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Extending Biomedical Concepts to Analysis

Presented by Kirsten Walther Langendorf, Partner at data4knowledge



#### **Meet the Speaker**

Kirsten Walther Langendorf

Title: Partner

Organization: data4knowledge

Worked 20 years in the pharmaceutical industry within programming, IT implementation, process improvement, standards implementation, and statistics.

Advices and implements better solutions and processes for use re-use of data and metadata.

Is motivated and driven by implementation of linked-data, metadatadriven solutions, and automated end-2-end process for clinical data.

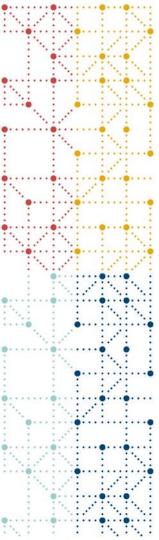
It should be simple and straight forward to use CDISC standards while working with clinical data.

Supports CDISC in the development of Biomedical Concepts.

#### **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. Background
- 2. End-2-end data flow with BCs extended to analysis
- 3. Derivation Concepts examples
- 4. Proof-of-concept implementation
- 5. Summary

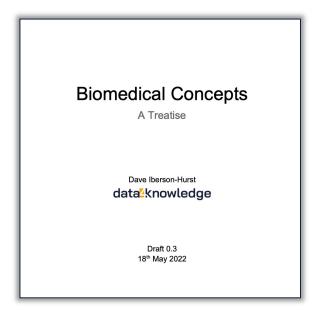


Background

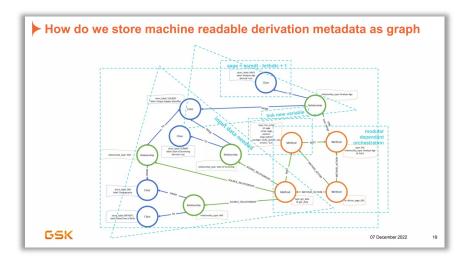
#### The input and inspiration

Paper by Dave Iberson-Hurst

Alexey Kuznetsov and Jorine Putter PhUSE EU 2022



https://www.slideshare.net/neo4j/how-will-knowledge-graphs-improve-clinical-reporting-workflows



https://github.com/data4knowledge/biomedical\_concepts/blob/main/docs/bc%20treatise/Biomedical%20Concepts%20Treatise.pdf

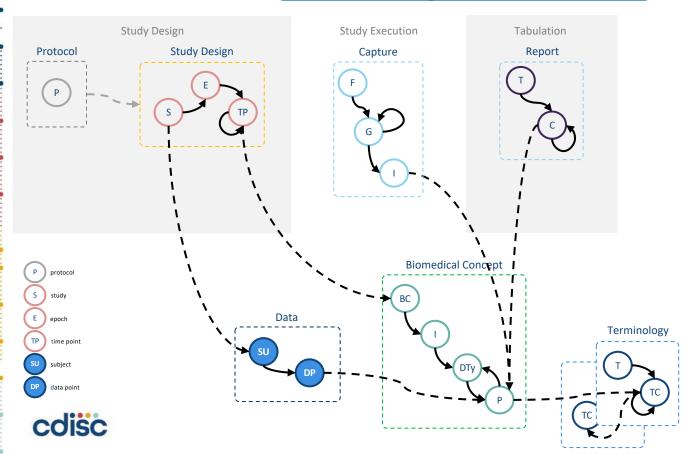




### **End-2-end data flow**

with Biomedical Concepts - extended to Analysis

# The end-to-end view – also focus for <u>CDISC Digital Data Flow</u>







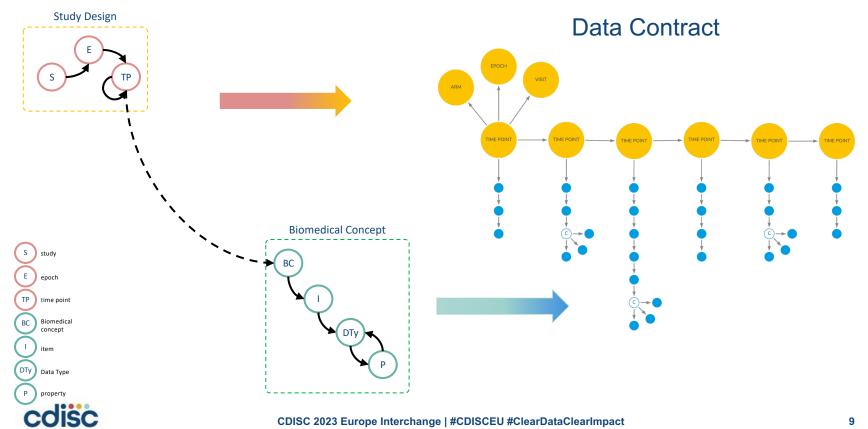


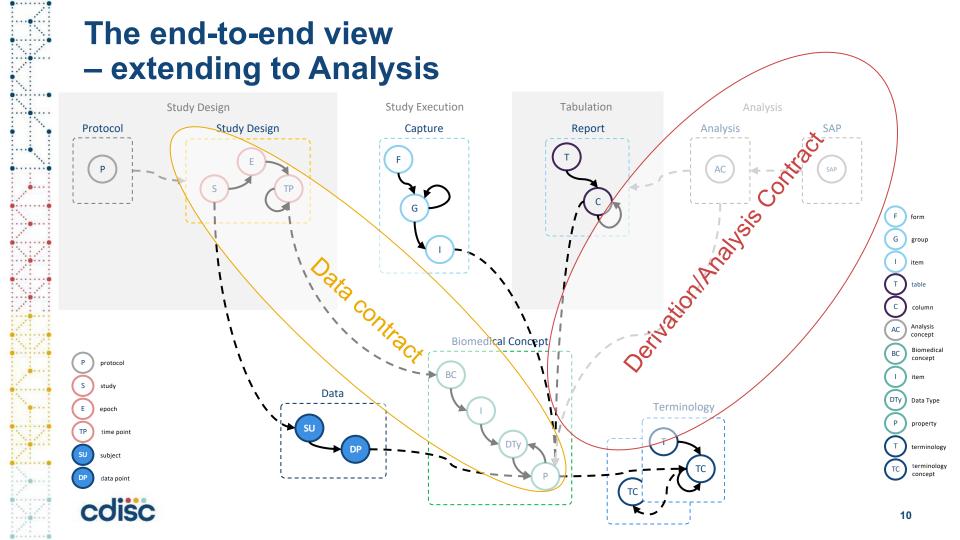


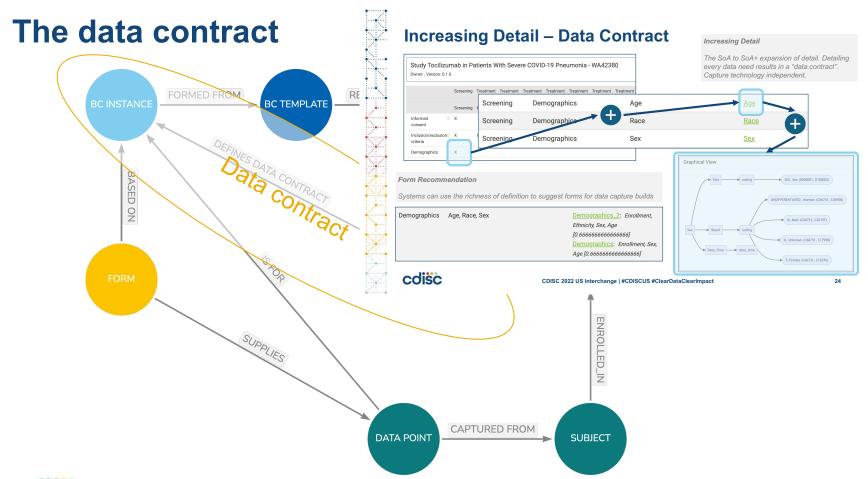




#### **BC** and the data contract

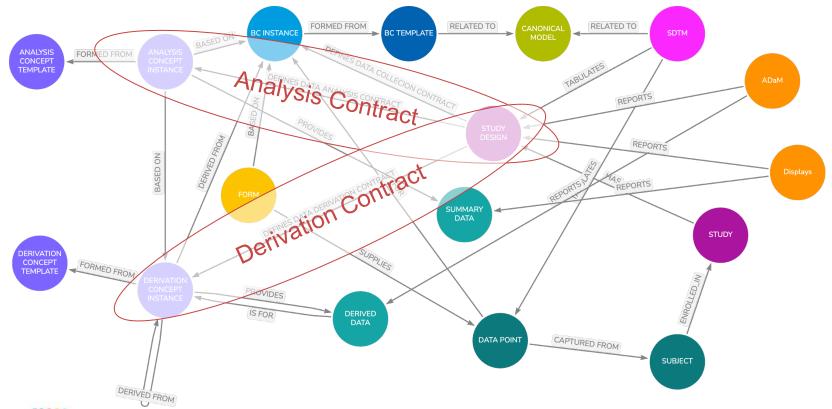








#### The data contract – extended to derivation/analysis

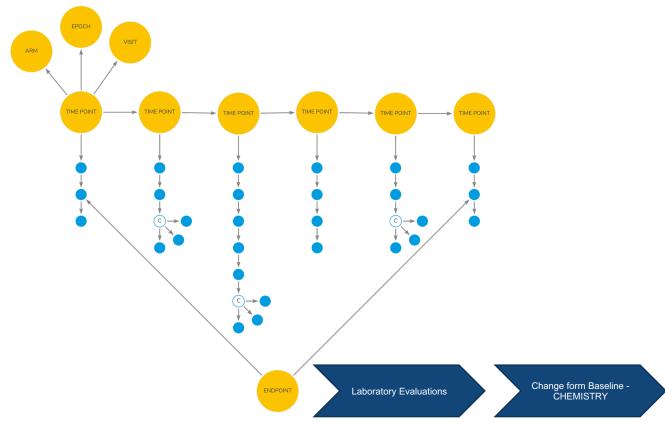




### **Derivation Concepts**

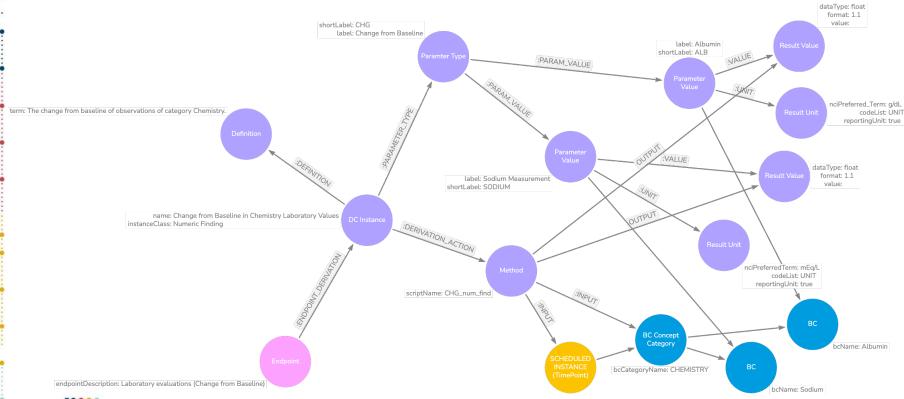
Examples and some initial design

### **Change from Baseline in Chemistry Laboratory**



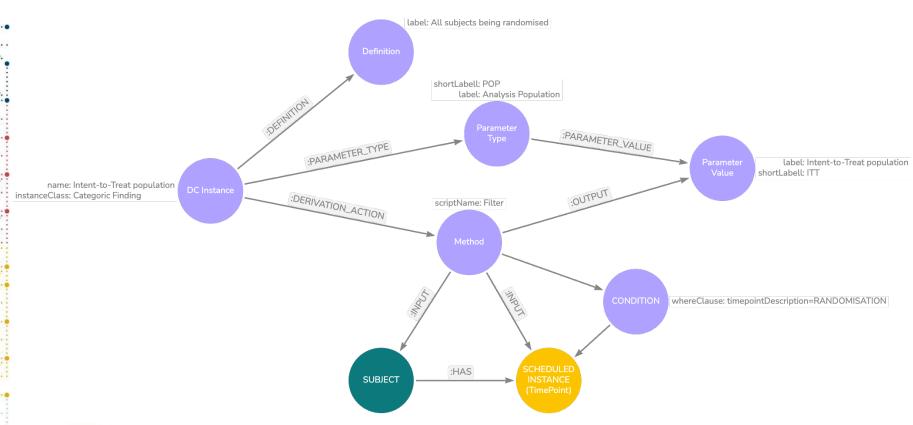


## Derivation concept – Change form Baseline in Chemistry Laboratory Values





### **Derivation concept – ITT Population**





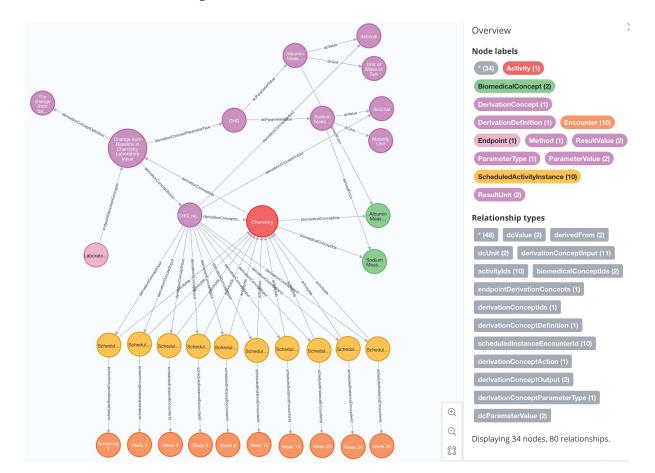


### **Proof of concept**

Derivation examples in DDF model

-using CDISC pilot study

#### A derivation concept metadata into DDF model





#### **Data contract**

- Library of Biomedical Concepts
- During study protocol setup the user selects Activities
- Next step is to detail which of the BCs (linked to Activity) is applicable for study

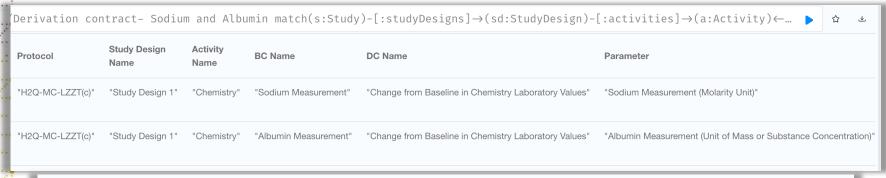
|\$ //Data contract- Sodium and Albumin match(s:Study)-[:studyDesigns] $\rightarrow$ (sd:StudyDesign)-[:activities] $\rightarrow$ (a:Activity) $\leftarrow$ [:activity]

Protocol	Study Design Name	Activity Name	BC Name	DC Name
"H2Q-MC-LZZT(c)"	"Study Design 1"	"Chemistry"	"Sodium Measurement"	"Change from Baseline in Chemistry Laboratory Values"
"H2Q-MC-LZZT(c)"	"Study Design 1"	"Chemistry"	"Albumin Measurement"	"Change from Baseline in Chemistry Laboratory Values"



#### **Derivation Contract**

- Library of Derivation Concepts these are linked to library BC concept
- During study protocol the user can pick relevant DCs based on the Activitties/BCs selected
- DCs can be linked to Endpoints



Protocol Objective		Endpoint	Activity Name DC Name		Parameter	
"H2Q-MC-LZZT(c)"	*To document the safety profile of the xanomeline TTS.*	"Laboratory evaluations (Change from Baseline)"	"Chemistry"	"Change from Baseline in Chemistry Laboratory Values"	"Sodium Measurement (Molarity Unit)"	
"H2Q-MC-LZZT(c)"	"To document the safety profile of the xanomeline TTS."	"Laboratory evaluations (Change from Baseline)"	"Chemistry"	"Change from Baseline in Chemistry Laboratory Values"	"Albumin Measurement (Unit of Mass or Substance Concentration)"	

## Data into DDF model – executing the derivation concept

df= get\_data('Sodium Measurement')



	USUBJID	PARAM	ENCOUNTER_NAME	scheduledInstanceId	VALUE
1	"01-718-1150"	"Sodium Measurement"	"Screening 1"	"ScheduledActivityInstance_1"	138.0
2	"01-718-1150"	"Sodium Measurement"	"Week 12"	"ScheduledActivityInstance_11"	138.0
3	"01-718-1150"	"Sodium Measurement"	"Week 16"	"ScheduledActivityInstance_13"	139.0
4	"01-718-1150"	"Sodium Measurement"	"Week 20"	"ScheduledActivityInstance_15"	139.0
5	"01-718-1150"	"Sodium Measurement"	"Week 24"	"ScheduledActivityInstance_17"	140.0
6	"01-718-1150"	"Sodium Measurement"	"Week 26"	"ScheduledActivityInstance_18"	144.0



## Data into DDF model – executing the derivation concept

```
df= get_data('Sodium Measurement')
df= CHG_num_find(df,'Screening 1')
```





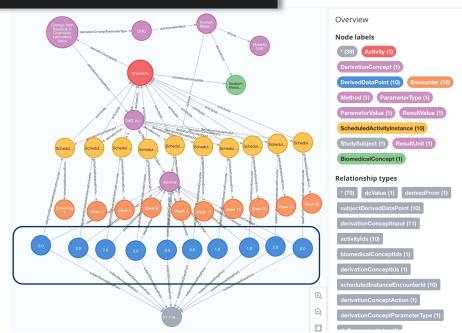
	USUBJID	PARAM	ENCOUNTER_NAME	scheduledInstanceId	VALUE	CHG
0	01-701-1015	Sodium Measurement	Screening 1	ScheduledActivityInstance_1	140.0	0.0
1	01-701-1015	Sodium Measurement	Week 12	ScheduledActivityInstance_11	139.0	-1.0
2	01-701-1015	Sodium Measurement	Week 16	<pre>ScheduledActivityInstance_13</pre>	141.0	1.0
3	01-701-1015	Sodium Measurement	Week 20	ScheduledActivityInstance_15	142.0	2.0
4	01-701-1015	Sodium Measurement	Week 24	ScheduledActivityInstance_17	140.0	0.0



### Data into DDF model – executing the derivation concept

```
df= get_data('Sodium Measurement')
df= CHG_num_find(df,'Screening 1')
save_derived_to_dc(df,'Sodium Measurement')
```

3 Save back to graph







#### **Summary**

- Presented an idea not a final model!
  - · More testing and alignment to DDF model is needed.
- To enable end-2-end metadata driven data flow we can introduce derivation concepts
- Introducing derivations concepts helps tracing how derived data was created and from where it was collected

#### **Future work**

- Further work needs to be done testing and modelling
- Need to cover the analysis concepts





#### data4knowledge

#### **Thank You!**

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