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Automate harmonization of study metadata for integrated define.xml





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Title: Statistical Programming Specialist

Organization: Novo Nordisk A/S

Nynne Storm joined Novo Nordisk A/S in 2018 and has worked as a Statistical Programmer on and across various projects and trials. Currently focusing on pooling of data for submission. She has a PhD from the Technical University of Denmark within the field of Geomagnetism.



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Matthew Phelps joined Novo Nordisk A/S to work on the newly formed Data Science Automation Team as a Statistical Programming Specialist in April 2022. Previously he worked as an epidemiologist at the Danish Heart foundation doing registry-based research. In addition to research, he had a lot of fun programming in R and developing public-facing Shiny apps.



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 or Novo Nordisk.
- The authors have no real or apparent conflicts of interest to report.





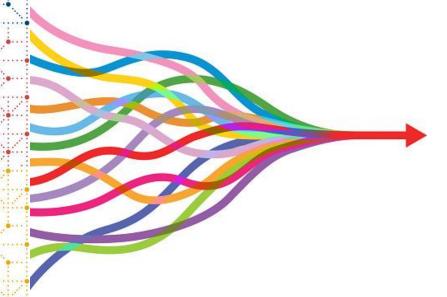
Agenda

- 1. Introduction
- 2. Challenges
- 3. Ambitions overview
- 4. Zoom in with features
- 5. Future

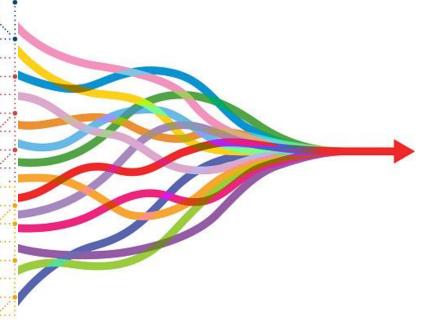
Introduction





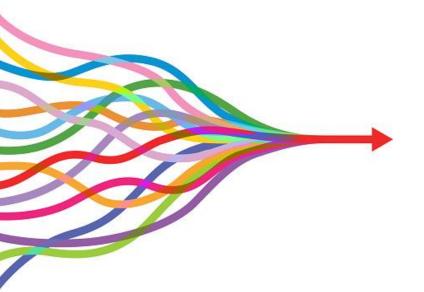






- Integrated database for submission
- Safety update database
- Exploratory database
- Etc.





- Integrated database for submission
- Safety update database
- Exploratory database
- Etc.

• Time consuming and tedious



Introduction

CST document - containing metadata about ADaM datasets and their variables, including CDISC ADaM IG information.

Dataset	Label	Class		
ADAE	Adverse Events Analysis	Occurence Data Structure	 	
ADSL	Subject-Level Analysis Dataset	Subject Level Analysis Dataset	 	



Introduction

CST document - containing metadata about ADaM datasets and their variables, including CDISC ADaM IG information.

Dataset	Label	Class		
ADAE	Adverse Events Analysis	Occurence Data Structure	 	
ADSL	Subject-Level Analysis Dataset	Subject Level Analysis Dataset	 	

Dataset	Variable	Label	Core	Origin	Description	
ADAE	USUBJID	Unique Subject Identifier	Req	Predecessor	A E. USUBJID	
ADAE	APERIOD	Period	Perm	Derived	xxx	

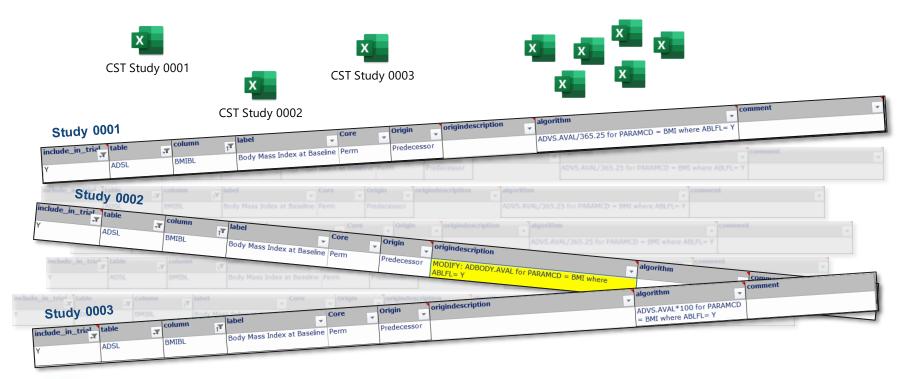




Study 0001

Study 0001		label	origin vigindesex-p	ADVS.AVAL/365.25 for PARAMCD = BMI with	nere ABLFL= 1
include_in_trial_table	column	Body Mass Index at Baseline Perm	Ddocessor		
ADSL	BMIBL	body 1122			





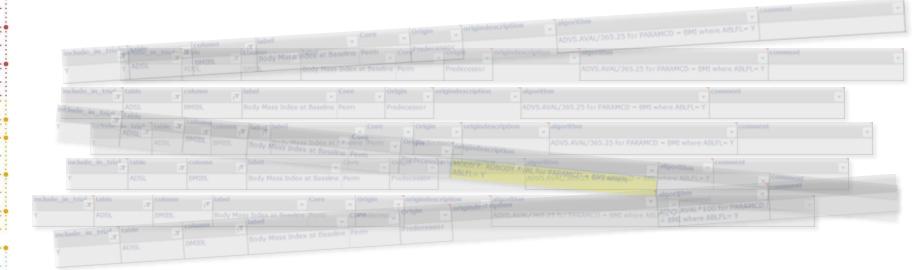
































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Only focuses on variables that are differently derived between studies











Only focuses on variables that are differently derived between studies

	trial 🍦	origindescription	$\stackrel{\triangle}{\triangledown}$	algorithm			
ADSL-BMIBL							
1	0001	blank		ADVS.AVAL*100 for PARAMCD = BMI where ABLFL= Υ			
2	0002	blank		ADVS.AVAL*100 for PARAMCD = BMI where ABLFL= Y			
3	0003	blank		ADVS.AVAL / 100 for PARAMCD = BMI where ABLFL= Υ			
Showing 1 to 3 of 3 entries							



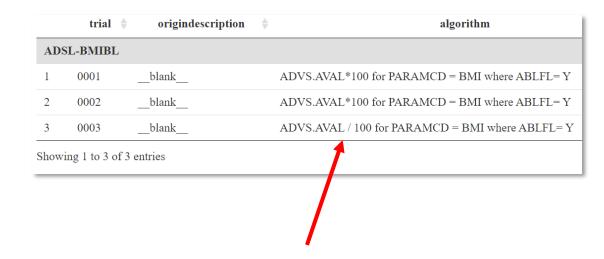




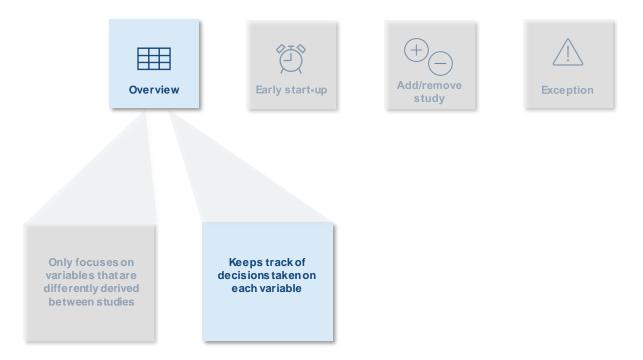




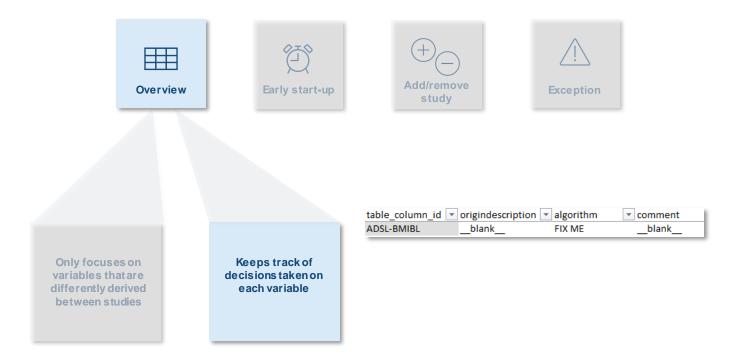
Only focuses on variables that are differently derived between studies



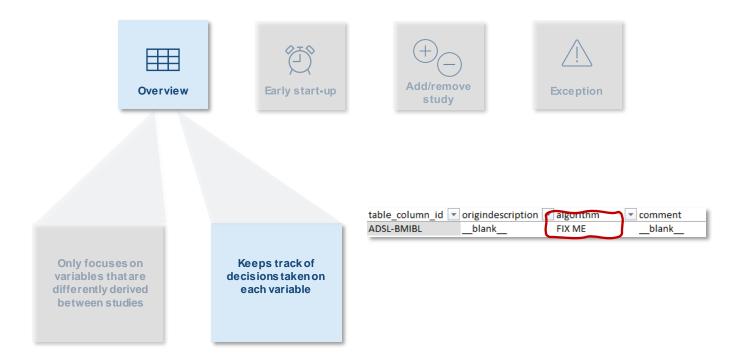




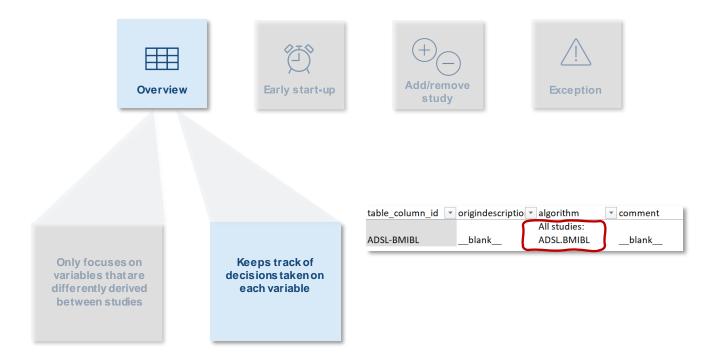


























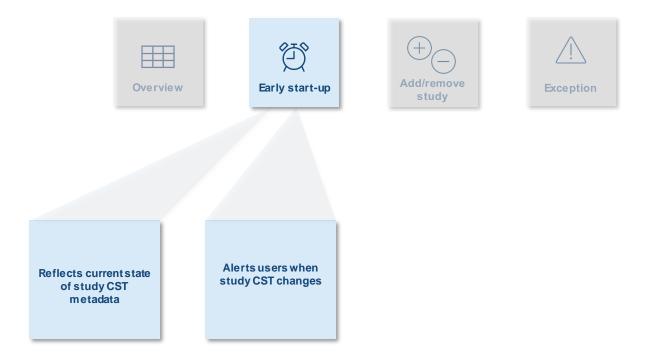




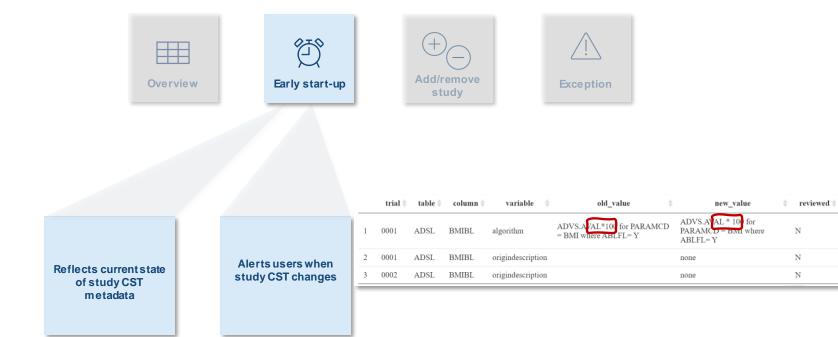


Reflects current state of study CST metadata

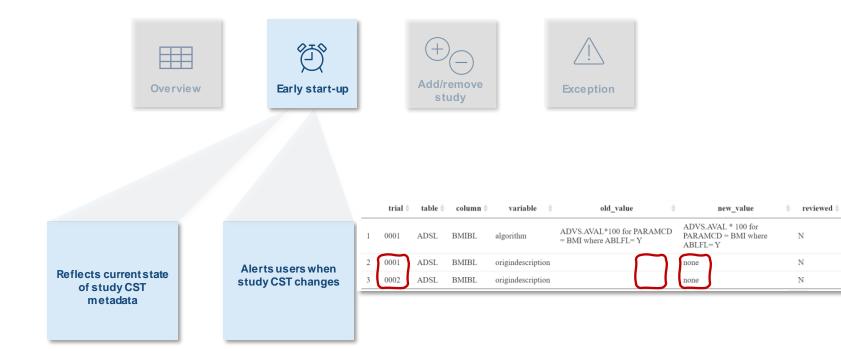




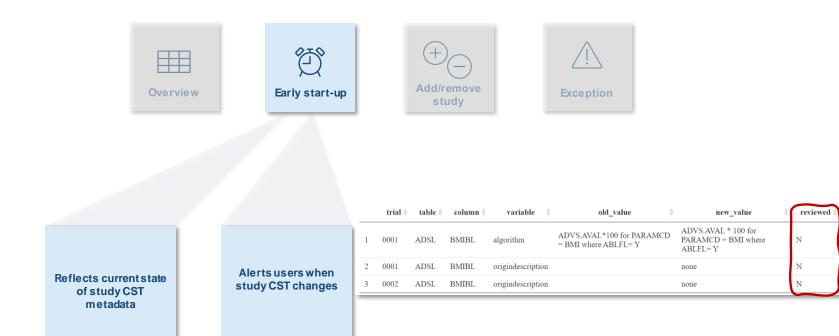
































New studies can easily be added





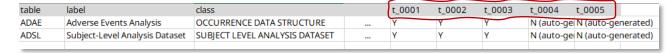






New studies can easily be added

trial	instance	stack
0001	current	Y
0002	csr_er	Υ
0003	current	Υ
0004	current	N
0005	current	N













New studies can easily be added

Studies to be included can be toggled on/off when ever needed

		$\overline{}$	١.
trial	instance	stack	
0001	current	Υ	
0002	csr_er	Υ	Г
0003	current	Υ	١
0004	current	N	ı
0005	current	N	L
_)























Specific exception in description for a study



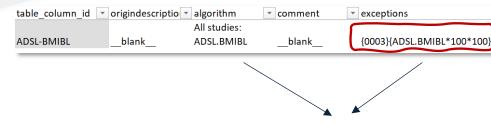








Specific exception in description for a study

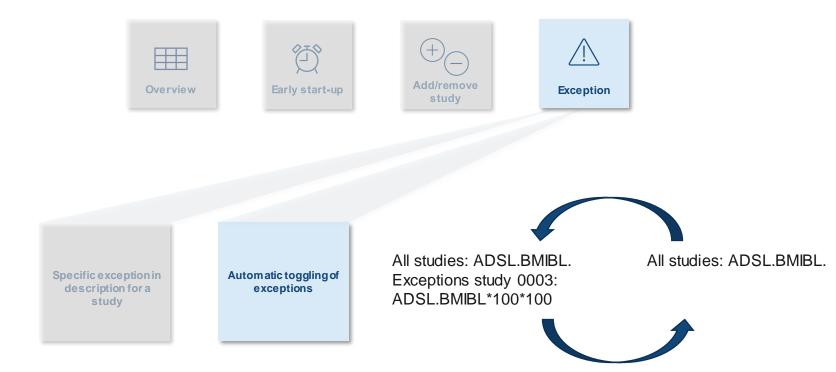


All studies: ADSL.BMIBL. Exceptions study 0003: ADSL.BMIBL*100*100



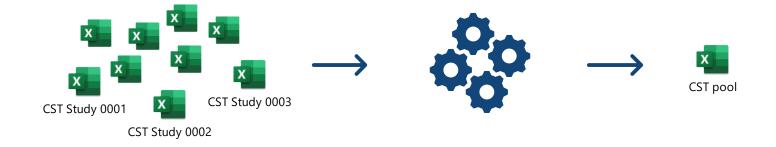
▼ stacked or pooled ▼

stacked





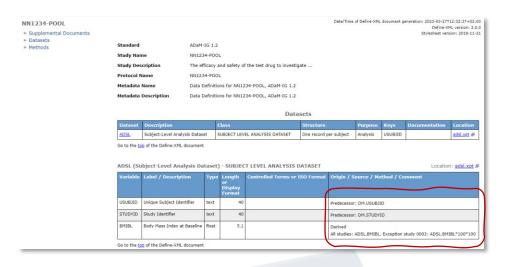
Output

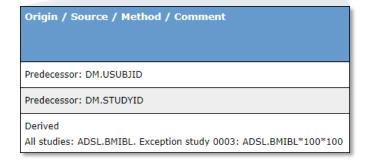




Output

 CST file with the descriptions for the pooled database – input for define.xml







Output

 CST file with the descriptions for the pooled database – input for define.xml

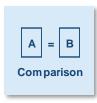
 Stacked study data – ready for data wrangling for final ADaM pool





Future

 Compare variables that are the same, but are named differently in the studies



• Compare studies across projects



"Smart" comparison of description text





Learnings

 Only doable due to standardization (CDISC & Novo Nordisk)

Work iteratively

Starting early > starting later



Thank You!

