Report

Graph And Al CA-1

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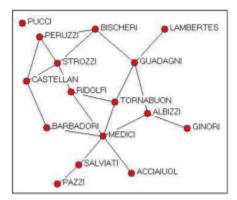
Excercise 1 Creating Nodes for Family Pseudo code Creating Relationship between families Pseudo code **Centrality Calculations Degree Centrality** Pseudo Code **Closeness Centrality** Pseudo Code **Betweenness Centrality** Pseudo Code **Eigenvector Centrality** Pseudo Code **PageRank** Pseudo Code **Final Calculations** Excercise 3 Personal Findings on Applying CRISP-DM Methodology using Neo4J **Business Understanding Data Understanding Data Preparation Data Cleaning Create Nodes Creating Relations Modeling Evaluation** Deployment **Note**

11010

References

Excercise 1

- Use Padgett's data on Florentine Families Marriage alliances
- Make Nodes Representing each Family
- Edge Denoting Connection in Marriage
- Load it in Neo4J Knowledge Graph
- Calculate Degree Centrality; Closeness Centrality; Betweenness Centrality; Eigenvector Centrality; PageRank



Creating Nodes for Family

Each Node Represents a Florentine Family, we create the node for each family one by one using the CREATE query of neo4j.

Pseudo code

// Creating the Family Nodes, the nodes will represent the family names, upon which relationships are built.

// Node PUCCI

CREATE (n:<NodeName> {name:'<NodeAttribute - eg.PUCCI >'});

```
// Node PUCCI
CREATE (n:FAMILY {name:'PUCCI'});
// Node PERUZZI
CREATE (n:FAMILY {name:'PERUZZI'});
// Node STROZZI
CREATE (n:FAMILY {name:'STROZZI'});
// Node BISCHERI
CREATE (n:FAMILY {name: 'BISCHERI'});
// Node LAMBERTES
CREATE (n:FAMILY {name:'LAMBERTES'});
// Node CASTELLAN
CREATE (n:FAMILY {name: 'CASTELLAN'});
//-Node-RIDOLF
CREATE (n:FAMILY {name:'RIDOLF'});
//-Node-TORNABUON
CREATE (n:FAMILY {name: 'TORNABUON'});
// Node GUADAGNI
CREATE (n:FAMILY {name: 'GUADAGNI'});
// Node ALBIZZI
CREATE (n:FAMILY {name: 'ALBIZZI'});
//-Node-BARBADORI
CREATE (n:FAMILY {name:'BARBADORI'});
// Node MEDICI
CREATE (n:FAMILY {name:'MEDICI'});
// Node GINORI
CREATE (n:FAMILY {name:'GINORI'});
// Node SALVIATI
CREATE (n:FAMILY {name:'SALVIATI'});
// Node ACCIAIUOL
CREATE (n:FAMILY {name:'ACCIAIUOL'});
// Node PAZZI
CREATE (n:FAMILY {name: 'PAZZI'});
```

Fig 1:- Nodes Created in Neo4J

Creating Relationship between families

After our Nodes are created we have a starting point of creating relations, each node represents a Family so the relations they are connected to and we can use to represent is `MARRIAGE`.

Pseudo code

// MEDICI - SALVIATI

MATCH (n1:FAMILY), (n2:FAMILY)

WHERE n1.name = 'MEDICI' AND n2.name = 'SALVIATI'

CREATE (n1)-[mrg:MARRAGE]->(n2);

On the above code **CREATE (n1)-[mrg:MARRAGE]->(n2)** is where we are creating the relationship, **n1** and **n2** are variables used to create this relationship which is declared **MATCH (n1:FAMILY)**, **(n2:FAMILY)** here.

To Create the Reverse Relationship

// SALVIATI - MEDICI

MATCH (n1:FAMILY), (n2:FAMILY)

WHERE n1.name = 'MEDICI' AND n2.name = 'SALVIATI'

CREATE (n2)-[mrg:MARRAGE]->(n1);

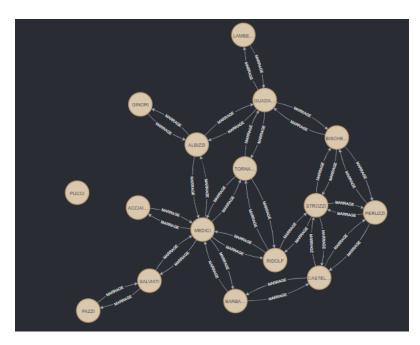


Fig 2: - Graph Visualised after Nodes added and Relationships Added.

Centrality Calculations

Calculations for Degree Centrality; Closeness Centrality; Betweenness Centrality; Eigenvector Centrality; PageRank.

Degree Centrality

It represents the number of edges the node has.

Pseudo Code

CALL gds.degree.write('MEDICIGraph', { writeProperty: 'degree' })

YIELD centralityDistribution, nodePropertiesWritten

RETURN centralityDistribution.min AS minimumScore, centralityDistribution.mean AS meanScore, nodePropertiesWritten;

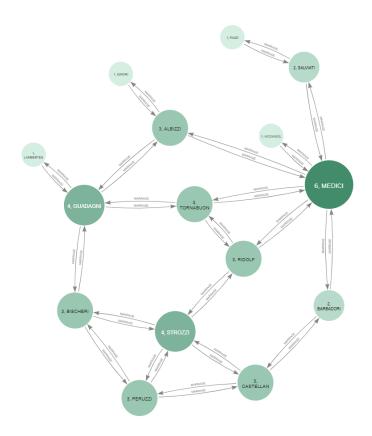


Fig 3: - Degree Centrality Visual Representation by Size and Color - Using Neo4j Bloom

Closeness Centrality

Closeness Centrality gives a numerical output representing how close is a particular node from other nodes.

Pseudo Code

//Using GDS to calculate closeness

CALL gds.alpha.closeness.write({

nodeProjection: 'FAMILY',

relationshipProjection: 'MARRAGE',

writeProperty: 'centrality'

)) YIELD nodes, writeProperty;

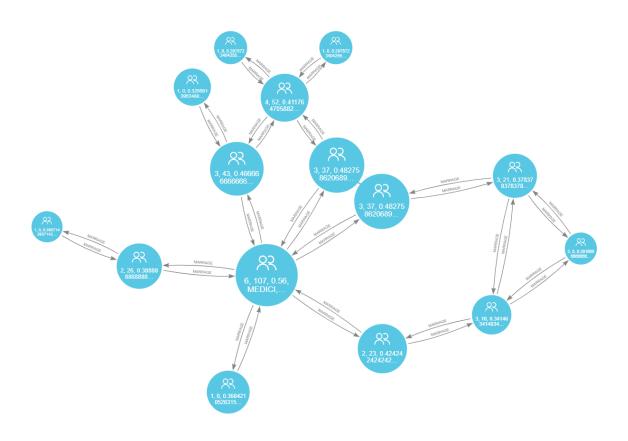


Fig 4: - Closeness Centrality Visual Representation by Size and Color - Using Neo4j Bloom

Betweenness Centrality

Betweenness Represents the shortest path in numerical terms

Pseudo Code

CALL gds.betweenness.write('MEDICIGraph', { writeProperty: 'betweenness' })

YIELD centralityDistribution, nodePropertiesWritten

RETURN centralityDistribution.min AS minimumScore, centralityDistribution.mean AS meanScore, nodePropertiesWritten;

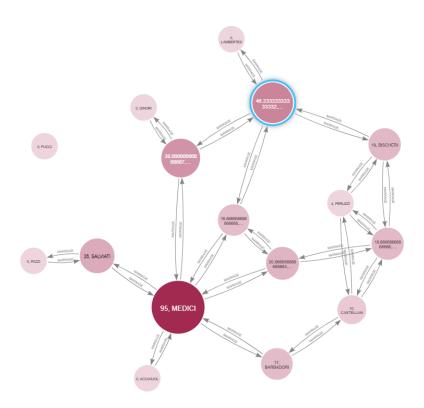


Fig 5 : - Betweenness Centrality Visual Representation by Size and Color - Using Neo4j Bloom

Eigenvector Centrality

Eigenvector Centrality represents Influence of the node in numerical terms

Pseudo Code

```
CALL gds.eigenvector.write('MEDICIGraph', {
   maxIterations: 20,
   writeProperty: 'Eigcentrality'
})
```

YIELD nodePropertiesWritten, ranlterations;

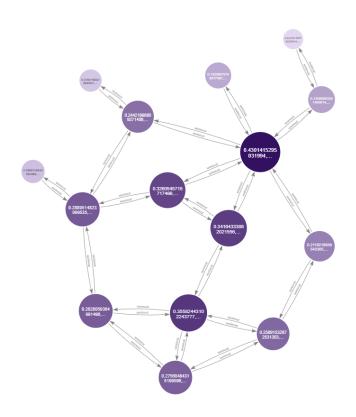


Fig 5 : - Eigenvector Centrality Visual Representation by Size and Color - Using Neo4j Bloom

PageRank

Pagerank sets Quality of node based on importance of the node

Pseudo Code

```
CALL gds.pageRank.write('MEDICIGraph', {
  maxIterations: 20,
  dampingFactor: 0.85,
  writeProperty: 'pagerank'
})
```

YIELD nodePropertiesWritten, ranIterations;

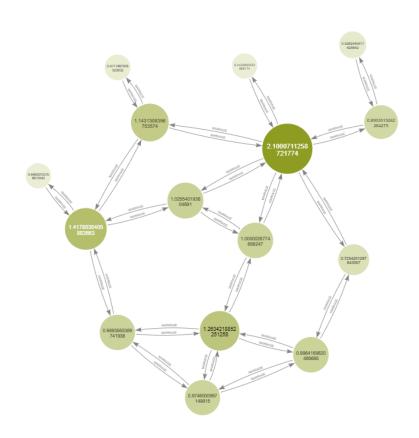


Fig 6 : -Pagerank Visual Representation by Size and Color - Using Neo4j Bloom

Final Calculations

ID	Name	Labels	Closeness	Eigenvector	Degree	Betweennes s	Pagerank
1	PERUZZI	FAMILY	0.368421052 631579	0.27560484 3151996	3	4	0.97460009 8714982
2	STROZZI	FAMILY	0.4375	0.355824431 022438	4	18.6666666 66667	1.263421885 22513
3	BISCHERI	FAMILY	0.4	0.28280593 649915	3	19	0.98935653 8974194
4	LAMBERTES	FAMILY	0.325581395 348837	0.08887088 39820869	1	0	0.449091007 58673
5	CASTELLAN	FAMILY	0.3888888 888889	0.258915328 725313	3	10	0.996416982 048869
6	RIDOLF	FAMILY	0.5	0.341643338 620216	3	20.6666666 666667	1.000002677 46882
7	TORNABUO N	FAMILY	0.48275862 0689655	0.326054571 57175	3	16.6666666 66667	1.025540193 60469
8	GUADAGNI	FAMILY	0.4666666 6666667	0.288951482 309953	4	46.33333333 33333	1.417903640 58837
9	ALBIZZI	FAMILY	0.48275862 0689655	0.244219988 092714	3	38.6666666 666667	1.1431308396 7536
10	BARBADORI	FAMILY	0.4375	0.2118218936 5433	2	17	0.725429128 784396
11	MEDICI	FAMILY	0.56	0.430141529 503199	6	95	2.1000711258 7218
12	GINORI	FAMILY	0.33333333 333333	0.07487599 27284355	1	0	0.471746783 802363
13	SALVIATI	FAMILY	0.3888888 888889	0.146089032 614658	2	26	0.890351526 226428
14	ACCIAIUOL	FAMILY	0.368421052 631579	0.132300727 485774	1	0	0.44529002 3396917
15	PAZZI	FAMILY	0.285714285 714286	0.044791707 7823384	1	0	0.526254581 762864

Excercise 3

Personal Findings on Applying CRISP-DM Methodology using Neo4J

Business Understanding

Movies comprise of many people, which includes directors, writers, producers actors, and many more, but when choosing a movie recommendation its extremely important to provide information about the movies the person wants to see, as it will reduce the time taken to search such movies by the customer and it will increase the business value of the platform that provides the service. Using IMDBs Top 250 Movies from the link below,

https://www.kaggle.com/datasets/jillanisofttech/imdb-top-250-eng-movies-dataset

We have visualized and modeled data for using algorithms like `adamicAdar` which helps us to find similar movies to recommend to customers.

Data Understanding

A Movie is directed in a Particular Year, in a particular Genre with specific people involved, hence we can better understand it by representing it in the form of nodes.

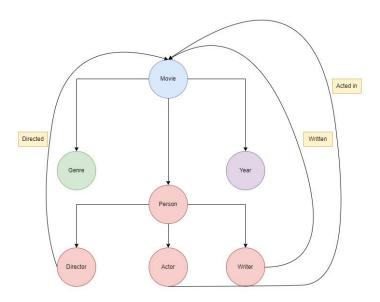


Fig 7: Relationships Defined as Nodes

As we can see above the entire movie process is related to each other by the people working on it and the genre and the year it was released on. The important aspect is the end goal, which is the rating of the movie that is to be determined on varying degrees of the likelihood of liking a movie, based on genre, person, and year, it might change from person to person.

Data Preparation

Data Cleaning

The data we have comprises 38 Columns, we selected the following columns due to their use in our project

- **Index**: To provide us with a starting point while we write the graph to neo4 js as nodes and relationships.
- **Title**: Movies are identified by titles, and that makes it one of the most important columns in our data set, as all the relationships will branch out of it.
- **Year**: Year at which the movie was released would create a link to the type of movies released depending on the community of the people present by that time.
- **Genre**: The column defines the theme of the movie, this will also determine what kind of similar movies the person would like.
- **Director**: People have affiliation with the movie directors as they also tend to choose the movies also directed by the same director which have a similar theme, like: "Christopher Nolan" Movie or "Martin Scorsese" Movie.
- **Writers**: The Entire story of the movie is written by the writers of the movie and that also determines how well the movie will do. Usually, Novel adaptations have a greater chance of success because of the depth of the story.
- **Actors**: The talent in the movies determines the screenplay quality and more or less the success of that screenplay and popularity of the talent itself to gain a pace of success, a movie with great acting and a mediocre story could also do ok in the industry.
- **Rating**: Rating reflects the end product of the movie, that is customer satisfaction, which is represented in the data on a scale of 1 10

All the columns above was important for the project, the rest was not necessary for our specific use-case, hence it was dropped.

The End Result looks something like this.



Fig 8 : - Data Set Viewing using Pandas Dataframe.

After the data is usable we need to load it as nodes in the NeO4j Graph Database.

Create Nodes

```
CREATE (m:Movie {id:$index,name: $Movie_Name})
SET
m.directors = $director,
m.Year = $year,
m.rating = $rating,
m.cast=$cast,
m.writers = $writer,
m.genre = $genre
```

Fig 9 : - Query String to create nodes

Using Py2Neo we can use the query to write to the graph database.

Creating Relations

Pseudo Code

//Match Query

MATCH (m:Movie)

//Where Clause

WHERE m.cast IS NOT NULL

WITH m

UNWIND split(m.cast, ',') AS actor

MERGE (p:Person (name: trim(actor)))

MERGE (p)-[r:ACTED_IN]->(m);

Following Relations have been Created for Our use

- 1. Genre
- 2. CREATED_ON
- 3. WORK_WITH
- 4. ACTED_IN
- 5. DIRECTED
- 6. NEXT

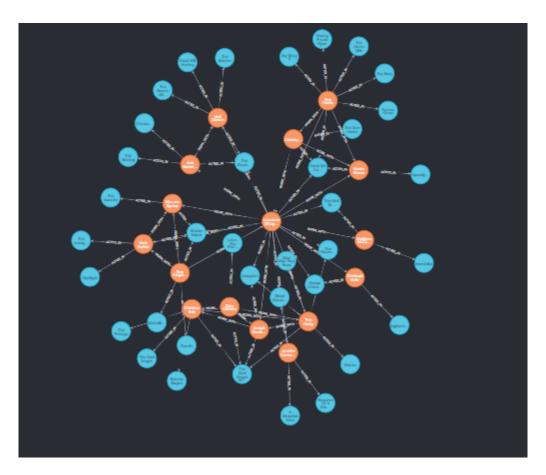


Fig 10: - Node Preview with relations

Modeling

Using Graph data Science Link Prediction Library Adamic Adar

The Algorithm **Adamic Adar** uses graph links we created above and determines the most similar nodes to it, provided we give a starting point, in this case, it will be our **Movie Name** in data we can leverage the prebuilt models in neo4j to find patterns in the data.

$$A(x,y) = \sum_{u \in N(x) \cap N(y)} \frac{1}{\log |N(u)|}$$

Fig 10: - Adamic Adar Equation

Adamic Adar takes the inverse of the logarithmic value of degree centrality, we run the above formula using the Graph Data Science library of Neo4j, it calculates recommendation on Node Strength and provides us the well-connected nodes with a score called Adamic Adar Index, The index shows what is the strength of the nodes relations and provides us with a numerical value for each node. It was generally developed for social network analysis.

Below is the implementation of Adamic Adar using Neo4j.

```
MATCH-(a:Movie-{name:$ptitle}-)-[*4]-(b:Movie)
WHERE-a-<>-b-AND-a.name-<-b.name
WITH-DISTINCT-a,b
RETURN-a.name-as-title,-b.name-as-recommendation,-gds.alpha.linkprediction.adamicAdar(a,-b)-AS-score
ORDER-BY-score-DESC
LIMIT-10
```

Fig 11: - Adamic Adar Neo4j Query

Using the graph object from Neo4J we can query the data as below

graph.run(request_link_prediction_movie,ptitle="Inception").to_data_frame()

We receive the output as a dictionary which contains the Movie Names that are similar to the ones we provided

	title	recommendation
0	Inception	The Dark Knight Rises
1	Inception	Mad Max: Fury Road
2	Inception	Interstellar
3	Inception	The Revenant
4	Inception	Spider-Man: Homecoming
5	Inception	The Prestige
6	Inception	Shutter Island
7	Inception	The Dark Knight
8	Inception	Toy Story 3
9	Inception	Warrior

Fig 12: - Recommendations eg

Evaluation

The Evaluation of Adamic Adar can be determined by the score provided by Adamic Adar, the higher the value of score the closer the nodes are to the recommendation, which can be seen below.

Title	Recommendation	Score	
	The Dark Knight Rises	1.65071083440955	
	Mad Max: Fury Road	1.17882767875969	
	Interstellar	1.06981187145939	
	The Revenant	0.882428994290986	
Inception	Spider-Man: Homecoming	0.832851422498494	
	The Prestige	0.824565809649557	
	Shutter Island	0.805105018589713	
	The Dark Knight	0.790836235778608	

Toy Story 3	0.759144404179581

Deployment

We can create a Py2Neo based Web API to get the recommender system of the Neo4j Working, Neo4j itself could be deployed as a cluster server which does the AI Part by itself when provided a pipeline for basic data management and machine learning.

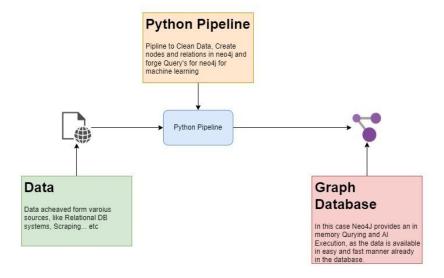


Fig 13: - Graph DB Pipeline

Note

Please find the cypher codes for Excercise 1 and Jupyter Notebook for Excercise 3 in the Zip File

References

- Neo4j Graph Data Platform. (n.d.). Adamic Adar Neo4j Graph Data Science. [online] Available at: https://neo4j.com/docs/graph-data-science/current/alpha-algorithms/adamic-adar/ [Accessed 21 Apr. 2022].
- Neo4j Graph Database Platform. (n.d.). *The Neo4j Graph Data Science Library Manual v1.8 Neo4j Graph Data Science*. [online] Available at: https://neo4j.com/docs/graph-data-science/current/.

 Wikipedia Contributors (2022). Adamic—Adar index. [online] Wikipedia. Available at: https://en.wikipedia.org/wiki/Adamic%E2%80%93Adar_index [Accessed 21 Apr. 2022].