Assessment Questions on 'Practicing Graph Analytics in Neo4j With Cypher'

9 questions

1.

[NOTE: The following questions apply to the results from the "Practicing Graph Analytics in Neo4j With Cypher" Assignment using the dataset titled 'gene_gene_associations_50k.csv'. They are based on the assumption you successfully completed the "Practicing Graph Analytics in Neo4j With Cypher" assignment using the dataset included in the download linked below.]

gene_gene_associations_50k.zip (http...

What is the number of nodes returned?

50,000

9656

O 9756

O 8673

2.

What's the number of edges?

50,000

0	49,834
0	46,621
0	None of the above
3.	
The nu	umber of loops in the graph is:
0	1035
0	1395
0	1221
0	1243
4. The qu gives u	the count of all non loop edges between every adjacent node
0	pair. the count of all edges between every adjacent node pair.
0	the count of all edges.
0	None of the above
=	uery match (n)-[r]->(m) where m <> n return distinct n, m, count(r) as unt order by myCount desc limit 1 produces what?
O	a random edge

0	two neighboring nodes, each with a high outdegree
0	the pair of nodes with the maximum number of multi-edges between them
_	uery match p=(n {Name:'BRCA1'})-[:AssociationType*2]->(m) return duces what?
0	The neighbors' neighbors of the node whose name is 'BRCA1'
0	The 2-neighborhood of the node whose name is 'BRCA1'
0	The neighbors whose distance is greater than 1 and less than 2 of the node whose name is 'BRCA1'
0	The neighbors of the node whose name is 'BRCA1'
	nany <u>non-directed</u> shortest paths are there between the node d 'BRCA1' and the node named 'NBR1'?
0	9
0	10
0	None of the above
8. The to	p 2 nodes with the highest outdegree are:
0	GRB2 and TP53
0	EP300 and BRCA1
0	MEPCE and EGFR

