Study of the operation of algorithms (Kruskal algorithm, Floyd-Warshall algorithm)

```
Import needed libraries
import random
import networkx as nx
import matplotlib.pyplot as plt
from itertools import combinations, groupby
import time
from tqdm import tqdm
import math
from typing import List, Tuple
Graph generation function
# You can use this function to generate a random graph with
'num of nodes' nodes
# and 'completeness' probability of an edge between any two nodes
# If 'directed' is True, the graph will be directed
# If 'draw' is True, the graph will be drawn
def gnp random connected graph(num of nodes: int,
                                completeness: int,
                                directed: bool = False,
                                draw: bool = False):
    Generates a random graph, similarly to an Erdős-Rényi
    graph, but enforcing that the resulting graph is conneted (in case
of undirected graphs)
    if directed:
        G = nx.DiGraph()
    else:
        G = nx.Graph()
    edges = combinations(range(num of nodes), 2)
    G.add nodes from(range(num of nodes))
    for , node edges in groupby(edges, key = lambda x: x[0]):
        node edges = list(node edges)
        random edge = random.choice(node edges)
        if random.random() < 0.5:</pre>
            random edge = random edge[::-1]
        G.add_edge(*random edge)
        for e in node edges:
            if random.random() < completeness:</pre>
                G.add edge(*e)
    for (u,v,w) in G.edges(data=True):
```

```
w['weight'] = random.randint(-5, 20)
if draw:
    plt.figure(figsize=(10,6))
    if directed:
        # draw with edge weights
        pos = nx.arf layout(G)
        nx.draw(G,pos, node color='lightblue',
                with labels=True,
                node size=500,
                arrowsize=20,
                arrows=True)
        labels = nx.get edge attributes(G,'weight')
        nx.draw networkx edge labels(G, pos,edge labels=labels)
    else:
        nx.draw(G, node color='lightblue',
            with_labels=True,
            node size=500)
return G
```

Kruskal algorithm

Task: write a function that will implement Kruskal's algorithm taking as an argument a graph generated using the module provided in the task (function gnp_random_connected_graph). Then compare developed algorithm with native one from imported library

```
#generate random graph
G kruskal = qnp random connected qraph(10, 1, False, False)
def kruskals algorithm(graph edges):
    Kruskal's algorithm impemetation
    sorted graph = sorted(graph edges, key= lambda x:
x[2].get('weight'))
    connected_nodes_groups = {}
    already connected nodes = []
    min_frame = []
    for edge in sorted graph:
        if not edge[0] in already connected nodes or not edge[1] in
already connected nodes: # checking if one of the nodes are not
connected
            if not edge[0] in already connected nodes and not edge[1]
in already connected nodes: # checking for both
                connected nodes groups[edge[0]] = [edge[0], edge[1]] #
connect two nodes with each other
```

```
connected nodes groups[edge[1]] = [edge[\theta], edge[1]]
             else:
                 if not connected nodes groups.get(edge[0]):
                      connected nodes groups[edge[1]].append(edge[0])
                      connected nodes groups[edge[0]] =
connected_nodes_groups[edge[1]]
                 else:
                      connected nodes groups[edge[0]].append(edge[1])
                      connected nodes groups[edge[1]] =
connected nodes groups[edge[0]]
             min frame.append(edge)
             already connected nodes.append(edge[0])
             already_connected_nodes.append(edge[1])
    for i in sorted graph:
        if i[0] in connected nodes groups[i[0]] and i[1] not in
connected nodes groups[i[0]]:
             min frame.append(i)
    return min_frame
Here you can see an example of using created Kruskal algorithm. Function returns List[List[int]]
(matrix). Full algorithm is implemented in kruskal_algorithm() function. It takes one argument -
List[Tuple[int, int, dict]] (graph edges list with information about weight as a dict). Function
returns List[Tuple[int, int, dict]] - minimum carcass for input graph
kruskals algorithm(list(G kruskal.edges(data=True)))
[(5, 8, {'weight': -5}),
 (7, 8, {'weight': -5}),
 (2, 8, {'weight': -4}),
 (4, 9, {'weight': -4}),
 (0, 4, {\text{weight': -3}}),
 (3, 6, {'weight': -2}),
 (1, 7, {\text{weight'}: -1}),
 (0, 8, {'weight': -2}),
 (6, 9, {'weight': -2}),
 (5, 9, {'weight': -1}),
 (1, 3, {'weight': 1}),
 (4, 7, {'weight': 1}),
 (0, 5, {'weight': 2}),
 (3, 8, {'weight': 2}),
 (0, 6, {'weight': 3}),
 (3, 7, {'weight': 4}),
 (4, 6, {'weight': 4}),
 (5, 6, {'weight': 4}),
 (6, 8, {'weight': 4}),
 (7, 9, {'weight': 4}),
 (3, 5, {\text{'weight': 6}}),
 (8, 9, {'weight': 6}),
 (6, 7, {'weight': 7}),
 (1, 4, {'weight': 8}),
 (4, 5, {'weight': 8}),
```

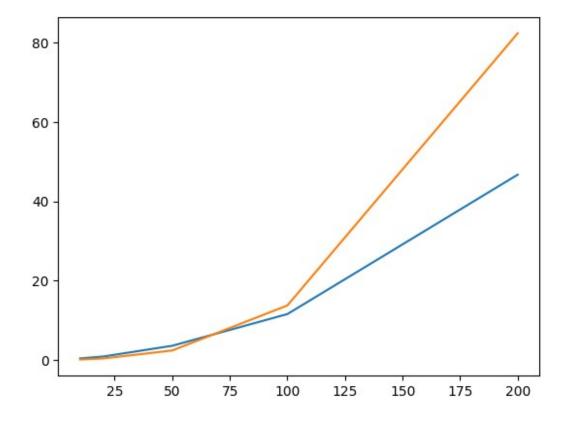
```
(4, 8, {'weight': 9}),
 (0, 3, {'weight': 10}),
 (0, 1, {'weight': 12}),
 (1, 6, {'weight': 13}),
 (0, 2, {'weight': 14}),
 (2, 3, {'weight': 15}),
 (3, 4, {'weight': 15}),
 (1, 9, {'weight': 17}),
 (2, 6, {'weight': 18}),
 (2, 4, {'weight': 18}),
 (0, 7, {'weight': 19}),
 (3, 9, {'weight': 19}),
 (5, 7, {'weight': 19}),
 (2, 9, {'weight': 20})]
Kruskal algrorithm time measuring
from networkx.algorithms import tree
nodes kruskal = [10, 20, 50, 100, 200]
NUM OF ITERATIONS = 1000
time taken imported = 0
kruskal native algorithm = []
#For imported kruskal algorithm for 10, 20, 50, 100 and 200 nodes
graph
for node in nodes kruskal:
    for i in tgdm(range(NUM OF ITERATIONS)):
        graph for imported = gnp random connected graph(node, 0.4,
False)
        #measure time while algorithm is working
        start = time.time() #start
        tree.minimum_spanning_tree(graph_for_imported,
algorithm="kruskal") #algorithm is working
        end = time.time() #end
        time taken imported += end - start
    kruskal native algorithm.append(time taken imported)
    time taken imported / NUM OF ITERATIONS
                 1000/1000 [00:00<00:00, 2052.22it/s]
100%|
100%|
                 1000/1000 [00:00<00:00, 1026.31it/s]
100%|
                 1000/1000 [00:05<00:00, 185.72it/s]
100%|
                 1000/1000 [00:17<00:00, 55.65it/s]
100%||
               | 1000/1000 [01:09<00:00, 14.30it/s]
time taken created = 0
kruskal own algorithm = []
#For created kruskal algorithm for 10, 20, 50, 100 and 200 nodes graph
for node in nodes kruskal:
    for i in tqdm(range(NUM OF ITERATIONS)):
        geaph_for_created = gnp random connected graph(node, 0.4,
False)
```

```
#measure time while algorithm is working
        start = time.time() #start
        kruskals algorithm(geaph for created.edges(data=True))
#algorithm is working
        end = time.time() #end
        time taken created += end - start
    kruskal own algorithm.append(time taken created)
    time taken created / NUM OF ITERATIONS
100%|
                 1000/1000 [00:00<00:00, 3224.02it/s]
100%
                 1000/1000 [00:00<00:00, 1282.23it/s]
100%|
                 1000/1000 [00:04<00:00, 218.55it/s]
                 1000/1000 [00:21<00:00, 47.23it/s]
100%|
                 1000/1000 [01:40<00:00, 9.93it/s]
100%|
```

Here you can see comparing native and created algorithms and a graph that shows difference in time

#comparing own and native kruskal algorithms by plotting
plt.plot(nodes kruskal, kruskal native algorithm)

plt.plot(nodes_kruskal, kruskal_own_algorithm)
plt.show()



As we see, created algorithm is a little faster than native when graph nodex < 100 and slower than native when nodex > 100. This is because the native algorithm is optimally written by professionals

Floyd-Warshall Algorithm

Task: write a function that will implement Floyd-Warshall's algorithm taking as an argument a graph generated using the module provided in the task (function gnp_random_connected_graph). Then compare developed algorithm with native one from imported library

```
G floyd = gnp random connected graph(7, 0.001, False, False)
G floyd.edges(data=True)
EdgeDataView([(0, 1, {\text{weight': -2}}), (1, 2, {\text{weight': 3}}), (2, 5, 
{'weight': 1}), (3, 5, {'weight': 12}), (4, 6, {'weight': -5}), (5, 6,
{'weight': 4})])
def create_table_for_floyd(graph, is_graph_directed):
    Function that creates table for
    floyd algorithm using generated
    graph
    N = len(graph.nodes)
    table = [[math.inf for in range(N)] for in range(N)]
    for x, y, w dict in list(graph.edges(data=True)):
        table[x][x] = 0
        table[y][y] = 0
        if is graph directed:
            table[x][y] = w_dict['weight']
        else:
            table[x][y] = w dict['weight']
            table[y][x] = w_dict['weight']
    return table
def floydworshal_algorithm(graph, is_graph_directed, table = None):
    Floyd-Warshall algorithm implementation
    Args:
        graph: format from generation function - graph
        nodes: int - the sum of nodes
        is graph directred: bool - is graph directed?
    Returns:
    if table == None:
```

```
table = create table for floyd(graph, is graph directed)
    N = len(graph.nodes)
    for k in range(N):
        for i in range(N):
             for j in range(N):
                 d = table[i][k] + table[k][j]
                 if i == j and d < 0:
                     return "Negative cycle detected!"
                 if table[i][j] > d:
                     table[i][i] = d
    return table
Here you can see an example of using created Floyd-Warshall algorithm. Function returns
List[List[int]] (matrix). Firstly function create table for floyd() creates the table for algorithm
using generated graph and then function floydworshall algorithm() implements algorithm
floydworshal algorithm(G floyd, False)
'Negative cycle detected!'
FLoyd-Warshall algrorithm time measuring
from networkx.algorithms import
floyd warshall predecessor and distance
nodes for floyd = [10, 20, 50]
NUM \overline{OF} ITERATIONS = 1000
time taken imported = 0
floyd native algorithm = []
#For imported kruskal algorithm for 10, 20, 50 nodes graph
for node in nodes for floyd:
    for i in tqdm(range(NUM OF ITERATIONS)):
        graph for imported = gnp random connected graph(node, 0.01,
False)
        #measure time while algorithm is working
        start = time.time() #start
        try:
             pred, dist =
floyd warshall predecessor and distance(graph for imported) #algorithm
is working
        except:
             pass
        end = time.time() #end
        time taken imported += end - start
    floyd_native_algorithm.append(time_taken imported)
    time_taken_imported / NUM_OF_ITERATIONS
                  1000/1000 [00:00<00:00, 1790.11it/s]
100%|
                  1000/1000 [00:03<00:00, 332.19it/s]
100%|
```

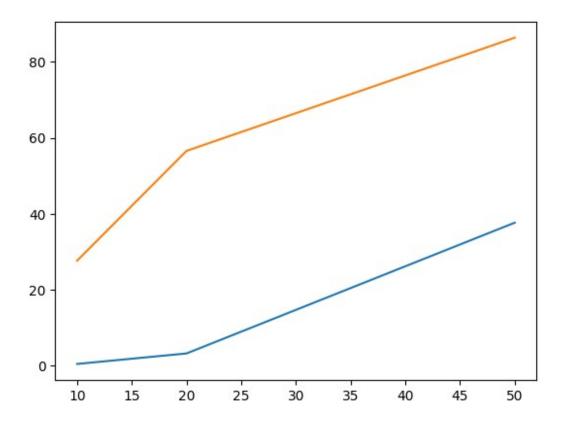
| 1000/1000 [00:35<00:00, 28.05it/s]

100%|

```
time taken created = 0
floyd own algorithm = []
#For created floyd algorithm for 10, 20, 50 nodes graph
for node in nodes for floyd:
    for i in tqdm(range(NUM OF ITERATIONS)):
        geaph for created = gnp random connected graph(node, 0.01,
False)
        #measure time while algorithm is working
        start = time.time() #start
        floydworshal algorithm(graph for imported, True)
        end = time.time() #end
        time taken created += end - start
    floyd own algorithm.append(time taken created)
    time taken created / NUM OF ITERATIONS
100%
                 1000/1000 [00:28<00:00, 35.64it/s]
100%|
                 1000/1000 [00:29<00:00, 33.93it/s]
100%|
                 1000/1000 [00:31<00:00, 32.15it/s]
```

Here you can see comparing native and created algorithms and a graph that shows difference in time

```
plt.plot(nodes_for_floyd, floyd_native_algorithm)
plt.plot(nodes_for_floyd, floyd_own_algorithm)
plt.show()
```



As you can see, created algorithm is always much slower than native one. This is because the native algorithm is optimally written by professionals

Conslusion

As we can see we are able to create our own algorithms, but they will not be so fast as we want. So, instead of creating, we can use built-in algorithms. Or, also, it is possible to raise programming skills and create as fast created algorithms as native ones.