



2023
EUROPE
INTERCHANGE
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**All you wanted to know about Blood Pressure
but were afraid to ask**

Presented by Marius Conjeaud, Neo4j
And by Nicolas de Saint Jorre, CEO, Quanticoft



Meet the Speakers

Nicolas de Saint Jorre

Title: CEO

Organization: Quanticsoft

More than 26 years of experience in the world of Data Management and Clinical Research, since 2000, working on EDC system. Since 2005: working with EvidentIQ (EDC with CDISC compliance). I contributed to the CDISC360 project since 2018 setting up a prototype of 'Study Builder' using a central library of metadata. Since 2019, working with Novo Nordisk on the OpenStudyBuilder.

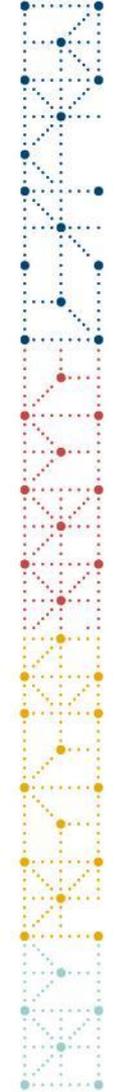


Marius Conjeaud

Title: Consulting Engineer & Team Leader

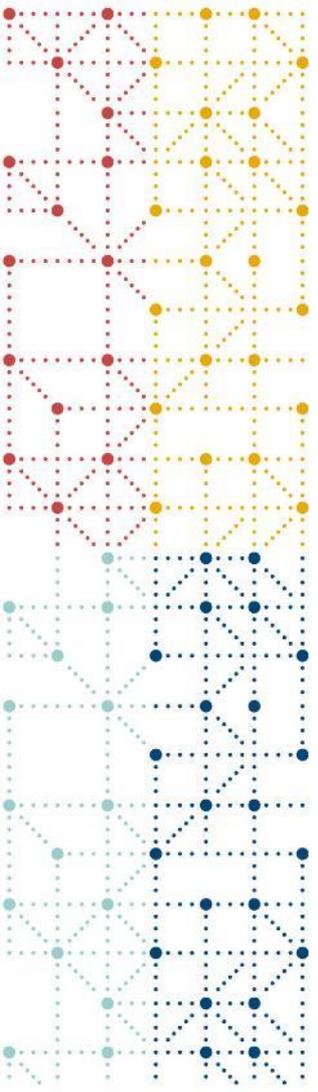
Organization: Neo4j

Experienced technical architect with a background in development, graph databases and visualization. As a consulting engineer, I help Neo4j's customers make sense of their data by leveraging the graph ecosystem - through graph data modeling, APIs, query and front-end development. I also have experience working with both business users and data actors - data engineers and data scientists



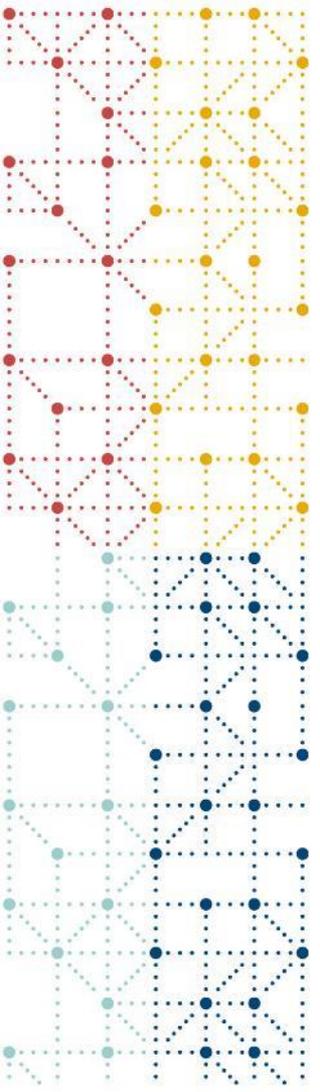
Disclaimer and Disclosures

- *The views and opinions expressed in this presentation are those of the author(s) and do not necessarily reflect the official policy or position of CDISC.*



Agenda

1. Metadata Repository? Where to start
2. The CT and the Models in a Graph database
3. Analyzing some examples
4. Biomedical Concept, 3D graphs and tools
5. Conclusion and future fiction



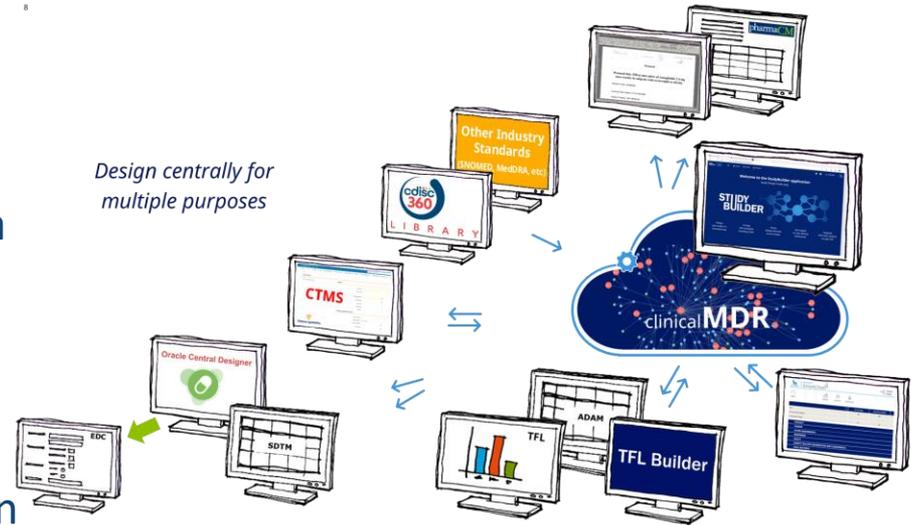
Metadata Repository – Where to start

From CDISC360 to the OpenStudyBuilder through Controlled Terminology and Models inside a Graph database...

CDISC360 – 2018, we started using Neo4j as a database

A Graph database to rule them all:

- Between 2018 and 2019: The CDISC360 project set a POC showing how to manage metadata from a Protocol to Reporting through CRF management and SDTM tabulation
- Neo4j was selected as a Graph database with an API and a Front in Python, using the Django framework



- Summary
- Objectives / Endpoints
- Derived Assessments ▲
- Collected Assessments ▲
- Schedule of Activities
- Schedule of Assessments**
- Data Collection ▲
- Tables, Figures and Listings ▲

CDISC360-2 / Schedule of Assessments

We have for this study the following visits and the following Assessments: i



Epoch		Screening	Treatment																
Activity	Assessment	Visit 1	Visit 2	Visit 3	Visit 4	Visit 5	Visit 6	Visit 7	Visit 8	Visit 9	Visit 10	Visit 11	Visit 12	Visit 13	Visit 14	Visit 15	Visit 16	Visit 17	
Randomisation	Randomisation Date	⊗	✔	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗
Body Measurement	Body Weight	✔	✔	⊗	✔	⊗	⊗	⊗	✔	⊗	✔	⊗	⊗	✔	⊗	⊗	⊗	⊗	⊗
Glucose metabolism	Hemoglobin A1C/Hemoglobin	✔	✔	✔	✔	✔	✔	✔	✔	✔	✔	✔	✔	✔	✔	✔	✔	✔	✔
	Glucose, Plasma	✔	✔	✔	✔	✔	✔	✔	✔	✔	✔	✔	✔	✔	✔	✔	✔	✔	✔
Lipids	HDL Cholesterol	✔	✔	✔	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗
	LDL Cholesterol	✔	✔	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗
	Total Cholesterol	✔	✔	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗
Biochemistry	Albumin	✔	✔	⊗	⊗	⊗	✔	✔	✔	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗
	Creatinine	✔	✔	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗
	Alanine Aminotransferase	✔	✔	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗
Haematology	Haemoglobin Blood	✔	✔	⊗	⊗	⊗	✔	⊗	⊗	✔	⊗	✔	⊗	⊗	⊗	⊗	⊗	⊗	⊗
	Haematocrit Blood	✔	✔	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗
	Thrombocytes	✔	✔	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗
Vital signs	Systolic Blood Pressure	✔	✔	⊗	⊗	✔	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗
	Diastolic Blood Pressure	✔	✔	⊗	⊗	✔	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗
	Pulse	✔	✔	⊗	⊗	✔	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗
	Body Temperature	✔	✔	⊗	⊗	✔	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗	⊗
Adverse Event	Adverse Event	✔	✔	✔	✔	✔	✔	✔	✔	✔	✔	✔	✔	✔	✔	✔	✔	✔	✔

- <<
- About Studies
- Process Overview
- Manage Studies
- Define Study
- Specification Overview
- Study Title
- Registry Identifiers
- Study Properties
- Study Structure
- Study Population
- Study Criteria
- Study Interventions
- Study Purpose
- Study Activities
- Terminology
- View Specifications
- View Listings

Studies / Define Study / Study Activities / Detailed Flowchart

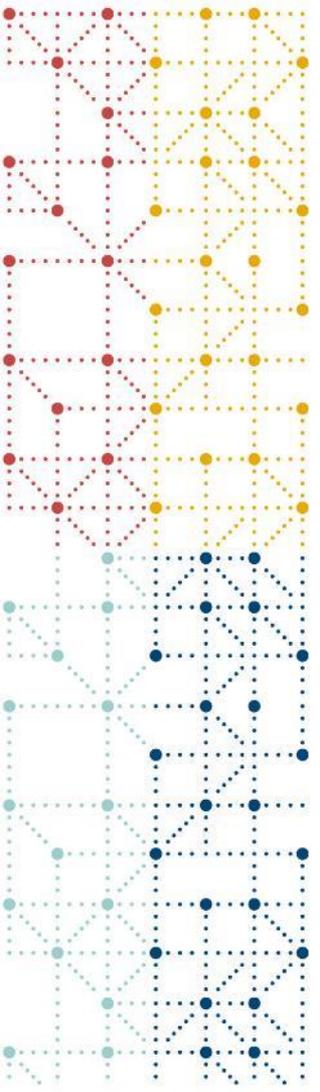
Study Activities (CDISC DEV-0)

List of Study Activities **Detailed Flowchart** Protocol Flowchart Activity Instructions

Expand table Collapse table Hide flowchart groups



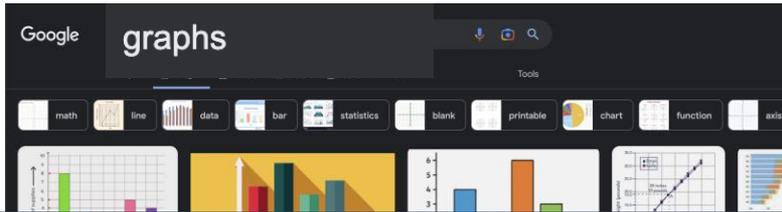
	Epoch	Screening	Treatment									Follow-up
	Visit	V1	V2	V3	V4	V5	V6	V7	V8	V9	V10	V11
	Day	-14	1	8	15	22	29	36	43	57	183	213
	Window	-13/+0	±0	-1/+1	-1/+1	-1/+1	-1/+1	-1/+1	-1/+1	-1/+1	-1/+1	0/+35
SUBJECT RELATED INFORMATION												
<input checked="" type="checkbox"/> General												
<input checked="" type="checkbox"/> Randomisation												
<input type="checkbox"/> Randomized		<input type="radio"/>	<input checked="" type="radio"/>	<input type="radio"/>								
<input checked="" type="checkbox"/> End of Study												
<input type="checkbox"/> End of Study		<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input checked="" type="radio"/>
<input checked="" type="checkbox"/> Physical Examination - early ph1												
<input type="checkbox"/> Cardiovascular System		<input checked="" type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>
<input type="checkbox"/> Abdomen		<input checked="" type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>	<input type="radio"/>



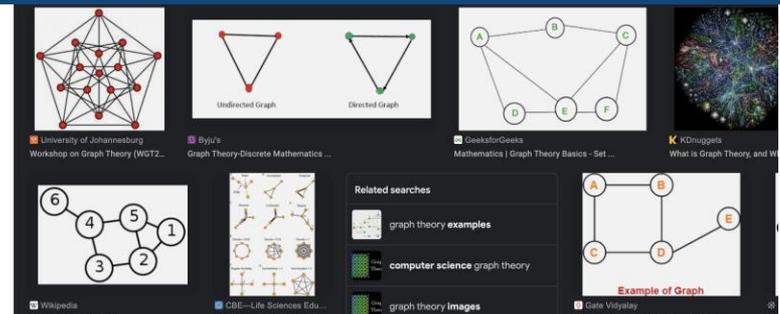
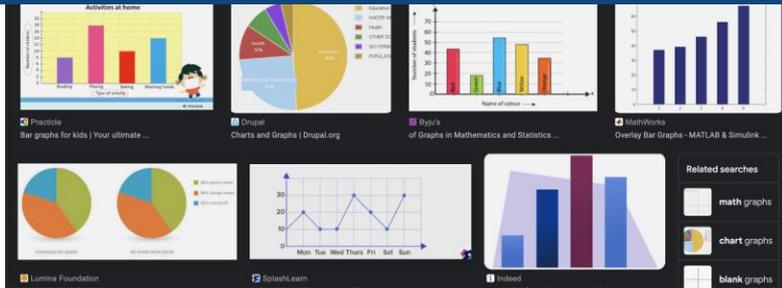
The CT and the Models in a Graph database

We are downloading the CT and the Models via the CDISC API into the Neo4j database without duplicating information!!!

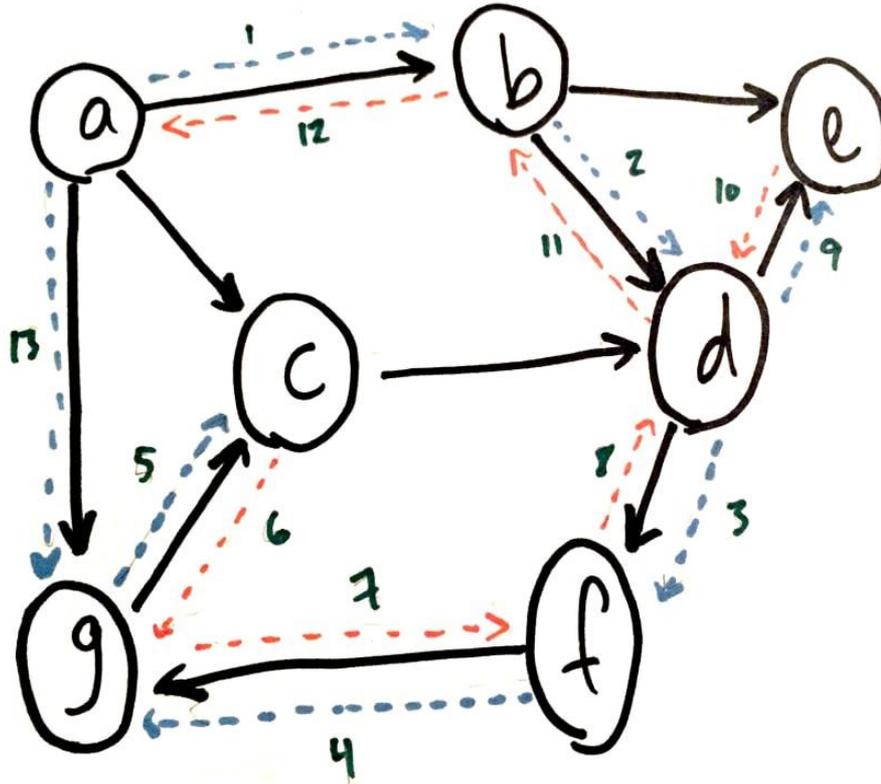
What is a Graph Database



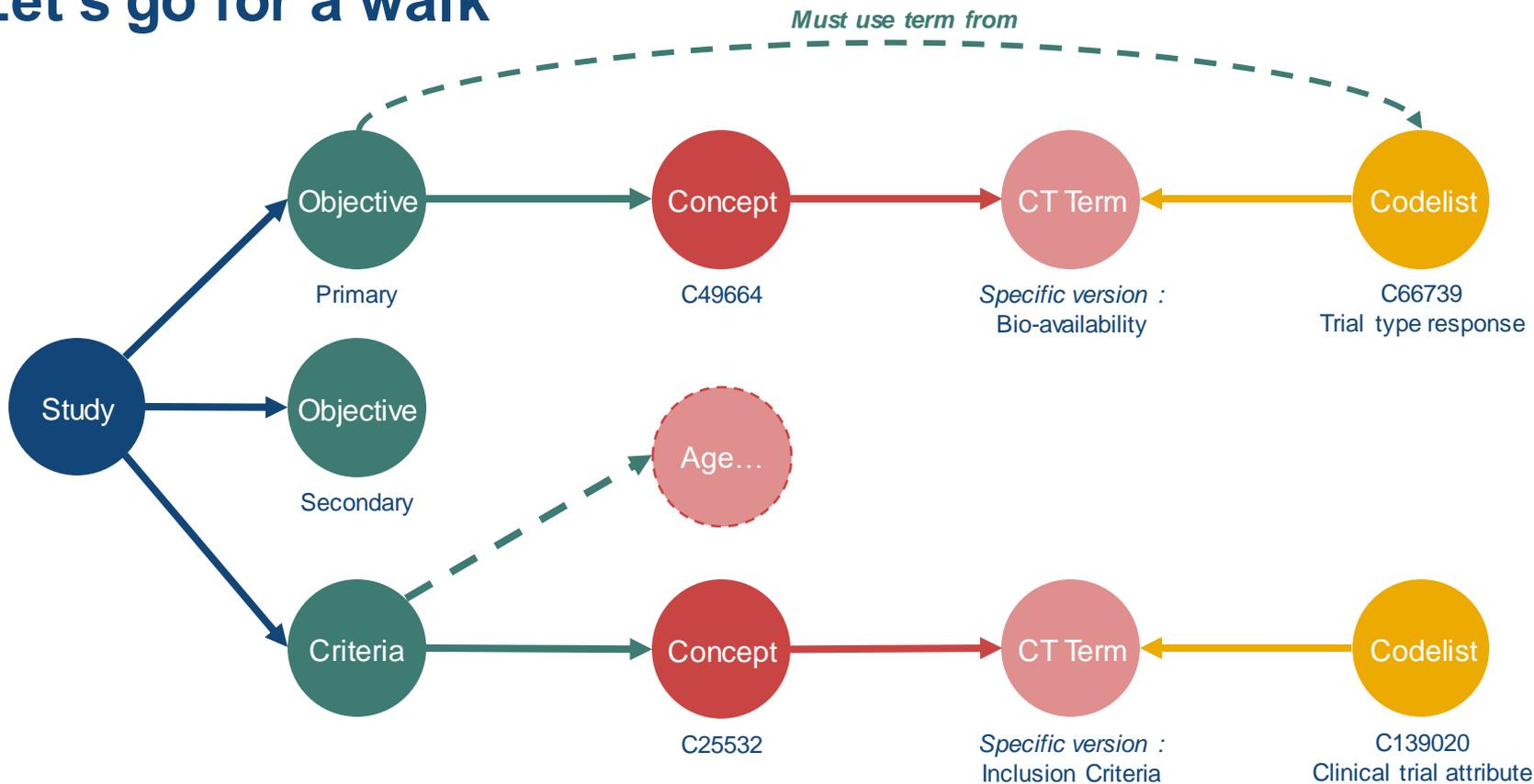
Where « relational » databases focus on data points, graph databases focus on relationships between these points

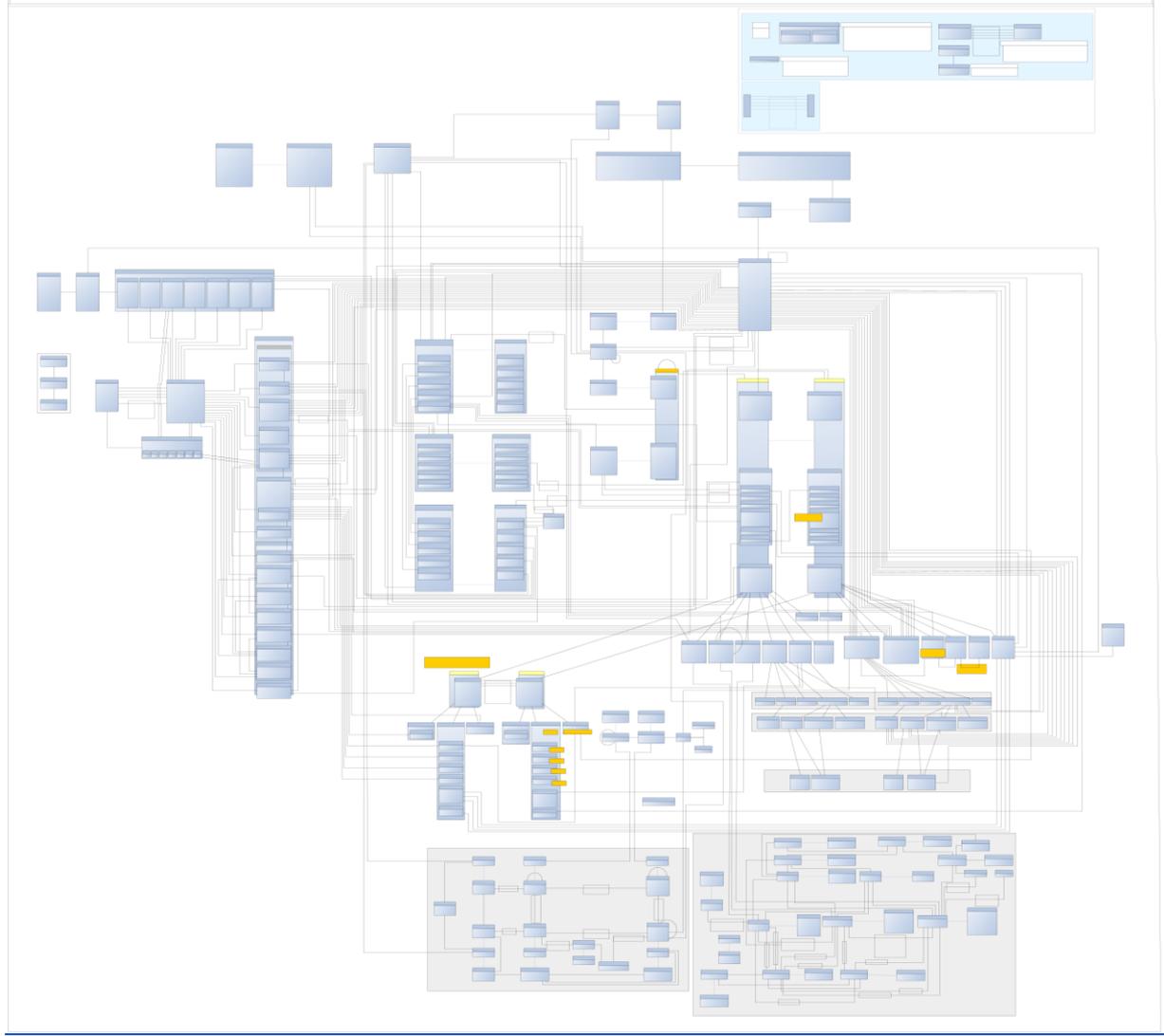
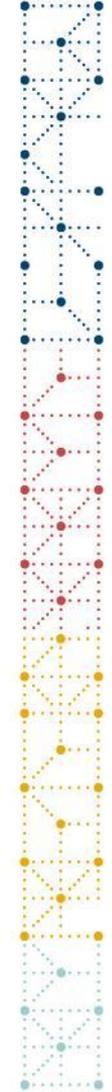


Graph Databases focus on relationships... and traversal

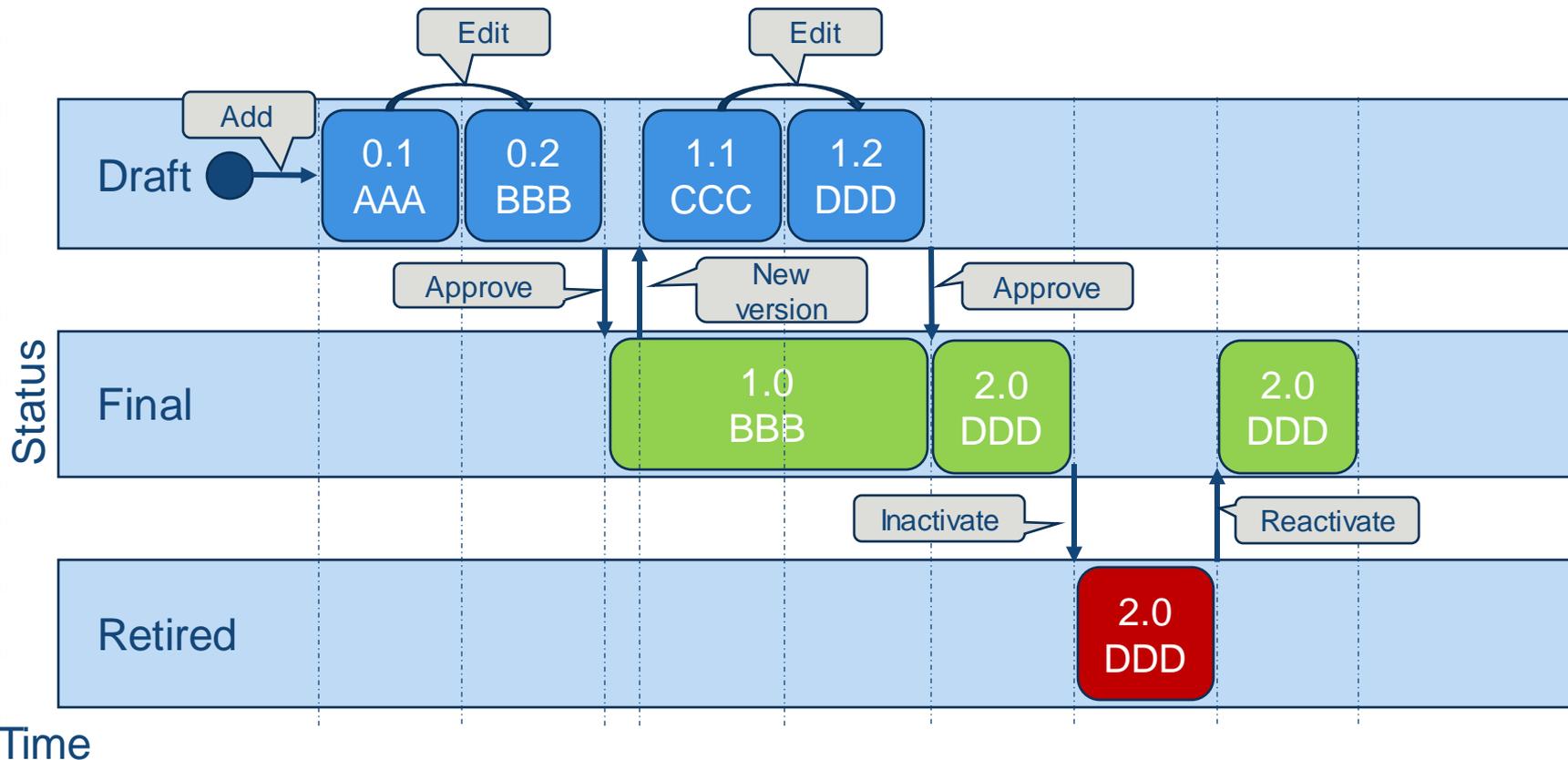


Let's go for a walk

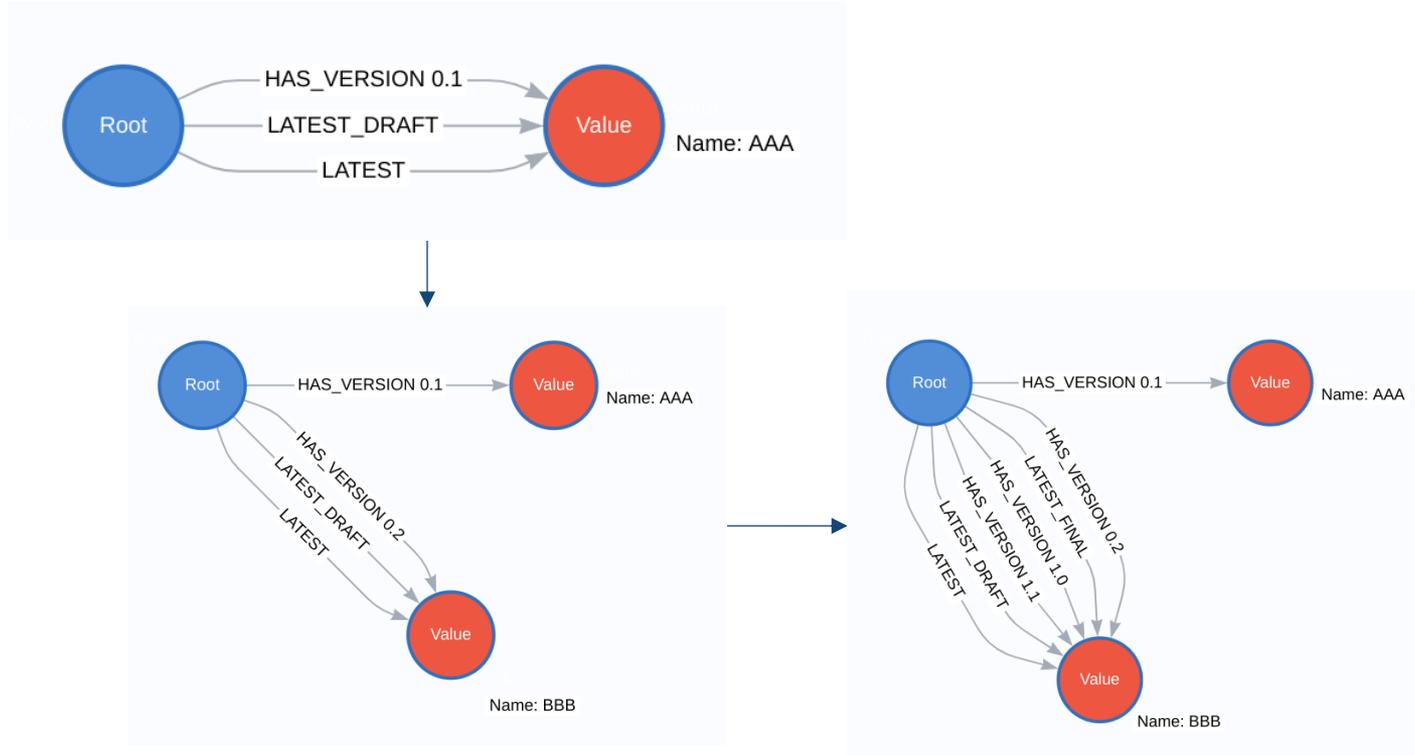




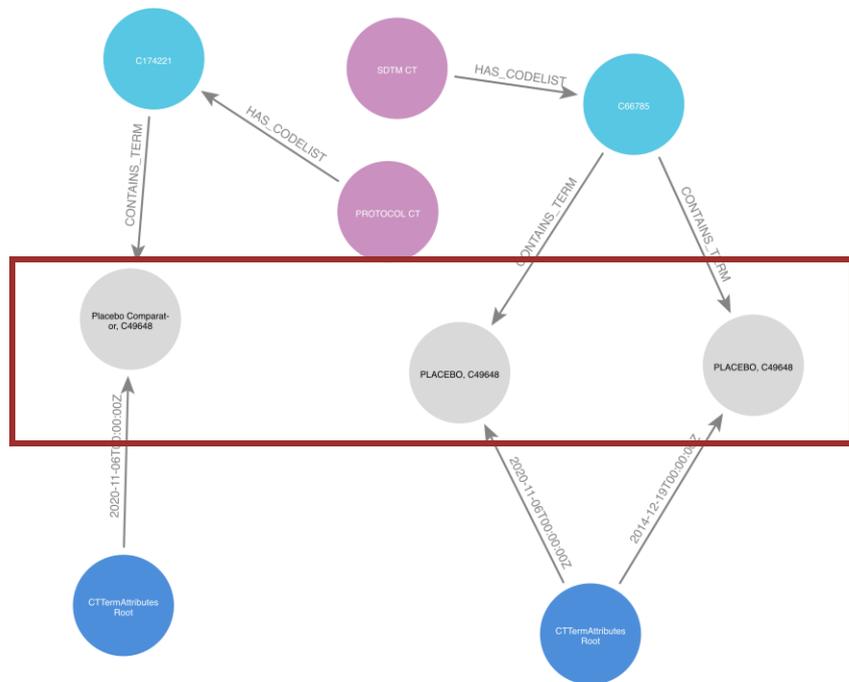
Version control concept for metadata



Version control around the CT and the Models in the Graph database



Very strange behavior?!



-  Catalogue
-  Codelist
-  Term
-  Term root

Terms having the same
ConceptID but with a
different Submission Value
In two Catalogues...

Things that we still need

Test-Case 17

JSON Input File(s)

Package "SDTM CT 2"

```
"conceptId": "C71620",
"name": "Unit",
"submissionValue": "UNIT"
```

Package "SDTM CT 2"

```
"conceptId": "C71620",
"name": "Unit",
"submissionValue": "UNIT"
```

Test-Case 18

JSON Input File(s)

Package "SDTM CT 2016-06-24"

```
"conceptId": "C66786",
"name": "Country",
"submissionValue": "COUNTRY"
```

This Co

Package "SDTM CT 2016-09-30"

```
"conceptId": "C66786",
"name": "Country",
"submissionValue": "COUNTRY"
```

Every T

Test-Case 19 Term deleted and replace in the next package by another

JSON Input File(s)

Package "SDTM CT 2014-09-26"

```
"conceptId": "C71153",
"name": "ECG Test Code",
"submissionValue": "EGTESTCD",
```

```
"conceptId": "C39779",
"submissionValue": "HRMAX",
"definition": "The minimum time between successive cycles of contraction and subsequent relaxation of the heart, usually expressed as beats per minute, obtained from a set of measurements of the heart rate. (NCI)"
"preferredTerm": "Maximum Heart Rate"
"synonyms": "Summary (Max) Heart Rate"
```

Code/Name codelist pair.

```
"conceptId": "C71152",
"name": "ECG Test Name",
"submissionValue": "EGTEST",
```

```
"conceptId": "C39779",
"submissionValue": "Summary (Max) Heart Rate",
"definition": "The minimum time between successive cycles of contraction and subsequent relaxation of the heart, usually expressed as beats per minute, obtained from a set of measurements of the heart rate. (NCI)"
"preferredTerm": "Maximum Heart Rate"
"synonyms": "Summary (Max) Heart Rate"
```

How are we going to track the replacement of the HRMAX by the EGHRMAX Tem???

Package "SDTM CT 2014-12-26"

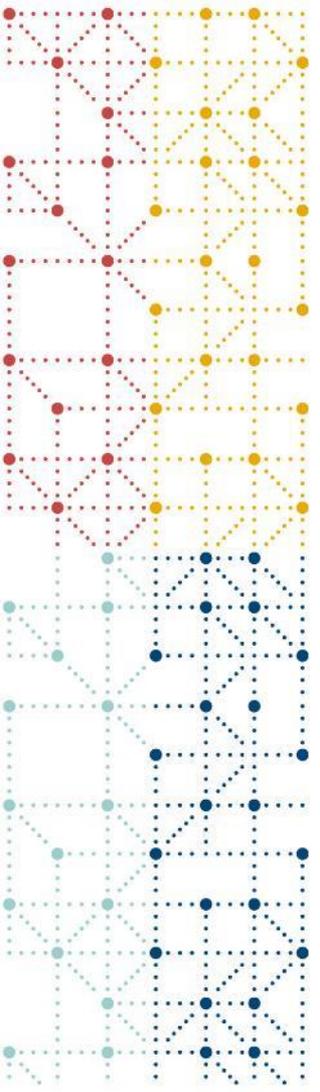
```
"conceptId": "C71153",
"name": "ECG Test Code",
"submissionValue": "EGTESTCD",
```

```
"conceptId": "C119257",
"submissionValue": "EGHRMAX",
"definition": "An electrocardiographic measurement of the maximum rate of depolarization of a specific region of the heart during an interval of time, usually expressed in beats per minute. Unless otherwise specified, this is usually the maximum ventricular rate."
"nameSubmissionValue": "ECG Maximum Heart Rate"
"preferredTerm": "Maximum Heart Rate by Electrocardiogram"
"synonyms": "ECG Maximum Heart Rate"
```

Code/Name codelist pair.

```
"conceptId": "C71152",
"name": "ECG Test Name",
"submissionValue": "EGTEST",
```

```
"conceptId": "C119257",
"submissionValue": "ECG Maximum Heart Rate",
"definition": "An electrocardiographic measurement of the maximum rate of depolarization of a specific region of the heart during an interval of time, usually expressed in beats per minute. Unless otherwise specified, this is usually the maximum ventricular rate."
"preferredTerm": "Maximum Heart Rate by Electrocardiogram"
"synonyms": "ECG Maximum Heart Rate"
```



Analyzing some examples

As we have a version control solution, let's see how this is helping us in managing our CDISC metadata with examples

Some Cypher examples

CT RACE Codelist:

```
// CT for the RACE - C74457 Codelist
```

```
MATCH (n0:CTCatalogue)
```

```
MATCH (n0)-[]->(n1:CTCodelistRoot)
```

```
MATCH (n1)-[]-
```

```
>(n2:CTCodelistAttributesRoot)-[]-
```

```
>(n3:CTCodelistAttributesValue)
```

```
MATCH (n1)-[]->(n4:CTTermRoot)
```

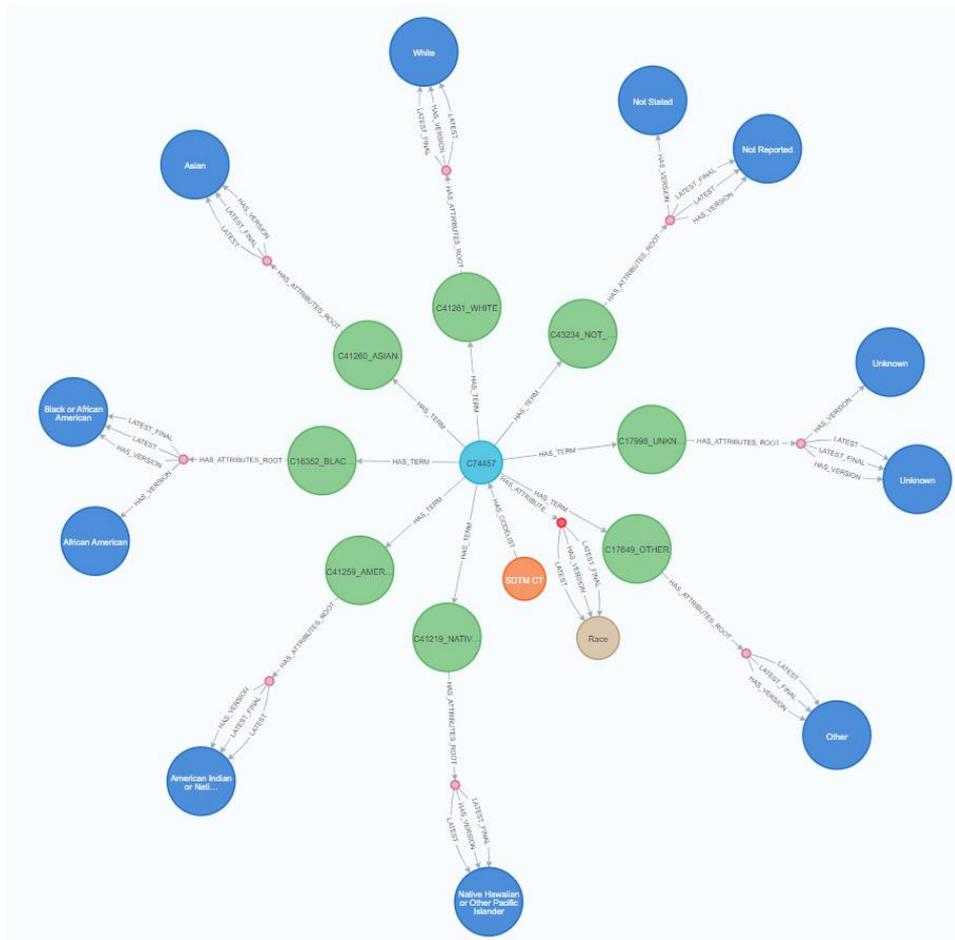
```
MATCH (n4)-[]-
```

```
>(n5:CTTermAttributesRoot)-[]-
```

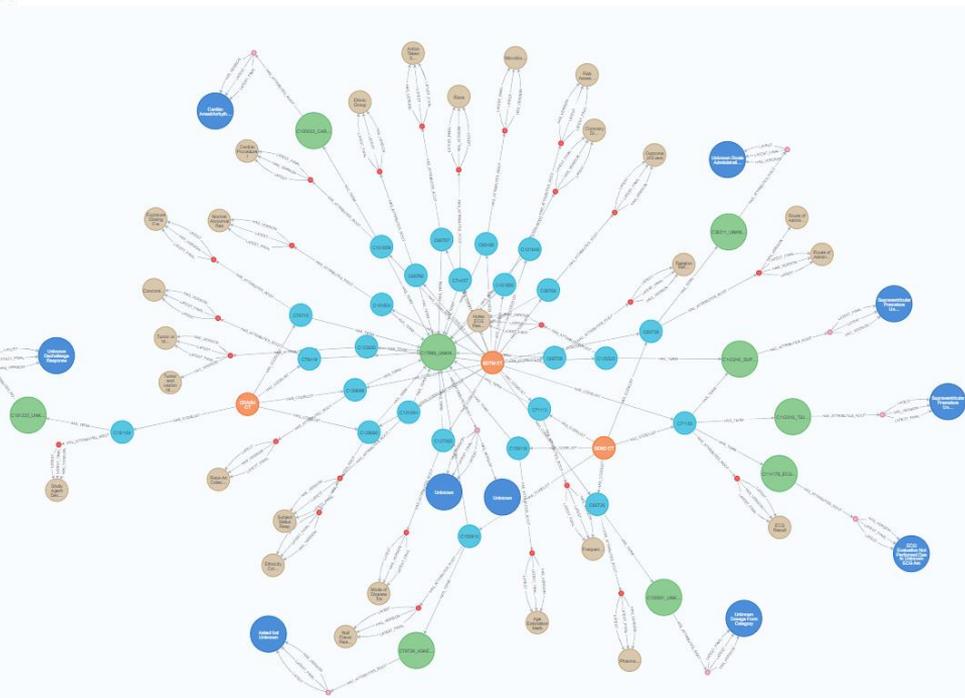
```
>(n6:CTTermAttributesValue)
```

```
WHERE n1.uid = 'C74457'
```

```
RETURN n0,n1,n2,n3,n4,n5,n6;
```



Some Cypher examples



CT UNKNOWN Terms...:

```
// CT with a UNK Term used
MATCH (n0:CTCatalogue)
MATCH (n0)-[]->(n1:CTCodelistRoot)
MATCH (n1)-[]->
>(n2:CTCodelistAttributesRoot)-[]->
>(n3:CTCodelistAttributesValue)
MATCH (n1)-[]->(n4:CTTermRoot)
MATCH (n4)-[]->
>(n5:CTTermAttributesRoot)-[]->
>(n6:CTTermAttributesValue)
WHERE n4.uid CONTAINS '_UNK'
RETURN n0,n1,n2,n3,n4,n5,n6;
```

Some Cypher examples

SDTMIG VS:

```
// SDTMIG VS Domain for Visit Number
```

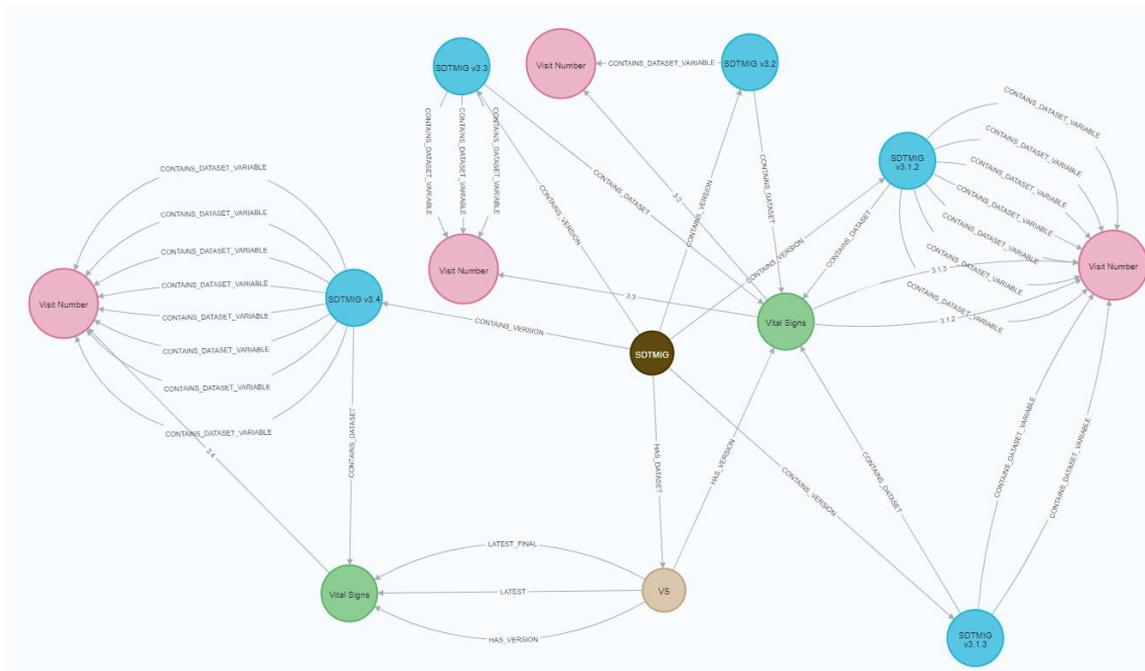
```
MATCH (n0:DataModelCatalogue)-[r1]->(n1:DatasetRoot)
```

```
MATCH (n1)-[r2]->(n2:DatasetValue)-[r3]->(n3:DatasetVariableValue)
```

```
MATCH (n3)<-[]-(n4:DataModelVersion)
```

```
WHERE n0.name = 'SDTMIG' AND  
n1.uid = 'VS' AND n4.name  
CONTAINS 'SDTMIG' AND n3.title =  
'Visit Number'
```

```
RETURN n0,n1,n2,n3,n4 ORDER BY  
n3.ordinal;
```



Some Cypher examples

SDTMIG VS:

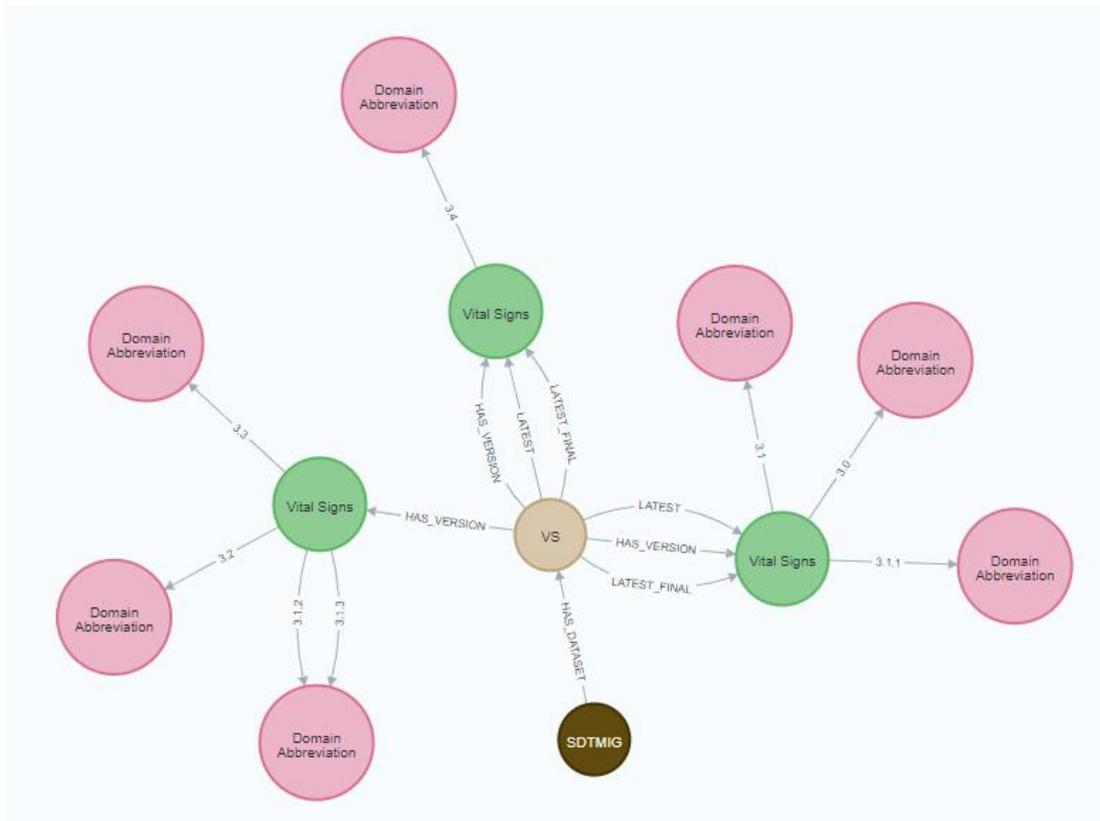
```
// SDTMIG VS Domain for Domain  
Abbreviation
```

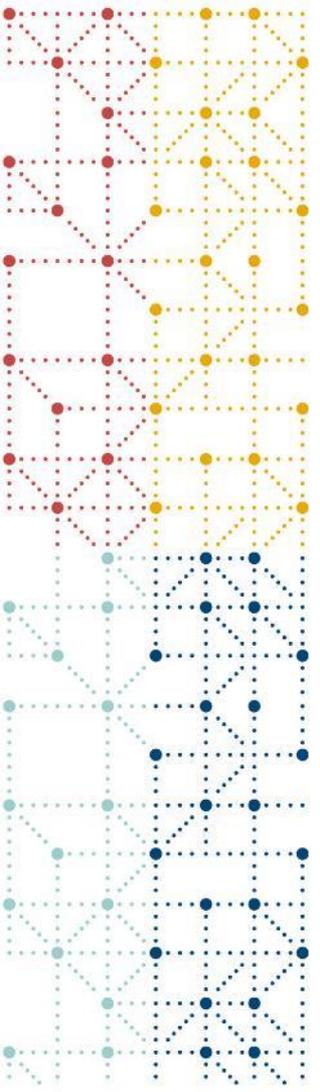
```
MATCH (n0:DataModelCatalogue)-  
[r1]->(n1:DatasetRoot)
```

```
MATCH (n1)-[r2]-  
>(n2:DatasetValue)-[r3]-  
>(n3:DatasetVariableValue)
```

```
WHERE n0.name = 'SDTMIG' AND  
n1.uid = 'VS' AND n3.title =  
'Domain Abbreviation'
```

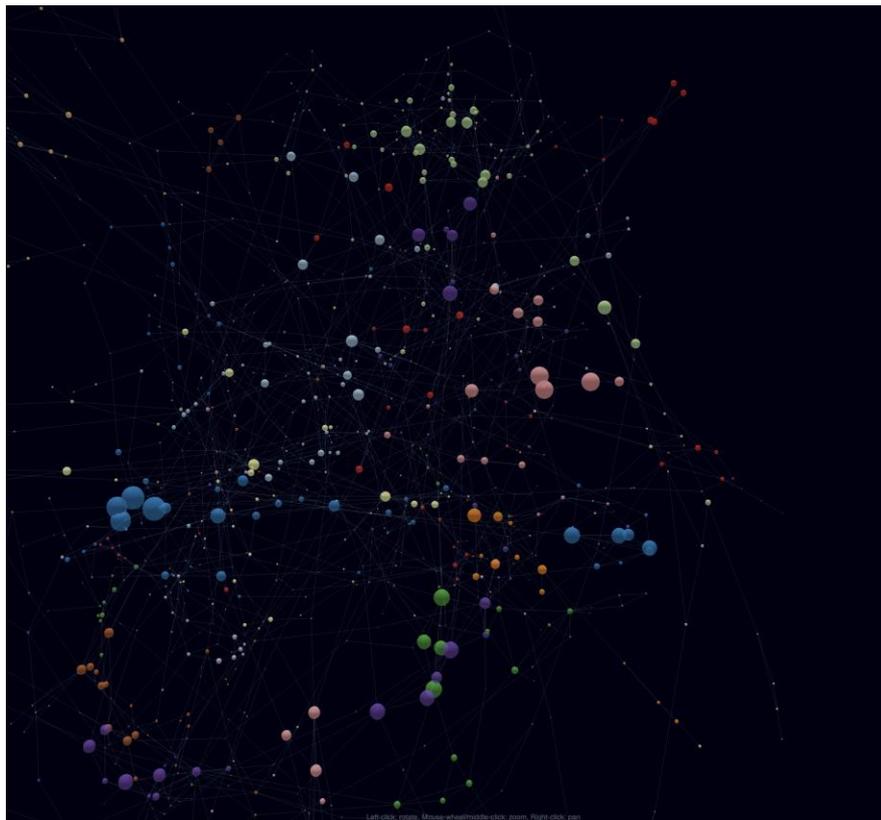
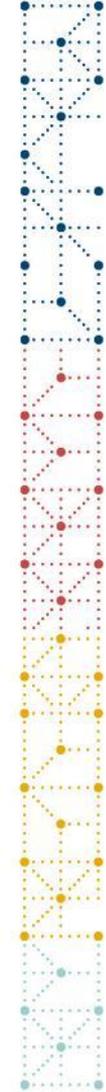
```
RETURN n0,n1,n2,n3 ORDER BY  
r3.ordinal;
```





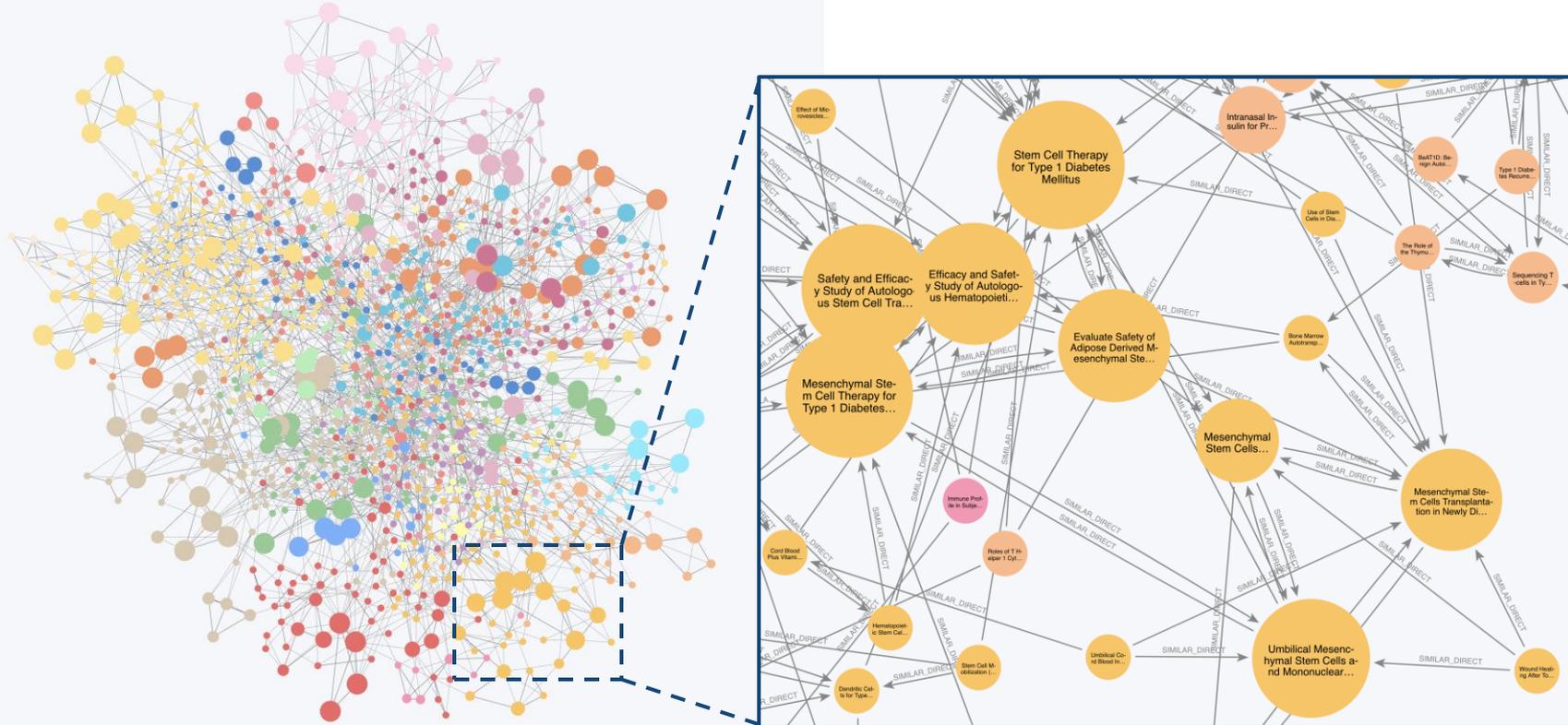
Biomedical Concept, 3D graphs and tools

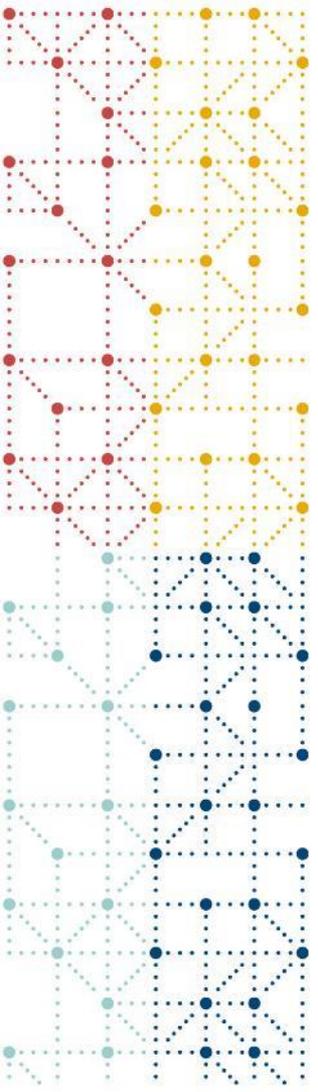
By adding some relationship between CT, Models, CRF elements, SDTM variables, we can create some BC.
A 3D view can help us in looking at those data, but we also have some other tools...



3D view to display our Neo4j data

Automatic, content-based grouping of studies



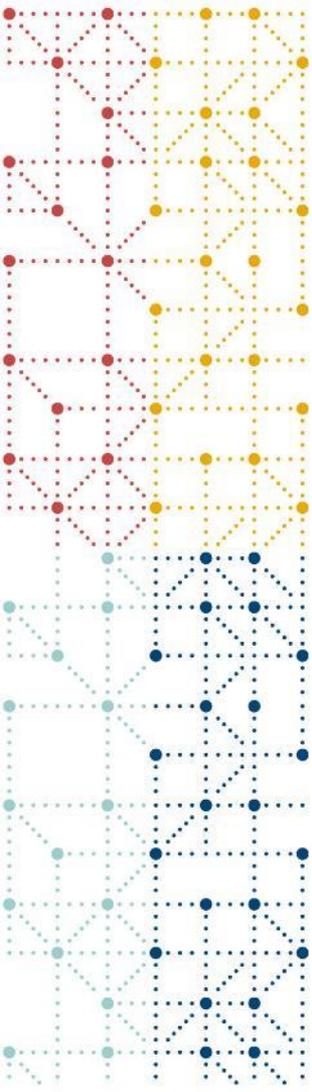


Conclusion and future fiction

Are we looking at new Data Management way of working?

Where is the AI around those tools?

Aren't we cutting the branch we are sitting on?



Thank You!

You can call us:

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