

```
import tensorflow as tf
tf.test.gpu_device_name()

from google.colab import drive
drive.mount('/content/drive/')

Mounted at /content/drive/

import networkx as nx
G = nx.read_edgelist("/content/drive/MyDrive/Colab Notebooks/facebook_combined.txt", nodetype=int)
```



Complete network using networkx library. Following Parameters we have of in all the Hops

1. degree of the node
2. sum of degrees of the neighbours (depending on the steps / hops)
3. Common connections in between neighbours
4. Average Degree Connectivity of the Node
5. Average Degree Connectivity of the Node's Network (hop step)
6. bridge count described as nodes which are end points which are not with any other node. (hop step)
7. K core components of the Node
8. K core components of the Node's Network (hop step)
9. Onion Layers of the Node
10. Onion Layers of the Node's Network (hop step)
11. Boundary expansion of the Node netwrok (hop step)
12. eccentricity of a node v is the maximum distance from v to all other nodes in G
13. Kemeny constant - measures the time needed for spreading across a graph
14. global efficiency of the Node's Network (hop step)
15. max\_weight\_matching
16. random\_probability -
17. Number of Triangles in the Network

Based on this data i have done 4 types of clustering

- \* K Means
- \* Heiarchial (Agglomeretic ) clustering
  
- \* DBScan
- \* Network X Clustering

For Score Calculation of metrics i have compared it to the NetworkX Clustering

Finally i have tried community detection through various algorithms

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```
print(G.degree(1))
```

```
#all nodes and node degree in one frame for two hop
#node data
import pandas as pd
nodeData = []
for i in range(len(G)):
    ndData = []

    # Initialize deg outside the loop to avoid overwriting
    deg = G.degree[i]
    node_data = [i, deg]
    ndData.append(node_data)

    #extend it to the node data
    nodeData.extend(ndData)
```

```
# Define column names
columns = ['Node', 'NodeDegree']
```

```
# Create a DataFrame
nodedf = pd.DataFrame(nodeData, columns=columns)
```

```
# Display the DataFrame
print(nodedf)
```

	Node	NodeDegree
0	0	347
1	1	17
2	2	10
3	3	17
4	4	10
...	...	...
4034	4034	2
4035	4035	1
4036	4036	2
4037	4037	4
4038	4038	9

```
[4039 rows x 2 columns]
```

common connections between neighbours of a node

```
import pandas as pd
import networkx as nx
from itertools import combinations

onehopcommonst_data = []

for node in range(len(G)):
    abData = []
    acData = sorted(nx.all_neighbors(G, node))

    # Use itertools.combinations to iterate over unique pairs of elements in acData
    for neighbor1, neighbor2 in combinations(acData, 2):
        # Check if the common neighbors have 1 unit distance with the original nodes
        N = nx.shortest_path_length(G, source=neighbor1, target=neighbor2)
```

```
# Count only when the shortest distance is 1
if N == 1:
    abData.append(1)
```

```
onehopcommonst_data.append(sum(abData)) # Append the sum of abData for the current node
```

```
# Create a DataFrame from the list of sums
onehopcommondf = pd.DataFrame(onehopcommonst_data, columns=["Common Connections"])
```

```
# Display the DataFrame
print(onehopcommondf)
```

	Common Connections
0	2519
1	57
2	40
3	86

```

4           39
...
4034        1
4035        0
4036        1
4037        4
4038       20

```

[4039 rows x 1 columns]

degree of the neighbours and eventually counting the number of suppose counting the number of 1 degree neighbours in the network

```

#near degree data
onedegree_data = []

for node in range(len(G)):
    abData = []
    acData = sorted(nx.all_neighbors(G, node))

    for j in acData:
        M = G.degree(j)
        abData.append(M)

    onedegree_data.append(abData) # Append a new list as a row to all_data

# Create a DataFrame from the list of lists
onehopdegreedf = pd.DataFrame(onedegree_data, columns= [f'{i} degree connections' for i in range(1, 1046)])

# Display the DataFrame
print(onehopdegreedf)

# Assuming you have a list of 100 values named values_to_count
values_to_count = range(1046) # Replace ... with the actual values

# Add columns for each value in values_to_count
for value in values_to_count:
    column_name = f'{value}_count'
    onehopdegreedf[column_name] = onehopdegreedf.apply(lambda row: row.eq(value).sum(), axis=1)

print(onehopdegreedf)

```

	1 degree connections	2 degree connections	3 degree connections	\
0	17	10.0	17.0	
1	347	22.0	31.0	
2	347	15.0	21.0	
3	347	57.0	69.0	
4	347	9.0	5.0	
...	...	...	...	
4034	59	18.0	NaN	
4035	59	NaN	NaN	
4036	59	4.0	NaN	
4037	59	14.0	12.0	
4038	59	6.0	10.0	
	4 degree connections	5 degree connections	6 degree connections	\
0	10.0	13.0	6.0	
1	8.0	10.0	20.0	
2	17.0	14.0	14.0	
3	68.0	76.0	24.0	
4	10.0	9.0	9.0	
...	...	...	...	
4034	NaN	NaN	NaN	
4035	NaN	NaN	NaN	
4036	NaN	NaN	NaN	
4037	8.0	NaN	NaN	
4038	6.0	12.0	8.0	
	7 degree connections	8 degree connections	9 degree connections	\
0	20.0	8.0	57.0	
1	21.0	62.0	7.0	
2	26.0	19.0	8.0	
3	14.0	63.0	43.0	
4	9.0	10.0	9.0	
...	...	...	...	
4034	NaN	NaN	NaN	
4035	NaN	NaN	NaN	
4036	NaN	NaN	NaN	
4037	NaN	NaN	NaN	
4038	18.0	7.0	11.0	

```
10 degree connections ... 1036 degree connections \
0          10.0 ...
1          18.0 ...
2          18.0 ...
3          46.0 ...
4           9.0 ...
...
4034        NaN ...
4035        NaN ...
4036        NaN ...
4037        NaN ...
4038        NaN ...
```

```
1037 degree connections 1038 degree connections \
0            NaN           NaN
1            NaN           NaN
2            NaN           NaN
3            NaN           NaN
4            NaN           NaN
```

```
df_dropped = onehopdegreedf.drop(onehopdegreedf.columns[:1045], axis=1)
print(df_dropped)
```

```
#dropping columns in degreedf
df_dropped = df_dropped.drop(df_dropped.columns[793:1045], axis=1)
df_dropped = df_dropped.drop(df_dropped.columns[756:792], axis=1)
df_dropped = df_dropped.drop(df_dropped.columns[548:755], axis=1)
df_dropped = df_dropped.drop(df_dropped.columns[348:547], axis=1)
df_dropped = df_dropped.drop(df_dropped.columns[295:347], axis=1)
df_dropped = df_dropped.drop(df_dropped.columns[292:294], axis=1)
df_dropped = df_dropped.drop(df_dropped.columns[255:291], axis=1)
df_dropped = df_dropped.drop(df_dropped.columns[246:254], axis=1)
df_dropped = df_dropped.drop(df_dropped.columns[236:245], axis=1)
df_dropped = df_dropped.drop(df_dropped.columns[232:234], axis=1)
df_dropped = df_dropped.drop(df_dropped.columns[230:231], axis=1)
df_dropped = df_dropped.drop(df_dropped.columns[227:229], axis=1)
df_dropped = df_dropped.drop(df_dropped.columns[225:226], axis=1)
# Display the DataFrame after dropping columns
print(df_dropped)
```

	0_count	1_count	2_count	3_count	4_count	5_count	6_count	7_count	...
0	0	14	27	22	17	12	15	18	
1	0	0	0	0	0	0	0	1	
2	0	0	0	0	0	0	0	0	
3	0	0	0	1	0	1	0	0	
4	0	0	0	0	0	1	0	0	
...	...	...	...	...	...	...	...	...	...
4034	0	0	0	0	0	0	0	0	0
4035	0	0	0	0	0	0	0	0	0
4036	0	0	0	0	1	0	0	0	0
4037	0	0	0	0	0	0	0	0	0
4038	0	0	0	0	0	0	2	1	
	8_count	9_count	...	235_count	245_count	254_count	291_count	...	
0	15	15	...	0	0	0	0	0	
1	1	0	...	0	0	0	0	0	
2	1	0	...	0	0	0	0	0	
3	0	0	...	0	0	0	0	0	
4	0	6	...	0	0	0	0	0	
...	...	...	...	...	...	...	...	...	...
4034	0	0	...	0	0	0	0	0	
4035	0	0	...	0	0	0	0	0	
4036	0	0	...	0	0	0	0	0	
4037	1	0	...	0	0	0	0	0	
4038	1	0	...	0	0	0	0	0	
	294_count	347_count	547_count	755_count	792_count	1045_count			
0	0	0	0	0	0	1			
1	0	1	0	0	0	0			
2	0	1	0	0	0	0			
3	0	1	0	0	0	0			
4	0	1	0	0	0	0			
...	...	...	...	...	...	...			
4034	0	0	0	0	0	0			
4035	0	0	0	0	0	0			
4036	0	0	0	0	0	0			
4037	0	0	0	0	0	0			
4038	0	0	0	0	0	0			

[4039 rows x 239 columns]

```

import pandas as pd
A = nx.average_neighbor_degree(G)
B = nx.node_degree_xy(G)
print(A)
average_degrees = [round(value, 2) for value in A.values()]

# Create a DataFrame with a single column
adcdf = pd.DataFrame({'Average Degree Connectivity for Node': average_degrees})

# Display the DataFrame
print(adcdf)

{0: 18.959654178674352, 1: 48.23529411764706, 2: 49.9, 3: 59.76470588235294, 4: 42.6, 5: 50.61538461538461, 6: 63.5, 7: 45.9, 8: 48.375,
   Average Degree Connectivity for Node
0                      18.96
1                      48.24
2                      49.90
3                      59.76
4                      42.60
...
4034                   ...
4035                   38.50
4035                   59.00
4036                   31.50
4037                   23.25
4038                   15.22

[4039 rows x 1 columns]

```



```

import pandas as pd
import networkx as nx

dcData = []

for node in G.nodes():
    neighbors = list(G.neighbors(node))

    # Include the node itself in the subgraph
    subgraph = G.subgraph([node] + neighbors)

    # Calculate the average neighbor degree for the subgraph
    average_neighbor_degree = nx.average_neighbor_degree(subgraph)

    # Round the values to 2 decimal places
    average_degrees = {k: round(v, 2) for k, v in average_neighbor_degree.items()}

    # Sum the average degrees
    B = sum(average_degrees.values())

    dcData.append(B)

# Create a DataFrame with a single column
oneadcdf = pd.DataFrame({'Average Degree Connectivity for network': dcData})

# Display the DataFrame
print(oneadcdf)

```

```

   Average Degree Connectivity for network
0                      28959.92
1                      183.50
2                      101.99
3                      232.98
4                      101.84
...
4034                   ...
4035                   6.00
4035                   2.00
4036                   6.00
4037                   17.34
4038                   68.15

```

[4039 rows x 1 columns]

```

node_bridge_counts = {}

# Iterate over nodes from 0 to (number of nodes - 1)
for node in range(len(G)):
    # Create a subgraph containing only the current node and its neighbors

```

```
subgraph = G.subgraph([node] + list(G.neighbors(node)))

# Calculate local bridges for the subgraph
local_bridges = list(nx.local_bridges(subgraph, with_span=False))

# Count the number of bridges for the current node
bridge_count = len(local_bridges)

# Store the result in the dictionary
node_bridge_counts[node] = bridge_count

# Convert the dictionary to a DataFrame
onehopbrdf = pd.DataFrame(list(node_bridge_counts.values()), columns=['Bridge_Count'])

# Display the DataFrame
print(onehopbrdf)

   Bridge_Count
0            14
1             0
2             0
3             0
4             0
...
4034          0
4035          1
4036          0
4037          0
4038          0

[4039 rows x 1 columns]

# Returns the core number for each vertex.

# A k-core is a maximal subgraph that contains nodes of degree k or more.

# The core number of a node is the largest value k of a k-core containing that node.
C = nx.core_number(G)
print(C)

coredf = pd.DataFrame(list(C.values()), columns=['Core_Number'])

# Display the DataFrame
print(coredf)

{0: 21, 1: 13, 2: 9, 3: 13, 4: 9, 5: 10, 6: 5, 7: 12, 8: 5, 9: 21, 10: 10, 11: 1, 12: 1, 13: 21, 14: 10, 15: 1, 16: 9, 17: 9, 18: 1, 19:
   Core_Number
0            21
1            13
2             9
3            13
4             9
...
4034           2
4035           1
4036           2
4037           4
4038           5

[4039 rows x 1 columns]
```

```

onekcore_counts = []

# Iterate over nodes from 0 to (number of nodes - 1)
for node in G.nodes():
    # Create a subgraph containing only the current node and its neighbors
    subgraph = G.subgraph([node] + list(G.neighbors(node)))

    # Calculate local bridges for the subgraph
    k_core = nx.core_number(subgraph)
    B = sum(list(k_core.values()))

    onekcore_counts.append(B)

# Convert the list to a DataFrame
onehopkcoredf = pd.DataFrame({'k_core for network': onekcore_counts})

# Display the DataFrame
print(onehopkcoredf)

      k_core for network
0                  3400
1                   106
2                    86
3                   170
4                   94
...
4034                  ...
4035                   2
4036                   6
4037                  14
4038                  44

[4039 rows x 1 columns]

# Returns the layer of each vertex in an onion decomposition of the graph.

# The onion decomposition refines the k-core decomposition by providing information on the internal organization of each k-shell.
# It is usually used alongside the core numbers.
O = nx.onion_layers(G)
print(O)
sorted_keys = sorted(O.keys())

# Create a DataFrame from the dictionary with sorted keys
oniondf = pd.DataFrame.from_dict({key: O[key] for key in sorted_keys}, orient='index', columns=['onion layers'])

# Display the DataFrame
print(oniondf)

{11: 1, 12: 1, 15: 1, 18: 1, 37: 1, 43: 1, 74: 1, 114: 1, 209: 1, 210: 1, 215: 1, 287: 1, 292: 1, 335: 1, 911: 1, 918: 1, 1096: 1, 1119: 1, 1120: 1, 1121: 1, 1122: 1, 1123: 1, 1124: 1, 1125: 1, 1126: 1, 1127: 1, 1128: 1, 1129: 1, 1130: 1, 1131: 1, 1132: 1, 1133: 1, 1134: 1, 1135: 1, 1136: 1, 1137: 1, 1138: 1, 1139: 1, 1140: 1, 1141: 1, 1142: 1, 1143: 1, 1144: 1, 1145: 1, 1146: 1, 1147: 1, 1148: 1, 1149: 1, 1150: 1, 1151: 1, 1152: 1, 1153: 1, 1154: 1, 1155: 1, 1156: 1, 1157: 1, 1158: 1, 1159: 1, 1160: 1, 1161: 1, 1162: 1, 1163: 1, 1164: 1, 1165: 1, 1166: 1, 1167: 1, 1168: 1, 1169: 1, 1170: 1, 1171: 1, 1172: 1, 1173: 1, 1174: 1, 1175: 1, 1176: 1, 1177: 1, 1178: 1, 1179: 1, 1180: 1, 1181: 1, 1182: 1, 1183: 1, 1184: 1, 1185: 1, 1186: 1, 1187: 1, 1188: 1, 1189: 1, 1190: 1, 1191: 1, 1192: 1, 1193: 1, 1194: 1, 1195: 1, 1196: 1, 1197: 1, 1198: 1, 1199: 1, 1200: 1, 1201: 1, 1202: 1, 1203: 1, 1204: 1, 1205: 1, 1206: 1, 1207: 1, 1208: 1, 1209: 1, 1210: 1, 1211: 1, 1212: 1, 1213: 1, 1214: 1, 1215: 1, 1216: 1, 1217: 1, 1218: 1, 1219: 1, 1220: 1, 1221: 1, 1222: 1, 1223: 1, 1224: 1, 1225: 1, 1226: 1, 1227: 1, 1228: 1, 1229: 1, 1230: 1, 1231: 1, 1232: 1, 1233: 1, 1234: 1, 1235: 1, 1236: 1, 1237: 1, 1238: 1, 1239: 1, 1240: 1, 1241: 1, 1242: 1, 1243: 1, 1244: 1, 1245: 1, 1246: 1, 1247: 1, 1248: 1, 1249: 1, 1250: 1, 1251: 1, 1252: 1, 1253: 1, 1254: 1, 1255: 1, 1256: 1, 1257: 1, 1258: 1, 1259: 1, 1260: 1, 1261: 1, 1262: 1, 1263: 1, 1264: 1, 1265: 1, 1266: 1, 1267: 1, 1268: 1, 1269: 1, 1270: 1, 1271: 1, 1272: 1, 1273: 1, 1274: 1, 1275: 1, 1276: 1, 1277: 1, 1278: 1, 1279: 1, 1280: 1, 1281: 1, 1282: 1, 1283: 1, 1284: 1, 1285: 1, 1286: 1, 1287: 1, 1288: 1, 1289: 1, 1290: 1, 1291: 1, 1292: 1, 1293: 1, 1294: 1, 1295: 1, 1296: 1, 1297: 1, 1298: 1, 1299: 1, 1300: 1, 1301: 1, 1302: 1, 1303: 1, 1304: 1, 1305: 1, 1306: 1, 1307: 1, 1308: 1, 1309: 1, 1310: 1, 1311: 1, 1312: 1, 1313: 1, 1314: 1, 1315: 1, 1316: 1, 1317: 1, 1318: 1, 1319: 1, 1320: 1, 1321: 1, 1322: 1, 1323: 1, 1324: 1, 1325: 1, 1326: 1, 1327: 1, 1328: 1, 1329: 1, 1330: 1, 1331: 1, 1332: 1, 1333: 1, 1334: 1, 1335: 1, 1336: 1, 1337: 1, 1338: 1, 1339: 1, 1340: 1, 1341: 1, 1342: 1, 1343: 1, 1344: 1, 1345: 1, 1346: 1, 1347: 1, 1348: 1, 1349: 1, 1350: 1, 1351: 1, 1352: 1, 1353: 1, 1354: 1, 1355: 1, 1356: 1, 1357: 1, 1358: 1, 1359: 1, 1360: 1, 1361: 1, 1362: 1, 1363: 1, 1364: 1, 1365: 1, 1366: 1, 1367: 1, 1368: 1, 1369: 1, 1370: 1, 1371: 1, 1372: 1, 1373: 1, 1374: 1, 1375: 1, 1376: 1, 1377: 1, 1378: 1, 1379: 1, 1380: 1, 1381: 1, 1382: 1, 1383: 1, 1384: 1, 1385: 1, 1386: 1, 1387: 1, 1388: 1, 1389: 1, 1390: 1, 1391: 1, 1392: 1, 1393: 1, 1394: 1, 1395: 1, 1396: 1, 1397: 1, 1398: 1, 1399: 1, 1400: 1, 1401: 1, 1402: 1, 1403: 1, 1404: 1, 1405: 1, 1406: 1, 1407: 1, 1408: 1, 1409: 1, 1410: 1, 1411: 1, 1412: 1, 1413: 1, 1414: 1, 1415: 1, 1416: 1, 1417: 1, 1418: 1, 1419: 1, 1420: 1, 1421: 1, 1422: 1, 1423: 1, 1424: 1, 1425: 1, 1426: 1, 1427: 1, 1428: 1, 1429: 1, 1430: 1, 1431: 1, 1432: 1, 1433: 1, 1434: 1, 1435: 1, 1436: 1, 1437: 1, 1438: 1, 1439: 1, 1440: 1, 1441: 1, 1442: 1, 1443: 1, 1444: 1, 1445: 1, 1446: 1, 1447: 1, 1448: 1, 1449: 1, 1450: 1, 1451: 1, 1452: 1, 1453: 1, 1454: 1, 1455: 1, 1456: 1, 1457: 1, 1458: 1, 1459: 1, 1460: 1, 1461: 1, 1462: 1, 1463: 1, 1464: 1, 1465: 1, 1466: 1, 1467: 1, 1468: 1, 1469: 1, 1470: 1, 1471: 1, 1472: 1, 1473: 1, 1474: 1, 1475: 1, 1476: 1, 1477: 1, 1478: 1, 1479: 1, 1480: 1, 1481: 1, 1482: 1, 1483: 1, 1484: 1, 1485: 1, 1486: 1, 1487: 1, 1488: 1, 1489: 1, 1490: 1, 1491: 1, 1492: 1, 1493: 1, 1494: 1, 1495: 1, 1496: 1, 1497: 1, 1498: 1, 1499: 1, 1500: 1, 1501: 1, 1502: 1, 1503: 1, 1504: 1, 1505: 1, 1506: 1, 1507: 1, 1508: 1, 1509: 1, 1510: 1, 1511: 1, 1512: 1, 1513: 1, 1514: 1, 1515: 1, 1516: 1, 1517: 1, 1518: 1, 1519: 1, 1520: 1, 1521: 1, 1522: 1, 1523: 1, 1524: 1, 1525: 1, 1526: 1, 1527: 1, 1528: 1, 1529: 1, 1530: 1, 1531: 1, 1532: 1, 1533: 1, 1534: 1, 1535: 1, 1536: 1, 1537: 1, 1538: 1, 1539: 1, 1540: 1, 1541: 1, 1542: 1, 1543: 1, 1544: 1, 1545: 1, 1546: 1, 1547: 1, 1548: 1, 1549: 1, 1550: 1, 1551: 1, 1552: 1, 1553: 1, 1554: 1, 1555: 1, 1556: 1, 1557: 1, 1558: 1, 1559: 1, 1560: 1, 1561: 1, 1562: 1, 1563: 1, 1564: 1, 1565: 1, 1566: 1, 1567: 1, 1568: 1, 1569: 1, 1570: 1, 1571: 1, 1572: 1, 1573: 1, 1574: 1, 1575: 1, 1576: 1, 1577: 1, 1578: 1, 1579: 1, 1580: 1, 1581: 1, 1582: 1, 1583: 1, 1584: 1, 1585: 1, 1586: 1, 1587: 1, 1588: 1, 1589: 1, 1590: 1, 1591: 1, 1592: 1, 1593: 1, 1594: 1, 1595: 1, 1596: 1, 1597: 1, 1598: 1, 1599: 1, 1600: 1, 1601: 1, 1602: 1, 1603: 1, 1604: 1, 1605: 1, 1606: 1, 1607: 1, 1608: 1, 1609: 1, 1610: 1, 1611: 1, 1612: 1, 1613: 1, 1614: 1, 1615: 1, 1616: 1, 1617: 1, 1618: 1, 1619: 1, 1620: 1, 1621: 1, 1622: 1, 1623: 1, 1624: 1, 1625: 1, 1626: 1, 1627: 1, 1628: 1, 1629: 1, 1630: 1, 1631: 1, 1632: 1, 1633: 1, 1634: 1, 1635: 1, 1636: 1, 1637: 1, 1638: 1, 1639: 1, 1640: 1, 1641: 1, 1642: 1, 1643: 1, 1644: 1, 1645: 1, 1646: 1, 1647: 1, 1648: 1, 1649: 1, 1650: 1, 1651: 1, 1652: 1, 1653: 1, 1654: 1, 1655: 1, 1656: 1, 1657: 1, 1658: 1, 1659: 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```

```

oneonion_counts = []

# Iterate over nodes from 0 to (number of nodes - 1)
for node in G.nodes():
    # Create a subgraph containing only the current node and its neighbors
    subgraph = G.subgraph([node] + list(G.neighbors(node)))

    # Calculate local bridges for the subgraph
    k_core = nx.onion_layers(subgraph)
    sorted_keys = sorted(k_core.values())

    B = sum(list(sorted_keys))

    oneonion_counts.append(B)

# Convert the list to a DataFrame
oneoniondf = pd.DataFrame({'onion layers for network': oneonion_counts})

# Display the DataFrame
print(oneoniondf)

```

	onion layers for network
0	8567
1	47
2	29
3	87
4	21
...	...
4034	3
4035	2
4036	3
4037	9
4038	31

[4039 rows x 1 columns]

#Returns the boundary expansion of the set S.

```
#The boundary expansion is the quotient of the size of the node boundary and the cardinality of S. [1]
oneboundary_expansion_list = []
```

```

# Iterate over all nodes in the graph
for node in G.nodes():
    # Create a subgraph with the current node and its neighbors
    subgraph = G.subgraph([node] + list(G.neighbors(node)))

    # Calculate the boundary expansion for the current node
    boundary_expansion = nx.boundary_expansion(G, subgraph)

    # Append the boundary expansion to the list
    oneboundary_expansion_list.append(boundary_expansion)

```

```
# Create a DataFrame from the list
oneboundddf = pd.DataFrame({'Boundary_Expansion': oneboundary_expansion_list})
```

```
# Display the DataFrame
print(oneboundddf)
```

	Boundary_Expansion
0	3.364943
1	18.333333
2	30.636364
3	18.333333
4	30.636364
...	...
4034	19.000000
4035	29.000000
4036	19.000000
4037	11.000000
4038	5.000000

[4039 rows x 1 columns]

#The eccentricity of a node v is the maximum distance from v to all other nodes in G.

```
ecc = nx.eccentricity(G, v=None, sp=None, weight=None)
eccdf = pd.DataFrame(list(ecc.values()), columns=['eccentricity'])
print(eccdf)
```

```
eccentricity
0           6
1           7
2           7
3           7
4           7
...
4034        8
4035        8
4036        8
4037        8
4038        8
```

[4039 rows x 1 columns]

```
#The Kemeny constant measures the time needed for spreading across a graph. Low values indicate a closely connected graph whereas high value
onekemeny_list = []
```

```
# Iterate over all nodes in the graph
for node in G.nodes():
    # Create a subgraph with the current node and its neighbors
    subgraph = G.subgraph([node] + list(G.neighbors(node)))

    # Calculate the boundary expansion for the current node
    kemeny = nx.kemeny_constant(subgraph)

    # Append the boundary expansion to the list
    onekemeny_list.append(kemeny)
```

```
# Create a DataFrame from the list
onekemenydf = pd.DataFrame({'kemeny constant for network': onekemeny_list})
```

```
# Display the DataFrame
print(onekemenydf)
```

```
kemeny constant for network
0           406.232390
1           17.493435
2           9.177039
3           16.649210
4           9.196660
...
4034        ...
4035        1.333333
4036        0.500000
4037        1.333333
4038        3.362500
4038        8.546052
```

[4039 rows x 1 columns]

```
#The average global efficiency of a graph is the average efficiency of all pairs of nodes
oneglobaleff_list = []
oneweighting_list = []
onerandom_list = []
panther_list = []
# Iterate over all nodes in the graph
for node in G.nodes():
    # Create a subgraph with the current node and its neighbors
    subgraph = G.subgraph([node] + list(G.neighbors(node)))

    # Calculate the boundary expansion for the current node
    globaleff = nx.global_efficiency(subgraph)

    weighteff = len(list(nx.max_weight_matching(subgraph, maxcardinality=False, weight='weight')))

    result = nx.non_randomness(subgraph, k=None, weight='weight')
    second_value = result[1]

    #panther = len(nx.panther_similarity(subgraph, 0, k=5, path_length=5, c=0.5, delta=0.1, eps=None, weight='weight'))

    # Append the boundary expansion to the list
    oneglobaleff_list.append(globaleff)
    oneweighting_list.append(weighteff)
    onerandom_list.append(second_value)
    #panther_list.append(panther)
# Create a DataFrame from the list
oneglobaldf = pd.DataFrame({
    'global_efficiency': oneglobaleff_list,
    'max_weight_matching': oneweighting_list,
    'random_probability': onerandom_list,
    # 'panther_similarity': panther_list # Uncomment if needed
})
# Display the DataFrame
print(oneglobaldf)

/usr/local/lib/python3.10/dist-packages/networkx/algorithms/non_randomness.py:94: RuntimeWarning: invalid value encountered in double_scalars
  nr_rd = (nr - ((n - 2 * k) * p + k)) / math.sqrt(2 * k * p * (1 - p))
/usr/local/lib/python3.10/dist-packages/networkx/algorithms/non_randomness.py:94: RuntimeWarning: divide by zero encountered in double_scalars
  nr_rd = (nr - ((n - 2 * k) * p + k)) / math.sqrt(2 * k * p * (1 - p))
   global_efficiency  max_weight_matching  random_probability
0           0.523734                  164      75.626117
1           0.741830                   9      0.814521
2           0.954545                  5      0.088836
3           0.836601                  9      1.395870
4           0.945455                  5      0.364554
...
4034        1.000000                  ...          ...
4035        1.000000                  1          NaN
4036        1.000000                  1          ...
4037        0.900000                  2      -0.135403
4038        0.822222                  5      0.266415

[4039 rows x 3 columns]
```

	global_efficiency	max_weight_matching	random_probability
0	0.523734	164	75.626117
1	0.741830	9	0.814521
2	0.954545	5	0.088836
3	0.836601	9	1.395870
4	0.945455	5	0.364554
...	...	...	...
4034	1.000000	1	-inf
4035	1.000000	1	NaN
4036	1.000000	1	-inf
4037	0.900000	2	-0.135403
4038	0.822222	5	0.266415

```
oneglcdf = oneglobaldf.fillna(0)
print(oneglcdf)

   global_efficiency  max_weight_matching  random_probability
0           0.523734                  164      75.626117
1           0.741830                   9      0.814521
2           0.954545                  5      0.088836
3           0.836601                  9      1.395870
4           0.945455                  5      0.364554
...
4034        1.000000                  ...          ...
4035        1.000000                  1          0.000000
4036        1.000000                  1          -inf
4037        0.900000                  2      -0.135403
4038        0.822222                  5      0.266415

[4039 rows x 3 columns]
```

```
import numpy as np
oneglcdf.replace([np.inf, -np.inf], 0, inplace=True)
print(oneglcdf)
```

	global_efficiency	max_weight_matching	random_probability
0	0.523734	164	75.626117
1	0.741830	9	0.814521
2	0.954545	5	0.088836
3	0.836601	9	1.395870
4	0.945455	5	0.364554
...	...	...	...
4034	1.000000	1	0.000000
4035	1.000000	1	0.000000
4036	1.000000	1	0.000000
4037	0.900000	2	-0.135403
4038	0.822222	5	0.266415

[4039 rows x 3 columns]

```
#dataset final
```

```
onehopdf = pd.concat([nodedf, df_dropped, onehopcommndf, adcdf, oneadcdf, onehopbrdf, coredf, onehopkcoredf, oniondf, oneoniondf, onebounddf])
print(onehopdf)
```

4035	4035	1	0	0	0	0	0
4036	4036	2	0	0	0	0	1
4037	4037	4	0	0	0	0	0
4038	4038	9	0	0	0	0	0

	6_count	7_count	...	Core_Number	k_core	for network	onion layers	\
0	15	18	...	21	3400		108	
1	0	1	...	13		106		51
2	0	0	...	9		86		30
3	0	0	...	13		170		49
4	0	0	...	9		94		30
...	...	...	...	...	...	...	...	...
4034	0	0	...	2		6		2
4035	0	0	...	1		2		1
4036	0	0	...	2		6		2
4037	0	0	...	4		14		8
4038	2	1	...	5		44		14

	onion layers	for network	Boundary_Expansion	\
0		8567	3.364943	
1		47	18.333333	
2		29	30.636364	
3		87	18.333333	
4		21	30.636364	
...	...	...	...	
4034		3	19.000000	
4035		2	29.000000	
4036		3	19.000000	
4037		9	11.000000	
4038		31	5.000000	

	kemeny constant	for network	global_efficiency	max_weight_matching	\
0	406.232390		0.523734	164	
1	17.493435		0.741830	9	
2	9.177039		0.954545	5	
3	16.649210		0.836601	9	
4	9.196660		0.945455	5	
...	...	...	...	...	
4034	1.333333		1.000000	1	
4035	0.500000		1.000000	1	
4036	1.333333		1.000000	1	
4037	3.362500		0.900000	2	
4038	8.546052		0.822222	5	

	random_probability	eccentricity
0	75.626117	6
1	0.814521	7
2	0.088836	7
3	1.395870	7
4	0.364554	7
...	...	...
4034	0.000000	8
4035	0.000000	8
4036	0.000000	8
4037	-0.135403	8
4038	0.266415	8

[4039 rows x 255 columns]

```
onehopdf.to_csv('/content/drive/MyDrive/Colab Notebooks/onehopdata.csv', index=False)
```

```
import pandas as pd
```

```
tri = nx.triangles(G, nodes=None)
tridf = pd.DataFrame(list(tri.values()), columns=['No of triangles with Node'])
print(tridf)
```

	No of triangles with Node
0	2519
1	57
2	40
3	86
4	39
...	...
4034	1
4035	0
4036	1
4037	4
4038	20

```
[4039 rows x 1 columns]
```

```
onetriangle_counts = []
```

```
# Iterate over nodes from 0 to (number of nodes - 1)
for node in G.nodes():
    # Create a subgraph containing only the current node and its neighbors
    subgraph = G.subgraph([node] + list(G.neighbors(node)))
```

```
# Calculate local bridges for the subgraph
triangle = nx.triangles(subgraph, nodes=None)
B = sum(list(triangle.values()))
```

```
onetriangle_counts.append(B)
```

```
# Convert the list to a DataFrame
onehopktriangledf = pd.DataFrame({'triangle for network': onetriangle_counts})
```

```
# Display the DataFrame
print(onehopktriangledf)
```

	triangle for network
0	39777
1	402
2	381
3	996
4	378
...	...
4034	3
4035	0
4036	3
4037	15
4038	114

```
[4039 rows x 1 columns]
```

```
path = '/content/drive/MyDrive/Colab Notebooks/onehopdata.csv'
onehop = pd.read_csv(path)
print(onehop)
```

```
#Final Dataset for Onehop step
onefinaldf = pd.concat([onehop, tridf, onehopktriangledf], axis=1)
print(onefinaldf)
```

4035	4035	1	0	0	0	0	0
4036	4036	2	0	0	0	1	0
4037	4037	4	0	0	0	0	0
4038	4038	9	0	0	0	0	0

6_count	7_count	...	onion layers	onion layers for network	\
0	15	18	...	108	8567
1	0	1	...	51	47

```

4038        2         1 ...      14      31
Boundary_Expansion kemény constant for network global_efficiency \
0           3.364943    406.232390   0.523734
1           18.333333   17.493435   0.741830
2           30.636364   9.177039   0.954545
3           18.333333   16.649210   0.836601
4           30.636364   9.196660   0.945455
...
...          ...
4034       19.000000   1.333333   1.000000
4035       29.000000   0.500000   1.000000
4036       19.000000   1.333333   1.000000
4037       11.000000   3.362500   0.900000
4038       5.000000    8.546052   0.822222

max_weight_matching random_probability eccentricity \
0           164        75.626117   6
1            9         0.814521   7
2            5         0.088836   7
3            9         1.395870   7
4            5         0.364554   7
...
...          ...
4034       1          0.000000   8
4035       1          0.000000   8
4036       1          0.000000   8
4037       2         -0.135403   8
4038       5         0.266415   8

No of triangles with Node triangle for network
0           2519        39777
1            57         402
2            40         381
3            86         996
4            39         378
...
...          ...
4034       1          3
4035       0          0
4036       1          3
4037       4          15
4038       20         114

```

[4039 rows x 257 columns]

K Means Clustering on the Data by determining the number of steps then plotting by the number of clusters which is determined by Elbow method and then plot accordingly

```

#determining the number of Steps for K means
import pandas as pd
from sklearn.cluster import KMeans
from sklearn.preprocessing import StandardScaler
import matplotlib.pyplot as plt

# Assuming df is your DataFrame with standardized features
# Step 1: Standardize the data
scaler = StandardScaler()
features_standardized = scaler.fit_transform(onefinaldf)

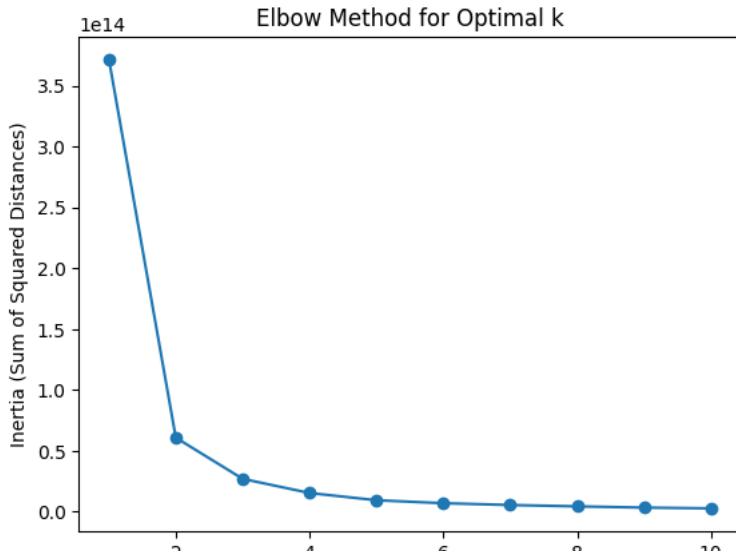
# Store the sum of squared distances for different values of k
inertia_values = []

# Define a range of k values to try
k_values = range(1, 11) # You can adjust the range based on your problem

# Apply k-means clustering for each k and store the inertia
for k in k_values:
    kmeans = KMeans(n_clusters=k, random_state=42)
    kmeans.fit(onefinaldf)
    inertia_values.append(kmeans.inertia_)

# Plot the elbow curve
plt.plot(k_values, inertia_values, marker='o')
plt.title('Elbow Method for Optimal k')
plt.xlabel('Number of Clusters (k)')
plt.ylabel('Inertia (Sum of Squared Distances)')
plt.show()

```



```
import pandas as pd
from sklearn.cluster import KMeans
from sklearn.preprocessing import StandardScaler
from sklearn.decomposition import PCA
import matplotlib.pyplot as plt

# Step 1: Standardize the data
scaler = StandardScaler()
features_standardized = scaler.fit_transform(onefinaldf)

# Step 3: Apply k-means clustering
k = 4 # Replace with your optimal k
kmeans = KMeans(n_clusters=k, random_state=42)
onefinaldf['k=4_Cluster'] = kmeans.fit_predict(features_standardized)

# Display the DataFrame with cluster assignments
print(onefinaldf)

# Step 4: Visualize the clusters (optional)
# This is just an example, adjust it based on your requirements
# Visualization is more challenging with a large number of features
# You may need to choose specific features for plotting or use advanced visualization techniques
plt.scatter(onefinaldf["NodeDegree"], onefinaldf['Common Connections'], c=onefinaldf['k=4_Cluster'], cmap='viridis')
plt.title('K-Means Clustering')
plt.show()
```

```
usr/local/lib/python3.10/dist-packages/sklearn/cluster/_kmeans.py:870: FutureWarning: The default value of `n_init` will change from
warnings.warn(
    Node  NodeDegree  0_count  1_count  2_count  3_count  4_count  5_count  \
    0      347       0       14      27      22      17      12
    1      17        0       0       0       0       0       0
    2      10        0       0       0       0       0       0
    3      17        0       0       0       0       1       0       1
    4      10        0       0       0       0       0       0       1
...
...     ...     ...     ...     ...     ...     ...
034  4034      2        0       0       0       0       0       0
035  4035      1        0       0       0       0       0       0
036  4036      2        0       0       0       0       0       1       0
037  4037      4        0       0       0       0       0       0       0
038  4038      9        0       0       0       0       0       0       0

  6_count  7_count  ... onion layers for network  Boundary_Expansion  \
    15      18     ...           8567          3.364943
    0      1     ...           47          18.333333
    0      0     ...           29          30.636364
    0      0     ...           87          18.333333
    0      0     ...           21          30.636364
...
...     ...     ...     ...
034  0        0     ...           3          19.000000
035  0        0     ...           2          29.000000
036  0        0     ...           3          19.000000
037  0        0     ...           9          11.000000
038  2        1     ...           31          5.000000

kemeny constant for network  global_efficiency  max_weight_matching  \
406.232390          0.523734          164
17.493435          0.741830          9
9.177039          0.954545          5
16.649210          0.836601          9
9.196660          0.945455          5
...
...     ...     ...
034  1.333333  1.000000          1
035  0.500000  1.000000          1
036  1.333333  1.000000          1
037  3.362500  0.900000          2
038  8.546052  0.822222          5

random_probability  eccentricity  No of triangles with Node  \
75.626117          6          2519
0.814521          7          57
0.088836          7          40
1.395870          7          86
0.364554          7          39
...
...     ...     ...
034  0.000000          8          1
035  0.000000          8          0
036  0.000000          8          1
037  -0.135403         8          4
038  0.266415          8          20

triangle for network  k=4_Cluster
39777          1
402            0
381            0
996            0
378            0
...
...     ...     ...
034  3            0
035  0            0
036  3            0
037  15           0
038  114          0
```

4039 rows x 258 columns]

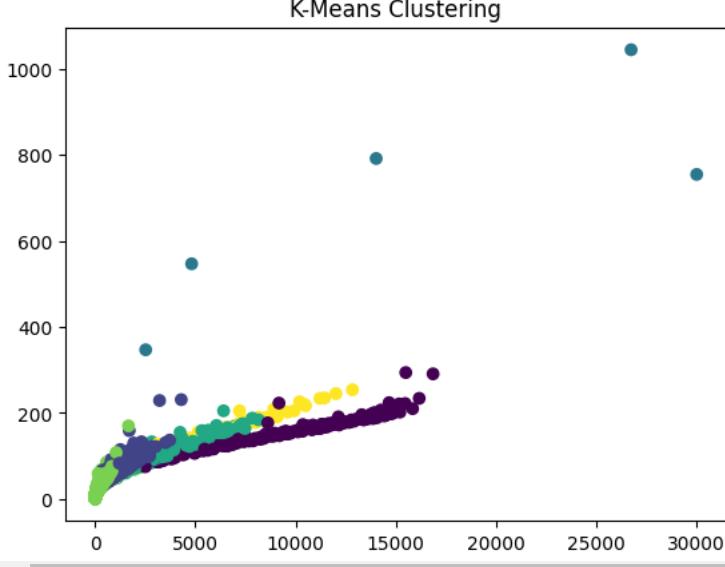
K-Means Clustering



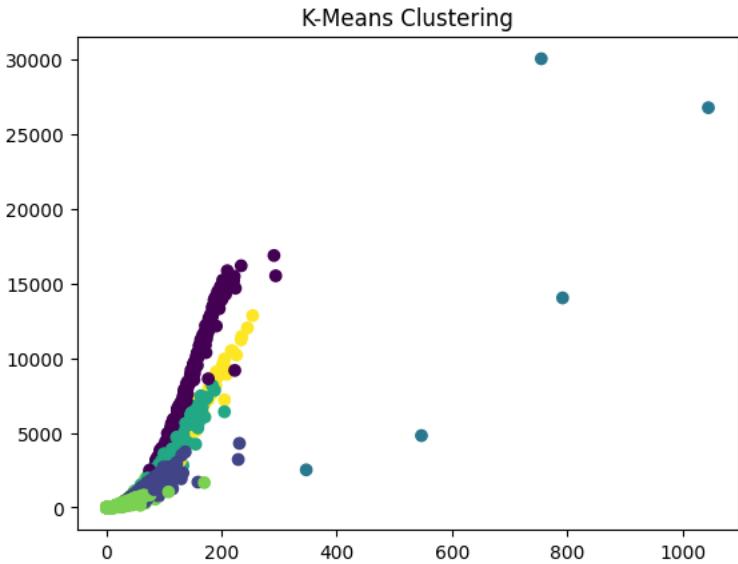
```
#Step 3: Apply k-means clustering
N = 6 # Replace with your optimal k
kmeans = KMeans(n_clusters=N, random_state=42)
onefinaldf['k = 6_Cluster'] = kmeans.fit_predict(features_standardized)

# Step 4: Visualize the clusters (optional)
# This is just an example, adjust it based on your requirements
# Visualization is more challenging with a large number of features
# You may need to choose specific features for plotting or use advanced visualization techniques
plt.scatter(onefinaldf["Common Connections"], onefinaldf['NodeDegree'], c=onefinaldf['k = 6_Cluster'], cmap='viridis')
plt.title('K-Means Clustering')
plt.show()
```

/usr/local/lib/python3.10/dist-packages/sklearn/cluster/\_kmeans.py:870: FutureWarning: The default value of `n\_init` will change from 10 to 20 in 0.22.0. The following code will raise a warning:



```
# You may need to choose specific features for plotting or use advanced visualization techniques
plt.scatter(onefinaldf["NodeDegree"], onefinaldf['Common Connections'], c=onefinaldf['k = 6_Cluster'], cmap='viridis')
plt.title('K-Means Clustering')
plt.show()
```



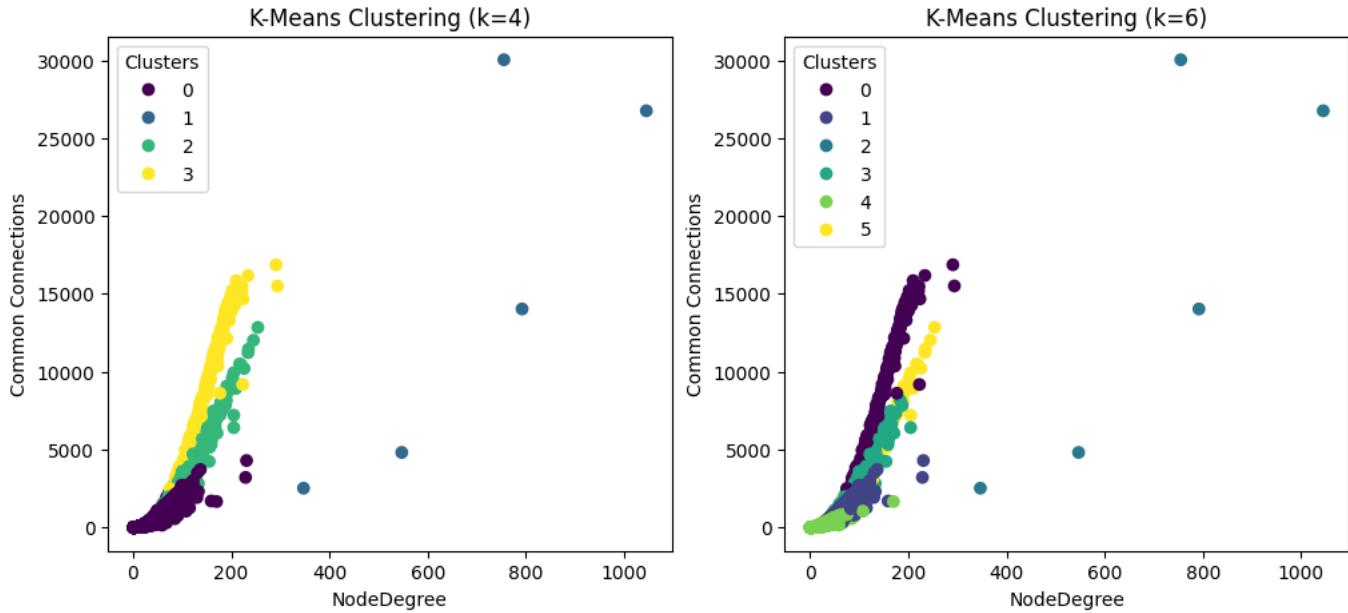
showing both for 4 clusters and 6 clusters in a same graph

```
# Step 4: Visualize the clusters
fig, axes = plt.subplots(nrows=1, ncols=2, figsize=(12, 5))

# Plot for k=4 clusters
scatter_4 = axes[0].scatter(onefinaldf["NodeDegree"], onefinaldf['Common Connections'], c=onefinaldf['k=4_Cluster'], cmap='viridis')
axes[0].set_title(f'K-Means Clustering (k=4)')
axes[0].set_xlabel('NodeDegree')
axes[0].set_ylabel('Common Connections')
axes[0].legend(*scatter_4.legend_elements(), title='Clusters')

# Plot for k=6 clusters
scatter_6 = axes[1].scatter(onefinaldf["NodeDegree"], onefinaldf['Common Connections'], c=onefinaldf['k = 6_Cluster'], cmap='viridis')
axes[1].set_title(f'K-Means Clustering (k=6)')
axes[1].set_xlabel('NodeDegree')
axes[1].set_ylabel('Common Connections')
axes[1].legend(*scatter_6.legend_elements(), title='Clusters')
```

&lt;matplotlib.legend.Legend at 0x7a199b5abe80&gt;



# visualize with kemeny constant - time and Onion layers of the network for clustering

```
# Step 4: Visualize the clusters
fig, axes = plt.subplots(nrows=1, ncols=2, figsize=(12, 5))
```

```
# Plot for k=4 clusters
scatter_4 = axes[0].scatter(onefinaldf["kemeny constant for network"], onefinaldf['onion layers for network'], c=onefinaldf['k=4_Cluster'], cmap='viridis')
axes[0].set_title(f'K-Means Clustering (k=4)')
axes[0].set_xlabel('kemeny constant for network')
axes[0].set_ylabel('onion layers for network')
axes[0].legend(*scatter_4.legend_elements(), title='Clusters')

# Plot for k=6 clusters
scatter_6 = axes[1].scatter(onefinaldf["kemeny constant for network"], onefinaldf['onion layers for network'], c=onefinaldf['k = 6_Cluster'], cmap='viridis')
axes[1].set_title(f'K-Means Clustering (k=6)')
axes[1].set_xlabel('kemeny constant for network')
axes[1].set_ylabel('onion layers for network')
axes[1].legend(*scatter_6.legend_elements(), title='Clusters')
```

&lt;matplotlib.legend.Legend at 0x7a198ff2f0a0&gt;

K-Means Clustering (k=4)



K-Means Clustering (k=6)



```
# visualize with No of triangle in the network and Onion layers of the network for clustering
# Step 4: Visualize the clusters
```

```
fig, axes = plt.subplots(nrows=1, ncols=2, figsize=(12, 5))
```

```
# Plot for k=4 clusters
```

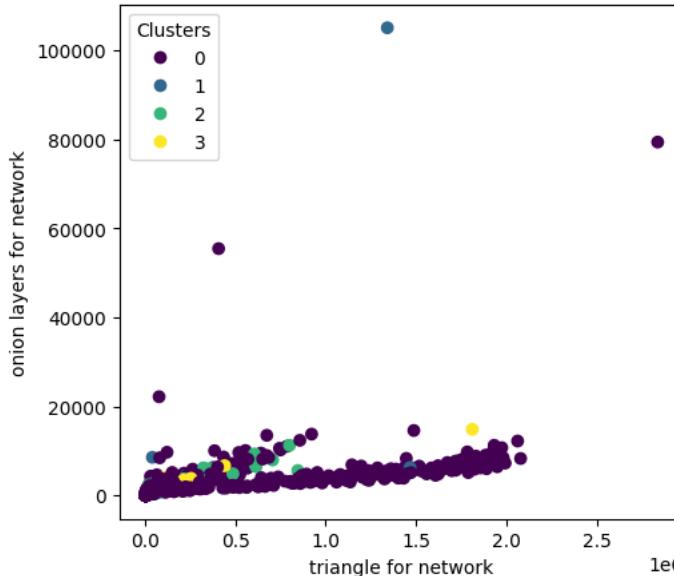
```
scatter_4 = axes[0].scatter(onefinaldf["triangle for network"], onefinaldf['onion layers for network'], c=onefinaldf['k=4_Cluster'], cmap='viridis')
axes[0].set_title(f'K-Means Clustering (k=4)')
axes[0].set_xlabel('triangle for network')
axes[0].set_ylabel('onion layers for network')
axes[0].legend(*scatter_4.legend_elements(), title='Clusters')
```

```
# Plot for k=6 clusters
```

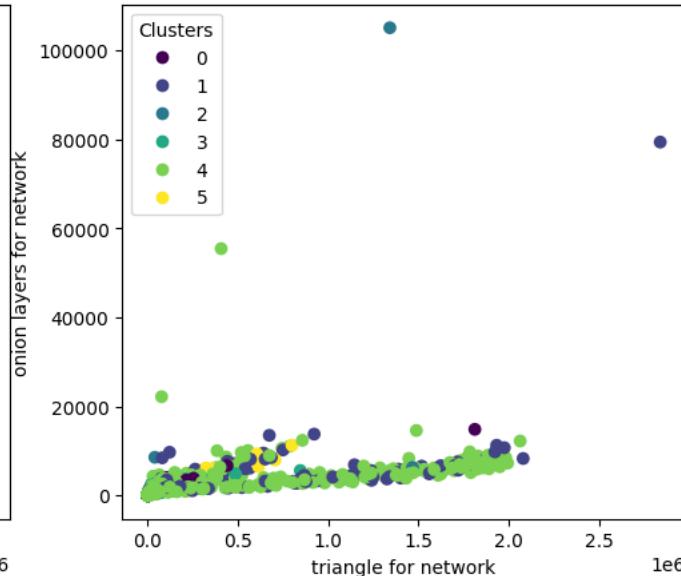
```
scatter_6 = axes[1].scatter(onefinaldf["triangle for network"], onefinaldf['onion layers for network'], c=onefinaldf['k = 6_Cluster'], cmap='viridis')
axes[1].set_title(f'K-Means Clustering (k=6)')
axes[1].set_xlabel('triangle for network')
axes[1].set_ylabel('onion layers for network')
axes[1].legend(*scatter_6.legend_elements(), title='Clusters')
```

&lt;matplotlib.legend.Legend at 0x7a199b411f00&gt;

K-Means Clustering (k=4)



K-Means Clustering (k=6)



```
# visualize with Boundary Expansion of network and Onion layers of the network for clustering
```

```
# Step 4: Visualize the clusters
```

```
fig, axes = plt.subplots(nrows=1, ncols=2, figsize=(12, 5))
```

```
# Plot for k=4 clusters
```

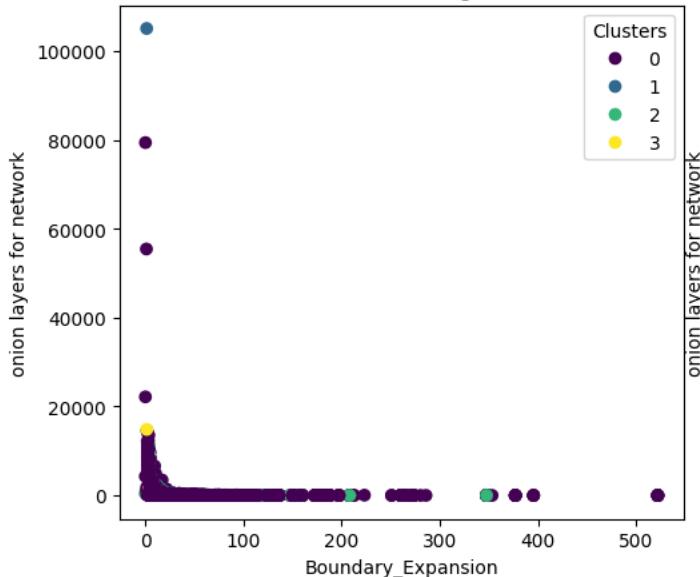
```
scatter_4 = axes[0].scatter(onefinaldf["Boundary_Expansion"], onefinaldf['onion layers for network'], c=onefinaldf['k=4_Cluster'], cmap='viridis')
axes[0].set_title(f'K-Means Clustering (k=4)')
axes[0].set_xlabel('Boundary_Expansion')
axes[0].set_ylabel('onion layers for network')
axes[0].legend(*scatter_4.legend_elements(), title='Clusters')
```

```
# Plot for k=6 clusters
```

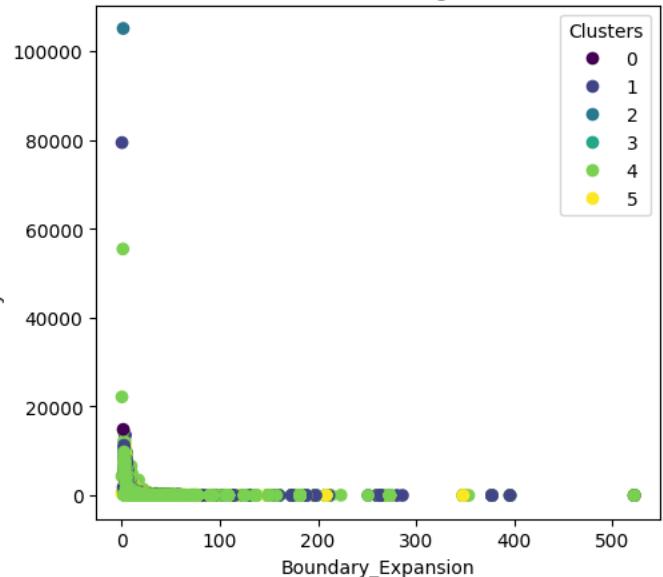
```
scatter_6 = axes[1].scatter(onefinaldf["Boundary_Expansion"], onefinaldf['onion layers for network'], c=onefinaldf['k = 6_Cluster'], cmap='viridis')
axes[1].set_title(f'K-Means Clustering (k=6)')
axes[1].set_xlabel('Boundary_Expansion')
axes[1].set_ylabel('onion layers for network')
axes[1].legend(*scatter_6.legend_elements(), title='Clusters')
```

&lt;matplotlib.legend.Legend at 0x7a199b675870&gt;

K-Means Clustering (k=4)



K-Means Clustering (k=6)



```

import seaborn as sns
def plot_clusters(graph, clusters, k_value):
    # Choose a colormap for clusters
    cluster_colors = sns.color_palette('Set1', n_colors=len(clusters.unique()))

    # Create a figure and axis
    fig, ax = plt.subplots(figsize=(10, 8))

    # Draw nodes with colors based on clusters
    nodes = nx.draw_networkx_nodes(graph, pos=nx.spring_layout(graph), node_color=clusters, cmap=plt.cm.get_cmap('Set1', len(clusters.unique)))

    # Draw edges
    nx.draw_networkx_edges(graph, pos=nx.spring_layout(graph), alpha=0.1)

    # Add a colorbar
    cbar = plt.colorbar(nodes, ticks=clusters.unique(), label='Cluster')
    cbar.set_ticklabels([f'Cluster {i}' for i in clusters.unique()])

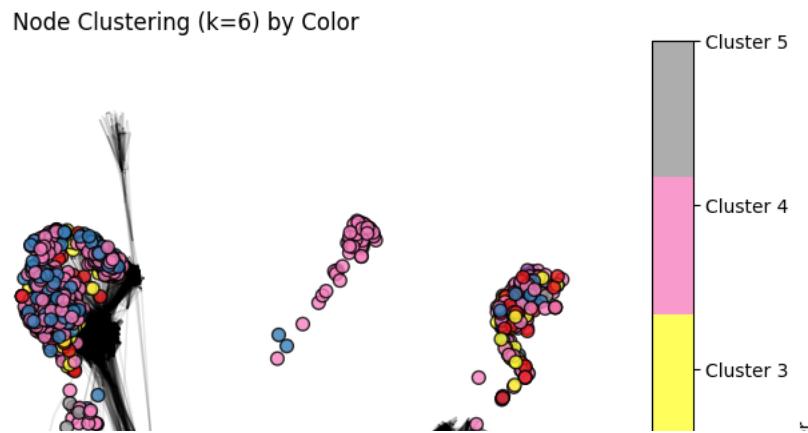
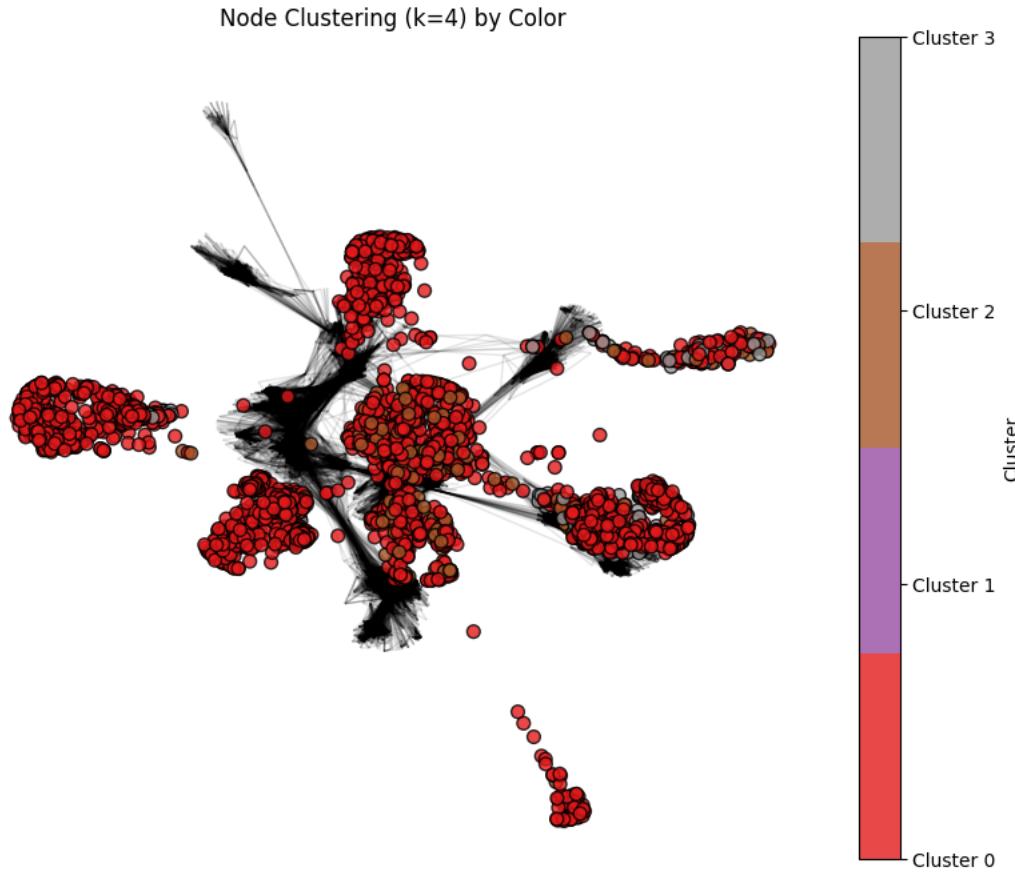
    plt.title(f'Node Clustering (k={k_value}) by Color')
    plt.axis('off')
    plt.show()

# Plot for k=4
plot_clusters(G, onefinaldf['k=4_Cluster'], 4)

# Plot for k=6
plot_clusters(G, onefinaldf['k = 6_Cluster'], 6)

```

```
<ipython-input-113-98016605a855>:10: MatplotlibDeprecationWarning: The get_cmap function was deprecated in Matplotlib 3.7 and will
nodes = nx.draw_networkx_nodes(graph, pos=nx.spring_layout(graph), node_color=clusters, cmap=plt.cm.get_cmap('Set1', len(clusters))
```



```
# Agglomerative Clustering
import pandas as pd
from sklearn.cluster import AgglomerativeClustering
from sklearn.preprocessing import StandardScaler
import matplotlib.pyplot as plt
from scipy.cluster.hierarchy import dendrogram, linkage

# Step 1: Standardize the data (optional but recommended for hierarchical clustering)
scaler = StandardScaler()
data_standardized = scaler.fit_transform(onefinaldf)

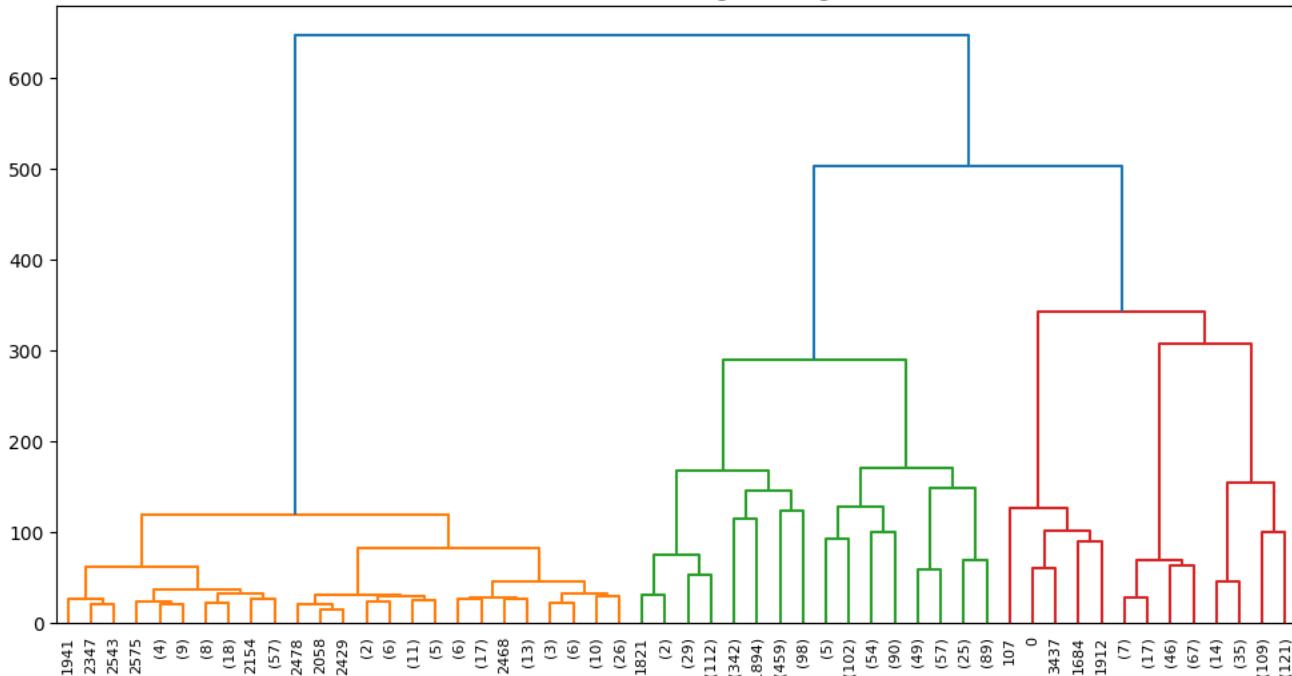
# Step 2: Determine the linkage matrix using the ward method
linkage_matrix = linkage(data_standardized, method='ward')

# Step 3: Plot the dendrogram (optional but useful for visualizing hierarchical clustering)
plt.figure(figsize=(12, 6))
dendrogram(linkage_matrix, leaf_rotation=90., leaf_font_size=8., truncate_mode='level', p=5)
plt.title('Hierarchical Clustering Dendrogram')
plt.show()

# Step 4: Apply Agglomerative Hierarchical Clustering
# Choose the appropriate number of clusters (n_clusters) based on the dendrogram
n_clusters = 4 # Replace with your desired number of clusters
hierarchical_clustering = AgglomerativeClustering(n_clusters=n_clusters, linkage='ward')
onefinaldf['agg4Cluster'] = hierarchical_clustering.fit_predict(data_standardized)

# Display the DataFrame with cluster assignments
print(onefinaldf)
```

## Hierarchical Clustering Dendrogram



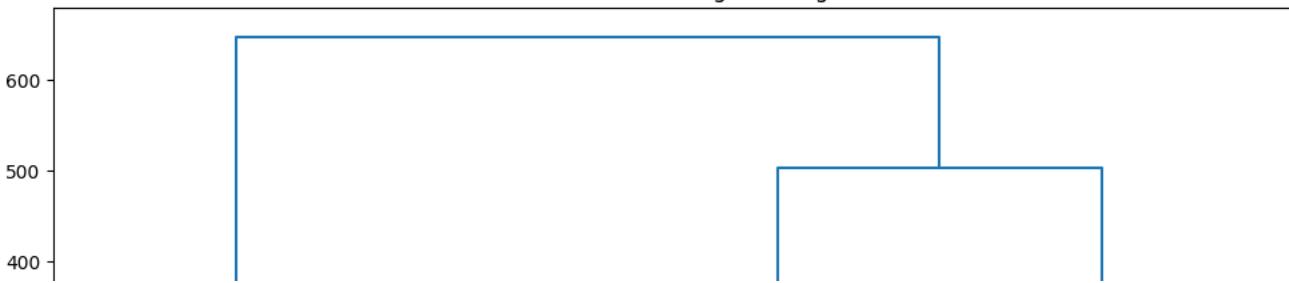
Node	NodeDegree	0_count	1_count	2_count	3_count	4_count	5_count	\
0	347	0	14	27	22	17	12	
1	17	0	0	0	0	0	0	
2	10	0	0	0	0	0	0	
3	17	0	0	0	1	0	1	
4	10	0	0	0	0	0	1	
...	...	...	...	...	...	...	...	
4034	4034	2	0	0	0	0	0	
4035	4035	1	0	0	0	0	0	
4036	4036	2	0	0	0	0	1	
4037	4037	4	0	0	0	0	0	
4038	4038	9	0	0	0	0	0	
6_count	7_count	...	kemeny constant for network	global_efficiency	\			
0	15	18	...	406.232390	0.523734			
1	0	1	...	17.493435	0.741830			
2	0	0	...	9.177039	0.954545			

```
linkage_matrix = linkage(data_standardized, method='ward')
```

```
# Step 3: Plot the dendrogram (optional but useful for visualizing hierarchical clustering)
plt.figure(figsize=(12, 6))
dendrogram(linkage_matrix, leaf_rotation=90., leaf_font_size=8., truncate_mode='level', p=7)
plt.title('Hierarchical Clustering Dendrogram')
plt.show()
```

```
# Step 4: Apply Agglomerative Hierarchical Clustering
# Choose the appropriate number of clusters (n_clusters) based on the dendrogram
n_clusters = 6 # Replace with your desired number of clusters
hierarchical_clustering = AgglomerativeClustering(n_clusters=n_clusters, linkage='ward')
onefinaldf['agg6Cluster'] = hierarchical_clustering.fit_predict(data_standardized)
```

## Hierarchical Clustering Dendrogram



```
print(onefinaldf)
```

	4035	4036	4037	4038	1	0	0	0	0	0	0
0	4035	4036	4037	4038	2	0	0	0	0	1	0
1					4	0	0	0	0	0	0
2					9	0	0	0	0	0	0

	6_count	7_count	...	global_efficiency	max_weight_matching	\
0	15	18	...	0.523734	164	
1	0	1	...	0.741830	9	
2	0	0	...	0.954545	5	
3	0	0	...	0.836601	9	
4	0	0	...	0.945455	5	
...	...	...	...	...	...	
4034	0	0	...	1.000000	1	
4035	0	0	...	1.000000	1	
4036	0	0	...	1.000000	1	
4037	0	0	...	0.900000	2	
4038	2	1	...	0.822222	5	

	random_probability	eccentricity	No of triangles with Node	\
0	75.626117	6	2519	
1	0.814521	7	57	
2	0.088836	7	40	
3	1.395870	7	86	
4	0.364554	7	39	
...	...	...	...	
4034	0.000000	8	1	
4035	0.000000	8	0	
4036	0.000000	8	1	
4037	-0.135403	8	4	
4038	0.266415	8	20	

	triangle for network	k=4_Cluster	k = 6_Cluster	agg4Cluster	\
0	39777	1	2	1	
1	402	0	4	2	
2	381	0	4	2	
3	996	0	4	2	
4	378	0	4	2	
...	...	...	...	...	
4034	3	0	4	2	
4035	0	0	4	2	
4036	3	0	4	2	
4037	15	0	4	2	
4038	114	0	4	2	

	agg6Cluster
0	1
1	2
2	2
3	2
4	2
...	...
4034	2
4035	2
4036	2
4037	2
4038	2

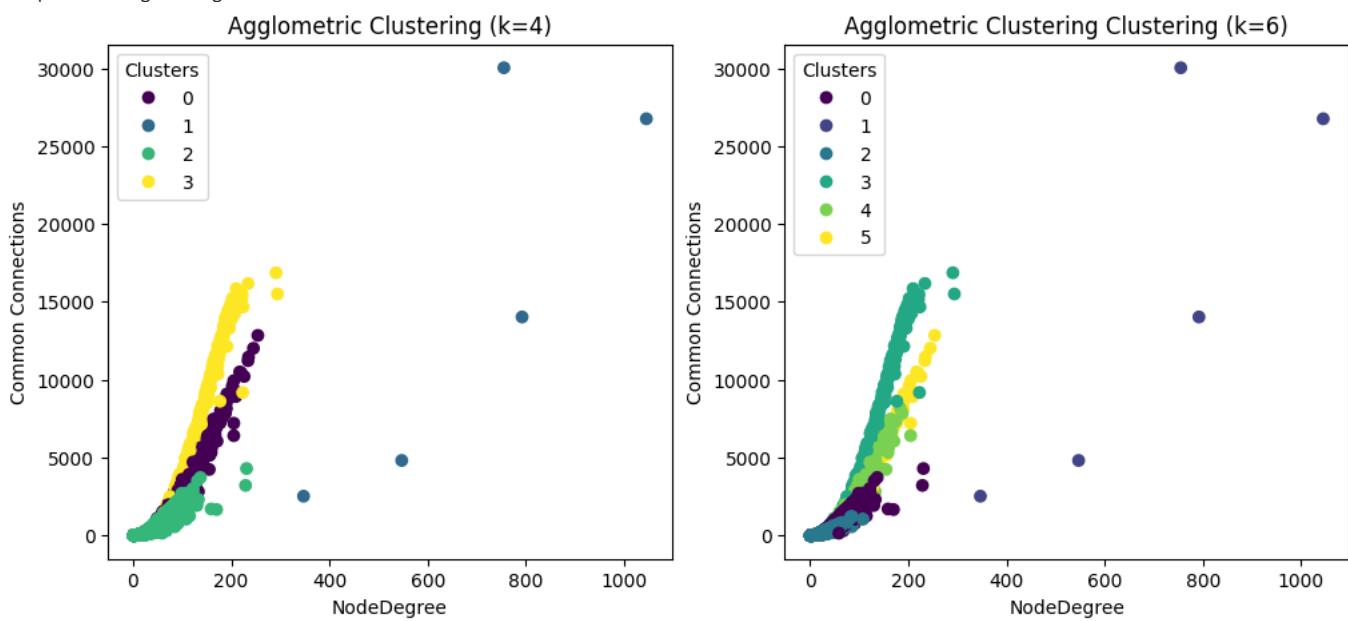
```
[4039 rows x 261 columns]
```

```
# Step 4: Visualize the clusters for agglomeretic clustering
fig, axes = plt.subplots(nrows=1, ncols=2, figsize=(12, 5))

# Plot for k=4 clusters
scatter_4 = axes[0].scatter(onefinaldf["NodeDegree"], onefinaldf['Common Connections'], c=onefinaldf['agg4Cluster'], cmap='viridis')
axes[0].set_title(f'Agglomeretic Clustering (k=4)')
axes[0].set_xlabel('NodeDegree')
axes[0].set_ylabel('Common Connections')
axes[0].legend(*scatter_4.legend_elements(), title='Clusters')

# Plot for k=6 clusters
scatter_6 = axes[1].scatter(onefinaldf["NodeDegree"], onefinaldf['Common Connections'], c=onefinaldf['agg6Cluster'], cmap='viridis')
axes[1].set_title(f'Agglomeretic Clustering Clustering (k=6)')
axes[1].set_xlabel('NodeDegree')
axes[1].set_ylabel('Common Connections')
axes[1].legend(*scatter_6.legend_elements(), title='Clusters')

<matplotlib.legend.Legend at 0x7a198e411f00>
```

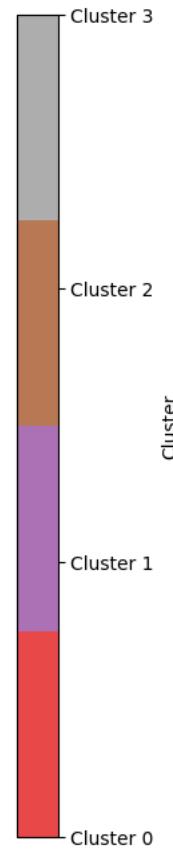


```
# Plot for k=4
plot_clusters(G, onefinaldf['agg4Cluster'], 4)

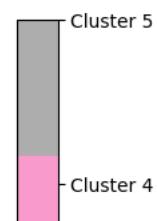
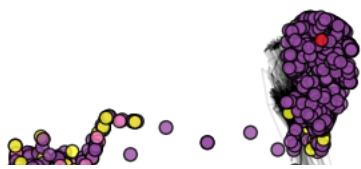
# Plot for k=6
plot_clusters(G, onefinaldf['agg6Cluster'], 6)
```

```
<ipython-input-113-98016605a855>:10: MatplotlibDeprecationWarning: The get_cmap function was deprecated in Matplotlib 3.7 and will
nodes = nx.draw_networkx_nodes(graph, pos=nx.spring_layout(graph), node_color=clusters, cmap=plt.cm.get_cmap('Set1', len(clusters))
```

Node Clustering (k=4) by Color



Node Clustering (k=6) by Color



```
# Step 4: Visualize the clusters for k-means and agglomerative clustering
fig, axes = plt.subplots(nrows=2, ncols=2, figsize=(12, 10))

# Plot for k=4 clusters in k-means
scatter_4_kmeans = axes[0, 0].scatter(onefinaldf["NodeDegree"], onefinaldf['Common Connections'], c=onefinaldf['k=4_Cluster'], cmap='viridis')
axes[0, 0].set_title('K-means Clustering (k=4)')
axes[0, 0].set_xlabel('NodeDegree')
axes[0, 0].set_ylabel('Common Connections')
axes[0, 0].legend(*scatter_4_kmeans.legend_elements(), title='Clusters')

# Plot for k=4 clusters in agglomerative clustering
scatter_4_agg = axes[0, 1].scatter(onefinaldf["NodeDegree"], onefinaldf['Common Connections'], c=onefinaldf['agg4Cluster'], cmap='viridis')
axes[0, 1].set_title('Agglomerative Clustering (k=4)')
axes[0, 1].set_xlabel('NodeDegree')
axes[0, 1].set_ylabel('Common Connections')
axes[0, 1].legend(*scatter_4_agg.legend_elements(), title='Clusters')

# Plot for k=6 clusters in k-means
scatter_6_kmeans = axes[1, 0].scatter(onefinaldf["NodeDegree"], onefinaldf['Common Connections'], c=onefinaldf['k = 6_Cluster'], cmap='viridis')
axes[1, 0].set_title('K-means Clustering (k=6)')
axes[1, 0].set_xlabel('NodeDegree')
axes[1, 0].set_ylabel('Common Connections')
axes[1, 0].legend(*scatter_6_kmeans.legend_elements(), title='Clusters')

# Plot for k=6 clusters in agglomerative clustering
scatter_6_agg = axes[1, 1].scatter(onefinaldf["NodeDegree"], onefinaldf['Common Connections'], c=onefinaldf['agg6Cluster'], cmap='viridis')
axes[1, 1].set_title('Agglomerative Clustering (k=6)')
axes[1, 1].set_xlabel('NodeDegree')
axes[1, 1].set_ylabel('Common Connections')
axes[1, 1].legend(*scatter_6_agg.legend_elements(), title='Clusters')

plt.tight_layout()
plt.show()
```

K-means Clustering (k=4)



Agglomerative Clustering (k=4)



```

import seaborn as sns

def plot_clustersname(graph, clusters, k_value, name, ax):
    # Choose a colormap for clusters
    cluster_colors = sns.color_palette('Set1', n_colors=len(clusters.unique()))

    # Draw nodes with colors based on clusters
    nodes = nx.draw_networkx_nodes(graph, pos=nx.spring_layout(graph), node_color=clusters, cmap=plt.cm.get_cmap('Set1', len(clusters.unique)))

    # Draw edges
    nx.draw_networkx_edges(graph, pos=nx.spring_layout(graph), alpha=0.1, ax=ax)

    # Add a colorbar
    cbar = plt.colorbar(nodes, ticks=clusters.unique(), label='Cluster', ax=ax)
    cbar.set_ticklabels([f'Cluster {i}' for i in clusters.unique()])

    ax.set_title(f'Node Clustering (k={k_value}) ({name}) by Color')
    ax.axis('off')

# Assuming G is your graph

# Create a 2x2 grid for subplots
fig, axes = plt.subplots(2, 2, figsize=(12, 10))

# Plot for k=4
plot_clustersname(G, onefinaldf['k=4_Cluster'], 4, 'kmeans', axes[0, 0])

# Plot for k=4 - agg
plot_clustersname(G, onefinaldf['agg4Cluster'], 4, 'aggclustering', axes[0, 1])

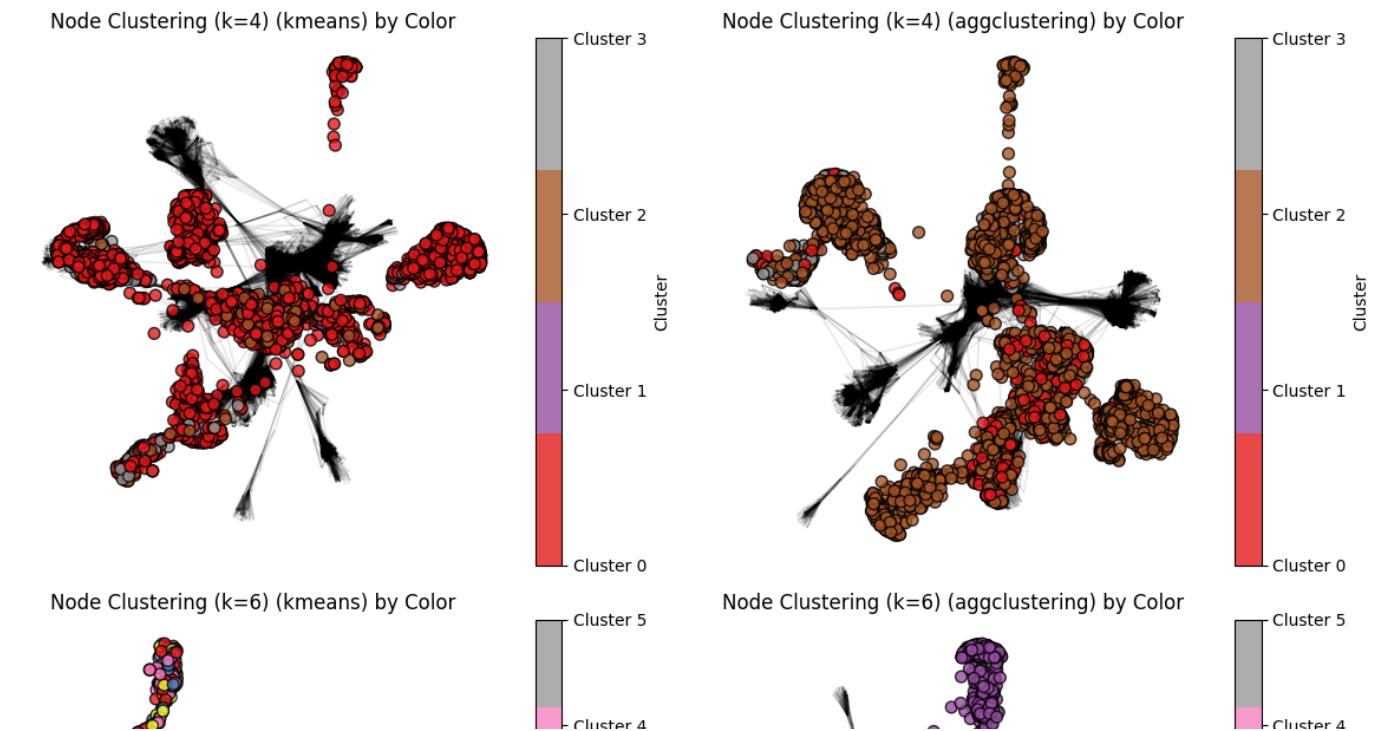
# Plot for k=6
plot_clustersname(G, onefinaldf['k = 6_Cluster'], 6, 'kmeans', axes[1, 0])

# Plot for k=6 - agg
plot_clustersname(G, onefinaldf['agg6Cluster'], 6, 'aggclustering', axes[1, 1])

# Adjust layout
plt.tight_layout()
plt.show()

```

```
<ipython-input-117-83f07bba3dce>:8: MatplotlibDeprecationWarning: The get_cmap function was deprecated in Matplotlib 3.7 and will be removed in 3.8.
```



```
# visualize with No of triangle in the network and Onion layers of the network for both of the Clustering
```

```
# Step 4: Visualize the clusters for k-means and agglomerative clustering
```

```
fig, axes = plt.subplots(nrows=2, ncols=2, figsize=(12, 10))
```

```
# Plot for k=4 clusters in k-means
```

```
scatter_4_kmeans = axes[0, 0].scatter(onefinaldf["triangle for network"], onefinaldf['onion layers for network'], c=onefinaldf['k=4_Cluster'])
axes[0, 0].set_title('K-means Clustering (k=4)')
axes[0, 0].set_xlabel('triangle for network')
axes[0, 0].set_ylabel('onion layers for network')
axes[0, 0].legend(*scatter_4_kmeans.legend_elements(), title='Clusters')
```

```
# Plot for k=4 clusters in agglomerative clustering
```

```
scatter_4_agg = axes[0, 1].scatter(onefinaldf["triangle for network"], onefinaldf['onion layers for network'], c=onefinaldf['agg4Cluster'])
axes[0, 1].set_title('Agglomerative Clustering (k=4)')
axes[0, 1].set_xlabel('triangle for network')
axes[0, 1].set_ylabel('onion layers for network')
axes[0, 1].legend(*scatter_4_agg.legend_elements(), title='Clusters')
```

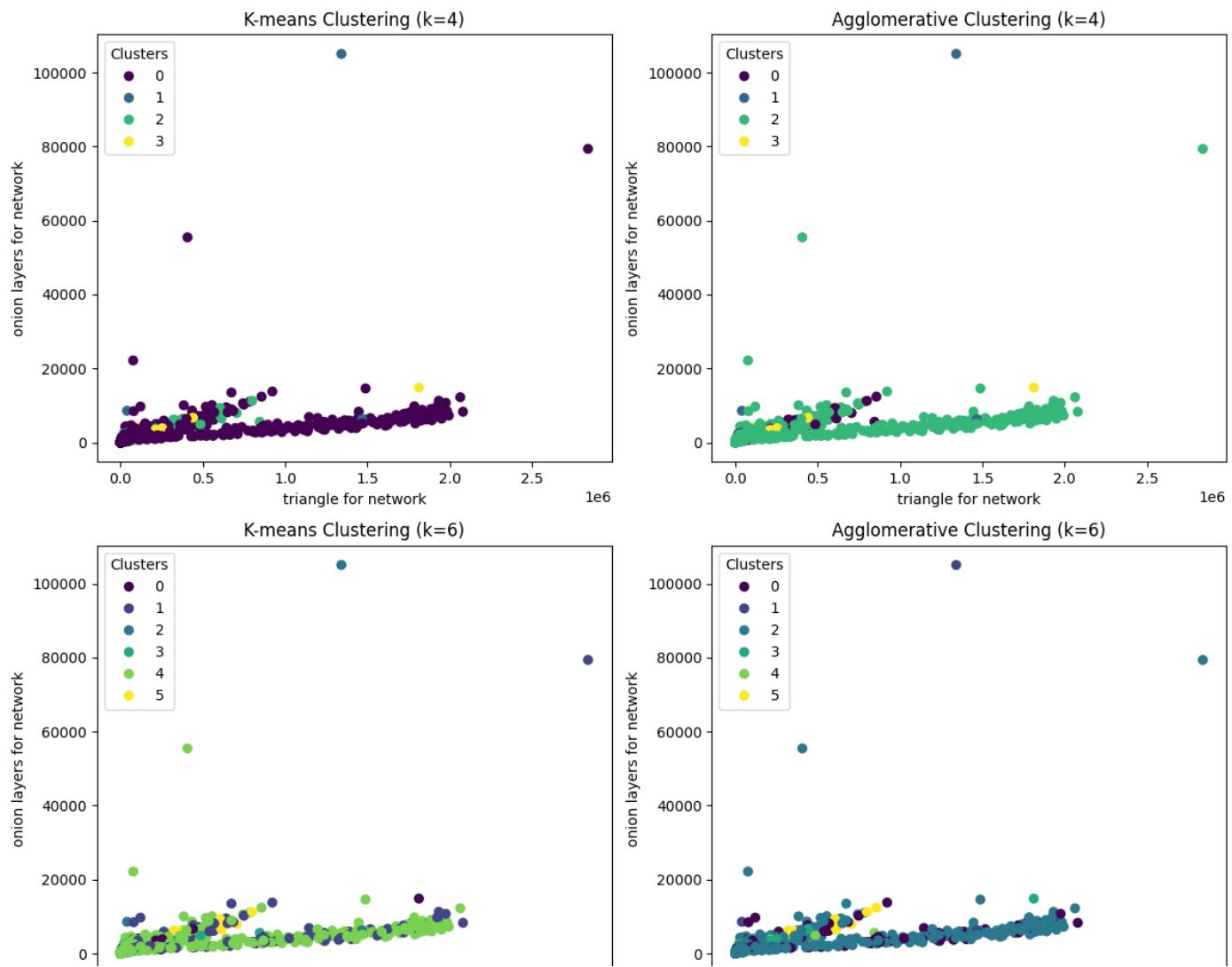
```
# Plot for k=6 clusters in k-means
```

```
scatter_6_kmeans = axes[1, 0].scatter(onefinaldf["triangle for network"], onefinaldf['onion layers for network'], c=onefinaldf['k = 6_Cluster'])
axes[1, 0].set_title('K-means Clustering (k=6)')
axes[1, 0].set_xlabel('triangle for network')
axes[1, 0].set_ylabel('onion layers for network')
axes[1, 0].legend(*scatter_6_kmeans.legend_elements(), title='Clusters')
```

```
# Plot for k=6 clusters in agglomerative clustering
```

```
scatter_6_agg = axes[1, 1].scatter(onefinaldf["triangle for network"], onefinaldf['onion layers for network'], c=onefinaldf['agg6Cluster'])
axes[1, 1].set_title('Agglomerative Clustering (k=6)')
axes[1, 1].set_xlabel('triangle for network')
axes[1, 1].set_ylabel('onion layers for network')
axes[1, 1].legend(*scatter_6_agg.legend_elements(), title='Clusters')
```

```
plt.tight_layout()
plt.show()
```



```
#dbscan algorithm
import pandas as pd
from sklearn.cluster import DBSCAN
from sklearn.preprocessing import StandardScaler
import matplotlib.pyplot as plt

# Assuming df is your DataFrame

# Step 1: Standardize the data (required for DBSCAN)
scaler = StandardScaler()
data_standardized = scaler.fit_transform(onefinaldf)

# Step 2: Apply DBSCAN
# Choose appropriate values for `eps` (maximum distance between two samples for one to be considered as in the neighborhood of the other)
# and `min_samples` (the number of samples in a neighborhood for a point to be considered as a core point)
dbscan = DBSCAN(eps=1, min_samples=20)
onefinaldf['dbs1Cluster'] = dbscan.fit_predict(data_standardized)

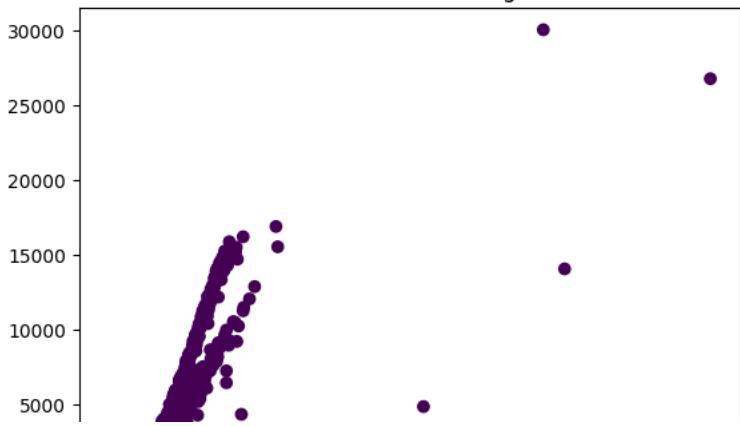
# Display the DataFrame with cluster assignments
print(onefinaldf)

# Step 3: Visualize the clusters (optional)
# This is just an example, adjust it based on your requirements
plt.scatter(onefinaldf['NodeDegree'], onefinaldf['Common Connections'], c=onefinaldf['dbs1Cluster'], cmap='viridis')
plt.title('DBSCAN Clustering')
plt.show()
```

	Node	NodeDegree	0_count	1_count	2_count	3_count	4_count	5_count	\
0	0	347	0	14	27	22	17	12	
1	1	17	0	0	0	0	0	0	
2	2	10	0	0	0	0	0	0	
3	3	17	0	0	0	1	0	1	
4	4	10	0	0	0	0	0	1	
...	...	...	...	...	...	...	...	...	
4034	4034	2	0	0	0	0	0	0	
4035	4035	1	0	0	0	0	0	0	
4036	4036	2	0	0	0	0	1	0	
4037	4037	4	0	0	0	0	0	0	
4038	4038	9	0	0	0	0	0	0	
			6_count	7_count	...	max_weight_matching	random_probability	\	
0		15	18	...		164	75.626117		
1	0	0	1	...		9	0.814521		
2	0	0	0	...		5	0.088836		
3	0	0	0	...		9	1.395870		
4	0	0	0	...		5	0.364554		
...	...	...	...	...		...	...		
4034	0	0	0	...		1	0.000000		
4035	0	0	0	...		1	0.000000		
4036	0	0	0	...		1	0.000000		
4037	0	0	0	...		2	-0.135403		
4038	2	1	1	...		5	0.266415		
			eccentricity	No of triangles with Node	triangle for network	\			
0		6		2519		39777			
1	7			57		402			
2	7			40		381			
3	7			86		996			
4	7			39		378			
...	...	...		...		...			
4034	8			1		3			
4035	8			0		0			
4036	8			1		3			
4037	8			4		15			
4038	8			20		114			
			k=4_Cluster	k = 6_Cluster	agg4Cluster	agg6Cluster	dbs1Cluster		
0		1		2	1	1	-1		
1	0		4	2	2	2	-1		
2	0		4	2	2	2	-1		
3	0		4	2	2	2	-1		
4	0		4	2	2	2	-1		
...	...	...	...	...	...	...	...		
4034	0		4	2	2	2	-1		
4035	0		4	2	2	2	-1		
4036	0		4	2	2	2	-1		
4037	0		4	2	2	2	-1		
4038	0		4	2	2	2	-1		

[4039 rows x 262 columns]

DBSCAN Clustering



we have to change eps and min samples value for the dbscan

v | — | |

```

from sklearn.neighbors import NearestNeighbors
import matplotlib.pyplot as plt

# Choose a suitable value for k (number of neighbors)
k = 4

# Compute pairwise distances between nodes
distances = nx.floyd_marshall_numpy(G)

# Choose a suitable value for k
k = 4

# Fit a nearest neighbors model
neigh = NearestNeighbors(n_neighbors=k)
neigh.fit(distances)

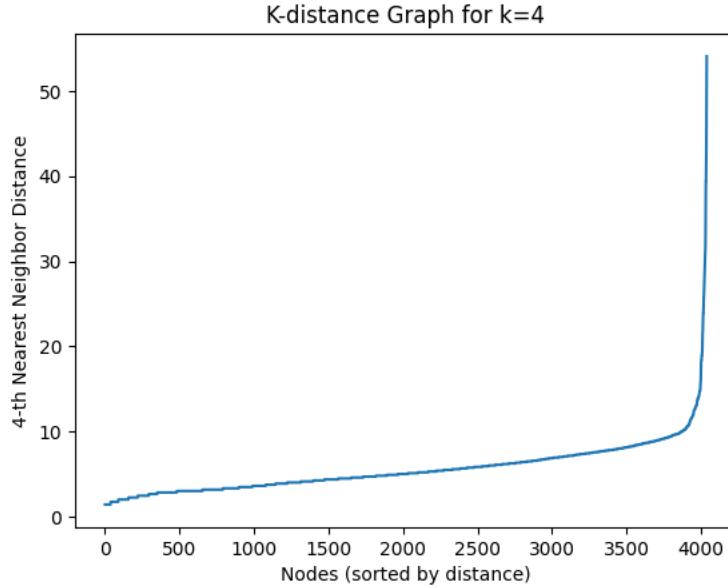
# Calculate distances to k-th nearest neighbors
distances_knn, _ = neigh.kneighbors(distances)

import numpy as np
import matplotlib.pyplot as plt

# Take the distances to the k-th nearest neighbors
distances_knn = np.max(distances_knn, axis=1)
distances_knn.sort()

# Plot the k-distance graph
plt.plot(distances_knn)
plt.xlabel('Nodes (sorted by distance)')
plt.ylabel(f'{k}-th Nearest Neighbor Distance')
plt.title(f'K-distance Graph for k={k}')
plt.show()

```



so it is increasing sharply near 11 , so i will take eps value as 11

```
# Compute pairwise distances between nodes (you may use other metrics based on your problem)
distances1 = nx.floyd_marshall_numpy(G)

# Choose a suitable range of values for min_samples
min_samples_values = range(1, 4040) # Adjust the range as needed

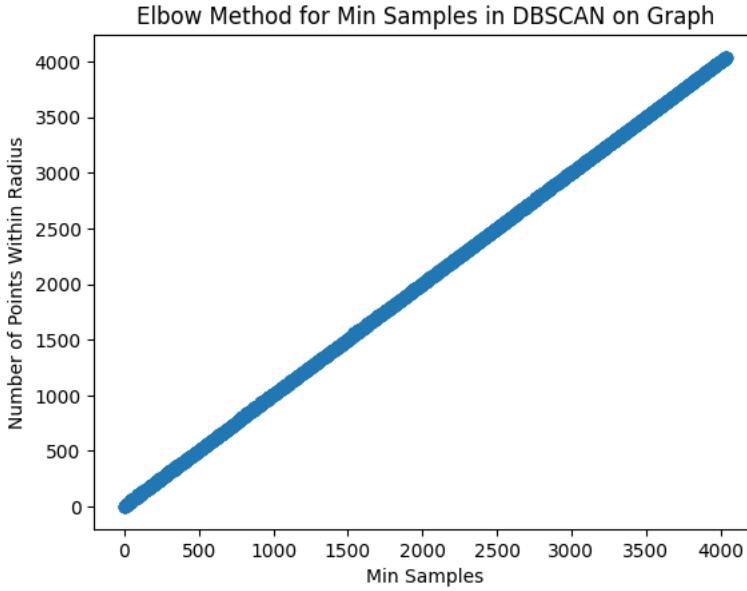
# Fit a nearest neighbors model
neigh = NearestNeighbors(n_neighbors=max(min_samples_values))
neigh.fit(distances1)

# Calculate distances to k-th nearest neighbors
distances_knn1, _ = neigh.kneighbors(distances)

# Sort the distances and calculate the number of points within each distance
distances_knn1 = np.max(distances_knn1, axis=1)
distances_knn1.sort()

num_points_within_radius = [np.sum(distances_knn1 <= d) for d in distances_knn1]

# Plot the elbow graph
plt.plot(min_samples_values, num_points_within_radius, marker='o')
plt.xlabel('Min Samples')
plt.ylabel('Number of Points Within Radius')
plt.title('Elbow Method for Min Samples in DBSCAN on Graph')
plt.show()
```



```
dbminvaluedf = onefinaldf.copy()
print(dbminvaluedf)

      Node  NodeDegree  0_count  1_count  2_count  3_count  4_count  5_count \
0         0        347       0      14      27      22      17      12
1         1        17       0       0       0       0       0       0
2         2        10       0       0       0       0       0       0
3         3        17       0       0       0       1       0       1
4         4        10       0       0       0       0       0       1
...
4034    4034        2       0       0       0       0       0       0
4035    4035        1       0       0       0       0       0       0
4036    4036        2       0       0       0       0       1       0
4037    4037        4       0       0       0       0       0       0
4038    4038        9       0       0       0       0       0       0

      6_count  7_count  ...  max_weight_matching  random_probability \
0        15     18   ...           164          75.626117
1        0      1   ...             9          0.814521
2        0      0   ...             5          0.088836
3        0      0   ...             9          1.395870
4        0      0   ...             5          0.364554
...
4034    0      0   ...             1          0.000000
4035    0      0   ...             1          0.000000
4036    0      0   ...             1          0.000000
4037    0      0   ...             2         -0.135403
4038    2      1   ...             5          0.266415
```

	eccentricity	No of triangles with Node	triangle for network	\
0	6	2519	39777	
1	7	57	402	
2	7	40	381	
3	7	86	996	
4	7	39	378	
...	...	...	...	
4034	8	1	3	
4035	8	0	0	
4036	8	1	3	
4037	8	4	15	
4038	8	20	114	

	k=4_Cluster	k = 6_Cluster	agg4Cluster	agg6Cluster	dbs1Cluster
0	1	2	1	1	-1
1	0	4	2	2	-1
2	0	4	2	2	-1
3	0	4	2	2	-1
4	0	4	2	2	-1
...	...	...	...	...	...
4034	0	4	2	2	-1
4035	0	4	2	2	-1
4036	0	4	2	2	-1
4037	0	4	2	2	-1
4038	0	4	2	2	-1

[4039 rows x 262 columns]

```

min_samples_values = range(2, 4040) # Starting from 2, as silhouette_score requires at least 2 clusters

# List to store silhouette scores for different min_samples values
silhouette_scores = []

# Loop over min_samples values
for min_samples in min_samples_values:
    # Apply DBSCAN for each min_samples value
    labels = DBSCAN(eps=11, min_samples=min_samples).fit_predict(dbminvaluedf)

    # Check if there is more than one unique label
    unique_labels = np.unique(labels)
    if len(unique_labels) > 1:
        # Calculate silhouette score
        silhouette_scores.append(silhouette_score(dbminvaluedf, labels))
    else:
        # If only one label, append a placeholder value (e.g., -1)
        silhouette_scores.append(-1)

# Plot silhouette scores
plt.plot(min_samples_values, silhouette_scores, marker='o')
plt.xlabel('Min Samples')
plt.ylabel('Silhouette Score')
plt.title('Silhouette Score for Different Min Samples in DBSCAN')
plt.show()

```

```

optimal_min_samples = min_samples_values[np.argmax(silhouette_scores)]
print(f'Optimal Min Samples: {optimal_min_samples}')

Optimal Min Samples: 2

#dbSCAN algorithm
import pandas as pd
from sklearn.cluster import DBSCAN
from sklearn.preprocessing import StandardScaler
import matplotlib.pyplot as plt

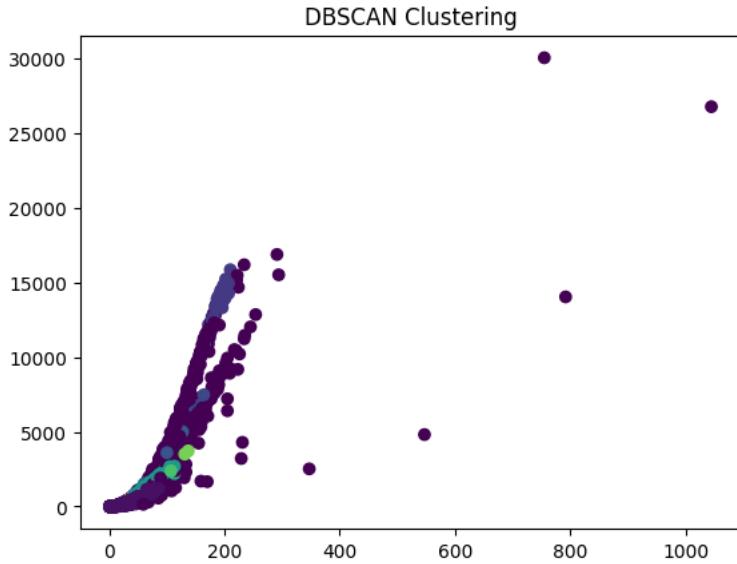
# Assuming df is your DataFrame

# Step 1: Standardize the data (required for DBSCAN)
scaler = StandardScaler()
data_standardized = scaler.fit_transform(onefinaldf)

# Step 2: Apply DBSCAN
# Choose appropriate values for `eps` (maximum distance between two samples for one to be considered as in the neighborhood of the other)
# and `min_samples` (the number of samples in a neighborhood for a point to be considered as a core point)
dbSCAN = DBSCAN(eps=11, min_samples=2)
onefinaldf['dbsCluster'] = dbSCAN.fit_predict(data_standardized)

# Step 3: Visualize the clusters (optional)
# This is just an example, adjust it based on your requirements
plt.scatter(onefinaldf['NodeDegree'], onefinaldf['Common Connections'], c=onefinaldf['dbsCluster'], cmap='viridis')
plt.title('DBSCAN Clustering')
plt.show()

```



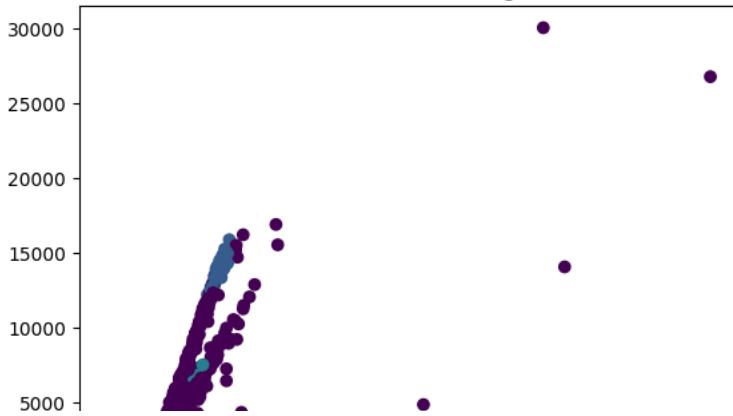
```

# Step 2: Apply DBSCAN
# Choose appropriate values for `eps` (maximum distance between two samples for one to be considered as in the neighborhood of the other)
# and `min_samples` (the number of samples in a neighborhood for a point to be considered as a core point)
dbSCAN = DBSCAN(eps=11, min_samples=4)
onefinaldf['dbs4Cluster'] = dbSCAN.fit_predict(data_standardized)

# Step 3: Visualize the clusters (optional)
# This is just an example, adjust it based on your requirements
plt.scatter(onefinaldf['NodeDegree'], onefinaldf['Common Connections'], c=onefinaldf['dbs4Cluster'], cmap='viridis')
plt.title('DBSCAN Clustering')
plt.show()

```

## DBSCAN Clustering



```
# Step 2: Apply DBSCAN
# Choose appropriate values for `eps` (maximum distance between two samples for one to be considered as in the neighborhood of the other)
# and `min_samples` (the number of samples in a neighborhood for a point to be considered as a core point)
dbscan = DBSCAN(eps=11, min_samples=6)
onefinaldf['dbs6Cluster'] = dbscan.fit_predict(data_standardized)

# Visualize clusters for k-means, agglomerative clustering, and DBSCAN in a 2x3 grid
fig, axes = plt.subplots(nrows=2, ncols=3, figsize=(18, 10))

# Plot for k=4 clusters in k-means
scatter_4_kmeans = axes[0, 0].scatter(onefinaldf["triangle for network"], onefinaldf['onion layers for network'], c=onefinaldf['k=4_Cluster'])
axes[0, 0].set_title('K-means Clustering (k=4)')
axes[0, 0].set_xlabel('triangle for network')
axes[0, 0].set_ylabel('onion layers for network')
axes[0, 0].legend(*scatter_4_kmeans.legend_elements(), title='Clusters')

# Plot for k=4 clusters in agglomerative clustering
scatter_4_agg = axes[0, 1].scatter(onefinaldf["triangle for network"], onefinaldf['onion layers for network'], c=onefinaldf['agg4Cluster'])
axes[0, 1].set_title('Agglomerative Clustering (k=4)')
axes[0, 1].set_xlabel('triangle for network')
axes[0, 1].set_ylabel('onion layers for network')
axes[0, 1].legend(*scatter_4_agg.legend_elements(), title='Clusters')

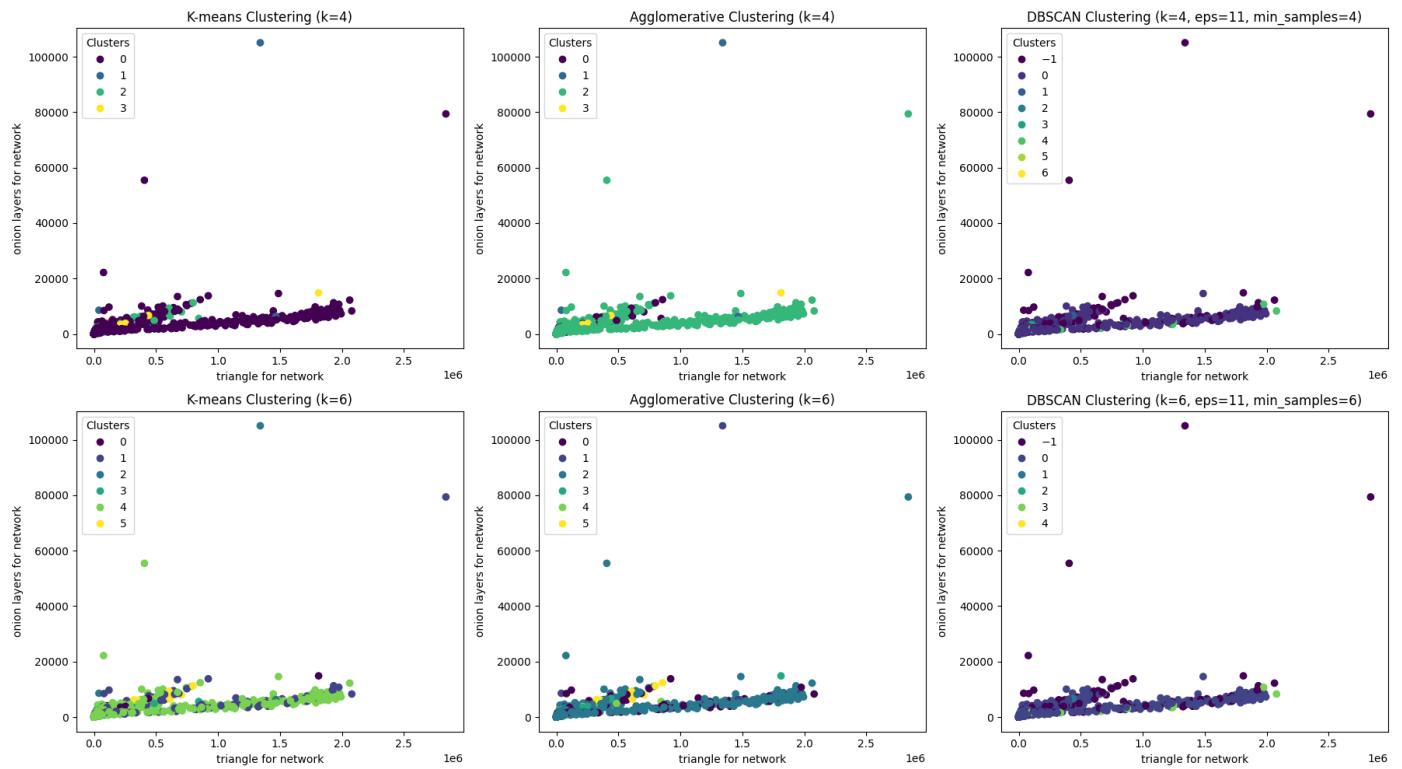
# Plot for k=6 clusters in k-means
scatter_6_kmeans = axes[1, 0].scatter(onefinaldf["triangle for network"], onefinaldf['onion layers for network'], c=onefinaldf['k = 6_Cluster'])
axes[1, 0].set_title('K-means Clustering (k=6)')
axes[1, 0].set_xlabel('triangle for network')
axes[1, 0].set_ylabel('onion layers for network')
axes[1, 0].legend(*scatter_6_kmeans.legend_elements(), title='Clusters')

# Plot for k=6 clusters in agglomerative clustering
scatter_6_agg = axes[1, 1].scatter(onefinaldf["triangle for network"], onefinaldf['onion layers for network'], c=onefinaldf['agg6Cluster'])
axes[1, 1].set_title('Agglomerative Clustering (k=6)')
axes[1, 1].set_xlabel('triangle for network')
axes[1, 1].set_ylabel('onion layers for network')
axes[1, 1].legend(*scatter_6_agg.legend_elements(), title='Clusters')

# Plot for DBSCAN clusters (k=4, eps=11, min_samples=5)
scatter_dbSCAN_4 = axes[0, 2].scatter(onefinaldf["triangle for network"], onefinaldf['onion layers for network'], c=onefinaldf['dbs4Cluster'])
axes[0, 2].set_title('DBSCAN Clustering (k=4, eps=11, min_samples=4)')
axes[0, 2].set_xlabel('triangle for network')
axes[0, 2].set_ylabel('onion layers for network')
axes[0, 2].legend(*scatter_dbSCAN_4.legend_elements(), title='Clusters')

# Plot for DBSCAN clusters (k=6, eps=0.8, min_samples=5)
scatter_dbSCAN_6 = axes[1, 2].scatter(onefinaldf["triangle for network"], onefinaldf['onion layers for network'], c=onefinaldf['dbs6Cluster'])
axes[1, 2].set_title('DBSCAN Clustering (k=6, eps=11, min_samples=6)')
axes[1, 2].set_xlabel('triangle for network')
axes[1, 2].set_ylabel('onion layers for network')
axes[1, 2].legend(*scatter_dbSCAN_6.legend_elements(), title='Clusters')

plt.tight_layout()
plt.show()
```



```
# Step 4: Visualize the clusters for k-means and agglomerative clustering
fig, axes = plt.subplots(nrows=2, ncols=3, figsize=(18, 10))

# Plot for k=4 clusters in k-means
scatter_4_kmeans = axes[0, 0].scatter(onefinaldf["NodeDegree"], onefinaldf['Common Connections'], c=onefinaldf['k=4_Cluster'], cmap='viridis')
axes[0, 0].set_title('K-means Clustering (k=4)')
axes[0, 0].set_xlabel('NodeDegree')
axes[0, 0].set_ylabel('Common Connections')
axes[0, 0].legend(*scatter_4_kmeans.legend_elements(), title='Clusters')

# Plot for k=4 clusters in agglomerative clustering
scatter_4_agg = axes[0, 1].scatter(onefinaldf["NodeDegree"], onefinaldf['Common Connections'], c=onefinaldf['agg4Cluster'], cmap='viridis')
axes[0, 1].set_title('Agglomerative Clustering (k=4)')
axes[0, 1].set_xlabel('NodeDegree')
axes[0, 1].set_ylabel('Common Connections')
axes[0, 1].legend(*scatter_4_agg.legend_elements(), title='Clusters')

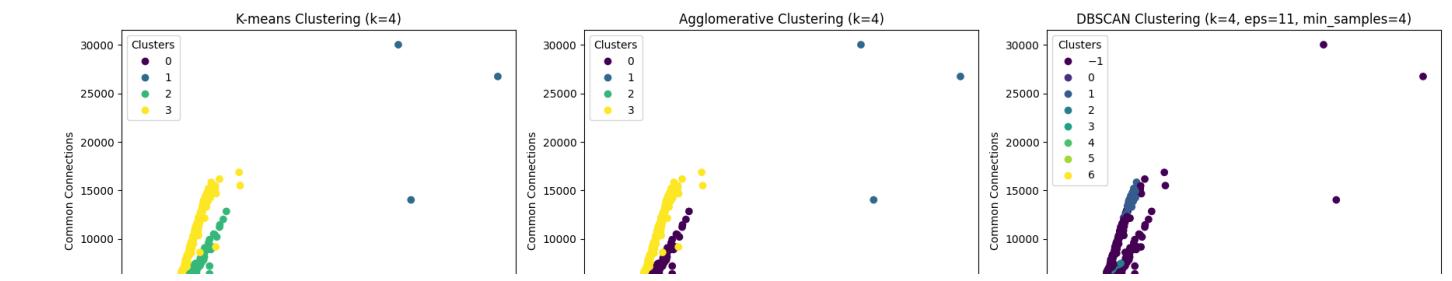
# Plot for DBSCAN clusters (k=4, eps=11, min_samples=5)
scatter_dbSCAN_4 = axes[0, 2].scatter(onefinaldf["NodeDegree"], onefinaldf['Common Connections'], c=onefinaldf['dbs4Cluster'], cmap='viridis')
axes[0, 2].set_title('DBSCAN Clustering (k=4, eps=11, min_samples=4)')
axes[0, 2].set_xlabel('NodeDegree')
axes[0, 2].set_ylabel('Common Connections')
axes[0, 2].legend(*scatter_dbSCAN_4.legend_elements(), title='Clusters')

# Plot for k=6 clusters in k-means
scatter_6_kmeans = axes[1, 0].scatter(onefinaldf["NodeDegree"], onefinaldf['Common Connections'], c=onefinaldf['k = 6_Cluster'], cmap='viridis')
axes[1, 0].set_title('K-means Clustering (k=6)')
axes[1, 0].set_xlabel('NodeDegree')
axes[1, 0].set_ylabel('Common Connections')
axes[1, 0].legend(*scatter_6_kmeans.legend_elements(), title='Clusters')

# Plot for k=6 clusters in agglomerative clustering
scatter_6_agg = axes[1, 1].scatter(onefinaldf["NodeDegree"], onefinaldf['Common Connections'], c=onefinaldf['agg6Cluster'], cmap='viridis')
axes[1, 1].set_title('Agglomerative Clustering (k=6)')
axes[1, 1].set_xlabel('NodeDegree')
axes[1, 1].set_ylabel('Common Connections')
axes[1, 1].legend(*scatter_6_agg.legend_elements(), title='Clusters')

# Plot for DBSCAN clusters (k=6, eps=0.8, min_samples=5)
scatter_dbSCAN_6 = axes[1, 2].scatter(onefinaldf["NodeDegree"], onefinaldf['Common Connections'], c=onefinaldf['dbs6Cluster'], cmap='viridis')
axes[1, 2].set_title('DBSCAN Clustering (k=6, eps=11, min_samples=6)')
axes[1, 2].set_xlabel('NodeDegree')
axes[1, 2].set_ylabel('Common Connections')
axes[1, 2].legend(*scatter_dbSCAN_6.legend_elements(), title='Clusters')

plt.tight_layout()
plt.show()
```



Metrics

```

import matplotlib.pyplot as plt
import numpy as np
from sklearn.metrics import silhouette_score, calinski_harabasz_score

# Extract features
X = onefinaldf[['triangle for network', 'onion layers for network']]

# Extract cluster labels
kmeans_labels_4 = onefinaldf['k=4_Cluster']
agg_labels_4 = onefinaldf['agg4Cluster']
dbSCAN_labels_4 = onefinaldf['dbs4Cluster']

kmeans_labels_6 = onefinaldf['k = 6_Cluster']
agg_labels_6 = onefinaldf['agg6Cluster']
dbSCAN_labels_6 = onefinaldf['dbs6Cluster']

# Calculate Silhouette Score
kmeans_silhouette_4 = silhouette_score(X, kmeans_labels_4)
agg_silhouette_4 = silhouette_score(X, agg_labels_4)
dbSCAN_silhouette_4 = silhouette_score(X, dbSCAN_labels_4)

kmeans_silhouette_6 = silhouette_score(X, kmeans_labels_6)
agg_silhouette_6 = silhouette_score(X, agg_labels_6)
dbSCAN_silhouette_6 = silhouette_score(X, dbSCAN_labels_6)

# Calculate Calinski-Harabasz Index
kmeans_calinski_harabasz_4 = calinski_harabasz_score(X, kmeans_labels_4)
agg_calinski_harabasz_4 = calinski_harabasz_score(X, agg_labels_4)
dbSCAN_calinski_harabasz_4 = calinski_harabasz_score(X, dbSCAN_labels_4)

kmeans_calinski_harabasz_6 = calinski_harabasz_score(X, kmeans_labels_6)
agg_calinski_harabasz_6 = calinski_harabasz_score(X, agg_labels_6)
dbSCAN_calinski_harabasz_6 = calinski_harabasz_score(X, dbSCAN_labels_6)

# Data for plotting
algorithms = ['K-means (k=4)', 'Agglomerative (k=4)', 'DBSCAN (k=4)', 'K-means (k=6)', 'Agglomerative (k=6)', 'DBSCAN (k=6)']
silhouette_scores = [kmeans_silhouette_4, agg_silhouette_4, dbSCAN_silhouette_4, kmeans_silhouette_6, agg_silhouette_6, dbSCAN_silhouette_6]
calinski_harabasz_scores = [kmeans_calinski_harabasz_4, agg_calinski_harabasz_4, dbSCAN_calinski_harabasz_4, kmeans_calinski_harabasz_6, agg_calinski_harabasz_6, dbSCAN_calinski_harabasz_6]

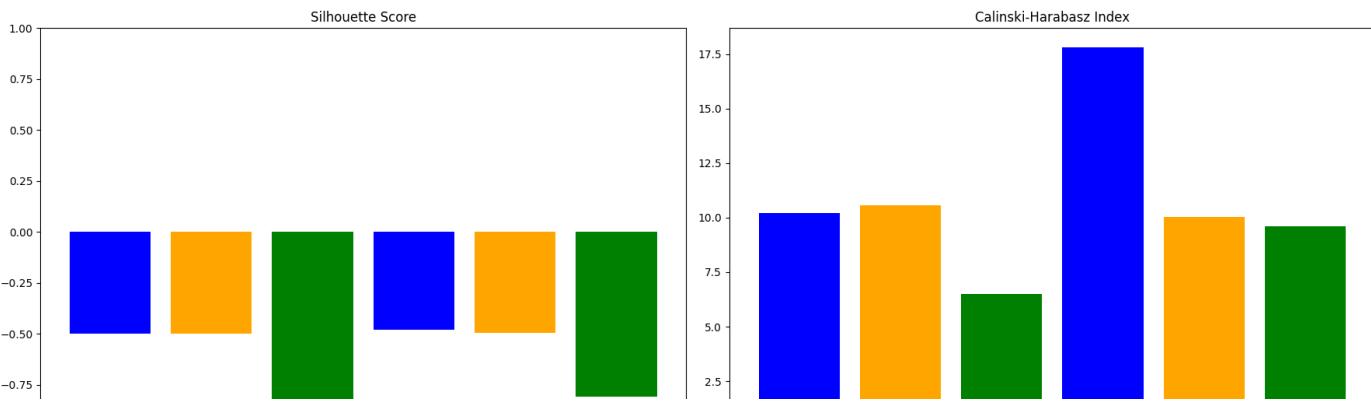
# Plot Silhouette Score and Calinski-Harabasz Index
fig, axes = plt.subplots(nrows=1, ncols=2, figsize=(18, 6))

# Plot Silhouette Score
axes[0].bar(algorithms, silhouette_scores, color=['blue', 'orange', 'green', 'blue', 'orange', 'green'])
axes[0].set_title('Silhouette Score')
axes[0].set_ylim(-1, 1)

# Plot Calinski-Harabasz Index
axes[1].bar(algorithms, calinski_harabasz_scores, color=['blue', 'orange', 'green', 'blue', 'orange', 'green'])
axes[1].set_title('Calinski-Harabasz Index')

plt.tight_layout()
plt.show()

```



```
#Clustering via networkx Library
Clus = nx.clustering(G)

Clust = [round(value, 2) for value in Clus.values()]

# Create a DataFrame with a single column
cludf = pd.DataFrame({'library Clustering': Clust})

# Display the DataFrame
print(cludf)
```

```
library Clustering
0 0.04
1 0.42
2 0.89
3 0.63
4 0.87
...
4034 1.00
4035 0.00
4036 1.00
4037 0.67
4038 0.56
```

[4039 rows x 1 columns]

```
onefinaldf = pd.concat([onefinaldf, cludf], axis = 1)
print(onefinaldf)
```

Node	NodeDegree	0_count	1_count	2_count	3_count	4_count	5_count	\
0	0	347	0	14	27	22	17	12
1	1	17	0	0	0	0	0	0
2	2	10	0	0	0	0	0	0
3	3	17	0	0	0	1	0	1
4	4	10	0	0	0	0	0	1
...	...	...	...	...	...	...	...	...
4034	4034	2	0	0	0	0	0	0
4035	4035	1	0	0	0	0	0	0
4036	4036	2	0	0	0	0	1	0
4037	4037	4	0	0	0	0	0	0
4038	4038	9	0	0	0	0	0	0

6_count	7_count	...	triangle for network	k=4_Cluster	k = 6_Cluster	\
0	15	18	...	39777	1	2
1	0	1	...	402	0	4
2	0	0	...	381	0	4
3	0	0	...	996	0	4
4	0	0	...	378	0	4
...	...	...	...	...	...	...
4034	0	0	...	3	0	4
4035	0	0	...	0	0	4
4036	0	0	...	3	0	4
4037	0	0	...	15	0	4
4038	2	1	...	114	0	4

agg4Cluster	agg6Cluster	dbs1Cluster	dbsCluster	dbs4Cluster	\
0	1	1	-1	-1	-1
1	2	2	-1	0	0
2	2	2	-1	0	0
3	2	2	-1	0	0
4	2	2	-1	0	0
...	...	...	...	...	...
4034	2	2	-1	0	0

4035	2	2	-1	0	0
4036	2	2	-1	0	0
4037	2	2	-1	0	0
4038	2	2	-1	0	0

dbs6Cluster	library	Clustering
0	-1	0.04
1	0	0.42
2	0	0.89
3	0	0.63
4	0	0.87
...	...	...
4034	0	1.00
4035	0	0.00
4036	0	1.00
4037	0	0.67
4038	0	0.56

[4039 rows x 266 columns]

```

#comparing scores with networkx library clustering
import matplotlib.pyplot as plt
import numpy as np
from sklearn.metrics import homogeneity_score, completeness_score, v_measure_score

# Extract cluster labels
kmeans_labels_4 = onefinaldf['k=4_Cluster']
agg_labels_4 = onefinaldf['agg4Cluster']
dbscan_labels_4 = onefinaldf['dbs4Cluster']

kmeans_labels_6 = onefinaldf['k = 6_Cluster']
agg_labels_6 = onefinaldf['agg6Cluster']
dbscan_labels_6 = onefinaldf['dbs6Cluster']

# Calculate scores for k=4 clusters
homogeneity_kmeans_4 = homogeneity_score(onefinaldf['library Clustering'], kmeans_labels_4)
completeness_kmeans_4 = completeness_score(onefinaldf['library Clustering'], kmeans_labels_4)
v_measure_kmeans_4 = v_measure_score(onefinaldf['library Clustering'], kmeans_labels_4)

homogeneity_agg_4 = homogeneity_score(onefinaldf['library Clustering'], agg_labels_4)
completeness_agg_4 = completeness_score(onefinaldf['library Clustering'], agg_labels_4)
v_measure_agg_4 = v_measure_score(onefinaldf['library Clustering'], agg_labels_4)

homogeneity_dbscan_4 = homogeneity_score(onefinaldf['library Clustering'], dbscan_labels_4)
completeness_dbscan_4 = completeness_score(onefinaldf['library Clustering'], dbscan_labels_4)
v_measure_dbscan_4 = v_measure_score(onefinaldf['library Clustering'], dbscan_labels_4)

# Calculate scores for k=6 clusters
homogeneity_kmeans_6 = homogeneity_score(onefinaldf['library Clustering'], kmeans_labels_6)
completeness_kmeans_6 = completeness_score(onefinaldf['library Clustering'], kmeans_labels_6)
v_measure_kmeans_6 = v_measure_score(onefinaldf['library Clustering'], kmeans_labels_6)

homogeneity_agg_6 = homogeneity_score(onefinaldf['library Clustering'], agg_labels_6)
completeness_agg_6 = completeness_score(onefinaldf['library Clustering'], agg_labels_6)
v_measure_agg_6 = v_measure_score(onefinaldf['library Clustering'], agg_labels_6)

homogeneity_dbscan_6 = homogeneity_score(onefinaldf['library Clustering'], dbscan_labels_6)
completeness_dbscan_6 = completeness_score(onefinaldf['library Clustering'], dbscan_labels_6)
v_measure_dbscan_6 = v_measure_score(onefinaldf['library Clustering'], dbscan_labels_6)

# Data for plotting
algorithms = ['K-means (k=4)', 'Agglomerative (k=4)', 'DBSCAN (k=4)', 'K-means (k=6)', 'Agglomerative (k=6)', 'DBSCAN (k=6)']
homogeneity_scores = [homogeneity_kmeans_4, homogeneity_agg_4, homogeneity_dbscan_4, homogeneity_kmeans_6, homogeneity_agg_6, homogeneity_dbscan_6]
completeness_scores = [completeness_kmeans_4, completeness_agg_4, completeness_dbscan_4, completeness_kmeans_6, completeness_agg_6, completeness_dbscan_6]
v_measure_scores = [v_measure_kmeans_4, v_measure_agg_4, v_measure_dbscan_4, v_measure_kmeans_6, v_measure_agg_6, v_measure_dbscan_6]

# Plot scores
fig, axes = plt.subplots(nrows=1, ncols=3, figsize=(18, 4))

# Plot Homogeneity Score
axes[0].bar(algorithms, homogeneity_scores, color=['blue', 'orange', 'green', 'blue', 'orange', 'green'])
axes[0].set_title('Homogeneity Score')

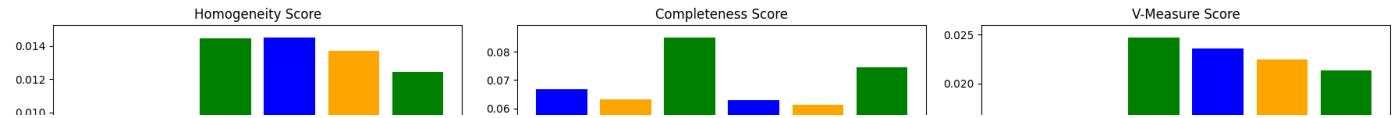
# Plot Completeness Score
axes[1].bar(algorithms, completeness_scores, color=['blue', 'orange', 'green', 'blue', 'orange', 'green'])
axes[1].set_title('Completeness Score')

# Plot V-Measure Score
axes[2].bar(algorithms, v_measure_scores, color=['blue', 'orange', 'green', 'blue', 'orange', 'green'])
axes[2].set_title('V-Measure Score')

plt.tight_layout()
plt.show()

```

```
/usr/local/lib/python3.10/dist-packages/sklearn/metrics/_cluster/_supervised.py:64: UserWarning: Clustering metrics expects discrete values
  warnings.warn(msg, UserWarning)
/usr/local/lib/python3.10/dist-packages/sklearn/metrics/_cluster/_supervised.py:64: UserWarning: Clustering metrics expects discrete values
  warnings.warn(msg, UserWarning)
/usr/local/lib/python3.10/dist-packages/sklearn/metrics/_cluster/_supervised.py:64: UserWarning: Clustering metrics expects discrete values
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  warnings.warn(msg, UserWarning)
/usr/local/lib/python3.10/dist-packages/sklearn/metrics/_cluster/_supervised.py:64: UserWarning: Clustering metrics expects discrete values
  warnings.warn(msg, UserWarning)
/usr/local/lib/python3.10/dist-packages/sklearn/metrics/_cluster/_supervised.py:64: UserWarning: Clustering metrics expects discrete values
  warnings.warn(msg, UserWarning)
/usr/local/lib/python3.10/dist-packages/sklearn/metrics/_cluster/_supervised.py:64: UserWarning: Clustering metrics expects discrete values
  warnings.warn(msg, UserWarning)
/usr/local/lib/python3.10/dist-packages/sklearn/metrics/_cluster/_supervised.py:64: UserWarning: Clustering metrics expects discrete values
  warnings.warn(msg, UserWarning)
/usr/local/lib/python3.10/dist-packages/sklearn/metrics/_cluster/_supervised.py:64: UserWarning: Clustering metrics expects discrete values
  warnings.warn(msg, UserWarning)
/usr/local/lib/python3.10/dist-packages/sklearn/metrics/_cluster/_supervised.py:64: UserWarning: Clustering metrics expects discrete values
  warnings.warn(msg, UserWarning)
/usr/local/lib/python3.10/dist-packages/sklearn/metrics/_cluster/_supervised.py:64: UserWarning: Clustering metrics expects discrete values
  warnings.warn(msg, UserWarning)
/usr/local/lib/python3.10/dist-packages/sklearn/metrics/_cluster/_supervised.py:64: UserWarning: Clustering metrics expects discrete values
  warnings.warn(msg, UserWarning)
/usr/local/lib/python3.10/dist-packages/sklearn/metrics/_cluster/_supervised.py:64: UserWarning: Clustering metrics expects discrete values
  warnings.warn(msg, UserWarning)
/usr/local/lib/python3.10/dist-packages/sklearn/metrics/_cluster/_supervised.py:64: UserWarning: Clustering metrics expects discrete values
  warnings.warn(msg, UserWarning)
/usr/local/lib/python3.10/dist-packages/sklearn/metrics/_cluster/_supervised.py:64: UserWarning: Clustering metrics expects discrete values
  warnings.warn(msg, UserWarning)
/usr/local/lib/python3.10/dist-packages/sklearn/metrics/_cluster/_supervised.py:64: UserWarning: Clustering metrics expects discrete values
  warnings.warn(msg, UserWarning)
```



```
onefinaldf.to_csv('/content/drive/MyDrive/Colab Notebooks/onefinaldata.csv', index=False)
```



```

import pandas as pd
import networkx as nx
from itertools import combinations

twohopcommon_data = []

for node in range(500):
    abData = []
    length = nx.single_source_shortest_path_length(G, node, cutoff=2)
    acData = sorted(list(length.keys()))

    if node in acData:
        acData.remove(node)

    # Use itertools.combinations to iterate over unique pairs of elements in acData
    for neighbor1, neighbor2 in combinations(acData, 2):
        # Check if the common neighbors have 1 unit distance with the original nodes

        N = nx.shortest_path_length(G, source=neighbor1, target=neighbor2)

        # Count only when the shortest distance is 1
        if N == 1:
            abData.append(1)

    twohopcommon_data.append(sum(abData)) # Append the sum of abData for the current node

```

```

# Create a DataFrame from the list of sums
twohopcommondf = pd.DataFrame(twohopcommon_data, columns=["Common Connections"])

# Display the DataFrame
print(twohopcommondf)

```

	Common Connections
0	33343
1	2849
2	2856
3	2849
4	2856
..	...
495	3811
496	8264
497	8377
498	3780
499	3438

[500 rows x 1 columns]

```
twohopcommondf.to_csv('/content/drive/MyDrive/Colab Notebooks/twohopcommonp1.csv', index=False)
```

```

path1 = '/content/drive/MyDrive/Colab Notebooks/twohopcommonp1.csv'
twohopcommon1 = pd.read_csv(path1)
print(twohopcommon1)

```

	Common Connections
0	33343
1	2849
2	2856
3	2849
4	2856
..	...
495	3811
496	8264
497	8377
498	3780
499	3438

[500 rows x 1 columns]

```

twohopcommon_data2 = []

for node in range(501, 1001):
    abData = []
    length = nx.single_source_shortest_path_length(G, node, cutoff=2)
    acData = sorted(list(length.keys()))

    if node in acData:
        acData.remove(node)

    # Use itertools.combinations to iterate over unique pairs of elements in acData
    for neighbor1, neighbor2 in combinations(acData, 2):
        # Check if the common neighbors have 1 unit distance with the original nodes

        N = nx.shortest_path_length(G, source=neighbor1, target=neighbor2)

        # Count only when the shortest distance is 1
        if N == 1:
            abData.append(1)

    twohopcommon_data2.append(sum(abData)) # Append the sum of abData for the current node

# Create a DataFrame from the list of sums
twohopcommondf2 = pd.DataFrame(twohopcommon_data2, columns=["Common Connections"])

# Display the DataFrame
print(twohopcommondf2)

   Common Connections
0              3440
1              4331
2              7810
3              4053
4              3660
..             ...
495             27916
496             27699
497             27786
498             27718
499             27779

[500 rows x 1 columns]

twohopcommondf2.to_csv('/content/drive/MyDrive/Colab Notebooks/twohopcommonp2.csv', index=False)

path2 = '/content/drive/MyDrive/Colab Notebooks/twohopcommonp2.csv'
twohopcommon2 = pd.read_csv(path2)
print(twohopcommon2)

   Common Connections
0              3440
1              4331
2              7810
3              4053
4              3660
..             ...
495             27916
496             27699
497             27786
498             27718
499             27779

[500 rows x 1 columns]

```

```

import pandas as pd
import networkx as nx
from itertools import combinations

twohopcommon_data3 = []

for node in range(1001, 1501):
    abData = []
    length = nx.single_source_shortest_path_length(G, node, cutoff=2)
    acData = sorted(list(length.keys()))

    if node in acData:
        acData.remove(node)

    # Use itertools.combinations to iterate over unique pairs of elements in acData
    for neighbor1, neighbor2 in combinations(acData, 2):
        # Check if the common neighbors have 1 unit distance with the original nodes

        N = nx.shortest_path_length(G, source=neighbor1, target=neighbor2)

        # Count only when the shortest distance is 1
        if N == 1:
            abData.append(1)

    twohopcommon_data3.append(sum(abData)) # Append the sum of abData for the current node

```

# Create a DataFrame from the list of sums  
twohopcommondf3 = pd.DataFrame(twohopcommon\_data3, columns=["Common Connections"])

# Display the DataFrame  
print(twohopcommondf3)

	Common Connections
0	29826
1	27791
2	27700
3	27675
4	27781
..	...
495	27932
496	27766
497	27789
498	27783
499	27766

[500 rows x 1 columns]

twohopcommondf3.to\_csv('/content/drive/MyDrive/Colab Notebooks/twohopcommonp3.csv', index=False)

path3 = '/content/drive/MyDrive/Colab Notebooks/twohopcommonp3.csv'  
twohopcommon3 = pd.read\_csv(path3)  
print(twohopcommon3)

	Common Connections
0	29826
1	27791
2	27700
3	27675
4	27781
..	...
495	27932
496	27766
497	27789
498	27783
499	27766

[500 rows x 1 columns]

```

import pandas as pd
import networkx as nx
from itertools import combinations
twohopcommon_data4 = []

for node in range(1501, 2001):
    abData = []
    length = nx.single_source_shortest_path_length(G, node, cutoff=2)
    acData = sorted(list(length.keys()))

    if node in acData:
        acData.remove(node)

    # Use itertools.combinations to iterate over unique pairs of elements in acData
    for neighbor1, neighbor2 in combinations(acData, 2):
        # Check if the common neighbors have 1 unit distance with the original nodes

        N = nx.shortest_path_length(G, source=neighbor1, target=neighbor2)

        # Count only when the shortest distance is 1
        if N == 1:
            abData.append(1)

    twohopcommon_data4.append(sum(abData)) # Append the sum of abData for the current node

# Create a DataFrame from the list of sums
twohopcommondf4 = pd.DataFrame(twohopcommon_data4, columns=["Common Connections"])

# Display the DataFrame
print(twohopcommondf4)

twohopcommondf4.to_csv('/content/drive/MyDrive/Colab Notebooks/twohopcommonp4.csv', index=False)

twohopcommondf4.to_csv('/content/drive/MyDrive/Colab Notebooks/twohopcommonp4.csv', index=False)

path4 = '/content/drive/MyDrive/Colab Notebooks/twohopcommonp4.csv'
twohopcommon4 = pd.read_csv(path4)
print(twohopcommon4)

   Common Connections
0              27777
1              27725
2              27767
3              27737
4              42496
..              ...
495             30776
496             30645
497             30851
498             30746
499             30747

[500 rows x 1 columns]

```

```

twohopcommon_data5 = []

for node in range(2001, 2501):
    abData = []
    length = nx.single_source_shortest_path_length(G, node, cutoff=2)
    acData = sorted(list(length.keys()))

    if node in acData:
        acData.remove(node)

    # Use itertools.combinations to iterate over unique pairs of elements in acData
    for neighbor1, neighbor2 in combinations(acData, 2):
        # Check if the common neighbors have 1 unit distance with the original nodes

        N = nx.shortest_path_length(G, source=neighbor1, target=neighbor2)

        # Count only when the shortest distance is 1
        if N == 1:
            abData.append(1)

    twohopcommon_data5.append(sum(abData)) # Append the sum of abData for the current node

# Create a DataFrame from the list of sums
twohopcommondf5 = pd.DataFrame(twohopcommon_data5, columns=["Common Connections"])

# Display the DataFrame
print(twohopcommondf5)

twohopcommondf5.to_csv('/content/drive/MyDrive/Colab Notebooks/twohopcommonp5.csv', index=False)

twohopcommondf5.to_csv('/content/drive/MyDrive/Colab Notebooks/twohopcommonp5.csv', index=False)

path5 = '/content/drive/MyDrive/Colab Notebooks/twohopcommonp5.csv'
twohopcommon5 = pd.read_csv(path5)
print(twohopcommon5)

   Common Connections
0              30845
1              30729
2              31117
3              30855
4              30625
..             ...
495             30796
496             30755
497             31239
498             30687
499             30599

[500 rows x 1 columns]

```

```
twohopcommon_data6 = []

for node in range(2501, 3001):
    abData = []
    length = nx.single_source_shortest_path_length(G, node, cutoff=2)
    acData = sorted(list(length.keys()))

    if node in acData:
        acData.remove(node)

    # Use itertools.combinations to iterate over unique pairs of elements in acData
    for neighbor1, neighbor2 in combinations(acData, 2):
        # Check if the common neighbors have 1 unit distance with the original nodes

        N = nx.shortest_path_length(G, source=neighbor1, target=neighbor2)

        # Count only when the shortest distance is 1
        if N == 1:
            abData.append(1)

    twohopcommon_data6.append(sum(abData)) # Append the sum of abData for the current node

# Create a DataFrame from the list of sums
twohopcommondf6 = pd.DataFrame(twohopcommon_data6, columns=["Common Connections"])

# Display the DataFrame
print(twohopcommondf6)
```

```
twohopcommondf6.to_csv('/content/drive/MyDrive/Colab Notebooks/twohopcommonp6.csv', index=False)
```

```
path6 = '/content/drive/MyDrive/Colab Notebooks/twohopcommonp6.csv'
twohopcommon6 = pd.read_csv(path6)
print(twohopcommon6)
```

```
Common Connections
0           30742
1           30897
2           30764
3           30629
4           30775
..          ...
495          14778
496          14803
497          14814
498          14826
499          14726
```

```
[500 rows x 1 columns]
```

```

twohopcommon_data7 = []

for node in range(3001, 3501):
    abData = []
    length = nx.single_source_shortest_path_length(G, node, cutoff=2)
    acData = sorted(list(length.keys()))

    if node in acData:
        acData.remove(node)

    # Use itertools.combinations to iterate over unique pairs of elements in acData
    for neighbor1, neighbor2 in combinations(acData, 2):
        # Check if the common neighbors have 1 unit distance with the original nodes

        N = nx.shortest_path_length(G, source=neighbor1, target=neighbor2)

        # Count only when the shortest distance is 1
        if N == 1:
            abData.append(1)

    twohopcommon_data7.append(sum(abData)) # Append the sum of abData for the current node

# Create a DataFrame from the list of sums
twohopcommondf7 = pd.DataFrame(twohopcommon_data7, columns=["Common Connections"])

# Display the DataFrame
print(twohopcommondf7)

twohopcommondf7.to_csv('/content/drive/MyDrive/Colab Notebooks/twohopcommonp7.csv', index=False)

path7 = '/content/drive/MyDrive/Colab Notebooks/twohopcommonp7.csv'
twohopcommon7 = pd.read_csv(path7)
print(twohopcommon7)

twohopcommon_data8 = []

for node in range(3501, len(G)):
    abData = []
    length = nx.single_source_shortest_path_length(G, node, cutoff=2)
    acData = sorted(list(length.keys()))

    if node in acData:
        acData.remove(node)

    # Use itertools.combinations to iterate over unique pairs of elements in acData
    for neighbor1, neighbor2 in combinations(acData, 2):
        # Check if the common neighbors have 1 unit distance with the original nodes

        N = nx.shortest_path_length(G, source=neighbor1, target=neighbor2)

        # Count only when the shortest distance is 1
        if N == 1:
            abData.append(1)

    twohopcommon_data8.append(sum(abData)) # Append the sum of abData for the current node

# Create a DataFrame from the list of sums
twohopcommondf8 = pd.DataFrame(twohopcommon_data8, columns=["Common Connections"])

# Display the DataFrame
print(twohopcommondf8)

twohopcommondf8.to_csv('/content/drive/MyDrive/Colab Notebooks/twohopcommonp8.csv', index=False)

```

```
twohopcommondf8.to_csv('/content/drive/MyDrive/Colab Notebooks/twohopcommonp8.csv', index=False)
```

```
path8 = '/content/drive/MyDrive/Colab Notebooks/twohopcommonp8.csv'
twohopcommon8 = pd.read_csv(path8)
print(twohopcommon8)
```

	Common Connections
0	5588
1	5352
2	5347
3	5352
4	5335
..	...
533	203
534	204
535	203
536	201
537	196

[538 rows x 1 columns]

```
twohopcommondf = pd.concat([twohopcommon1, twohopcommon2, twohopcommon3, twohopcommon4, twohopcommon5, twohopcommon6, twohopcommon7, twohopcommon8])
print(twohopcommondf)
```

	Common Connections
0	33343
1	2849
2	2856
3	2849
4	2856
..	...
4033	203
4034	204
4035	203
4036	201
4037	196

[4038 rows x 1 columns]

```
import pandas as pd
#near degree data
twodegree_data = []

for node in range(len(G)):
    abData = []
    length = nx.single_source_shortest_path_length(G, node, cutoff=2)
    acData = sorted(list(length.keys()))

    if node in acData:
        acData.remove(node)

    for j in acData:
        M = G.degree(j)
        abData.append(M)

    twodegree_data.append(abData) # Append a new list as a row to all_data

# Create a DataFrame from the list of lists
twodegreedf = pd.DataFrame(twodegree_data, columns= [f'{i} degree connections' for i in range(1, 2916)])

# Display the DataFrame
print(twodegreedf)
```

	1 degree connections	2 degree connections	3 degree connections	4 degree connections	5 degree connections	6 degree connections
0	17	10	17	10	13	6
1	347	10	17	8	8	8
2	347	17	17	59	59	8
3	347	17	10	59	59	8
4	347	17	10	59	59	8
..	...	...	...	...	...	...
4034	8	59	8	8	8	8
4035	8	59	8	8	8	8
4036	8	59	8	8	8	8
4037	8	59	8	8	8	8
4038	8	59	8	8	8	8

```

1          10          13          6
2          10          13          6
3          10          13          6
4          17          13          6
...
...          ...          ...
4034        14          2          1
4035        14          2          1
4036        14          2          1
4037        14          2          1
4038        14          2          1

    7 degree connections  8 degree connections  9 degree connections \
0            20          8          57
1            20          8          57
2            20          8          57
3            20          8          57
4            20          8          57
...
...          ...
4034        6          9          2
4035        6          9          2
4036        6          9          2
4037        6          9          2
4038        6          9          2

    10 degree connections ...  2906 degree connections \
0           10 ...          NaN
1           10 ...          NaN
2           10 ...          NaN
3           10 ...          NaN
4           10 ...          NaN
...
...          ...
4034        7 ...          NaN
4035        7 ...          NaN
4036        7 ...          NaN
4037        7 ...          NaN
4038        7 ...          NaN

    2907 degree connections  2908 degree connections \
0             NaN          NaN
1             NaN          NaN
2             NaN          NaN
3             NaN          NaN
4             NaN          NaN

#transforming degree data by adding all the degree and counting for a specific degree

# Assuming you have a list of 100 values named values_to_count
values_to_count = range(1046) # Replace ... with the actual values

# Add columns for each value in values_to_count
for value in values_to_count:
    column_name = f'{value}_count'
    twodegreedf[column_name] = twodegreedf.apply(lambda row: row.eq(value).sum(), axis=1)

print(twodegreedf)

```

	4035	4036	4037	4038
7 degree connections	14	14	14	14
8 degree connections	2	2	2	2
9 degree connections	1	1	1	1

7 degree connections 8 degree connections 9 degree connections \

```

4036      / ...   0   0   0
4037      7 ...   0   0   0
4038      7 ...   0   0   0

  1039_count  1040_count  1041_count  1042_count  1043_count  1044_count \
0          0          0          0          0          0          0
1          0          0          0          0          0          0
2          0          0          0          0          0          0
3          0          0          0          0          0          0
4          0          0          0          0          0          0
...
...       ...     ...     ...     ...     ...
4034      0          0          0          0          0          0
4035      0          0          0          0          0          0
4036      0          0          0          0          0          0
4037      0          0          0          0          0          0
4038      0          0          0          0          0          0

  1045_count
0          1
1          1
2          1
3          1
4          1
...
...       ...
4034      0
4035      0
4036      0
4037      0
4038      0

```

[4039 rows x 3961 columns]

```

twohopdegreedf = twodegreedf.drop(twodegreedf.columns[:2915], axis=1)
#dropping columns in degreedf
twohopdegreedf = twohopdegreedf.drop(twohopdegreedf.columns[793:1045], axis=1)
twohopdegreedf = twohopdegreedf.drop(twohopdegreedf.columns[756:792], axis=1)
twohopdegreedf = twohopdegreedf.drop(twohopdegreedf.columns[548:755], axis=1)
twohopdegreedf = twohopdegreedf.drop(twohopdegreedf.columns[348:547], axis=1)
twohopdegreedf = twohopdegreedf.drop(twohopdegreedf.columns[295:347], axis=1)
twohopdegreedf = twohopdegreedf.drop(twohopdegreedf.columns[292:294], axis=1)
twohopdegreedf = twohopdegreedf.drop(twohopdegreedf.columns[255:291], axis=1)
twohopdegreedf = twohopdegreedf.drop(twohopdegreedf.columns[246:254], axis=1)
twohopdegreedf = twohopdegreedf.drop(twohopdegreedf.columns[236:245], axis=1)
twohopdegreedf = twohopdegreedf.drop(twohopdegreedf.columns[232:234], axis=1)
twohopdegreedf = twohopdegreedf.drop(twohopdegreedf.columns[230:231], axis=1)
twohopdegreedf = twohopdegreedf.drop(twohopdegreedf.columns[227:229], axis=1)
twohopdegreedf = twohopdegreedf.drop(twohopdegreedf.columns[225:226], axis=1)
# Display the DataFrame after dropping columns
print(twohopdegreedf)

```

	0_count	1_count	2_count	3_count	4_count	5_count	6_count	7_count	\
0	0	25	45	38	35	38	32	30	
1	0	14	27	22	17	12	15	18	
2	0	14	27	22	17	12	15	18	
3	0	14	27	22	17	12	15	18	
4	0	14	27	22	17	12	15	18	
...	...	...	...	...	...	...	...	...	
4034	0	7	9	4	9	1	4	5	
4035	0	6	10	4	9	1	5	4	
4036	0	7	9	4	9	1	4	5	
4037	0	7	10	4	8	1	4	5	
4038	0	7	10	4	9	1	4	5	
8_count	9_count	...	235_count	245_count	254_count	291_count	...		
0	33	39	...	1	1	1	0		
1	15	15	...	0	0	0	0		
2	15	15	...	0	0	0	0		
3	15	15	...	0	0	0	0		
4	15	15	...	0	0	0	0		
...	...	...	...	...	...	...	...		
4034	4	8	...	0	0	0	0		
4035	4	7	...	0	0	0	0		
4036	4	8	...	0	0	0	0		
4037	5	6	...	0	0	0	0		
4038	4	6	...	0	0	0	0		
294_count	347_count	547_count	755_count	792_count	1045_count				
0	1	0	0	1	1	1			
1	0	1	0	0	0	1			
2	0	1	0	0	0	1			
3	0	1	0	0	0	1			
4	0	1	0	0	0	1			
...	...	...	...	...	...	...			

4034	0	0	0	0	0	0
4035	0	0	0	0	0	0
4036	0	0	0	0	0	0
4037	0	0	0	0	0	0
4038	0	0	0	0	0	0

[4039 rows x 239 columns]

```
path8 = '/content/drive/MyDrive/Colab Notebooks/twohopdegreedf.csv'
twohopdegreedf = pd.read_csv(path8)
print(twohopdegreedf)

import pandas as pd
import networkx as nx

twodcData = []

for node in G.nodes():
    length = nx.single_source_shortest_path_length(G, node, cutoff=2)
    acData = sorted(list(length.keys()))

    # Include the node itself in the subgraph
    subgraph = G.subgraph(acData)

    # Calculate the average neighbor degree for the subgraph
    average_neighbor_degree = nx.average_neighbor_degree(subgraph)

    # Round the values to 2 decimal places
    average_degrees = {k: round(v, 2) for k, v in average_neighbor_degree.items()}

    # Sum the average degrees
    B = sum(average_degrees.values())

    twodcData.append(B)

# Create a DataFrame with a single column
twoadcdf = pd.DataFrame({'Average Degree Connectivity for network': twodcData})

# Display the DataFrame
print(twoadcdf)

twohopdegreedf.to_csv('/content/drive/MyDrive/Colab Notebooks/twohopdegreedf.csv', index=False)
twoadcdf.to_csv('/content/drive/MyDrive/Colab Notebooks/twoadcdf.csv', index=False)

path8 = '/content/drive/MyDrive/Colab Notebooks/twoadcdf.csv'
twoadcdf = pd.read_csv(path8)
print(twoadcdf)
```

```

twonode_bridge_counts = {}

# Iterate over nodes from 0 to (number of nodes - 1)
for node in range(len(G)):
    # Create a subgraph containing only the current node and its neighbors
    length = nx.single_source_shortest_path_length(G, node, cutoff=2)
    acData = sorted(list(length.keys()))

    # Include the node itself in the subgraph
    subgraph = G.subgraph(acData)

    # Calculate local bridges for the subgraph
    local_bridges = list(nx.local_bridges(subgraph, with_span=False))

    # Count the number of bridges for the current node
    bridge_count = len(local_bridges)

    # Store the result in the dictionary
    twonode_bridge_counts[node] = bridge_count

# Convert the dictionary to a DataFrame
twohopbrdf = pd.DataFrame(list(twonode_bridge_counts.values()), columns=['Bridge_Count'])

# Display the DataFrame
print(twohopbrdf)

```

	Bridge_Count
0	26
1	14
2	14
3	14
4	14
...	...
4034	7
4035	7
4036	7
4037	7
4038	7

[4039 rows x 1 columns]

```
twohopbrdf.to_csv('/content/drive/MyDrive/Colab Notebooks/twohopbrdf.csv', index=False)
```

```

path8 = '/content/drive/MyDrive/Colab Notebooks/twohopbrdf.csv'
twohopbrdf = pd.read_csv(path8)
print(twohopbrdf)

```

	Bridge_Count
0	26
1	14
2	14
3	14
4	14
...	...
4034	7
4035	7
4036	7
4037	7
4038	7

[4039 rows x 1 columns]

```
#Returns the boundary expansion of the set S.

#The boundary expansion is the quotient of the size of the node boundary and the cardinality of S. [1]
twoboundary_expansion_list = []

# Iterate over all nodes in the graph
for node in G.nodes():
    # Create a subgraph with the current node and its neighbors
    length = nx.single_source_shortest_path_length(G, node, cutoff=2)
    acData = sorted(list(length.keys()))

    # Include the node itself in the subgraph
    subgraph = G.subgraph(acData)
    # Calculate the boundary expansion for the current node
    boundary_expansion = nx.boundary_expansion(G, subgraph)

    # Append the boundary expansion to the list
    twoboundary_expansion_list.append(boundary_expansion)

# Create a DataFrame from the list
twoboundddf = pd.DataFrame({'Boundary_Expansion': twoboundary_expansion_list})

# Display the DataFrame
print(twoboundddf)
twoboundddf.to_csv('/content/drive/MyDrive/Colab Notebooks/twoboundddf.csv', index=False)
```

	Boundary_Expansion
0	1.146807
1	3.364943
2	3.364943
3	3.364943
4	3.364943
...	...
4034	0.066667
4035	0.066667
4036	0.066667
4037	0.066667
4038	0.066667

[4039 rows x 1 columns]

```
path8 = '/content/drive/MyDrive/Colab Notebooks/twoboundddf.csv'
twoboundddf = pd.read_csv(path8)
print(twoboundddf)
```

	Boundary_Expansion
0	1.146807
1	3.364943
2	3.364943
3	3.364943
4	3.364943
...	...
4034	0.066667
4035	0.066667
4036	0.066667
4037	0.066667
4038	0.066667

[4039 rows x 1 columns]

```
#The average global efficiency of a graph is the average efficiency of all pairs of nodes
import pandas as pd
import networkx as nx
from itertools import combinations

twoglobaleff_list = []
two_weight_matching_list = []
two_random_list = []
panther_list = []
# Iterate over all nodes in the graph
for node in G.nodes():
    # Create a subgraph with the current node and its neighbors
    length = nx.single_source_shortest_path_length(G, node, cutoff=2)
    acData = sorted(list(length.keys()))

    # Include the node itself in the subgraph
    subgraph = G.subgraph(acData)

    # Calculate the boundary expansion for the current node
    globaleff = nx.global_efficiency(subgraph)

    weighteff = len(list(nx.max_weight_matching(subgraph, maxcardinality=False, weight='weight')))

    result = nx.non_randomness(subgraph, k=None, weight='weight')
    second_value = result[1]

    #panther = len(nx.panther_similarity(subgraph, 0, k=5, path_length=5, c=0.5, delta=0.1, eps=None, weight='weight'))

    # Append the boundary expansion to the list
    twoglobaleff_list.append(globaleff)
    two_weight_matching_list.append(weighteff)
    two_random_list.append(second_value)
    #panther_list.append(panther)
# Create a DataFrame from the list
twoglobaldf = pd.DataFrame({
    'global_efficiency': twoglobaleff_list,
    'max_weight_matching': two_weight_matching_list,
    'random_probability': two_random_list,
    # 'panther_similarity': panther_list # Uncomment if needed
})
# Display the DataFrame
print(twoglobaldf)
```

```
twoglobaldf.to_csv('/content/drive/MyDrive/Colab Notebooks/twoglobaldf.csv', index=False)
```

	global_efficiency	max_weight_matching	random_probability
0	0.435346	744	-66.852262
1	0.523734	164	75.626117
2	0.523734	164	75.626117
3	0.523734	164	75.626117
4	0.523734	164	75.626117
...	...	...	...
4034	0.557910	26	10.139742
4035	0.557910	26	10.139742
4036	0.557910	26	10.139742
4037	0.557910	26	10.139742
4038	0.557910	26	10.139742

[4039 rows x 3 columns]

```
path8 = '/content/drive/MyDrive/Colab Notebooks/twoglobaldf.csv'
twoglobaldf = pd.read_csv(path8)
print(twoglobaldf)
```

	global_efficiency	max_weight_matching	random_probability
0	0.435346	744	-66.852262
1	0.523734	164	75.626117
2	0.523734	164	75.626117
3	0.523734	164	75.626117
4	0.523734	164	75.626117
...	...	...	...
4034	0.557910	26	10.139742
4035	0.557910	26	10.139742
4036	0.557910	26	10.139742
4037	0.557910	26	10.139742
4038	0.557910	26	10.139742

[4039 rows x 3 columns]

```
#The Kemeny constant measures the time needed for spreading across a graph. Low values indicate a closely connected graph whereas high value
```

```
twokemeny_list = []
```

```
# Iterate over all nodes in the graph
```

```
for node in G.nodes():
```

```
    # Create a subgraph with the current node and its neighbors
```

```
    length = nx.single_source_shortest_path_length(G, node, cutoff=2)
```

```
    acData = sorted(list(length.keys()))
```

```
# Include the node itself in the subgraph
```

```
subgraph = G.subgraph(acData)
```

```
# Calculate the boundary expansion for the current node
```

```
kemeny = nx.kemeny_constant(subgraph)
```

```
# Append the boundary expansion to the list
```

```
twokemeny_list.append(kemeny)
```

```
# Create a DataFrame from the list
```

```
twokemenydf = pd.DataFrame({'kemeny constant for network': twokemeny_list})
```

```
# Display the DataFrame
```

```
print(twokemenydf)
```

```
twokemenydf.to_csv('/content/drive/MyDrive/Colab Notebooks/twokemenydf.csv', index=False)
```

```
    kemeny constant for network
```

```
0           2030.583502
```

```
1           406.232390
```

```
2           406.232390
```

```
3           406.232390
```

```
4           406.232390
```

```
...
...
```

```
4034        68.707931
```

```
4035        68.707931
```

```
4036        68.707931
```

```
4037        68.707931
```

```
4038        68.707931
```

```
[4039 rows x 1 columns]
```

```
path8 = '/content/drive/MyDrive/Colab Notebooks/twokemenydf.csv'
```

```
twokemenydf = pd.read_csv(path8)
```

```
print(twokemenydf)
```

```
    kemeny constant for network
```

```
0           2030.583502
```

```
1           406.232390
```

```
2           406.232390
```

```
3           406.232390
```

```
4           406.232390
```

```
...
...
```

```
4034        68.707931
```

```
4035        68.707931
```

```
4036        68.707931
```

```
4037        68.707931
```

```
4038        68.707931
```

```
[4039 rows x 1 columns]
```

```

twokcore_counts = []

# Iterate over nodes from 0 to (number of nodes - 1)
for node in G.nodes():
    # Create a subgraph containing only the current node and its neighbors
    length = nx.single_source_shortest_path_length(G, node, cutoff=2)
    acData = sorted(list(length.keys()))

    # Include the node itself in the subgraph
    subgraph = G.subgraph(acData)

    # Calculate local bridges for the subgraph
    k_core = nx.core_number(subgraph)
    B = sum(list(k_core.values()))

    twokcore_counts.append(B)

```

```

# Convert the list to a DataFrame
twohopkkcoredf = pd.DataFrame({'K-core for network': twokcore_counts})

```

```

# Display the DataFrame
print(twohopkkcoredf)

```

K-core for network	
0	40653
1	3400
2	3400
3	3400
4	3400
...	...
4034	255
4035	255
4036	255
4037	255
4038	255

```
[4039 rows x 1 columns]
```

```
twotriangle_counts = []
```

```

# Iterate over nodes from 0 to (number of nodes - 1)
for node in G.nodes():
    # Create a subgraph containing only the current node and its neighbors
    length = nx.single_source_shortest_path_length(G, node, cutoff=2)
    acData = sorted(list(length.keys()))

```

```

    # Include the node itself in the subgraph
    subgraph = G.subgraph(acData)

    # Calculate local bridges for the subgraph
    triangle = nx.triangles(subgraph, nodes=None)
    B = sum(list(triangle.values()))

```

```
twotriangle_counts.append(B)
```

```

# Convert the list to a DataFrame
twohopktriangledf = pd.DataFrame({'triangle for network': twotriangle_counts})

```

```

# Display the DataFrame
print(twohopktriangledf)

```

triangle for network	
0	1514841
1	39777
2	39777
3	39777
4	39777
...	...
4034	942
4035	942
4036	942
4037	942
4038	942

```
[4039 rows x 1 columns]
```

```
#getting all the necessary data from one hop for the Node Data
selected_columns = ['Node', 'NodeDegree', 'Average Degree Connectivity for Node', 'Core_Number', 'onion layers', 'eccentricity', 'No of triangles with Node']

# Creating a new DataFrame with the selected columns
commonhopdf = onefinaldf[selected_columns]

# Display the new DataFrame
print(commonhopdf)

   Node  NodeDegree  Average Degree Connectivity for Node  Core_Number \
0      0          347                           18.96           21
1      1          17                            48.24           13
2      2          10                            49.90            9
3      3          17                            59.76           13
4      4          10                            42.60            9
...     ...         ...
4034  4034          2                            38.50           2
4035  4035          1                            59.00           1
4036  4036          2                            31.50           2
4037  4037          4                            23.25           4
4038  4038          9                            15.22           5

   onion layers  eccentricity  No of triangles with Node
0            108             6                  2519
1             51              7                  57
2             30              7                  40
3             49              7                  86
4             30              7                  39
...           ...
4034            2              8                  1
4035            1              8                  0
4036            2              8                  1
4037            8              8                  4
4038           14              8                  20
```

[4039 rows x 7 columns]

```
twoonion_counts = []

# Iterate over nodes from 0 to (number of nodes - 1)
for node in G.nodes():
    # Create a subgraph containing only the current node and its neighbors
    length = nx.single_source_shortest_path_length(G, node, cutoff=2)
    acData = sorted(list(length.keys()))

    # Include the node itself in the subgraph
    subgraph = G.subgraph(acData)

    # Calculate local bridges for the subgraph
    k_core = nx.onion_layers(subgraph)
    sorted_keys = sorted(k_core.values())

    B = sum(list(sorted_keys))

    twoonion_counts.append(B)

# Convert the list to a DataFrame
twooniondf = pd.DataFrame({'onion layers for network': twoonion_counts})

# Display the DataFrame
print(twooniondf)

   onion layers for network
0                143440
1                 8567
2                 8567
3                 8567
4                 8567
...               ...
4034                447
4035                447
4036                447
4037                447
4038                447
```

[4039 rows x 1 columns]

```
#Adding the twohop data
twohopdatafd = pd.concat([commonhopdf, twohopcommndf, twohopdegreedf, twoadcdf, twooniondf, twohopbrdf, twobounddf, twoglobaldf, twokemenydf
print(twohopdatafd)
```

	Node	NodeDegree	Average Degree Connectivity for Node	Core_Number	\
0	0	347	18.96	21	
1	1	17	48.24	13	
2	2	10	49.90	9	
3	3	17	59.76	13	
4	4	10	42.60	9	
...	...	...	...	...	
4034	4034	2	38.50	2	
4035	4035	1	59.00	1	
4036	4036	2	31.50	2	
4037	4037	4	23.25	4	
4038	4038	9	15.22	5	
	onion layers	eccentricity	No of triangles with Node		\
0	108	6	2519		
1	51	7	57		
2	30	7	40		
3	49	7	86		
4	30	7	39		
...	...	...	...	...	
4034	2	8	1		
4035	1	8	0		
4036	2	8	1		
4037	8	8	4		
4038	14	8	20		
	Common Connections	0_count	1_count	...	\
0	33343.0	0	25	...	
1	2849.0	0	14	...	
2	2856.0	0	14	...	
3	2849.0	0	14	...	
4	2856.0	0	14	...	
...	...	...	...	...	
4034	204.0	0	7	...	
4035	203.0	0	6	...	
4036	201.0	0	7	...	
4037	196.0	0	7	...	
4038	NaN	0	7	...	
	Average Degree Connectivity for network	onion layers	for network		\
0	174898.08		143440		
1	28959.92		8567		
2	28959.92		8567		
3	28959.92		8567		
4	28959.92		8567		
...	...		...	...	
4034	1481.20		447		
4035	1481.20		447		
4036	1481.20		447		
4037	1481.20		447		
4038	1481.20		447		
	Bridge_Count	Boundary_Expansion	global_efficiency		\
0	26	1.146807	0.435346		
1	14	3.364943	0.523734		
2	14	3.364943	0.523734		
3	14	3.364943	0.523734		
4	14	3.364943	0.523734		

```
#finalizing two hop data by adding networkx library clustering coefficient data to check with the metrics
twofinaldf = pd.concat([twohopdatafd, cludf], axis = 1)
print(twofinaldf)
```

	Node	NodeDegree	Average Degree Connectivity for Node	Core_Number	\
0	0	347	18.96	21	
1	1	17	48.24	13	
2	2	10	49.90	9	
3	3	17	59.76	13	
4	4	10	42.60	9	
...	...	...	...	...	
4034	4034	2	38.50	2	
4035	4035	1	59.00	1	
4036	4036	2	31.50	2	
4037	4037	4	23.25	4	
4038	4038	9	15.22	5	
	onion layers	eccentricity	No of triangles with Node		\
0	108	6	2519		
1	51	7	57		

```

2          30      7          40
3          49      7          86
4          30      7          39
...
...          ...      ...
4034         2      8          1
4035         1      8          0
4036         2      8          1
4037         8      8          4
4038        14      8          20

    Common_Connections  0_count  1_count  ... onion_layers_for_network \
0            33343.0      0     25  ...                      143440
1            2849.0       0     14  ...                      8567
2            2856.0       0     14  ...                      8567
3            2849.0       0     14  ...                      8567
4            2856.0       0     14  ...                      8567
...
...          ...      ...
4034         204.0       0      7  ...                      447
4035         203.0       0      6  ...                      447
4036         201.0       0      7  ...                      447
4037         196.0       0      7  ...                      447
4038         NaN        0      7  ...                      447

    Bridge_Count  Boundary_Expansion  global_efficiency \
0              26           1.146807      0.435346
1              14           3.364943      0.523734
2              14           3.364943      0.523734
3              14           3.364943      0.523734
4              14           3.364943      0.523734
...
...          ...
4034         7           0.066667      0.557910
4035         7           0.066667      0.557910
4036         7           0.066667      0.557910
4037         7           0.066667      0.557910
4038         7           0.066667      0.557910

    max_weight_matching  random_probability  kemeny_constant_for_network \
0                  744           -66.852262      2030.583502
1                  164            75.626117      406.232390
2                  164            75.626117      406.232390
3                  164            75.626117      406.232390
4                  164            75.626117      406.232390

```

```
twofinaldf.replace([np.inf, -np.inf], 0, inplace=True)
```

```
twofinaldata = twofinaldf.fillna(0)
```

```
#determining the number of Steps for K means
import pandas as pd
from sklearn.cluster import KMeans
from sklearn.preprocessing import StandardScaler
import matplotlib.pyplot as plt

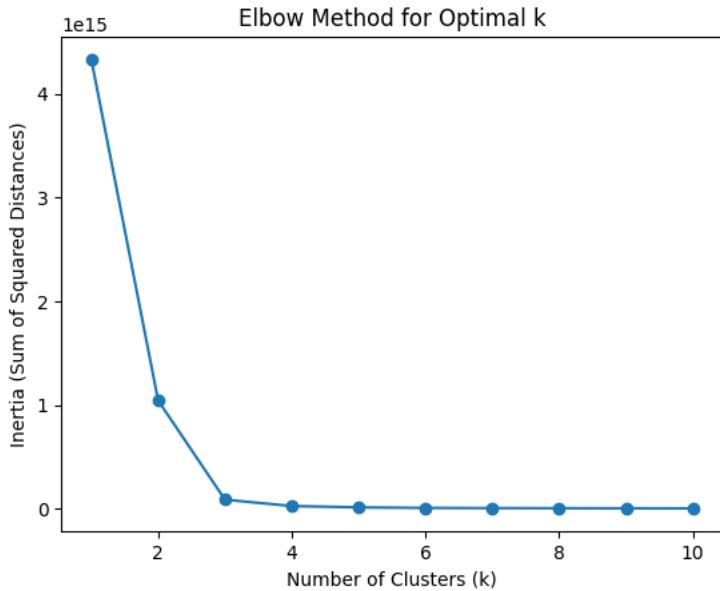
# Assuming df is your DataFrame with standardized features
# Step 1: Standardize the data
scaler = StandardScaler()
features_standardized = scaler.fit_transform(twofinaldata)

# Store the sum of squared distances for different values of k
inertia_values = []

# Define a range of k values to try
k_values = range(1, 11) # You can adjust the range based on your problem

# Apply k-means clustering for each k and store the inertia
for k in k_values:
    kmeans = KMeans(n_clusters=k, random_state=42)
    kmeans.fit(twofinaldata)
    inertia_values.append(kmeans.inertia_)

# Plot the elbow curve
plt.plot(k_values, inertia_values, marker='o')
plt.title('Elbow Method for Optimal k')
plt.xlabel('Number of Clusters (k)')
plt.ylabel('Inertia (Sum of Squared Distances)')
plt.show()
```



```
import pandas as pd
from sklearn.cluster import KMeans
from sklearn.preprocessing import StandardScaler
from sklearn.decomposition import PCA
import matplotlib.pyplot as plt

# Step 1: Standardize the data
scaler = StandardScaler()
features_standardized = scaler.fit_transform(twofinaldata)

# Step 3: Apply k-means clustering
k = 4 # Replace with your optimal k
kmeans = KMeans(n_clusters=k, random_state=42)
twofinaldata['k=4_Cluster'] = kmeans.fit_predict(features_standardized)

# Display the DataFrame with cluster assignments
print(twofinaldata)

# Step 4: Visualize the clusters (optional)
# This is just an example, adjust it based on your requirements
# Visualization is more challenging with a large number of features
# You may need to choose specific features for plotting or use advanced visualization techniques
plt.scatter(twofinaldata["NodeDegree"], twofinaldata['Common Connections'], c=twofinaldata['k=4_Cluster'], cmap='viridis')
plt.title('K-Means Clustering')
plt.show()
```

```
/usr/local/lib/python3.10/dist-packages/scikit-learn/_kmeans.py:870: FutureWarning: The default value of `n_init` will change from
```

```
warnings.warn(
    Node  NodeDegree  Average Degree Connectivity for Node  Core_Number \
0      0          347                      18.96        21
1      1          17                       48.24        13
2      2          10                       49.90         9
3      3          17                       59.76        13
4      4          10                       42.60         9
...
4034  4034          2                      38.50        2
4035  4035          1                      59.00        1
4036  4036          2                      31.50        2
4037  4037          4                      23.25        4
4038  4038          9                      15.22        5
```

```
onion layers  eccentricity  No of triangles with Node \
0           108          6                      2519
1           51           7                      57
2           30           7                      40
3           49           7                      86
4           30           7                      39
...
4034          2           8                      1
4035          1           8                      0
4036          2           8                      1
4037          8           8                      4
4038         14           8                     20
```

```
Common Connections  0_count  1_count  ...  Bridge_Count \
0            33343.0       0       25  ...        26
1            2849.0        0       14  ...        14
2            2856.0        0       14  ...        14
3            2849.0        0       14  ...        14
4            2856.0        0       14  ...        14
...
4034          204.0        0       7  ...        7
4035          203.0        0       6  ...        7
4036          201.0        0       7  ...        7
4037          196.0        0       7  ...        7
4038          0.0          0       7  ...        7
```

```
Boundary_Expansion  global_efficiency  max_weight_matching \
0             1.146807       0.435346        744
1             3.364943       0.523734        164
2             3.364943       0.523734        164
3             3.364943       0.523734        164
4             3.364943       0.523734        164
...
4034          0.066667       0.557910        26
4035          0.066667       0.557910        26
4036          0.066667       0.557910        26
4037          0.066667       0.557910        26
4038          0.066667       0.557910        26
```

```
random_probability  kemeny constant for network  K-core for network \
0            -66.852262     2030.583502      40653
1            75.626117      406.232390      3400
2            75.626117      406.232390      3400
3            75.626117      406.232390      3400
4            75.626117      406.232390      3400
...
4034          10.139742      68.707931      255
4035          10.139742      68.707931      255
4036          10.139742      68.707931      255
4037          10.139742      68.707931      255
4038          10.139742      68.707931      255
```

```
triangle for network  library Clustering  k=4_Cluster
0            1514841        0.04        2
1            39777         0.42        3
2            39777         0.89        3
3            39777         0.63        3
4            39777         0.87        3
...
4034          942          1.00        3
4035          942          0.00        3
4036          942          1.00        3
4037          942          0.67        3
4038          942          0.56        3
```

[4039 rows x 259 columns]

### K-Means Clustering

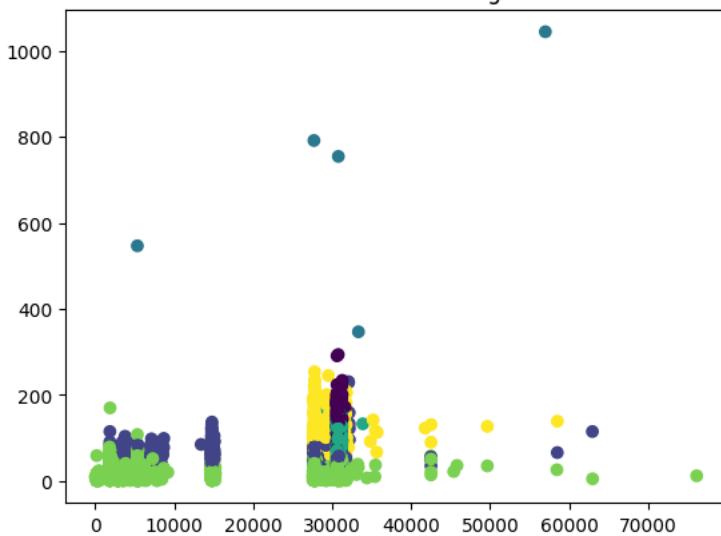


```
#Step 3: Apply k-means clustering
N = 6 # Replace with your optimal k
kmeans = KMeans(n_clusters=N, random_state=42)
twofinaldata['k = 6_Cluster'] = kmeans.fit_predict(features_standardized)

# Step 4: Visualize the clusters (optional)
# This is just an example, adjust it based on your requirements
# Visualization is more challenging with a large number of features
# You may need to choose specific features for plotting or use advanced visualization techniques
plt.scatter(twofinaldata["Common Connections"], twofinaldata['NodeDegree'], c=onefinaldf['k = 6_Cluster'], cmap='viridis')
plt.title('K-Means Clustering')
plt.show()
```

/usr/local/lib/python3.10/dist-packages/scikit-learn/cluster/\_kmeans.py:870: FutureWarning: The default value of `n\_init` will change from 10 to 100 in 0.23. To silence this warning, you can either set `n\_init` explicitly or set the environment variable `SKLEARN\_N\_INIT` to 100.

K-Means Clustering



```
# You may need to choose specific features for plotting or use advanced visualization techniques
plt.scatter(twofinaldata["NodeDegree"], twofinaldata['Common Connections'], c=twofinaldata['k = 6_Cluster'], cmap='viridis')
plt.title('K-Means Clustering')
plt.show()
```

### K-Means Clustering

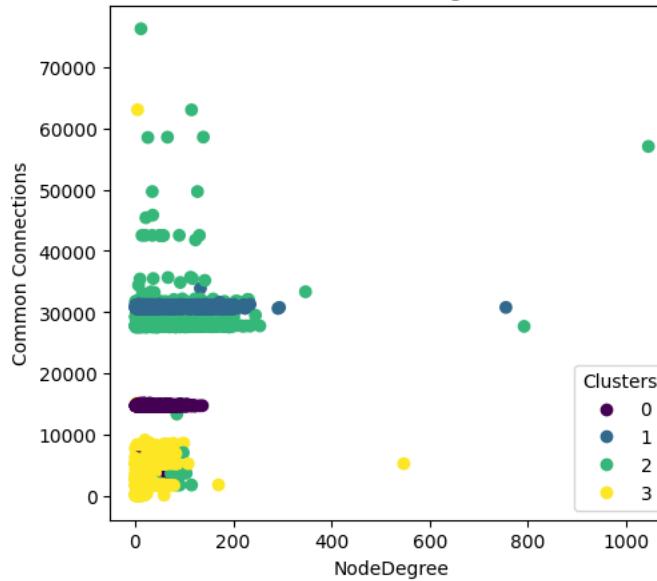
```
# Step 4: Visualize the clusters
fig, axes = plt.subplots(nrows=1, ncols=2, figsize=(12, 5))

# Plot for k=4 clusters
scatter_4 = axes[0].scatter(twofinaldata["NodeDegree"], twofinaldata['Common Connections'], c=twofinaldata['k=4_Cluster'], cmap='viridis')
axes[0].set_title(f'K-Means Clustering (k=4)')
axes[0].set_xlabel('NodeDegree')
axes[0].set_ylabel('Common Connections')
axes[0].legend(*scatter_4.legend_elements(), title='Clusters')

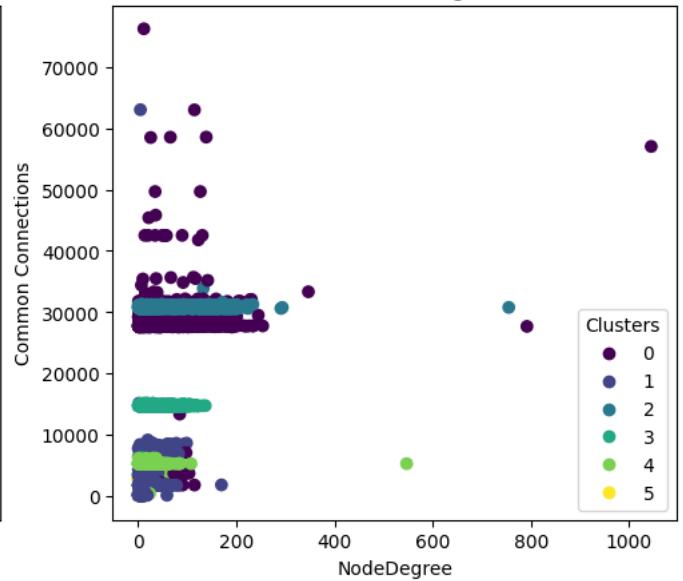
# Plot for k=6 clusters
scatter_6 = axes[1].scatter(twofinaldata["NodeDegree"], twofinaldata['Common Connections'], c=twofinaldata['k = 6_Cluster'], cmap='viridis')
axes[1].set_title(f'K-Means Clustering (k=6)')
axes[1].set_xlabel('NodeDegree')
axes[1].set_ylabel('Common Connections')
axes[1].legend(*scatter_6.legend_elements(), title='Clusters')
```

&lt;matplotlib.legend.Legend at 0x7a199aae6c20&gt;

### K-Means Clustering (k=4)



### K-Means Clustering (k=6)



# visualize with kemény constant - time and Onion layers of the network for clustering

# Step 4: Visualize the clusters

fig, axes = plt.subplots(nrows=1, ncols=2, figsize=(12, 5))

# Plot for k=4 clusters

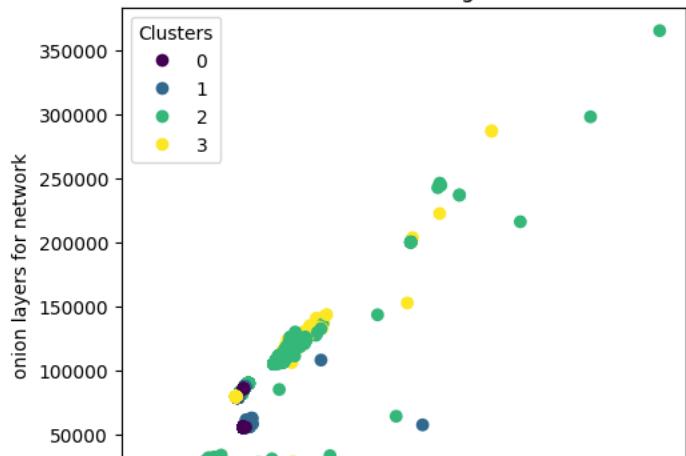
```
scatter_4 = axes[0].scatter(twofinaldata["kemeny constant for network"], twofinaldata['onion layers for network'], c=twofinaldata['k=4_Clust
axes[0].set_title(f'K-Means Clustering (k=4)')
axes[0].set_xlabel('kemeny constant for network')
axes[0].set_ylabel('onion layers for network')
axes[0].legend(*scatter_4.legend_elements(), title='Clusters')
```

# Plot for k=6 clusters

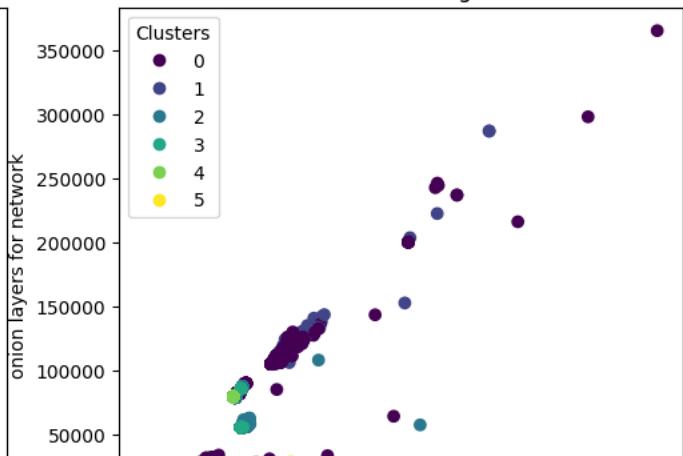
```
scatter_6 = axes[1].scatter(twofinaldata["kemeny constant for network"], twofinaldata['onion layers for network'], c=twofinaldata['k = 6_Cl
axes[1].set_title(f'K-Means Clustering (k=6)')
axes[1].set_xlabel('kemeny constant for network')
axes[1].set_ylabel('onion layers for network')
axes[1].legend(*scatter_6.legend_elements(), title='Clusters')
```

&lt;matplotlib.legend.Legend at 0x7a198d744a30&gt;

K-Means Clustering (k=4)



K-Means Clustering (k=6)



```
# Agglomerative Clustering
import pandas as pd
from sklearn.cluster import AgglomerativeClustering
from sklearn.preprocessing import StandardScaler
import matplotlib.pyplot as plt
from scipy.cluster.hierarchy import dendrogram, linkage
```

```
# Step 1: Standardize the data (optional but recommended for hierarchical clustering)
scaler = StandardScaler()
data_standardized = scaler.fit_transform(twofinaldata)

# Step 2: Determine the linkage matrix using the ward method
linkage_matrix = linkage(data_standardized, method='ward')

# Step 3: Plot the dendrogram (optional but useful for visualizing hierarchical clustering)
plt.figure(figsize=(12, 6))
dendrogram(linkage_matrix, leaf_rotation=90., leaf_font_size=8., truncate_mode='level', p=5)
plt.title('Hierarchical Clustering Dendrogram')
plt.show()

# Step 4: Apply Agglomerative Hierarchical Clustering
# Choose the appropriate number of clusters (n_clusters) based on the dendrogram
n_clusters = 4 # Replace with your desired number of clusters
hierarchical_clustering = AgglomerativeClustering(n_clusters=n_clusters, linkage='ward')
twofinaldata['agg4Cluster'] = hierarchical_clustering.fit_predict(data_standardized)
```

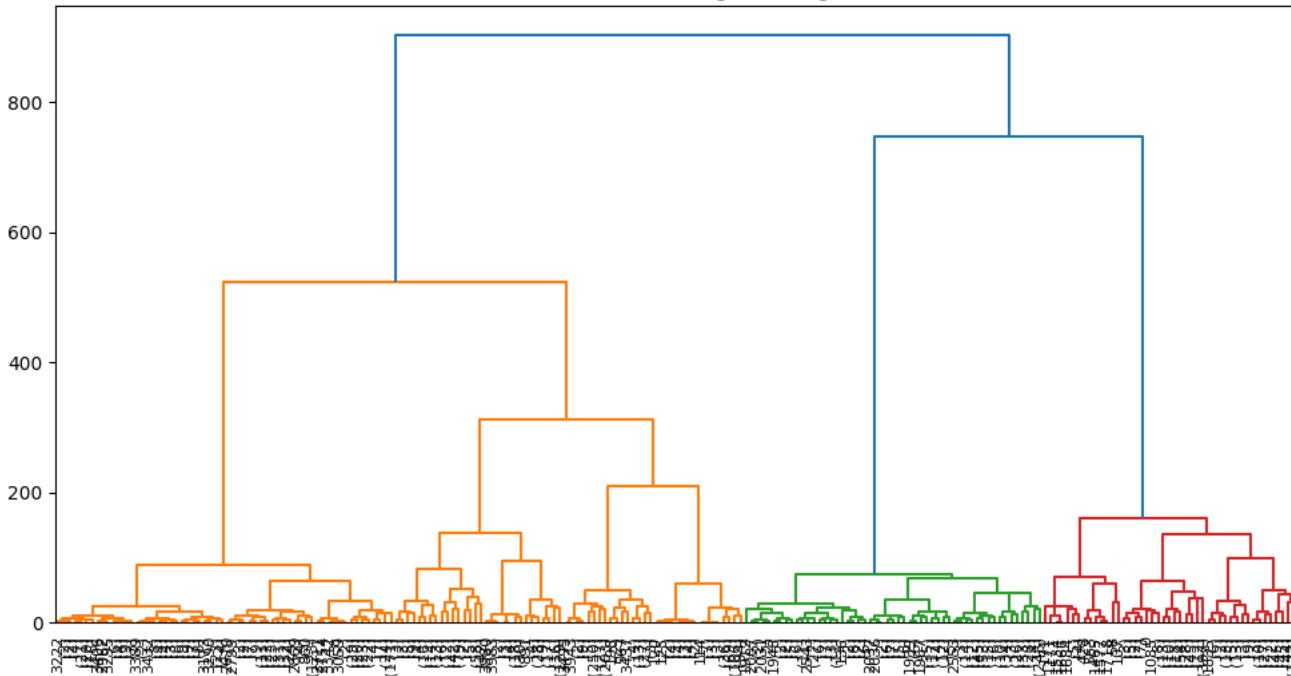
### Hierarchical Clustering Dendrogram

```
linkage_matrix = linkage(data_standardized, method='ward')

# Step 3: Plot the dendrogram (optional but useful for visualizing hierarchical clustering)
plt.figure(figsize=(12, 6))
dendrogram(linkage_matrix, leaf_rotation=90., leaf_font_size=8., truncate_mode='level', p=7)
plt.title('Hierarchical Clustering Dendrogram')
plt.show()

# Step 4: Apply Agglomerative Hierarchical Clustering
# Choose the appropriate number of clusters (n_clusters) based on the dendrogram
n_clusters = 6 # Replace with your desired number of clusters
hierarchical_clustering = AgglomerativeClustering(n_clusters=n_clusters, linkage='ward')
twofinaldata['agg6Cluster'] = hierarchical_clustering.fit_predict(data_standardized)
```

### Hierarchical Clustering Dendrogram



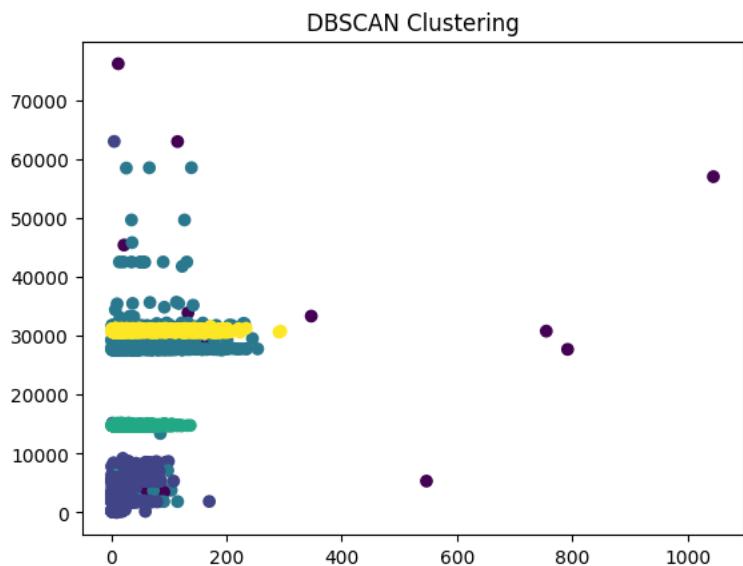
```
#DBSCAN
#dbscan algorithm
import pandas as pd
from sklearn.cluster import DBSCAN
from sklearn.preprocessing import StandardScaler
import matplotlib.pyplot as plt

# Assuming df is your DataFrame

# Step 1: Standardize the data (required for DBSCAN)
scaler = StandardScaler()
data_standardized = scaler.fit_transform(twofinaldata)

# Step 2: Apply DBSCAN
# Choose appropriate values for `eps` (maximum distance between two samples for one to be considered as in the neighborhood of the other)
# and `min_samples` (the number of samples in a neighborhood for a point to be considered as a core point)
dbscan = DBSCAN(eps=11, min_samples=4)
twofinaldata['dbs4Cluster'] = dbscan.fit_predict(data_standardized)

# Step 3: Visualize the clusters (optional)
# This is just an example, adjust it based on your requirements
plt.scatter(twofinaldata['NodeDegree'], twofinaldata['Common Connections'], c=twofinaldata['dbs4Cluster'], cmap='viridis')
plt.title('DBSCAN Clustering')
plt.show()
```



```
# Step 2: Apply DBSCAN
# Choose appropriate values for `eps` (maximum distance between two samples for one to be considered as in the neighborhood of the other)
# and `min_samples` (the number of samples in a neighborhood for a point to be considered as a core point)
dbscan = DBSCAN(eps=11, min_samples=6)
twofinaldata['dbs6Cluster'] = dbscan.fit_predict(data_standardized)
```

```
# Visualize clusters for k-means, agglomerative clustering, and DBSCAN in a 2x3 grid
fig, axes = plt.subplots(nrows=2, ncols=3, figsize=(18, 10))

# Plot for k=4 clusters in k-means
scatter_4_kmeans = axes[0, 0].scatter(twofinaldata["triangle for network"], twofinaldata['onion layers for network'], c=twofinaldata['k=4_Clusters'])
axes[0, 0].set_title('K-means Clustering (k=4)')
axes[0, 0].set_xlabel('triangle for network')
axes[0, 0].set_ylabel('onion layers for network')
axes[0, 0].legend(*scatter_4_kmeans.legend_elements(), title='Clusters')

# Plot for k=4 clusters in agglomerative clustering
scatter_4_agg = axes[0, 1].scatter(twofinaldata["triangle for network"], twofinaldata['onion layers for network'], c=twofinaldata['agg4Clusters'])
axes[0, 1].set_title('Agglomerative Clustering (k=4)')
axes[0, 1].set_xlabel('triangle for network')
axes[0, 1].set_ylabel('onion layers for network')
axes[0, 1].legend(*scatter_4_agg.legend_elements(), title='Clusters')

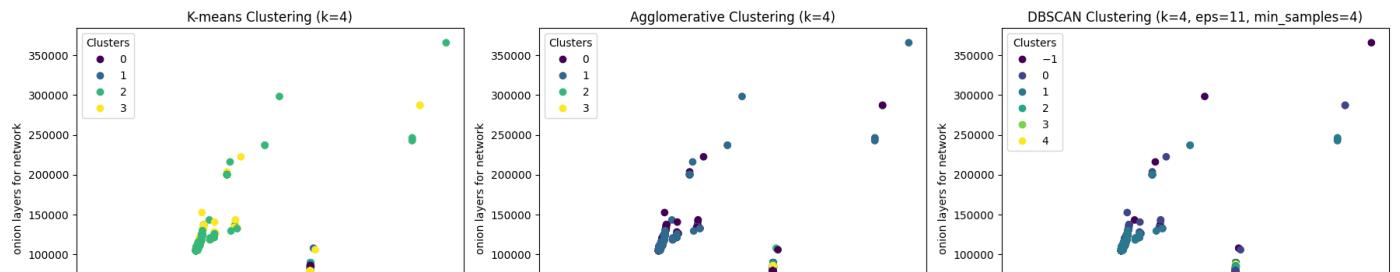
# Plot for k=6 clusters in k-means
scatter_6_kmeans = axes[1, 0].scatter(twofinaldata["triangle for network"], twofinaldata['onion layers for network'], c=twofinaldata['k = 6_Clusters'])
axes[1, 0].set_title('K-means Clustering (k=6)')
axes[1, 0].set_xlabel('triangle for network')
axes[1, 0].set_ylabel('onion layers for network')
axes[1, 0].legend(*scatter_6_kmeans.legend_elements(), title='Clusters')

# Plot for k=6 clusters in agglomerative clustering
scatter_6_agg = axes[1, 1].scatter(twofinaldata["triangle for network"], twofinaldata['onion layers for network'], c=twofinaldata['agg6Clusters'])
axes[1, 1].set_title('Agglomerative Clustering (k=6)')
axes[1, 1].set_xlabel('triangle for network')
axes[1, 1].set_ylabel('onion layers for network')
axes[1, 1].legend(*scatter_6_agg.legend_elements(), title='Clusters')

# Plot for DBSCAN clusters (k=4, eps=11, min_samples=5)
scatter_dbSCAN_4 = axes[0, 2].scatter(twofinaldata["triangle for network"], twofinaldata['onion layers for network'], c=twofinaldata['dbs4Clusters'])
axes[0, 2].set_title('DBSCAN Clustering (k=4, eps=11, min_samples=4)')
axes[0, 2].set_xlabel('triangle for network')
axes[0, 2].set_ylabel('onion layers for network')
axes[0, 2].legend(*scatter_dbSCAN_4.legend_elements(), title='Clusters')

# Plot for DBSCAN clusters (k=6, eps=0.8, min_samples=5)
scatter_dbSCAN_6 = axes[1, 2].scatter(twofinaldata["triangle for network"], twofinaldata['onion layers for network'], c=twofinaldata['dbs6Clusters'])
axes[1, 2].set_title('DBSCAN Clustering (k=6, eps=11, min_samples=6)')
axes[1, 2].set_xlabel('triangle for network')
axes[1, 2].set_ylabel('onion layers for network')
axes[1, 2].legend(*scatter_dbSCAN_6.legend_elements(), title='Clusters')

plt.tight_layout()
plt.show()
```



```
# Step 4: Visualize the clusters for k-means and agglomerative clustering
fig, axes = plt.subplots(nrows=2, ncols=3, figsize=(18, 10))
```

```
# Plot for k=4 clusters in k-means
scatter_4_kmeans = axes[0, 0].scatter(twofinaldata["NodeDegree"], twofinaldata['Common Connections'], c=twofinaldata['k=4_Cluster'], cmap='viridis')
axes[0, 0].set_title('K-means Clustering (k=4)')
axes[0, 0].set_xlabel('NodeDegree')
axes[0, 0].set_ylabel('Common Connections')
axes[0, 0].legend(*scatter_4_kmeans.legend_elements(), title='Clusters')

# Plot for k=4 clusters in agglomerative clustering
scatter_4_agg = axes[0, 1].scatter(twofinaldata["NodeDegree"], twofinaldata['Common Connections'], c=twofinaldata['agg4Cluster'], cmap='viridis')
axes[0, 1].set_title('Agglomerative Clustering (k=4)')
axes[0, 1].set_xlabel('NodeDegree')
axes[0, 1].set_ylabel('Common Connections')
axes[0, 1].legend(*scatter_4_agg.legend_elements(), title='Clusters')

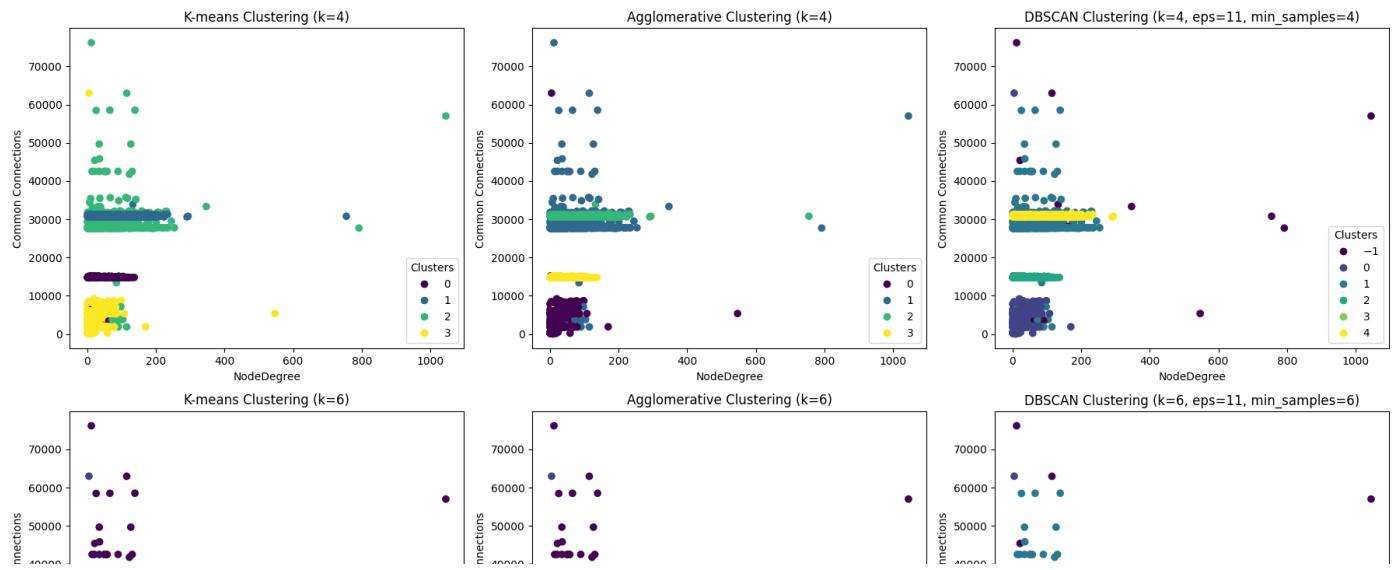
# Plot for DBSCAN clusters (k=4, eps=11, min_samples=5)
scatter_dbSCAN_4 = axes[0, 2].scatter(twofinaldata["NodeDegree"], twofinaldata['Common Connections'], c=twofinaldata['dbs4Cluster'], cmap='viridis')
axes[0, 2].set_title('DBSCAN Clustering (k=4, eps=11, min_samples=4)')
axes[0, 2].set_xlabel('NodeDegree')
axes[0, 2].set_ylabel('Common Connections')
axes[0, 2].legend(*scatter_dbSCAN_4.legend_elements(), title='Clusters')

# Plot for k=6 clusters in k-means
scatter_6_kmeans = axes[1, 0].scatter(twofinaldata["NodeDegree"], twofinaldata['Common Connections'], c=twofinaldata['k = 6_Cluster'], cmap='viridis')
axes[1, 0].set_title('K-means Clustering (k=6)')
axes[1, 0].set_xlabel('NodeDegree')
axes[1, 0].set_ylabel('Common Connections')
axes[1, 0].legend(*scatter_6_kmeans.legend_elements(), title='Clusters')

# Plot for k=6 clusters in agglomerative clustering
scatter_6_agg = axes[1, 1].scatter(twofinaldata["NodeDegree"], twofinaldata['Common Connections'], c=twofinaldata['agg6Cluster'], cmap='viridis')
axes[1, 1].set_title('Agglomerative Clustering (k=6)')
axes[1, 1].set_xlabel('NodeDegree')
axes[1, 1].set_ylabel('Common Connections')
axes[1, 1].legend(*scatter_6_agg.legend_elements(), title='Clusters')

# Plot for DBSCAN clusters (k=6, eps=0.8, min_samples=5)
scatter_dbSCAN_6 = axes[1, 2].scatter(twofinaldata["NodeDegree"], twofinaldata['Common Connections'], c=twofinaldata['dbs6Cluster'], cmap='viridis')
axes[1, 2].set_title('DBSCAN Clustering (k=6, eps=11, min_samples=6)')
axes[1, 2].set_xlabel('NodeDegree')
axes[1, 2].set_ylabel('Common Connections')
axes[1, 2].legend(*scatter_dbSCAN_6.legend_elements(), title='Clusters')

plt.tight_layout()
plt.show()
```



```
#generate graphs
# Create a 2x4 grid for subplots
fig, axes = plt.subplots(2, 4, figsize=(9, 5))

# Plot for k=4
plot_clustersname(G, twofinaldata['k=4_Cluster'], 4, 'kmeans', axes[0, 0])

# Plot for k=4 - agg
plot_clustersname(G, twofinaldata['agg4Cluster'], 4, 'aggclustering', axes[0, 1])

# Plot for k=4 - dbSCAN
plot_clustersname(G, twofinaldata['dbs4Cluster'], 4, 'dbSCAN4sample', axes[0, 2])

#plot for networkx - library
plot_clustersname(G, twofinaldata['library Clustering'], 6, 'networkx', axes[0,3])

# Plot for k=6
plot_clustersname(G, twofinaldata['k = 6_Cluster'], 6, 'kmeans', axes[1, 0])

# Plot for k=6 - agg
plot_clustersname(G, twofinaldata['agg6Cluster'], 6, 'aggclustering', axes[1, 1])

# Plot for k=6 - dbSCAN
plot_clustersname(G, twofinaldata['dbs6Cluster'], 4, 'dbSCAN6sample', axes[1, 2])

#plot for networkx - library
plot_clustersname(G, twofinaldata['library Clustering'], 6, 'networkx', axes[1,3])

# Adjust layout
plt.tight_layout()
plt.show()
```

```
<ipython-input-117-83f07bba3dce>:8: MatplotlibDeprecationWarning: The get_cmap function was deprecated in Matplotlib 3.7 and will be removed in 3.8.
```



Metrics



```
import matplotlib.pyplot as plt
import numpy as np
from sklearn.metrics import silhouette_score, calinski_harabasz_score

# Extract features
X = twofinaldata[['triangle for network', 'onion layers for network']]

# Extract cluster labels
kmeans_labels_4 = twofinaldata['k=4_Cluster']
agg_labels_4 = twofinaldata['agg4Cluster']
dbSCAN_labels_4 = twofinaldata['dbs4Cluster']

kmeans_labels_6 = twofinaldata['k = 6_Cluster']
agg_labels_6 = twofinaldata['agg6Cluster']
dbSCAN_labels_6 = twofinaldata['dbs6Cluster']

# Calculate Silhouette Score
kmeans_silhouette_4 = silhouette_score(X, kmeans_labels_4)
agg_silhouette_4 = silhouette_score(X, agg_labels_4)
dbSCAN_silhouette_4 = silhouette_score(X, dbSCAN_labels_4)

kmeans_silhouette_6 = silhouette_score(X, kmeans_labels_6)
agg_silhouette_6 = silhouette_score(X, agg_labels_6)
dbSCAN_silhouette_6 = silhouette_score(X, dbSCAN_labels_6)

# Calculate Calinski-Harabasz Index
kmeans_calinski_harabasz_4 = calinski_harabasz_score(X, kmeans_labels_4)
agg_calinski_harabasz_4 = calinski_harabasz_score(X, agg_labels_4)
dbSCAN_calinski_harabasz_4 = calinski_harabasz_score(X, dbSCAN_labels_4)

kmeans_calinski_harabasz_6 = calinski_harabasz_score(X, kmeans_labels_6)
agg_calinski_harabasz_6 = calinski_harabasz_score(X, agg_labels_6)
dbSCAN_calinski_harabasz_6 = calinski_harabasz_score(X, dbSCAN_labels_6)

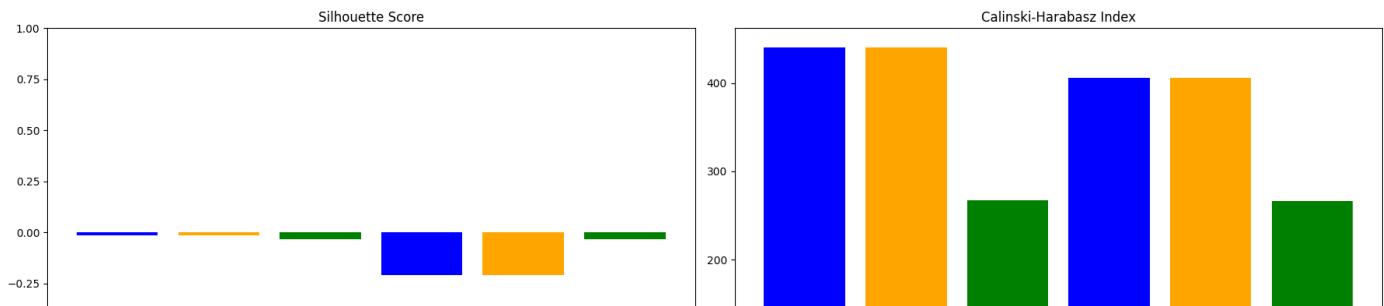
# Data for plotting
algorithms = ['K-means (k=4)', 'Agglomerative (k=4)', 'DBSCAN (k=4)', 'K-means (k=6)', 'Agglomerative (k=6)', 'DBSCAN (k=6)']
silhouette_scores = [kmeans_silhouette_4, agg_silhouette_4, dbSCAN_silhouette_4, kmeans_silhouette_6, agg_silhouette_6, dbSCAN_silhouette_6]
calinski_harabasz_scores = [kmeans_calinski_harabasz_4, agg_calinski_harabasz_4, dbSCAN_calinski_harabasz_4, kmeans_calinski_harabasz_6, agg_calinski_harabasz_6, dbSCAN_calinski_harabasz_6]

# Plot Silhouette Score and Calinski-Harabasz Index
fig, axes = plt.subplots(nrows=1, ncols=2, figsize=(18, 6))

# Plot Silhouette Score
axes[0].bar(algorithms, silhouette_scores, color=['blue', 'orange', 'green', 'blue', 'orange', 'green'])
axes[0].set_title('Silhouette Score')
axes[0].set_ylim(-1, 1)

# Plot Calinski-Harabasz Index
axes[1].bar(algorithms, calinski_harabasz_scores, color=['blue', 'orange', 'green', 'blue', 'orange', 'green'])
axes[1].set_title('Calinski-Harabasz Index')

plt.tight_layout()
plt.show()
```



as calinski score is high for no of triangles in network to onion layers, i am trying to check the score for no of common connections and kemeny

-----