

#### Session 4, Track C: CDISC 360, Part II



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6.

Q & A session

2. PoC for Study Design and Configuration using CDISC 360 Concept-based Standards – Mikkel and Nicolas

3. Automation of SDTM & ADaM Generation and Artifacts using CDISC 360 Enriched Metadata – Bhavin and Jimmy

4. Automation of TFL Generation using CDISC 360 Enriched Metadata – Bhavin, Prasanna & Stuart

5. Concluding Remarks and Next Steps – Bhavin and Mikkel

5 mins

10 mins

15 mins

Introduction, Future State, Process and



Mikkel Traun, *Novo Nordisk*Nicolas de Saint Jorre, *XClinical* 

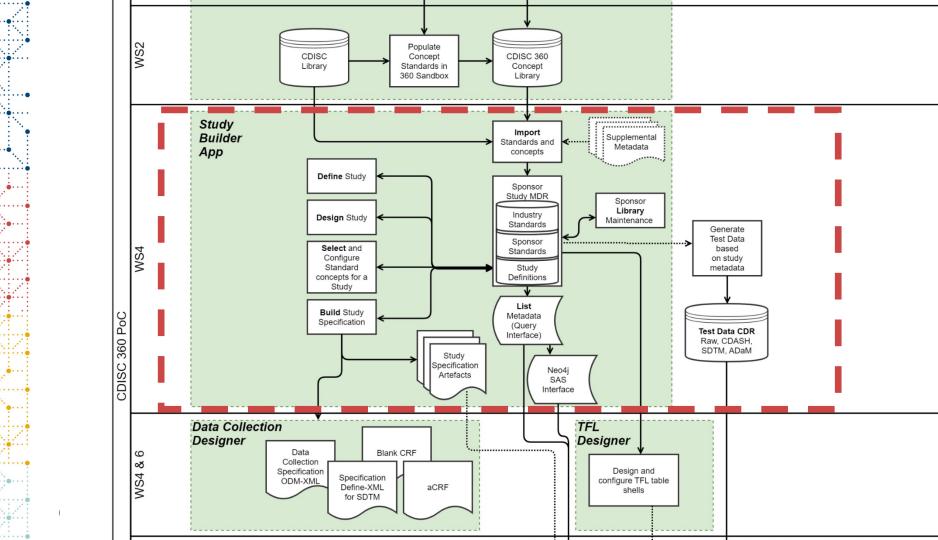
CDISC US Interchange, October 2020





### Agenda

- General Introduction to Study Builder App & MDR
  - Demo
    - Define Design Select Build
    - List and interface study metadata
  - BC's for Activities and Assessment
    - Linked Graph Data Model
  - API for Sponsor Study MDR
  - Neo4j to SAS Interface



#### Key features in the Study Designer App

#### Library

Import definitions from external libraries.

Manage sponsor defined selections and definitions. **Define** Identifiers and a general set of trial summary parameters for the study

**Design** Study design parameters as well as defining study arms, elements, epochs and visits

**Select** Search and select concept-based standards and define schedule of activities and assessments

**Build** Generate study specification artefacts that support automation of study setup and execution

#### List

Extract study metadata in tabular format for down stream usage.

From within the App with export to multiple format as well as direct from SAS.



#### **Import Concept Based Standards**

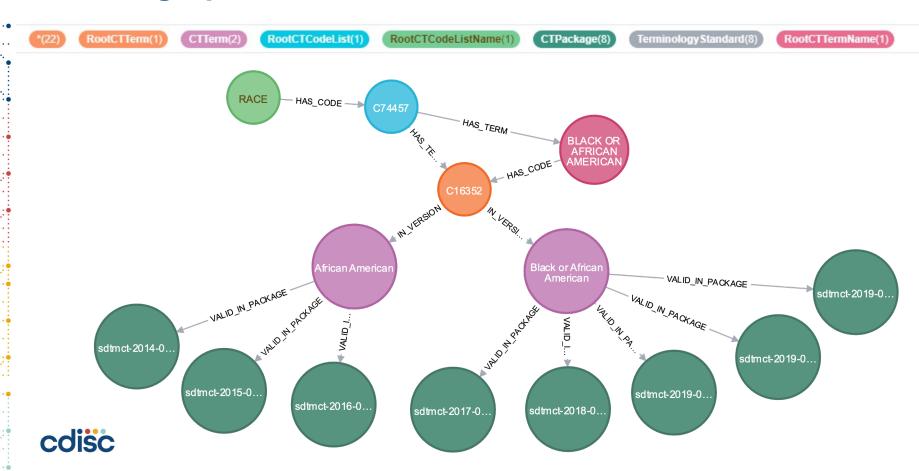
- Currently the Concept Based Standards are imported as a combination of data from
  - Current CDISC Library
  - Supplemental Metadata

- This is done in Cypher program scripts loading data into the Neo4j based Study Metadata Library
- Each CT term is stored once and Neo4j enable version tracking over time

"https://library.cdisc.org/api/mdr/ct/packages"

```
// Load Scope of CT packages
CALL apoc.load.jsonParams("https://library.cdisc.org/api/mdr/ct/packages",{Authorization:
   "Basic Y2xxx", Accept: "application/json"}, null) YIELD value AS link
UNWIND link._links.packages AS package
WITH DISTINCT SPLIT(package.title, ' ')[0] AS model
MERGE (mdl:Model {name: model})
RETURN mdl.name;
```

#### Linked graph domain model for CDISC CT





#### Demo

Library – Import and manage sponsor standards

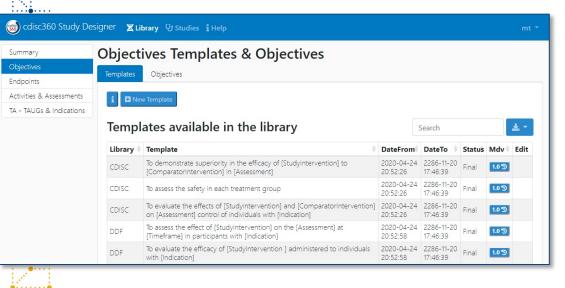
Define – Design – Select – Build

List and interface study metadata

How do you work with a 360 enabled Sponsor Study MDR



#### **Study Designer App - Library**

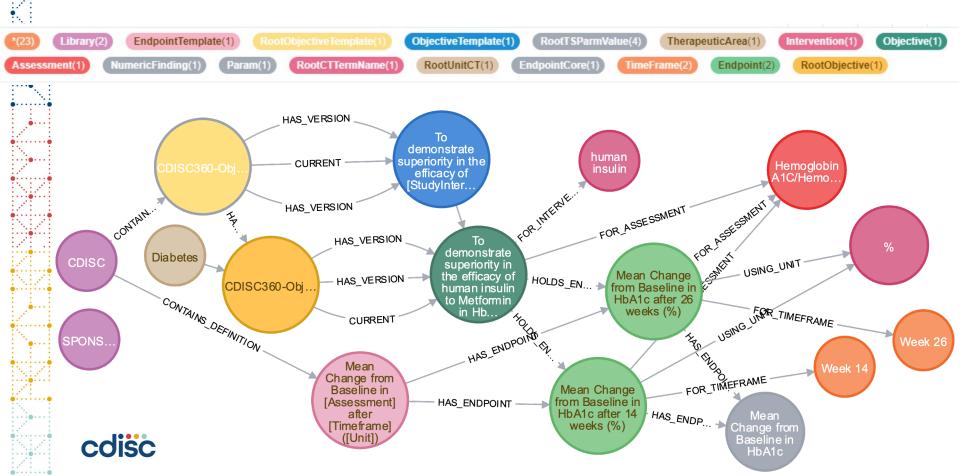


On the **Library** menu the user

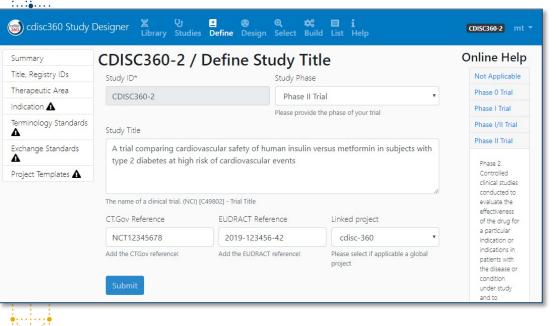
- Create additional templates for Objectives and Endpoints
- Create instantiations of imported or sponsor defined templates
- Instances of Objectives and Endpoints include reference to dependent parameters



#### Linked graph domain model for Library



#### **Study Designer App - Define**

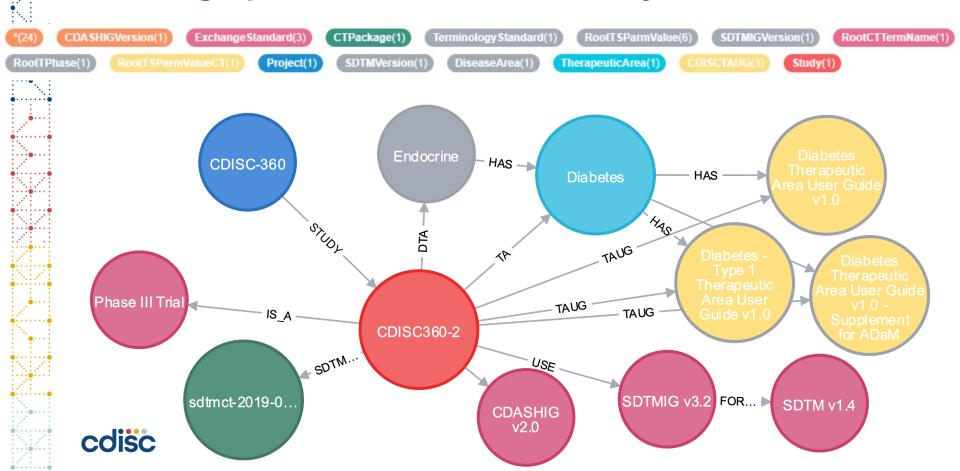


On the **Define** menu the user

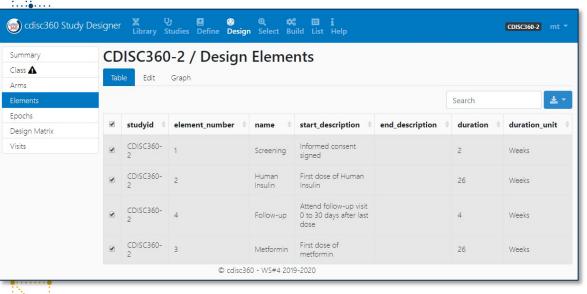
- Enter the basic description of the trial like the study phase, title, registry identifiers
- Therapeutic Area of the study and CDISC TAUGs used
- Version of terminology standards
- Version of exchange standards



#### Linked graph domain model for Study Define



#### **Study Designer App - Design**

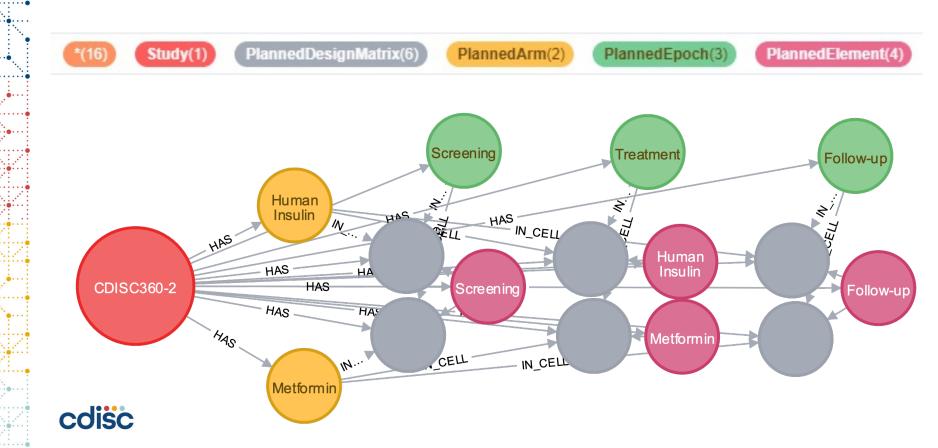


On the **Design** menu the user

- Make basic selection of trial design related trial summary parameters like Intervention Type, Intervention Model etc.
- Define the Trial Arms, Epochs, Elements and the Design matrix
- Define the visit schedule
- Define the planned interventions



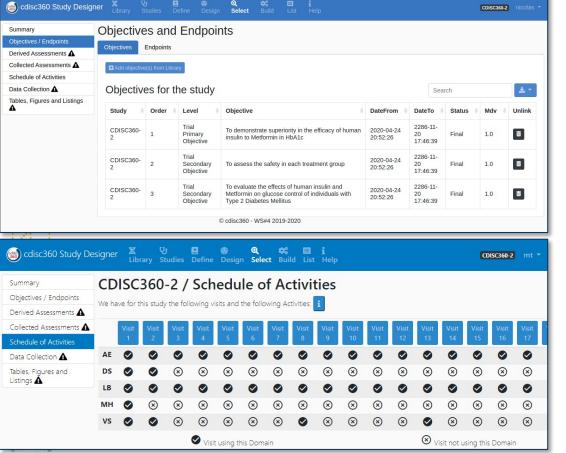
#### Linked graph domain model for Study Design





#### **Study Designer App - Select**

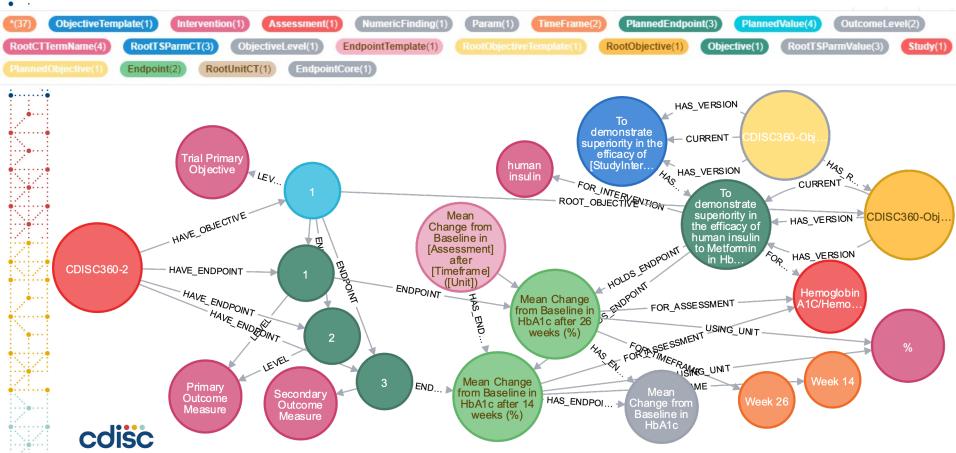
CDISC360-2



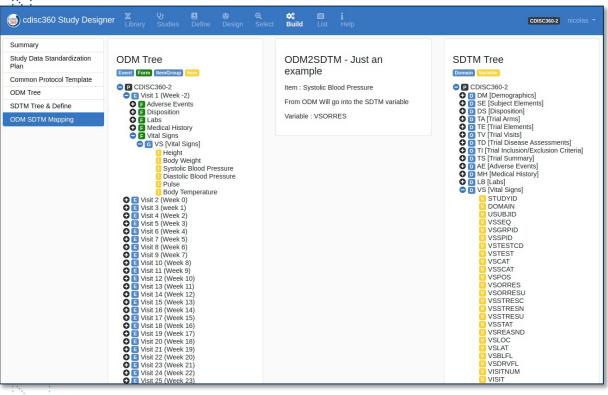
#### On the **Select** menu the user

- Selects the concept based standards from the libraries that are to be used in the study
  - These can be based on templates that are instantiated in the local library
- Objectives and Endpoints
- Activities and Assessments
- Schedule of Activities and **Assessments**
- TFL metadata

#### Linked graph domain model for Study Select



#### **Study Designer App - Build**



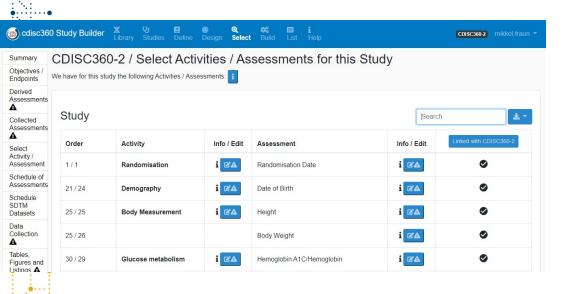
On the **Build** menu the user can generate:

- Study data standards plan
- Protocol metadata report
  - To be copy paste into CPT
  - As XML to be imported into eCPT
  - As tables that can be exported
- Data collection specification
  - ODM-XML
  - Blank CRF, techCRF and aCRF
- Tabulation Specification
  - Define-XML specification
- Analysis Specification





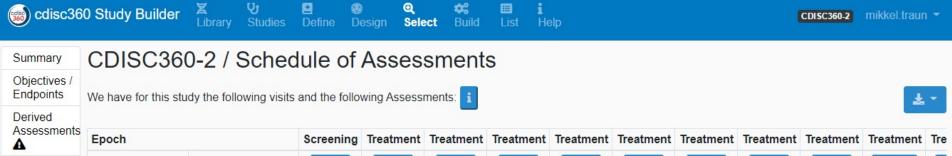
#### **Study Designer App – Select Activities & Assessments**



On the **Select** menu the user

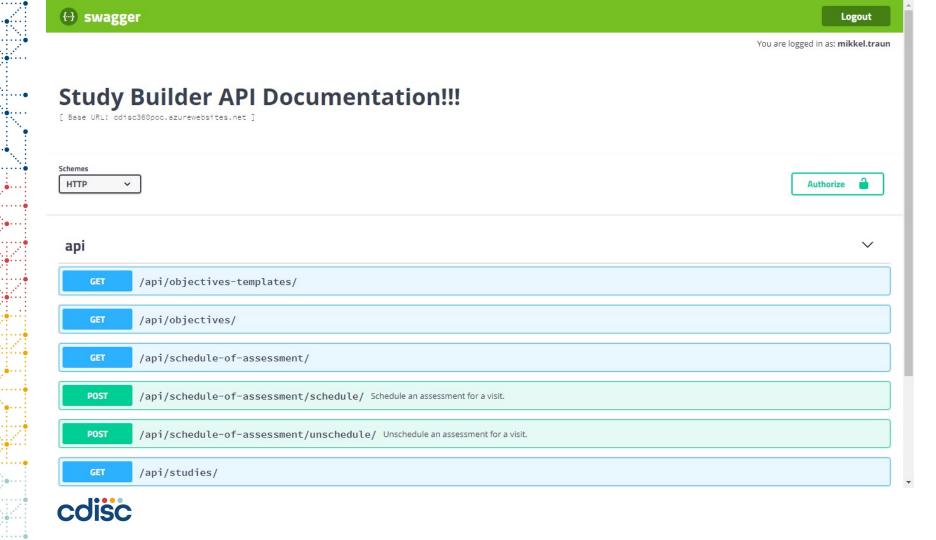
- Selects BC's in the form of Activities and Assessments
- Configure these in context of the study
- Schedule the Activities and Assessments in Study Design



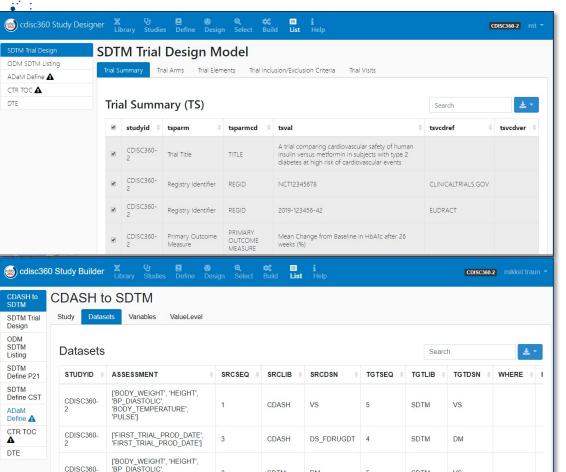


A	Epoch		Screening	Treatment	Treatment	Treatment	Treatment	Treatment	Treatment	Treatment	Treatment	Treatment	Tre
Collected	Activity	Assessment	Visit 1	Visit 2	Visit 3	Visit 4	Visit 5	Visit 6	Visit 7	Visit 8	Visit 9	Visit 10	V
Assessments  A	Randomisation	Randomisation Date	(X)	•	⊗	<b>⊗</b>	⊗	<b>⊗</b>	<b>×</b>	<b>⊗</b>	⊗	<b>×</b>	
Select Activity / Assessment Schedule of	Demography	Date of Birth	_	<b>×</b>	<b>⊗</b>	<b>(X)</b>	<b>⊗</b>	<b>×</b>	<b>⊗</b>	<b>×</b>	<b>⊗</b>	<b>×</b>	
	Vital signs	Systolic Blood Pressure	~	0	<b>⊗</b>	<b>×</b>	<b>⊗</b>	<b>⊗</b>	<b>×</b>	<b>⊗</b>	<b>⊗</b>	<b>×</b>	
Assessments Schedule SDTM Datasets		Diastolic Blood Pressure		<b>Ø</b>	<b>⊗</b>	<b>⊗</b>	<b>×</b>	<b>⊗</b>	×	<b>×</b>	<b>⊗</b>	×	
		Pulse	_	•	⊗	⊗	<b>⊗</b>	⊗	<b>⊗</b>	⊗	⊗	<b>⊗</b>	
Data  Collection		Body Temperature	•	•	⊗	⊗	<b>⊗</b>	$\otimes$	<b>⊗</b>	$\otimes$	⊗	<b>⊗</b>	
	Glucose metabolism	Hemoglobin A1C/Hemoglobin		•	•	<b>Ø</b>	<b>⊗</b>	⊗	<b>⊗</b>	⊗	•	⊗	
Tables, Figures and		Glucose, Plasma	_	<b>⊗</b>	<b>Ø</b>	<b>Ø</b>	<b>Ø</b>	<b>Ø</b>	<b>Ø</b>	<b>Ø</b>	<b>Ø</b>	•	





## **Study Designer App - List**



On the **List** menu the user can generate:

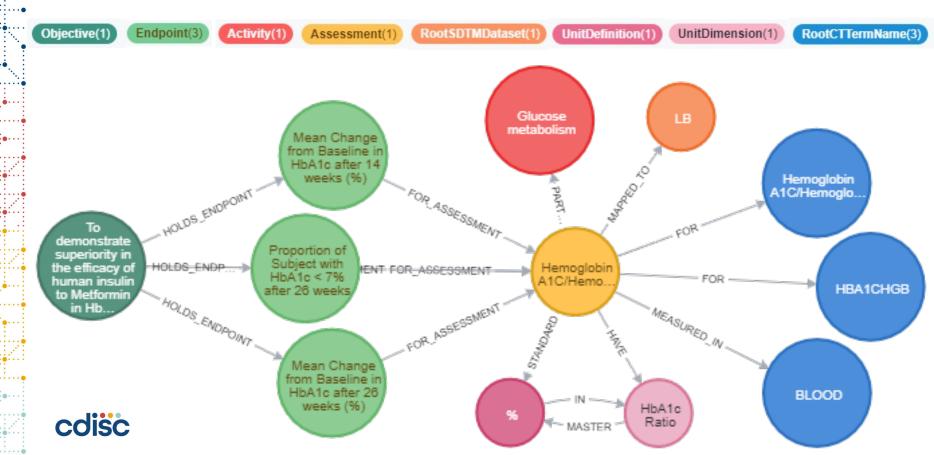
- Browse all study metadata in tabular form
- Export these into various file formats
- Will correspond to the SAS based interface to the Study Metadata Library enabling extract of study metadata into SAS datasets
- This include CDASH2SDTM and SDTM2ADaM Bindings



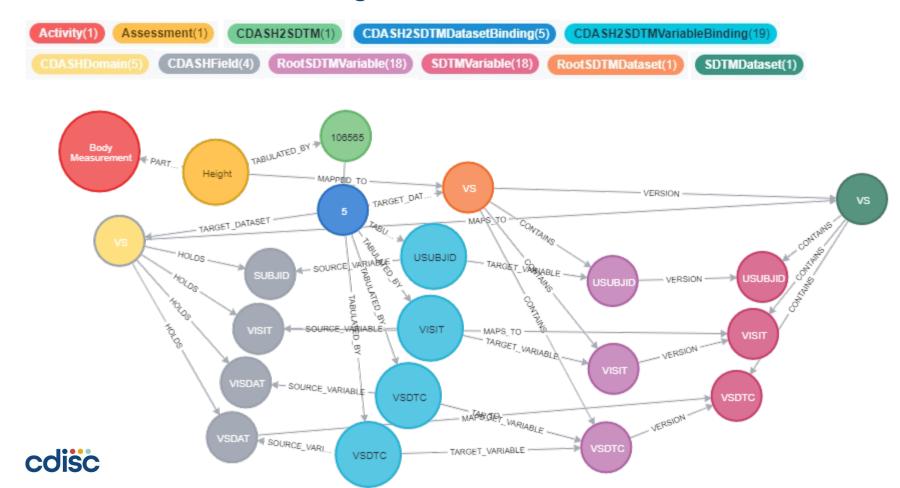
#### **BC's for Activities and Assessment**

Linked Graph Data Model

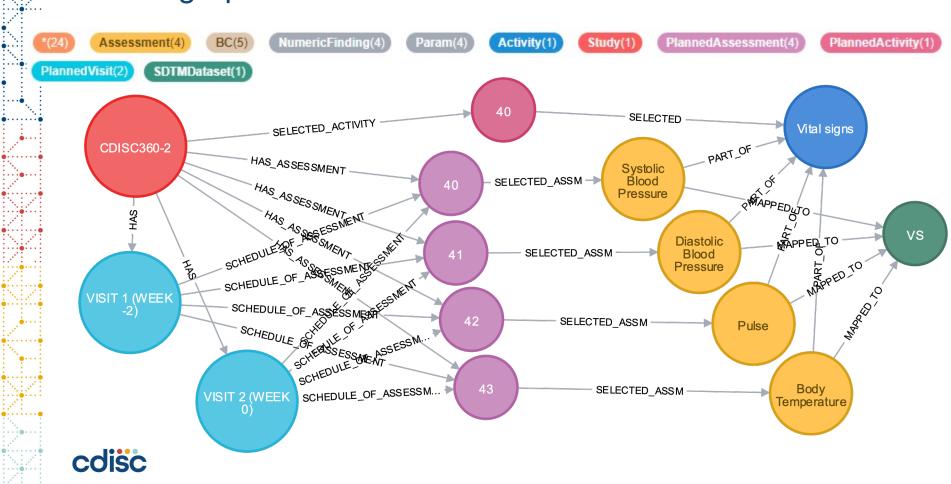
# Linked graph domain model for Activities and Assessments The Concept Definition



#### The CDISC 360 Bindings from CDASH -> SDTM

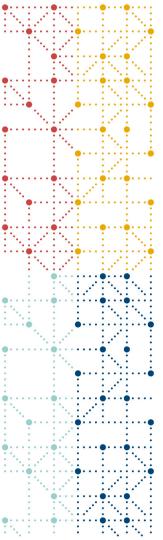


#### Linked graph domain model for Schedule of Assessment



#### Sample Cypher Query Schedule of Assessments

	study_id	activity	assessment	visitnum	pa.order	pm.order
1	"CDISC360-2"	"Vital signs"	"Systolic Blood Pressure"	100	40	40
2	"CDISC360-2"	"Vital signs"	"Diastolic Blood Pressure"	100	40	41
3	"CDISC360-2"	"Vital signs"	"Pulse"	100	40	42
4	"CDISC360-2"	"Vital signs"	"Body Temperature"	100	40	43
5	"CDISC360-2"	"Vital signs"	"Systolic Blood Pressure"	200	40	40

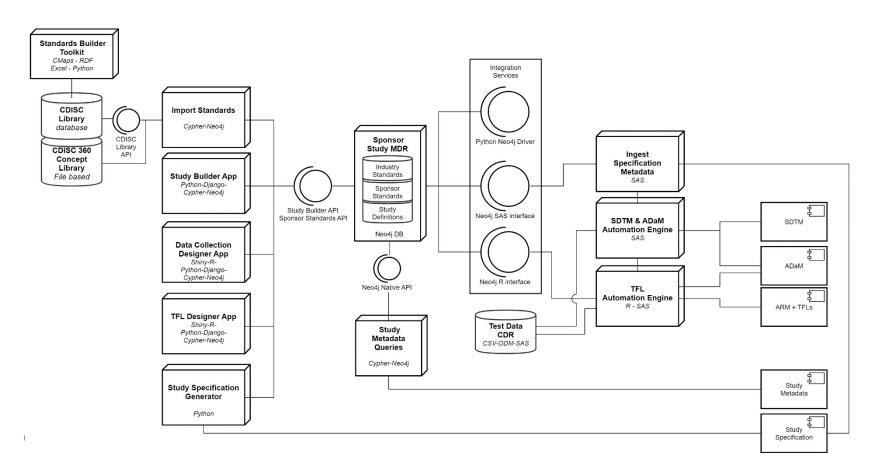


#### **API for Sponsor Study MDR**

PoC implementation of API endpoints

#### API for Sponsor MDR managing the Study Metadata

This component diagram will be described in the CDISC 360 PoC Yellow-paper





#### Neo4j to SAS Interface

All metadata in Sponsor Study MDR Neo4j database can be access directly from SAS

### Access Neo4j Sponsor Study MDR from SAS

- The Neo4j transactional HTTP endpoint allows you to execute Cypher statements
- Using SAS PROC LUA to easily interface with the Neo4j REST API from SAS
- Neo4j to SAS Interface manage SAS Dataset metadata

["FIRST TRIAL PROD DATE".

"FIRST TRIAL PROD DATE"1

ASSESSMENT	SRCSEQ	SRCLIB	SRCDSN	SRCVAR	SRCTYPE	ORIGIN	METHOD	CODELIST	TGTLIB
["BODY_WEIGHT", "HEIGHT", "BP_DIASTOLIC", "BODY_TEMPERATURE", "PULSE"]	1	CDASH	VS	SUBJID	text	Assigned	ALL.USUBJID		SDTM
["BODY_WEIGHT", "HEIGHT", "BP_DIASTOLIC", "BODY_TEMPERATURE", "PULSE"]	1	CDASH	VS	VISIT	text	Predecessor		VISIT	SDTM
["BODY_WEIGHT", "HEIGHT", "BP_DIASTOLIC", "BODY_TEMPERATURE", "PULSE"]	1	CDASH	VS	VSDAT	text	Assigned	VS.VSDTC		SDTM
["BODY_WEIGHT", "HEIGHT", "BP_DIASTOLIC", "BODY_TEMPERATURE", "PULSE"]	1	CDASH	VS	VISDAT	text	Assigned	VS.VSDTC		SDTM
["BODY_WEIGHT", "HEIGHT", "BP_DIASTOLIC", "BODY_TEMPERATURE", "PULSE"]	1	CDASH	DM	SUBJID	text	Assigned	ALL.USUBJID		SDTM
["BODY_WEIGHT", "HEIGHT", "BP_DIASTOLIC", "BODY_TEMPERATURE", "PULSE"]	1	CDASH	DM	RACE	text	Predecessor		RACE	SDTM

DS FDRUGDT DSSTDAT

SDTM



#### **Reflections from WS4**

What have we accomplished in the cdisc360 PoC

#### Machine-readable Study Specification Metadata

#### Study Design

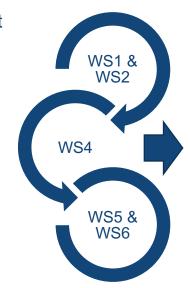
 Implemented basic Study Definition and Design in Study Repository linked to Schedule of Assessment

#### CDASH & SDTM

- Implemented Assessment BC's in a Label Property Graph Model linked with versioned metadata from the CDISC Library
- Including sample CDASH2SDTM bindings

#### ADaM

- Implemented basic ADaM Sponsor model in a Label Property Graph Model linked with versioned metadata from the CDISC Library
- Including sample SDTM2ADaM bindings
- Study Builder & Sponsor Study MDR
  - Standard API based to enable tool and vendor agnostic system integrations







# **Thank You!**

Mikkel Traun, *Novo Nordisk*Nicolas de Saint Jorre, *XClinical* 

