Introduction



Please find two other learning partners,

- form a standing group and
- ▶ tell them what you already know about
 - graphs,
 - graph databases and
 - ▶ Neo4j.



Graph Data - Modelling and Quering with Neo4j and Cypher

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Agenda

What are graphs?

Definition

Typical use cases

Not so typical use cases

Starting with Neo4j and Cypher

Starting the database

Brief look at the configuration

CRUD operations with Cypher

Node operations

Relation operations

Quering for paths and patterns

Using graph algorithms

apoc library

algo library

Importing data

Basic loading command

Performance issues and dealing with big data



Graph



Definition

Graph is an ordered pair G = (V, E) comprising a set V of vertices, nodes or points together with a set E of edges, arcs or lines, which are 2-element subsets of V.

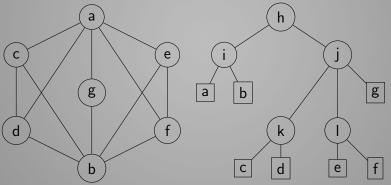
¹en.wikipedia.org/wiki/Graph_(discrete_mathematics)

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- Networks
 - Social networks



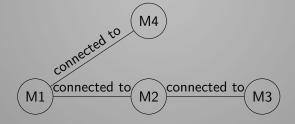


Networks

Social networks

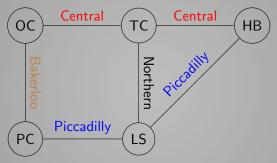


Computer networks





- Networks
 - Transport networks



OC = Oxford Circus

TC = Tottenham Court Road

HB = Holborn

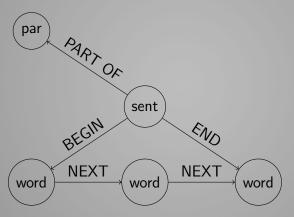
LS = Leicester Square

PC = Piccadilly Circus



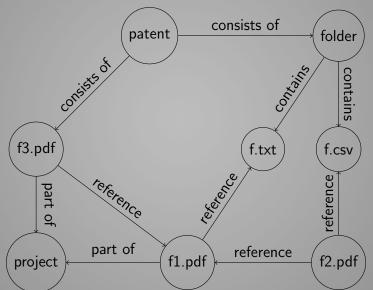


Natural Language Processing



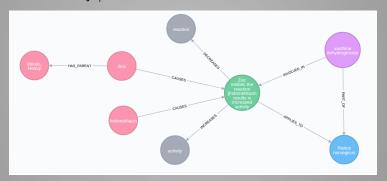


Document management





▶ Biochemistry / Genomics





Find the right installation file for your OS at neo4j-training-files/neo4j on the flash drive and install the software.



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- Copy the neo4j-training-files/data/odsc.db folder into your NEO4J_HOME/data/databases/ directory

Starting Neo4j



Start the database with
NEO4J_HOME/bin/neo4j start

Starting Neo4j



- Start the database with
 NEO4J_HOME/bin/neo4j start
- ► Go to http://localhost:7474 within you browser

Demonstration



Important configuration entries dbms.active_database=odsc.db dbms.security.auth_enabled=false dbms.security.procedures.unrestricted=algo.*,apoc.* apoc.import.file.enabled=true



- create node CREATE (c:Chemical {name: 'Helium'}) RETURN c
- update node
 MERGE (c:Chemical {name: 'Helium'}) SET c.symbol =
 'He' RETURN c
- delete node
 - without relations
 MATCH (c:Chemical {name:'Helium'}) DELETE c
 MATCH (c:Chemical)
 WHERE c.name = 'Helium'
 DELETE c
 - with existing relations
 MATCH (c:Chemical {name:'Helium'})
 DETACH DELETE c



- create relation
 - between new nodes

 CREATE (c:Chemical chemicalName:'Helium')[:BELONGS_TO]-¿(g:ChemicalGroup groupName:'Noble gases') RETURN c,g
 - between existing nodes
 MATCH (g:ChemicalGroup groupName:'Noble gases'),
 (p:ChemicalGroup groupName:'Gases') CREATE
 (g)-[:HAS_PARENT]-¿(p) RETURN g,p
- update relation
 MATCH ()-[r:BELONGS_TO]-() SET r.updateTime =
 timestamp() RETURN r
- delete relation MATCH ()-[r:BELONGS_TO]-() DELETE r

Examples:

► MATCH (g:Gene) WHERE g.geneSymbol = 'CTSD' RETURN g

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- MATCH (g:Gene)_i-[:ASSOCIATED_WITH]-(d:Disease) WHERE g.geneSymbol = 'CTSD' RETURN g, d
- MATCH (g:Gene)_i-[:ASSOCIATED_WITH]-(d:Disease)WHERE g.geneSymbol = 'CTSD' RETURN g, count(d)
- ► MATCH (g:Gene)_i-[:ASSOCIATED_WITH]-(d:Disease) WITH g, count(d) as diseases WHERE diseases ¿ 50 RETURN g.geneName, g.geneSymbol, diseases ORDER BY diseases DESC
- MATCH (g:Gene)_i-[:ASSOCIATED_WITH]-(d:Disease)-[:ASSOCIATED_WITH]-(otherGene:Gene) WHERE g.geneSymbol = 'CTSD' AND d.diseaseName = 'Osteoarthritis' RETURN otherGene.geneName, otherGene.geneSymbol

-) where
- MATCH p = (c:Chemical)-[*2]-(d:Disease) where d.diseaseName STARTS WITH 'Osteo' RETURN p LIMIT 20
- MATCH (c:Chemical)_i-[:HAS_PARENT*3..4]-(descendant:Chemical) WITH c, count(descendant) AS descendants, collect(descendant.chemicalName) as names ORDER BY descendants DESC LIMIT 10 RETURN c.chemicalName, names[1..10], descendants
- MATCH (c:Chemical) WHERE c.chemicalName = 'Zinc Acetate' MATCH (d:Disease) WHERE d.diseaseName = 'Alzheimer Disease' MATCH p = (c)-[*1..3]-(d) RETURN p LIMIT 20
- MATCH (c:Chemical chemicalName:'Zinc Acetate'),
 (d:Disease diseaseName:'Alzheimer Disease') MATCH
 path = allShortestPaths((c)-[*..3]-(d)) RETURN path

Calling procedures



- ▶ CALL db.schema
- CALL dbms.procedures
- call dbms.functions
- CALL apoc.help('dijkstra')



Definition

In a connected graph, the normalized closeness centrality (or closeness) of a node is the average length of the shortest path between the node and all other nodes in the graph. Thus the more central a node is, the closer it is to all other nodes.²

Closeness Centrality Example

MATCH (node:Chemical) WHERE node.chemicalName CONTAINS 'Vitamin' WITH collect(node) AS nodes CALL apoc.algo.closeness(['HAS_PARENT'],nodes,'BOTH') YIELD node, score RETURN node, score ORDER BY score DESC

²en.wikipedia.org/wiki/Centrality#Closeness_centrality



Definition

Betweenness centrality quantifies the number of times a node acts as a bridge along the shortest path between two other nodes 3

MATCH (node:Disease) WHERE node.diseaseName CONTAINS 'deficiency' WITH collect(node) AS nodes CALL apoc.algo.betweenness(['HAS_PARENT'],nodes,'BOTH') YIELD node, score RETURN node, diseaseName, score ORDER BY score DESC LIMIT 10

³en.wikipedia.org/wiki/Centrality#Betweenness_centrality



PageRank example

call algo.pageRank.stream('InteractionType', 'HAS_PARENT',iterations:20) YIELD node, score WITH * ORDER BY score DESC LIMIT 5 RETURN node.typeName, node.code, score;



MATCH (n:Movie) CALL apoc.create.addLabels(id(n), [n.genre]) YIELD node REMOVE node.genre RETURN node CALL apoc.periodic.iterate("MATCH (p:Person) WHERE (p)-[:ACTED_IN]- $\dot{\epsilon}$ () RETURN p", "SET p:Actor", batchSize:10000, parallel:true) call apoc.refactor.setType(rel, 'NEW-TYPE')