

Tuesday, April 19, 2016, 1–2 PM Eastern

Hosted by Abel Kho, MD

Facilitated by Shelley Rusincovitch and Michelle Smerek



Agenda

- Welcome and announcements
- Review of the GitHub CDM issue tracker and errata
- Recent work with the lab result reference table
- Discussion of the CONDITION table and "PCORnet-defined cohort algorithm" category
- Wrap up



Announcements



Upcoming data characterization office hours

Why it may be of interest to this group:

 Offers an opportunity for in-depth discussion with the team about your questions

April 20, 2-3 ET

1-650-479-3207 Call-in toll number (US/Canada)

Access code: 738 585 443

April 26, 2-3 ET

1-650-479-3207 Call-in toll number (US/Canada)

Access code: 734 755 577

May 6, 12-1 ET

1-650-479-3207 Call-in toll number (US/Canada)

Access code: 736 965 276



Research Using a DRN Forum on April 25

Why it may be of interest to this group:

- This is a new forum with focus on Distributed Research Network (DRN) research methods
- Topic for April 25: Flexible Analytic Tool Development: A SAS based perspective, presented by Jennifer Popovic

Research Using a DRN Forum Monday April 25th, 11am-12pm ET Online:

https://dukemed.webex.com/dukemed/j.php?MTID=mca9d81877417a4558450 558d80cc5e22

Phone: 1-855-244-8681 / Access code: 736 510 205



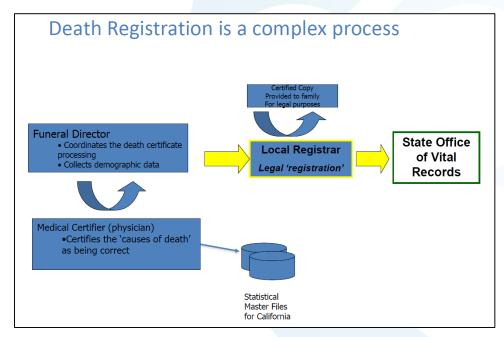
Recap: CDM Forum from March 14

Why it may be of interest to this group:

- Mike Hogarth presented on dearth event registration and mortality data
- Session prompted considerable interest and attention, even a request to rebroadcast recording with internal team

Recording, slides, and summary:

https://pcornet.imeetcentra I.com/p/ZgAAAAAACTEB





Recap: AMIA presentation on CDM v3.0

Why it may be of interest to this group:

 Shelley Rusincovitch presented on the Common Data Model v3.0 design at the AMIA Joint Summits in March

Slides and abstract:

https://github.com/CDMFO RUM/CDM-GUIDANCE/wiki/CDMrelated-Abstracts

Iterative Design and Modeling for the PCORnet Common Data Model v3.0

Shelley A. Rusincovitch, Jeffrey S. Brown, PhD, Keith Marsolo, PhD, Jenny Ibarra, RN, MSN, Abel N. Kho, MD, MS, Daniella Meeker, PhD, Lesley H. Curtis, PhD

S36: Infrastructure for Data-driven Translational Science **Thursday, March 24, 2016**







Recap: AMIA presentation on lab mapping strategies

Why it may be of interest to this group:

 Michelle Smerek presented at the AMIA Joint Summits in March; a product of the Lab Mappings Interest Group; includes survey results

Slides and abstract:

https://github.com/CDMFO RUM/CDM-GUIDANCE/wiki/CDMrelated-Abstracts

Factors and Approaches to Mapping Laboratory Results in PCORnet

Michelle M. Smerek, Elisa Priest, Dr.PH, S. Trent Rosenbloom, MD, MPH, FACMI,

Jon E. Puro, MPA, Pedro Rivera, MSCS, Shelley A. Rusincovitch, Rahul Jain, MPH, CPHIMS, and Keith Marsolo, PhD

AMIA 2016 Joint Summits on Translational Science March 21-24, 2016, San Francisco, California





GitHub and Issue Tracker Review

Some content repeated from DRN OC Forum on March 7, 2016



What is GitHub?

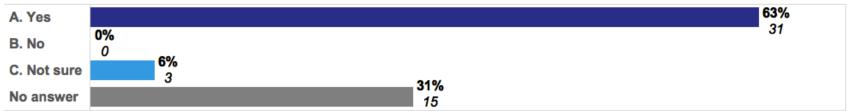
- Web-based repositories for storing software program code
- Largest platform of its kind with >12 million people, >31 million projects (https://github.com/about)
- Often used for open-source projects
- Used by initiatives such as:
 - openFDA: https://github.com/FDA/openfda
 - precisionFDA: https://github.com/FDA/precisionFDA
 - OHDSI: https://github.com/OHDSI
- Used by several PCORnet-related groups, including:
 - LHSNet: https://github.com/LHSNet
 - SCILHS: https://github.com/SCILHS
 - PEDSnet: https://github.com/PEDSnet



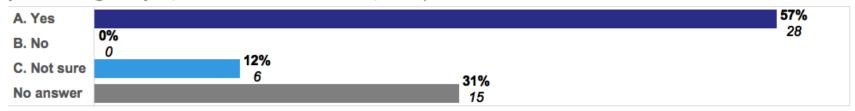


Poll results from CDM Forum on November 11, 2015

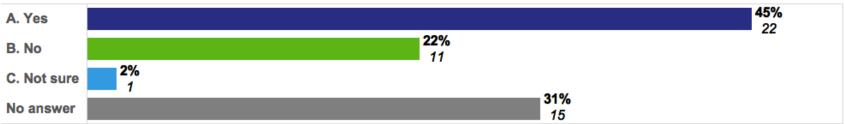
1. Would you (or your team) potentially have interest in USING or IMPROVING UPON PCORnet-related code shared by others? ("Code" might include, but is not limited to, database scripts, processing scripts, client-based utilities, etc.)



2. Would you (or your team) potentially have interest in SHARING PCORnet-related code from your team with others? ("Code" might include, but is not limited to, database scripts, processing scripts, client-based utilities, etc.)



3. Do you (or your team) have experience with a code repository tool (like GitHub or BitBucket)?



Summary of GitHub progression

3 phases of development and soft launches of GitHub repositories by the DRN OC:

(Planning stage: Discussion and assessment, primarily through the CDM Forum in fall 2015)

- 1. ADAPTABLE base phenotype (December 2015)
- 2. CDM Errata (January 2016)
- 3. Diagnostic Query and Data Characterization packages (March 2016)



Using the GitHub issue tracker functionality

- These areas are often brought to our attention by data partners, with many thanks
 - Multiple mechanisms include the CDM forum, CDM forum interest groups, data characterization office hours, and site liaison activity
- For the Common Data Model, a separation between errata (errors and typos in the current specification) and non-errata (guidance and potential enhancements)



Issue Tracker review (live)

CDM errata issue tracker:

https://github.com/CDMFORUM/CDM-ERRATA/issues

CDM guidance issue tracker:

https://github.com/CDMFORUM/CDM-GUIDANCE/issues



Future thoughts

- Many different PCORnet-related resources exist
 - Already a number of repositories, and expected to grow
 - Different organizations represented, and many desire to brand and manage their own resources
 - A challenge will be connecting and indexing this "web of resources"
- We look forward to the Data Committee's strategic assessment and recommendations
- Your suggestions and thoughts are appreciated!



Lab Result Reference Table

Michelle Smerek



From presentation at AMIA TBI/CRI Summit: March 2016

DEMOGRAPHIC

PATID
BIRTH_DATE
BIRTH_TIME
SEX
HISPANIC
RACE
BIOBANK FLAG

Fundamental basis

ENROLLMENT

PATID
ENR_START_DATE
ENR_END_DATE
CHART
ENR BASIS

DISPENSING

DISPENSINGID
PATID
PRESCRIBINGID (optional)
DISPENSE_DATE
NDC
DISPENSE_SUP

DISPENSE AMT

DEATH

PATID
DEATH_DATE
DEATH_DATE_IMPUTE
DEATH_SOURCE
DEATH MATCH CONFIDENCE

DEATH CAUSE

PATID
DEATH_CAUSE
DEATH_CAUSE_CODE
DEATH_CAUSE_TYPE
DEATH_CAUSE_SOURCE
DEATH_CAUSE_CONFIDENCE

Data captured from processes associated with healthcare delivery

http://www.pcornet.org/pcornet-common-data-model/

PCORnet Common Data Model v3.0

VITAL

VITALID

PATID
ENCOUNTERID (optional)
MEASURE_DATE
MEASURE_TIME
VITAL_SOURCE
HT
WT
DIASTOLIC
SYSTOLIC
ORIGINAL_BMI
BP_POSITION
SMOKING
TOBACCO
TOBACCO TYPE

CONDITION

CONDITIONID

PATID
ENCOUNTERID (optional)
REPORT_DATE
RESOLVE_DATE
ONSET_DATE
CONDITION_STATUS
CONDITION

PRO CM

PRO CM ID

CONDITION TYPE

PATID
ENCOUNTERID (optional)
PRO_ITEM
PRO_LOINC
PRO_DATE
PRO_TIME
PRO_RESPONSE
PRO_METHOD
PRO_MODE
PRO_CAT

Data captured within multiple contexts: healthcare delivery, registry activity, or directly from patients

ENCOUNTER

ENCOUNTERID
PATID
ADMIT_DATE
ADMIT_TIME
DISCHARGE_DATE
DISCHARGE_TIME
PROVIDERID
FACILITY_LOCATION
ENC_TYPE
FACILITYID
DISCHARGE_DISPOSIT

DISCHARGE_DISPOSITION DISCHARGE_STATUS DRG

DRG_TYPE

ADMITTING_SOURCE

DIAGNOSIS

DIAGNOSISID

PATID

ENCOUNTERID
ENC_TYPE (replicated)
ADMIT_DATE (replicated)
PROVIDERID (replicated)
DX
DX_TYPE
DX_SOURCE
PDX

PROCEDURES

PROCEDURESID

PATID
ENCOUNTERID
ENC_TYPE (replicated)
ADMIT_DATE (replicated)
PROVIDERID (replicated)
PX_DATE
PX
PX_TYPE
PX_SOURCE

LAB_RESULT_CM LAB_RESULT_CM_ID

PATID

ENCOUNTERID (optional) LAB NAME SPECIMEN SOURCE LAB LOINC PRIORITY RESULT LOC LAB PX LAB PX TYPE LAB ORDER DATE SPECIMEN_DATE SPECIMEN TIME RESULT_DATE RESULT TIME RESULT QUAL RESULT_NUM RESULT MODIFIER RESULT UNIT NORM_RANGE_LOW NORM MODIFIER LOW NORM RANGE HIGH NORM_MODIFIER_HIGH ABN IND

PRESCRIBING

PRESCRIBINGID
PATID
ENCOUNTERID (optional)
RX_PROVIDERID
RX_ORDER_DATE
RX_ORDER_TIME
RX_START_DATE
RX_END_DATE
RX_QUANTITY
RX_REFILLS
RX_DAYS_SUPPLY
RX_FREQUENCY
RX_BASIS
RXNORM CUI

PCORNET TRIAL

PATID
TRIALID
PARTICIPANTID
TRIAL_SITEID
TRIAL_ENROLL_DATE
TRIAL_END_DATE
TRIAL_WITHDRAW_DATE
TRIAL_INVITE_CODE

Associations with PCORnet clinical trials

HARVEST

NETWORKID NETWORK NAME DATAMARTID DATAMART_NAME DATAMART PLATFORM CDM VERSION DATAMART_CLAIMS DATAMART EHR BIRTH DATE MGMT ENR START DATE MGMT ENR_END_DATE_MGMT ADMIT DATE MGMT DISCHARGE DATE MGMT PX_DATE_MGMT RX_ORDER_DATE_MGMT RX_START_DATE_MGMT RX_END_DATE_MGMT DISPENSE DATE MGMT LAB_ORDER_DATE_MGMT SPECIMEN_DATE_MGMT RESULT DATE MGMT MEASURE_DATE_MGMT ONSET_DATE_MGMT REPORT DATE MGMT RESOLVE_DATE_MGMT PRO DATE MGMT REFRESH DEMOGRAPHIC DATE REFRESH ENROLLMENT DATE REFRESH ENCOUNTER DATE REFRESH_DIAGNOSIS_DATE REFRESH PROCEDURES DATE REFRESH VITAL DATE REFRESH_DISPENSING_DATE REFRESH LAB RESULT CM DATE REFRESH_CONDITION_DATE REFRESH PRO CM DATE REFRESH_PRESCRIBING_DATE REFRESH PCORNET TRIAL DATE REFRESH DEATH DATE REFRESH DEATH CAUSE DATE

Process-related data

Data captured from healthcare delivery, direct encounter basis





LAB_RESULT_CM: contents

LAB_RESULT_CM

LAB_RESULT_CM_ID
PATID

ENCOUNTERID (optional)

LAB NAME

SPECIMEN SOURCE

LAB LOINC

PRIORITY

RESULT_LOC

LAB_PX

LAB_PX_TYPE

LAB_ORDER_DATE

SPECIMEN_DATE

SPECIMEN_TIME

RESULT_DATE

RESULT_TIME

RESULT QUAL

RESULT_NUM

RESULT_MODIFIER

RESULT_UNIT

NORM_RANGE_LOW

NORM_MODIFIER_LOW

NORM_RANGE_HIGH

NORM_MODIFIER_HIGH

ABN_IND

The "PCORnet 11"

Hemoglobin A1c

Creatine kinase total

Creatine kinase MB

Creatine kinase MB/creatine kinase total

Creatinine

Hemoglobin

Low-density lipoprotein

International normalized ratio

Troponin I cardiac

Troponin T cardiac (qualitative)

Troponin T cardiac (quantitative)





Lab result reference table 1, v1

Reference Table 1: Laboratory Results and LOINC Codes

This table is intended to be a guide and does not represent a complete list of codes for each laboratory test. Regenstrief Institute, the organization that has developed and maintains the LOINC system, has completed a partial mapping of LOINC to CPT codes. The mapping is publicly available on the US National Library of Medicine's webpage at the following link (valid as of October 15, 2014):

1	http://www.nlm	.nih.gov/research/un	nls/mapping projects	loine to ept	map.html	

	ov/research/umls/mapping projects/loinc to c		in (valia as of October 15, 2014).
LAB NAME	SPECIMEN SOURCE	LOINC	Comments
CK MB	SERUM, PLASMA, or SR PLS	2154-3	
CK MB	SERUM, PLASMA, or SR PLS	32673-6	·
CK MB	BLOOD	49551-5	•
CK MBI	SERUM, PLASMA, or SR PLS	12187-1	
CK MBI	SERUM, PLASMA, or SR_PLS	12189-7	·
CK_MBI	SERUM, PLASMA, or SR_PLS	20569-0	•
CK_MBI	SERUM, PLASMA, or SR_PLS	49136-5	Rarely used
CK_MBI	_	·	Do not use LOINC code 15049-0, as this is a ratio for CK-MM instead of CK-MB.
CREATININE	SERUM, PLASMA, or SR_PLS	14682-9	•
CREATININE	BLOOD	21232-4	
CREATININE	SERUM, PLASMA, or SR_PLS	2160-0	,
CREATININE	SERUM, PLASMA, or SR_PLS	35203-9	This code is "discouraged" (guidance added in v3.0)
CREATININE	BLOOD	38483-4	
CREATININE	SERUM, PLASMA, or SR_PLS	44784-7	·
CREATININE	SERUM	54052-6	HEDIS 2009 code.
CREATININE	BLOOD	59826-8	
HGB	BLOOD	14775-1	•
HGB	BLOOD	20509-6	
HGB	BLOOD	24360-0	HGB and HCT panel - keep only the HGB results, e.g., those with units "g/dl" instead of "%".
HGB	BLOOD	30313-1	•
HGB	BLOOD	30350-3	
HGB	BLOOD	30351-1	<u> </u>
HGB	BLOOD	30352-9	·
HGB	BLOOD	55782-7	
HGB	BLOOD	59260-0	·
HGB	BLOOD	718-7	•
LDL	SERUM, PLASMA, or SR PLS	47213-4	
INR	BLOOD	34714-6	
INR	BLOOD	46418-0	

"This table is intended to be a guide and does not represent a complete list of codes for each laboratory test."

http://www.pcornet.org/pcornet-common-data-model/



From Lab mapping Interest Group: Lab mapping survey

- Survey launched November 11, 2015
- 34 responses received

How many unique lab result names does your site have?

24%: 1,001 - 5,000

32%: 5,001 - 10,000

36%: > 10,000 unique lab result names





From lab mapping survey

How many different synonyms does your site have for "hemoglobin?"

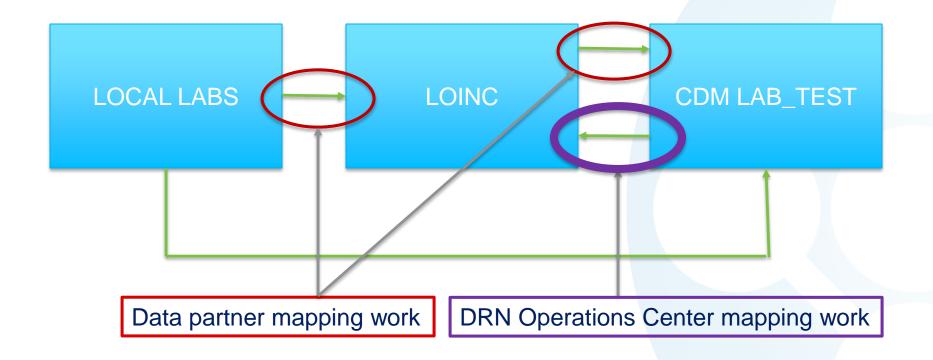
64%: > 7 synonyms for "hemoglobin" $48\% \ge 10$ synonyms





Multiple Mapping Steps

Mapping from Local Labs to CDM "common measure" labs







Lab result reference table 1, v2

This table is intended to be a dynamic resource containing LOINC codes that may map to PCORnet laboratory result name common measures.

Categorical value for PCORnet LAB_NAME variable

LOINC unique identifier for the measure

The information contained in this table can be used to search for laboratory results in local data sources that may map to PCORnet laboratory result name common measures.

Attribute that distinguishes between different kinds of quantities relating to the same substance

72563-0 Creatine kinase.MB/Creatine kinase.total [Pure catalytic fraction] in Serum or Plasma by Electrophoresis

Creatine kinase.MB/Creatine kinase.total [Ratio] in Serum or Plasma

This table is a living document that will be updated when/if necessary. We welcome feedback from PCORnet data partners regarding questions or suggestions for additional codes that should be considered for inclusion.

Rows highlighted in yellow indicate LOINC codes that may not correspond to the PCORnet lab result name common measure listed. They are included for data partners to determine whether or not they are a good fit.

The LOINC® codes, LOINC® table (regardless of format), LOINC® Release Notes, LOINC® Changes File, and LOINC® Users' Guide are copyright @ 1995-2015, Regenstrief Institute, Inc. and the Logical Observation Identifiers Names and Codes (LOINC) Committee. All rights reserved.

Timing Attribute that describes whether the measurement was for a point-in-time or time interval									
System	Type of sample the measurement was made on. This attribute consists of two subparts; the first part names the system, the optional second part, delimited with a "^", indicates the super system source of the sample if it is not the patient								
Scale	Scale of the	Scale of the measure							
Method	Method by	y which the test was performed. Methods need only be expressed as part of the name when they provide a distir	nction between tests that measure the same component (analyte) but w	which have differe	ent clinical	significance or have a differe	ant clinica	al reference ranges.	'
ExUnits	Example un								7
PCORnet LAB_NAME	LOINC code	le Category or Name	Component/analyte	Property	Timing	g System	Scale	e Method	ExUnits
	48425-3	Troponin T.cardiac [Mass/volume] in Blood	Troponin T.cardiac	MCnc	Pt	Bld	Qn		ug/L
Transpir T cardiac (quantitative)	6597-9	Troponin T.cardiac [Mass/volume] in Venous blood	Troponin T.cardiac	MCnc	Pt	BldV	Qn		ug/L
Troponin T cardiac (quantitative)	6598-7	Troponin T.cardiac [Mass/volume] in Serum or Plasma	Troponin	MCnc	Pt	Ser/Plas	Qn		ug/L
<u> </u>	67151-1	Troponin T.cardiac [Mass/volume] in Serum or Plasma by Detection limit <= 5 ng/L	Troponin	MCnc	Pt	Ser/Plas	Qn	Detection limit <= 5 ng/L	
	41995-2	Hemoglobin A1c [Mass/volume] in Blood	oglobii c	MCnc	Pt	Bld	Qn		g/dL
	4548-4	Hemoglobin A1c/Hemoglobin.total in Blood	He lobin /Hemoglobin.total	MFr	Pt	Bld	Qn		%
	17855-8	Hemoglobin A1c/Hemoglobin.total in Blood by calculation	Hemoglobin.total	MFr	Pt	Bld	Qn	Calculated	%
	4549-2	Hemoglobin A1c/Hemoglobin.total in Blood by Electrophoresis	Hemograpin A1c/Hemoglobin.total	MFr	Pt	Bld	Qn	Electrophoresis	%
Hemoglobin A1c	17856-6	Hemoglobin A1c/Hemoglobin.total in Blood by HPLC	Hemoglobin A1c/Hemoglobin.total	MFr	Pt	Bld	Qn	HPLC	%
	62388-4	Hemoglobin A1c/Hemoglobin.total in Blood by JDS/JSCC protocol	Hemoglobin A1c/Hemoglobin.total	MFr	Pt	Bld	Qn	JDS/JSCC	%
1	71875-9	Hemoglobin A1c/Hemoglobin.total [Pure mass fraction] in Blood	Hemoglobin A1c/Hemoglobin.total	MFr.DF	Pt	Bld	Qn		
	59261-8	Hemoglobin A1c/Hemoglobin.total in Blood by IFCC protocol	Hemoglobin A1c/Hemoglobin.total	SFr	Pt	Bld	Qn	IFCC	mmol/mol
1		- · · · · · · · · · · · · · · · · · · ·	-	·					 ,
1	49551-5	Creatine kinase.MB [Mass/volume] in Blood	Creatine kinase.MB	MCnc	Pt	Bld	Qn		ng/mL
4	51506-4	Creatine kinase.MB [Enzymatic activity/volume] in Cerebral spinal fluid by Electrophoresis	Creatine kinase.MB	CCnc	Pt	CSF	Qn	Electrophoresis	U/L
Creatine kinase MB	38482-6	Creatine kinase.MB [Presence] in Serum or Plasma	Creatine kinase.MB	ACnc	Pt	Ser/Plas	Ord		4
Creatine Kinase IVID	32673-6	Creatine kinase.MB [Enzymatic activity/volume] in Serum or Plasma	Creatine kinase.MB	CCnc	Pt	Ser/Plas	Qn		U/L
4	2154-3	Creatine kinase.MB [Enzymatic activity/volume] in Serum or Plasma by Electrophoresis	Creatine kinase.MB	CCnc	Pt	Ser/Plas	Qn	Electrophoresis	U/L
4	13969-1	Creatine kinase.MB [Mass/volume] in Serum or Plasma	Creatine kinase.MB	MCnc	Pt	Ser/Plas	Qn		ng/mL; ug/L
<u> </u>									
4	2156-8	Creatine kinase [Enzymatic activity/volume] in Amniotic fluid	Creatine kinase	CCnc	Pt	Amnio fld	Qn		U/L
4	50756-6	Creatine kinase [Mass/volume] in Blood	Creatine kinase	MCnc	Pt	Bld	Qn		ng/mL
4	34160-2	Creatine kinase [Presence] in Body fluid	Creatine kinase	ACnc	Pt	Body fld	Ord		
Creatine kinase total	16688-4	Creatine kinase [Enzymatic activity/volume] in Body fluid	Creatine kinase	CCnc	Pt	Body fld	Qn		U/L
4	2151-9	Creatine kinase [Enzymatic activity/volume] in Cerebral spinal fluid	Creatine kinase	CCnc	Pt	CSF	Qn		U/L
4	53433-9	Creatine kinase [Enzymatic activity/volume] in Dialysis fluid	Creatine kinase	CCnc	Pt	Dial fld	Qn		U/L
4	2157-6	Creatine kinase [Enzymatic activity/volume] in Serum or Plasma	Creatine kinase	CCnc	Pt	Ser/Plas	Qn		U/L
	20569-0	Creatine kinase.MB/Creatine kinase.total in Serum or Plasma	Creatine kinase.MB/Creatine kinase.total	CFr	Pt	Ser/Plas	Qn		%
	12189-7	Creatine kinase.MB/Creatine kinase.total in Serum or Plasma by calculation	Creatine kinase.MB/Creatine kinase.total	CFr	Pt	Ser/Plas	Qn	Calculated	%
Creatine kinase MB/creatine kinase total	12187-1	Creatine kinase.MB/Creatine kinase.total in Serum or Plasma by Electrophoresis	Creatine kinase.MB/Creatine kinase.total	CFr	Pt	Ser/Plas	Qn	Electrophoresis	%
Creatine kinase MB/Creatine kinase total	72564-8	Creatine kinase.MB/Creatine kinase.total [Pure catalytic fraction] in Serum or Plasma by calculation	Creatine kinase.MB/Creatine kinase.total	CFr.DF	Pt	Ser/Plas	Qn	Calculated	

(1) the principal name (e.g., the name of the analyte or the measurement); (2) the challenge or provocation, if relevant, including the time delay, substance of challenge, amount administered, and route of administration; and (3) any standardization or adjustment

https://github.com/CDMFORUM/CDM-GUIDANCE/wiki/Lab-Mapping-Resources

Creatine kinase.MB/Creatine kinase.total

Creatine kinase.MB/Creatine kinase.total

CFr.DF Pt

Ser/Plas

On Electrophoresis



PCORnet LAB NAME

Category or Name Component/analyte

LOINC code

Property

WHAT IT IS

- A dynamic resource containing LOINC codes that may map to PCORnet laboratory result name common measures.
- Information can be used to search for laboratory results in local data sources that may map to PCORnet laboratory result name common measures.
- © Rows highlighted in yellow indicate LOINC codes that may not correspond to the PCORnet lab result name common measure listed. They are included for data partners to determine whether or not they are a good fit.



WE NEED YOU!

- We welcome (NEED) feedback from PCORnet data partners about additional codes that should be considered for inclusion
- Also need feedback about codes that don't seem to match the lab result name common measure listed
 - Provide feedback via email: michelle.smerek@duke.edu
- Anyone interested can join the Lab Mapping Interest Group call in May.
 - Please send email to <u>michelle.smerek@duke.edu</u> if interested!



WHAT IT IS NOT

- An exhaustive list of all the LOINC codes that have ever, or could ever correspond to the PCORnet lab result common measures
 - New lab names will be created in source systems
 - New LOINC codes will be created by Regenstrief
- A static resource that will make all lab data in PCORnet usable for all potential analytic purposes



Next Steps

- Cab mapping interest group to meet and discuss in May
- Once initial round of feedback has been received/reviewed, "draft" designation will be removed, and it will be available to data partners in GitHub as a living document



CONDITION table and "PCORnet-defined cohort algorithm" category

Some content repeated from CDM v3.0 development phase in 2015 for context





Wednesday, April 29, 2015



CDM v3.0 Development Processes

- Feedback cycle: April 7-20, 2015
- 236 discreet comments received
 - With many thanks!
- Each comment assessed and tagged into 23 thematic categories
- Comments spreadsheet with response to every comment will be finalized and posted after second stakeholder meeting

v3.0 CDM draft on Central Desktop:

https://pcornet.centraldesktop.com/taskforces/file/39183449/

v3.0 comments spreadsheet on Central Desktop:

https://pcornet.centraldesktop.com/taskforces/file/39897440/



12 Themes Distilled from Tagging System

- Death
- Expectation
- Dates
- PCORNET_TRIAL
- Harvest
- SAS
- Global
- Phenotypes
- Other
- Medications
- DISPENSING
- PRESCRIBING



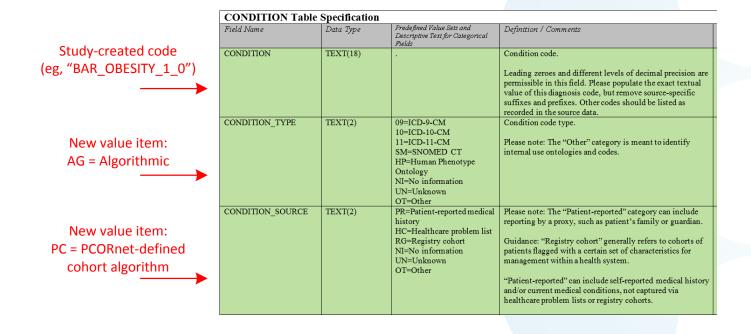
Tag CONDITION DATETIME DEATH DEMOGRAPHIC DIAGNOSIS DISPENSING ENCOUNTER ENROLLMENT EXPECTATION GLOBAL HARVEST HIT LAB RESULT CM MEDICATIONS NULLS OVERVIEW PCORNET TRIAL PHENOTYPE PRESCRIBING PRO_CM PROCEDURES RELATIONAL

SAS

VITAL

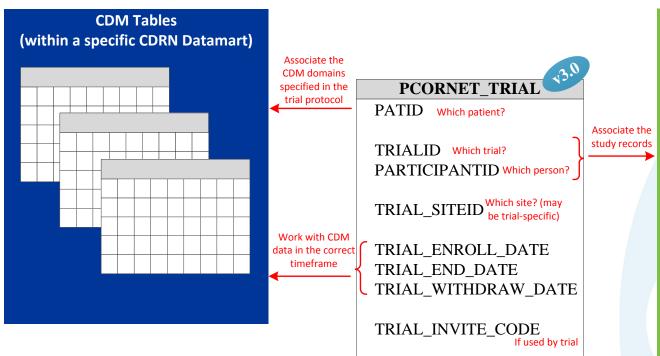
Theme: Phenotypes

Process of developing computable algorithms to recognize/define disease states and conditions

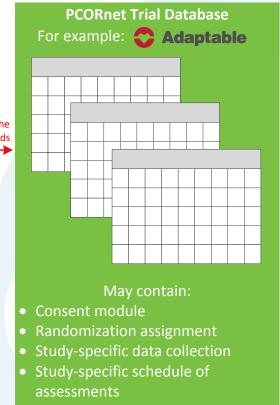




The PCORNET_TRIAL table serves as a connector and filter for CDM data within the parameters of a given trial protocol:



Any given PCORnet datamart may contain data from a large number of patients. The PCORNET_TRIAL table assists the ADAPTABLE Data Core in knowing which of these patients have consented and been enrolled in the ADAPTABLE trial for querying purposes.





Wrap up



Next CDM Forum: May 11, 2016

PCORnet Common Data Model (CDM) Implementation Forum

Wednesday, May 11, 2016, 1:00 - 2:00 PM Eastern time

Hosted by Michael Matheny, MD; facilitated by Shelley Rusincovitch and Michelle Smerek

ONLINE:

https://dukemed.webex.com/dukemed/j.php?MTID=m 501d523454c97d76bb86dec256c4853a

PHONE: 1-650-479-3207 / Access code: 739 305 302



These slides will also be posted on GitHub:

https://github.com/CDMFORUM/CDM-GUIDANCE/wiki



Summary of links

iMeetCentral resources:

- PCORnet main blog (includes weekly announcements): https://pcornet.imeetcentral.com/pcornetmain/blog/
- DRN OC home: https://pcornet.imeetcentral.com/p/aQAAAAAB6T9b
 - Data Characterization home: https://pcornet.imeetcentral.com/p/aQAAAAACjjsH
 - Query Fulfillment home: https://pcornet.imeetcentral.com/p/aQAAAAACIFkW
 - DRN OC blog: https://pcornet.imeetcentral.com/drnoc-workgroups/blog/
- Collaborative threads on iMeetCentral: https://pcornet.imeetcentral.com/p/ZgAAAAAAPgOl

GitHub:

- ADAPTABLE base phenotype : https://github.com/ADAPTABLETRIAL/PHENOTYPE
- CDM errata issue tracker: https://github.com/CDMFORUM/CDM-ERRATA/issues
- CDM guidance issue tracker: https://github.com/CDMFORUM/CDM-GUIDANCE/issues
- PCORnet diagnostic query package: https://github.com/PCORnet-DRN-OC/PCORnet-Diagnostic-Query
- PCORnet data characterization query package: https://github.com/PCORnet-DRN-OC/PCORnet-Data-Characterization
- PCORnet Data Committee on GitHub: https://github.com/PCORnet/DataCommittee

