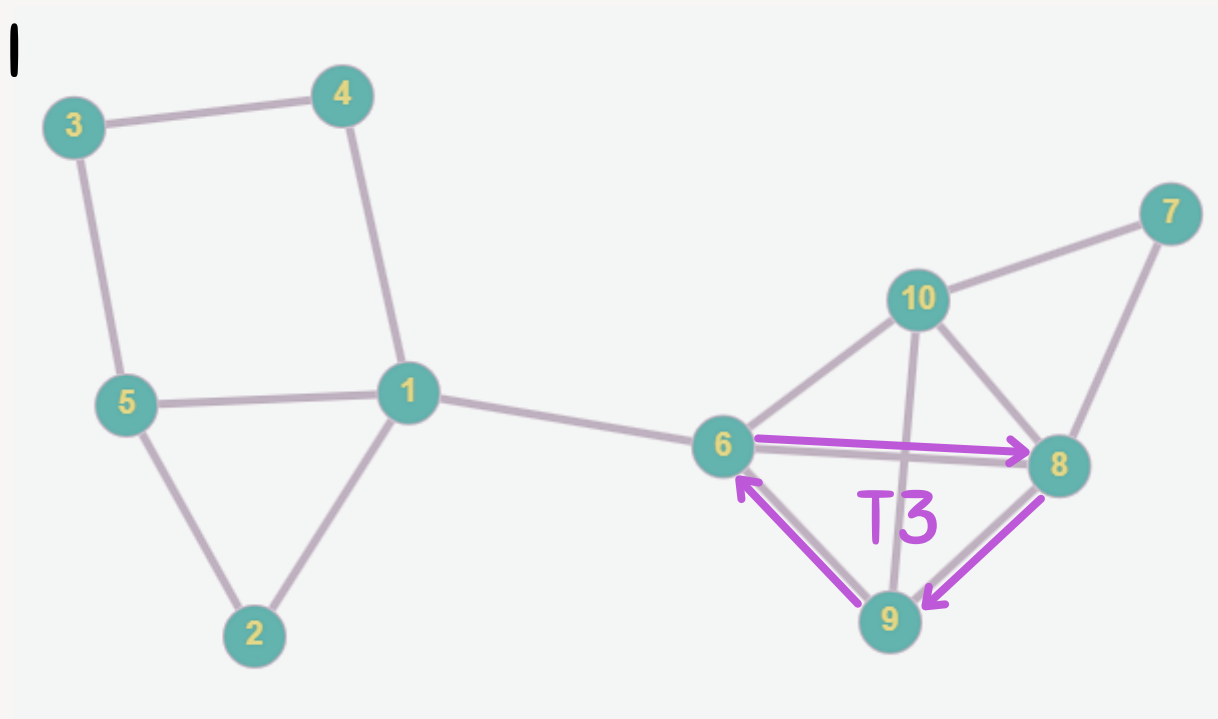
$$c_u = \frac{2T(u)}{\deg(u)(\deg(u) - 1)}$$

Coeficiente de agrupamento



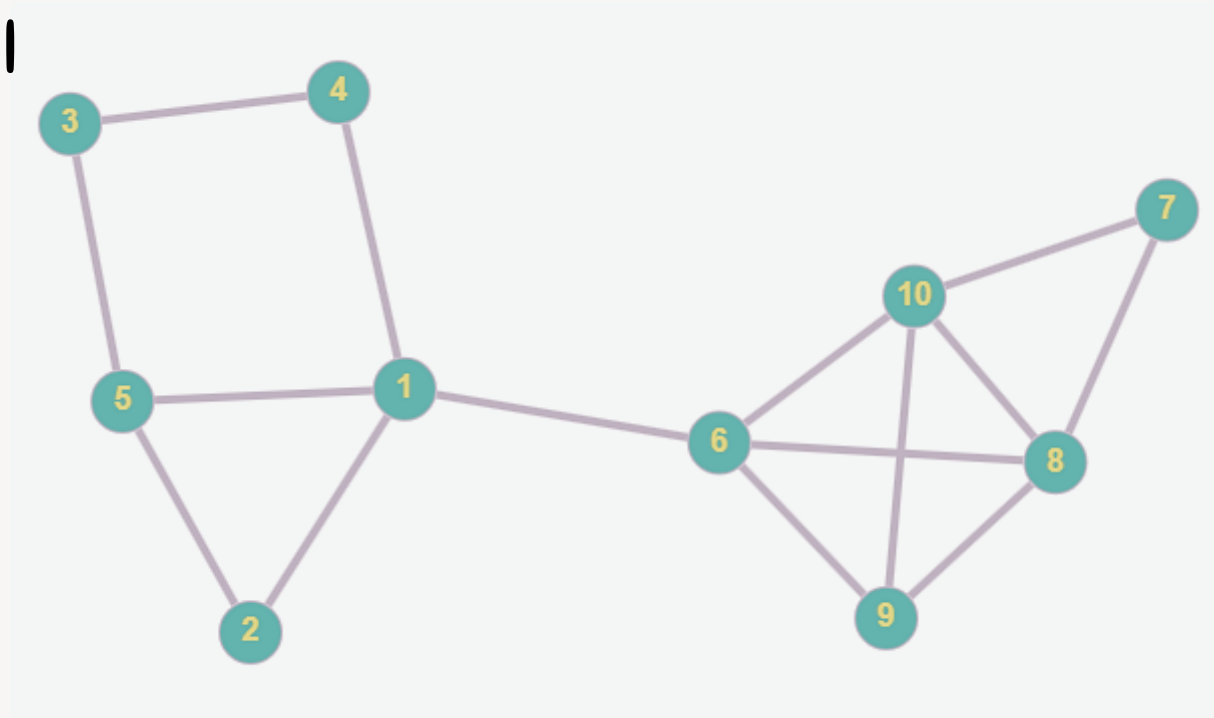
$$c_u = \frac{2T(u)}{\deg(u)(\deg(u) - 1)}$$

Coeficiente de agrupamento



$$c_u = \frac{2T(u)}{\deg(u)(\deg(u) - 1)}$$

≡ Manipulação de grafos no NetworkX

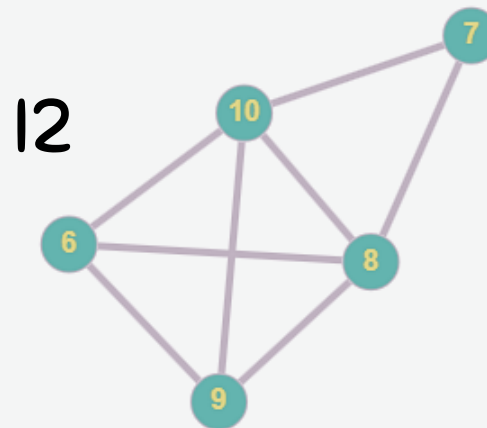
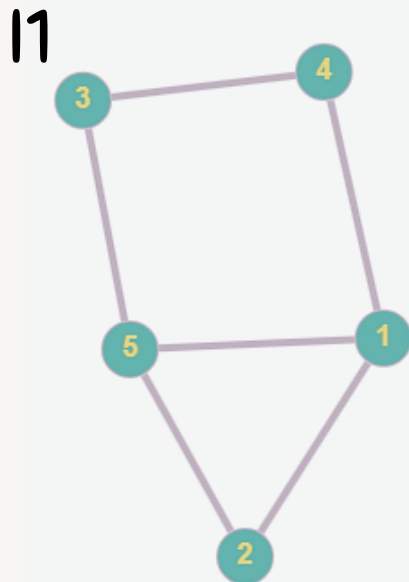


$$c_u = \frac{2T(u)}{\deg(u)(\deg(u) - 1)}$$

```
>>> nx.clustering(l, 6)  
0.5
```

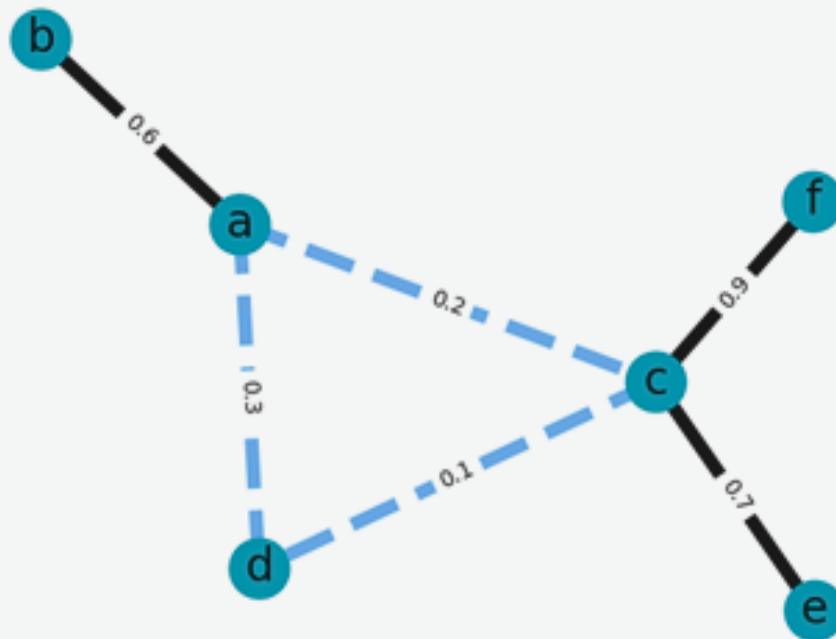
≡ Manipulação de grafos no NetworkX

```
>>> from networkx.algorithms.community centrality
import girvan_newman
>>> comp = girvan_newman(l)
>>> tuple(sorted(c) for c in next(comp))
([1, 2, 3, 4, 5], [6, 7, 8, 9, 10])
```



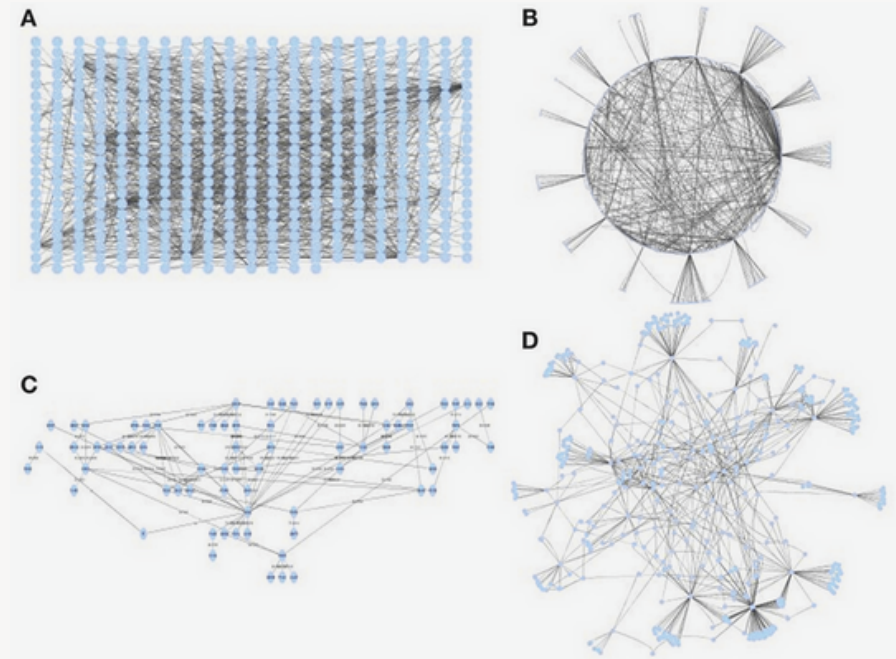
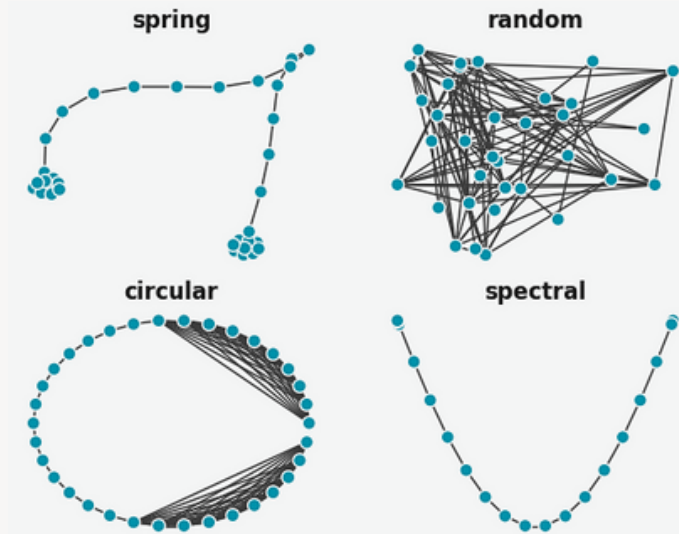
≡ Modelagem de grafos no NetworkX

```
>>> nx.draw(G)
>>> nx.draw_networkx_nodes(G, pos)
>>> nx.draw_networkx_edges(G, pos)
>>> nx.draw_networkx_labels(G, pos)
>>> nx.draw_networkx_edge_labels(G, pos)
```



≡ Modelagem de grafos no NetworkX

Layout da rede



JOURNAL ARTICLE

PyWGCNA: a Python package for weighted gene co-expression network analysis

Narges Rezaie, Farilie Reese, Ali Mortazavi ✉

Bioinformatics, Volume 39, Issue 7, July 2023, btad415,

<https://doi.org/10.1093/bioinformatics/btad415>

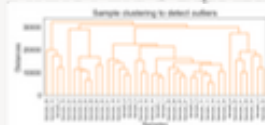
Published: 03 July 2023 Article history ▼

1) PyWGCNA object



2) Preprocessing

1. removed low expressed genes
2. removed samples with too many missing values
3. removed outliers sample by clustering



3) Find modules

1. calculating correlation matrix

	Gene A	Gene B	Gene C	Gene D	Gene E
Gene A	0	0.98	1	0.31	0.31
Gene B	0.98	0	0.98	0.22	0.22
Gene C	1	0.98	0	0.31	0.31
Gene D	0.31	0.22	0.31	0	0.99
Gene E	0.31	0.22	0.31	0.99	0

2. calculating adjacency matrix

$|cor(t(matrix))|^{\beta}$, $\beta = \text{high power}$

	Gene A	Gene B	Gene C	Gene D	Gene E
Gene A	0	0.90	1	0	0
Gene B	0.90	0	0.90	0	0
Gene C	1	0.90	0	0	0
Gene D	0	0	0	0	0.98
Gene E	0	0	0	0.98	0

3. calculating dissimilarity matrix and hierarchical clustering

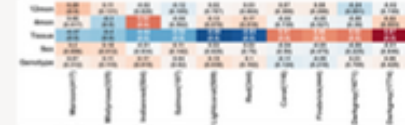


4. Identifying gene modules



4) Study modules

1. module trait relationship heatmap



2. module eigen-gene



3. module GO term



4. module network

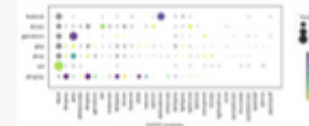


5) Compare PyWGCNAs

1. comparison table

IsxP4D	IsxP4D	IsxP4D_xxx	IsxP4D_xxx	fraction(%)	P_value
black	dimgray	6746	5080	97	1.5e-0006
black	black	6746	8157	2457	30.611743
black	dimgray	6746	2988	180	5.39e-071
black	darkgray	6746	479	44	9.18e-004
black	darkgray	6746	907	40	7.88e-046

2. confusion matrix



\$ pip install pyWGCNA