

### **COVID-19 User Guide Project Review**

### Presented by:

David R. Bobbitt, President and CEO

Kit Howard, Sr. Director, Standards Development and Education

Dr. Diane Wold, Sr. Director, Standards Development

Bess Leroy, Head of Standards Development

Amy Palmer, Head of Standards Development

Jon Neville, Sr. Standards Developer

Dr. Erin Muhlbradt, Clincal/Biomedical Information Specialist, US NCI Enterprise Vocabulary Services [C]



### 6 MAY 2020



## Today's Agenda

- 1. Housekeeping
- 2. Presenter Introductions
- 3. Feature Presentations
- 4. Question & Answer Session
- 5. Upcoming Learning Opportunities + Resources



### Housekeeping



### Housekeeping

- You will remain on **mute** for the entirety of the call
- There will be a Q&A after all of the presentations are finished
- Audio issues? Shut down and restart the GoToWebinar app
- The slides from the presentation and a recording of this webinar will be available in the Members Only section of the CDISC website
  - To access make sure that you create a login for the CDISC website if you haven't already
  - If you are employed by a CDISC member organization, please ensure you use your employer-issued email address with your employer's domain name, so we can verify membership for the purpose of applying discounts to purchasing event tickets, online courses, and more!





### **Content Disclaimer**

- The purpose of this webinar is to provide examples of implementation and should not be considered official recommendations by CDISC unless otherwise stated in the presentation.
- This webinar is not an authorized CDISC course, is not developed or delivered under CDISC Operating Procedures, and should not replace a published standard. Please refer to the latest published standards for the most authoritative implementation information.





### **Our Presenters**

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### 6 MAY 2020



## **COVID-19 Task Force** Being the global community

David R. Bobbitt, MSc, MBA President and CEO





### [Adversity] doesn't change who you are. It reveals who you are.

-Michelle Obama

## Thank you to the Task Force members

Jennifer Alf Rebecca Baker Cathy Bezek Dana Booth Assia Bouhadouza **Stephanie Chen** Karen Fanouillere Nikki Flores Nate Freimark Praveen Garg Tom Guinter Ajay Gupta **Brian Harris** 

Keith Hibbetts Kit Howard Chris Kaiser Smitha Karra Kalynn Kennon **Bess LeRoy** Laura Merson Erin Muhlbradt Jon Neville Amy Palmer **David Parkinson** Nik Pemble Chris Price

Heather Ribaudo Justin Ritz Lauren Shinaberry **Trisha Simpson** Lorraine P. Spencer Will Stevens Alana St. Clair Sarah Strobino Helena Sviglin Peter Van Reusel **Robin White Diane Wold** Jennifer Xio





















Johnson & Johnson





NATIONAL CANCER INSTITUTE Enterprise Vocabulary Services



National Institute of Allergy and Infectious Diseases













# Thank you to our community



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### **CDISC** Response to COVID-19

6 May 2020



### Agenda

- 1. Project Background
- 2. Scope and Content Development
- 3. Interim User Guide for COVID-19
  - Diagnostics and Virology
- 4. Guidance for Ongoing Studies Disrupted by the COVID-19 Pandemic
  - Relationships to COVID-19
- 5. Resources for Public Health Researchers
- 6. Controlled Terminology COVID-19 Package 41a Publication

### **Project Background**

- Need identified for a CDISC Interim User Guide for COVID-19
- CDISC convened a Task Force in Late March 2020
  - Industry stakeholders
  - Regulatory
  - Academia
  - Key CDISC data standards staff
- Development did not follow formal Standards Development Process (CDISC COP-001)
- Development of the Guide on CDISC wiki
- Published on CDISC Website on 21 April 2020
  - <u>https://www.cdisc.org/standards/therapeutic-areas/covid-19</u>

### NOW AVAILABLE: Interim User Guide for COVID-19



# Defining and Refining the Scope

- Quickly realized that we had two areas of focus beyond an Interim User Guide for COVID-19 for new studies
  - Guidance for Ongoing Studies Disrupted by the COVID-19 Pandemic
  - Resources for Public Health Researchers WHO Annotated Forms and Mapping Spreadsheet
- QRS Supplement in development for National Early Warning Scale 2 (NEWS2)
- Off-cycle COVID-19 Controlled Terminology Release scheduled for 8 May 2020





### Content Development

- Task Force identified scope and provided use-cases and CRFs
- CDISC staff members developed modeling
- Reused existing approved modeling approaches when available, leveraging previous TA work and SDTMIG examples
  - Influenza
  - Tuberculosis
  - Malaria
  - Ebola
  - HIV
  - Virology
  - Vaccines

- Chronic Kidney Disease
- Duchenne Muscular Dystrophy
- Asthma
- COPD
- SDTMIG v3.4
- CDASHIG v2.1





### **Meeting Schedule**

- Task Force met weekly
- Ongoing Studies Team met twice weekly
- CDISC Standards Development Team met daily
- QRS development occurred during QRS CT and Subteam meetings
- CT development involved ongoing coordination with NCI-EVS



### **Review Process and Timelines**

- All development on CDISC wiki and publicly viewable for transparency
- Public could create JIRA issues to provide comments throughout the development process
- Batches of examples were sent to Task Force daily and feedback requested within 3 days
- CDISC staff resolved comments



### Interim User Guide for COVID-19

### Interim User Guide Topics Covered

- Risk Factors
  - Pre-existing Medical Conditions
  - Personal Protective Equipment (PPE)
  - Travel
  - Contacts
  - Substance Use
  - Exposure to Animals
- Onset of Disease
- Signs and Symptoms
- Laboratory Test Results
- Diagnostics and Virology
  - Virus Identification
  - Antibody Testing
  - SARS-CoV-2 Viral Load
- Vital Signs and Urine Output

- Concomitant Medications
- Respiratory Findings
  - Imaging
  - Pulmonary Function Tests
- Cardiac Events/Findings
- Hospitalization
- Procedures
  - Assisted Ventilation and Oxygen Treatments
  - Renal Treatment
- Vaccines
- Questionnaires, Ratings, and Scales



### Environmental and Social Factors (ER) Draft Domain

- The Environmental and Social Factors (ER) domain represents data that was collected to assess the factors that might influence a subject's disease or medical condition via environmental contact or through participation in activities associated with increased or decreased risk.
- Used in COVID-19 Interim User Guide to represent data on travel, contacts, personal protective equipment, and exposure to animals
- This domain went through public review during the TB v2.0 TAUG public review. It also went through public review as part of SDTMIG v3.3 batch 3.
  - Held out from publication due to maturity concerns
    - Unclear if a single observation class (Events) is adequate to cover all use-cases
  - Publicly available on the CDISC Wiki



### **Diagnostics and Virology**

### **Diagnostics and Virology Section Overview**

- This section includes the following examples:
  - Virus identification- testing for the presence of SARS-CoV-2 in a subject sample
  - Antibody Testing- detection of IgG and IgM (terminology also exists for testing of SARS-Cov-2 IgA antibody)
  - Viral load testing
    - Quantification of SARS-CoV-2 RNA by quantitative PCR
    - Threshold Cycle value

### Sources of input

- Published literature
- Task Force member feedback
- Prior examples from existing CDISC therapeutic-area user guides



### **Virus Identification**

• The example follows SDTMIG conventions (v3.2-3.3) by representing these data in the MB domain

		1										
US	UBJID	MBSEQ	MBREFID	MBGPRID	MBTESTCD	MBTEST	MBTSTDTL	MBORRES	MBSTRESC	MBSPEC	MBLOC	MBMETHOD
A	3C-01-	1	60101	1	SARSCOV2	Severe Acute Resp Syndrome	DETECTION	DOSITIVE	DOCITIVE	ENDOTRACHEAL		QUANTITATIVE REVERSE TRANSCRIPTASE
	601		00101	00101 I SA		Coronavirus 2	DETECTION	POSITIVE	POSITIVE	FLUID		POLYMERASE CHAIN REACTION
A	3C-01-	2	72201	1	SARSCOVA	Severe Acute Resp Syndrome	DETECTION	NEGATIVE	NECATIVE	SWABBED		QUANTITATIVE REVERSE TRANSCRIPTASE
	722 2		72201	'	SARSCOV2	Coronavirus 2	DETECTION	NEGATIVE	NEGATIVE	MATERIAL	INKOAI	POLYMERASE CHAIN REACTION
		1										

- MBTEST/MBTESTCD represent the name of the virus
- Since we are only interested in detecting the presence of virus (versus quantifying it) MBTSTDTL=DETECTION
- Results are expressed as POSITIVE / NEGATIVE



### Antibody Testing

• As of SDTMIG v3.3, this concept is still represented in MB

DOMAIN	USUBJID	MBSEQ	MBREFID	MBTESTCD	MBTEST	MBTSTDTL	MBORRES	MBSTRESC	MBSPEC
MB	COVID-ABC-011	1	13668	SAR2IGM	SARS-CoV-2 IgM Antibody	DETECTION	POSITIVE	POSITIVE	SERUM
MB	COVID-ABC-011	2	13668	SAR2IGG	SARS-CoV-2 IgG Antibody	DETECTION	NEGATIVE	NEGATIVE	SERUM
MB	COVID-ABC-022	1	23433	SAR2IGGM	SARS-CoV-2 IgG/IgM Antibody	DETECTION	POSITIVE	POSITIVE	SERUM

- Based on the 3 most common antibody detection tests: IgG, IgM, and combination IgG/IgM
- Since we are not quantifying antibodies, the modeling approach is similar to virus detection with regard to MBTEST, MBTSTDTL, and results of POSITIVE / NEGATIVE
- If the assay were designed quantify antibodies, we would use MBTSTDTL=QUANTIFICATION, with numeric results and appropriate units



### Viral Load

	MPCPDID	MODEEID	MOTECTOD	MOTECT	MOTOTOTI	MROBBEC	MPOPPECII	MOCTOCC	MOCTOCON	MOCTOCCU	MPCDEC	MPMETHOD
•	IVIDGRPID	WIDKEFID	WIDTESTED	IVIDIESI	WIDISIDIL	WIDUKKES	WIDUKKESU	IVIDSTRESC	IVIDO I REDIN	IVIDO I REOU	IVIDSPEC	IVIDIVIETHOD
	1	001-02	CARORNIA	SARS-CoV-		2.0	log 10	2.0	2.0	log 10	SDUTUM	QUANTITATIVE REVERSE TRANSCI
	· · · ·	001-02	SANZNINA	2 RNA	VINAL LOAD	5.9	copies/mL	2.9	5.9	copies/mL	SPOTOW	POLYMERASE CHAIN REACTI
•	1	001.00	CARODNIA	SARS-CoV-	THRESHOLD CYCLE	27.42		27.42	27.42		COLITUM	QUANTITATIVE REVERSE TRANSCI
		001-02	SANZINIA	2 RNA		27.43		27.43	27.43		SPUTUM	POLYMERASE CHAIN REACTI
	2	001 02	CARODNIA	SARS-CoV-		47	log 10	47	4.7	log 10	COLITUM	QUANTITATIVE REVERSE TRANSCI
	2	001-05	SARZNINA	2 RNA	VIRAL LOAD	4.7	copies/mL	4.7	4.7	copies/mL	SPUTUIVI	POLYMERASE CHAIN REACTI
:	2	001-02	SARORNIA	SARS-CoV-	THRESHOLD	22.11		22.11	22.11		CDUITUM	QUANTITATIVE REVERSE TRANSCI
	2	001-05	SARZNINA	2 RNA	CYCLE	25.11		25.11	25.11		SPUTUIVI	POLYMERASE CHAIN REACTI
	2	001-04	SARORNIA	SARS-CoV-		4.5	log 10	4.5	4.5	log 10	SDUTUM	QUANTITATIVE REVERSE TRANSCI
	5	001-04	JANZINNA	2 RNA	VINAL LOAD	4.5	copies/mL	4.5	4.5	copies/mL	SPOTOW	POLYMERASE CHAIN REACTI
•	2	001-04	SAPOPNIA	SARS-CoV-	THRESHOLD	22.22		22.22	22.22		CDUITUM	QUANTITATIVE REVERSE TRANSCI
	5	001-04	SANZINIA	2 RNA	CYCLE	23.22		23,22	23.22		SPOTOW	POLYMERASE CHAIN REACTI
:		001-05	SAPOPNIA	SARS-CoV-		TARGET NOT		TARGET NOT			SDUTUM	QUANTITATIVE REVERSE TRANSCI
		001-05	SANZINIA	2 RNA		DETECTED		DETECTED			SPOTOW	POLYMERASE CHAIN REACTI
		001-06	SAPOPNIA	SARS-CoV-		TARGET NOT		TARGET NOT			SDUTUM	QUANTITATIVE REVERSE TRANSCI
:		001-00	SANZINIA	2 RNA		DETECTED		DETECTED			SPOTOW	POLYMERASE CHAIN REACTI
		001-07	SAP2PNIA	SARS-CoV-		TARGET NOT		TARGET NOT			SDUTUM	QUANTITATIVE REVERSE TRANSCI
		001-07	SANZNINA	2 RNA		DETECTED		DETECTED			3-010101	POLYMERASE CHAIN REACTI

- The example follows one subject over 6 visits
- All records: MBTEST= SARS-CoV-2 RNA
- Test may quantify RNA (viral load) and/or Threshold Cycle; MBTSTDTL distinguishes these, and when both are present, GRPID is used to group both by subject-visit
- When SARS-CoV-2 RNA isn't detected in the subject sample, MBTSTDTL is null



### Quantitative Reverse-Transcriptase PCR (qRT-PCR)

Threshold Cycle  $(C_t)$  – The PCR cycle at which the signal from the amplified sample crosses the threshold



(1) For the qRT-PCR assay illustrated in this example, conversion from the Ct readout to a viral load readout requires the use of a known concentration of reference standard sample of the target RNA to be run parallel with the subject sample. It is important to note that in the case of emerging pathogens like SARS-CoV-2 such reference standards may not be available. In such cases only the C+ value will be available and a quantitative viral load expressed as number of copies per unit volume will not be reported.





### Summary

- All concepts represented in MB (LB is not appropriate for these concepts)
- Pay attention to the value of MBTSTDTL based on what the test is reporting
  - DETECTION, QUANITIFICATION, VIRAL LOAD, THRESHOLD CYCLE
- Always check controlled terminology. More controlled terminology exists
  than what is shown in the examples
  - SARS-CoV-2 IgA Antibody
  - "copies/mL" is also valid for viral load



# Guidance for Ongoing Studies Disrupted by the COVID-19 Pandemic

# Guidance for Ongoing Studies Disrupted by the COVID-19 Pandemic

- Listing of COVID-19 Related Impacts as Part of CSR
- Relationships to COVID-19
- Protocol Deviations
- Disposition
- Missed Visits
- Missed Assessments
- Changes to Drug Accountability
- Changes to Adverse Event Data Collection
- Changes in Exposure
- Transfer to Another Site
- Trial Summary to Provide Pandemic Relationship



**GUIDANCE DOCUMENT** 

# FDA Guidance on Conduct of Clinical Trials of Medical Products during COVID-19 Public Health Emergency

Guidance for Industry, Investigators, and Institutional Review Boards

**MARCH 2020** 

	Download	the Final Guida	ance Docum	ent
		Final		
f Share	y Tweet	in Linkedin	🔀 Email	🔒 Print

https://www.fda.gov/regulatory-information/search-fda-guidancedocuments/fda-guidance-conduct-clinical-trials-medical-products-during-covid-19-public-health-emergency





### **Multiple Approaches**

- It may not be possible to update case report forms (CRFs) for ongoing trials
- Collection of relevant information can be performed multiple ways
  - Key words such as "COVID-19" in existing text fields
  - Comments on the CRF
  - Documentation outside the regular CRF
- Instead of showing one way to model the data in SDTM, multiple options were shown
- Data about impacts of the epidemic, whether from CRFs or other sources, may be represented different ways
  - Standard SDTM variables,
  - Non-standard variables (NSVs),
  - Custom domains
  - Flag SDTM dataset records that document epidemic impacts using indicator NSVs



### **Relationships to COVID-19**
#### Direct and Indirect Relationships to COVID-19





#### Which study subjects contracted COVID-19?

- COVID-19 as an Adverse Event
  - Coding in MedDRA depends on MedDRA version
- Adverse Events related to COVID-19
  - Use non-standard variable (NSV) Epi/Pandemic Related Indicator
- What treatments were given for COVID-19?
  - Use existing standard variable Indication (--INDC)



# **Possible Disruptions**

- Scheduled activities missed
  - Missed visits
  - Missed assessments
  - Missed study drug treatments
- Scheduled activities modified
  - Remote visits and assessments
  - Assessments at an alternate facility
  - Remote study drug distribution and return
  - Changes in study drug administration
- Protocol Deviations
  - Subject-level deviations
  - Site-level deviations
- Early subject withdrawals
- Site shut-downs and subject transfers



#### Why did these disruptions occur? Which reasons were related to COVID-19?

#### • Use "reason" variables

- Standard variables such as Reason Not Done, Indication, Reason for Adjustment, Reason for Discontinuation, DSTERM in the Disposition Domain
- Established non-standard variables such as Reason (e.g., for a protocol deviation), Reason for Occur Value (e.g., reason a treatment was not given), Reason for Interruption
- New non-standard variables as needed
- If desired, use additional non-standard indicator variables to flag reasons that were related to COVID-19.



#### Modifications to Scheduled Activities

- For activities originally scheduled as in-person, use NSV Contact Mode
  - For remote visits
  - For study treatment shipped directly to subjects
  - For data on amount of study treatment used collected by phone
- For tests and examination, use --NAM and--EVAL to record a different lab or a different evaluator





#### **Missed Visits**

- May be recorded in existing Protocol Deviations domain
  - Missing an originally scheduled visit may not be a deviation under a later protocol amendment
  - NSV: Reason for Deviation
  - NSV: Epi/Pandemic Related Indicator
- Custom Visit Events domain can be used to record both visits that occurred and those that were missed.
  - NSV: Reason for Occur Value (i.e., reason visit did not occur)
  - NSV: Epi/Pandemic Related Indicator
  - NSV: Contact Mode





#### **Site-Level Protocol Deviations**

- Identify subjects affected by the deviation
- Choose one of the following approaches
  - Record the deviation for each affected subject
  - Use POOLDEF to assign an identifier to the affected subjects as a pool, then record the deviation using POOLID





#### Site Transfers

- Choose one of the following approaches:
  - Custom domain Site Transfers
  - Record site transfers in the Disposition domain using DSCAT of "OTHER EVENT"
- In either case, record the additional site as supplemental qualifier in the Demographics domain.



#### Resources for Public Health Researchers

#### **Resources for Public Health Researchers**

- Novel Coronavirus (nCoV) Acute Respiratory Infection Clinical Characterisation Data Tool
  - Developed by the World Health Organization (WHO) and the International Severe Acute Respiratory and Emerging Infection Consortium (ISARIC)
  - Data tool is being used as the foundation for many COVID-19 research studies globally by more that 40 countries
- SDTM and CDASH Annotations and Excel Mapping File created by CDISC team published on CDISC website



#### WHO / ISARIC / IDDO Collaboration











#### CDISC's Participation: Annotated CRFs

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CORE CASE RECORD FORM
DEMOGRAPHICS
Clinical centre name: DM.SITEID Country: DM.COUNTRY {ISO-3166-1 Alpha-3}
Enrolmentdate: [_D_][_D_]/[_M_][_M_]/[_2_][_0_][_Y_][_Y_] DM.RFICDTC
Ethnic group (check all that apply): Arab Black East Asian South Asian West Asian Latin American White
Aboriginal/First Nations Other:Unknown RACE {RACE} RACE1, RACE2, etc. {RACEC}
Employed as a Healthcare Worker? DYES <u>NO</u> N/A ERTESTCD {ERTESTCD} ERORRES
Employed in a microbiology laboratory?     YES     NON/A     ERTESTCD {ERTESTCD}     ERORRES
Sex at Birth:  Male Female Not specified SCTESTCD {SCTESTCD} SCORRES
Estimated Age [][_]years OR [_][_]months AGE AGEU {AGEU}
Pregnant?  YES NO Unknown N/A If YES: Gestational weeks assessment: [][] weeks
MHTERM (MHPRESP)/MHOCCUR {YN} RPTESTCD {RPTESTCD}="Estimated Gestational Age" on mother's I
RPORRES RPORRESU {AGEU}



#### CDISC's Participation: Annotated CRFs

CORE CASE RECORD FORM

	DEMOGRAPHICS
	Clinical centre name: DM.SITEID Country: DM.COUNTRY {ISO-3166-1 Alpha-3}
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DIW	Ethnic group <i>(check all that apply):</i> <u>Arab</u> Black East Asian South Asian West Asian Latin American White Aboriginal/First Nations Other:Unknown RACE {RACE}
	Employed as a Healthcare Worker?  IYES INO IN/A ERTESTCD {ERTESTCD} ERORRES
-	Employed in a microbiology laboratory?     Image: YES     Image: N/A     ERTESTED {ERTESTED}     ERORRES
ļ	Sex at Birth:  Male_ Female  Not specified  SCTESTCD {SCTESTCD} SCORRES
DM	Estimated Age [][]years OR [_][]months AGE AGEU {AGEU}
	Pregnant?       YES       NO       Unknown       N/A If YES: Gestational weeks assessment:       []]       ] weeks         MHTERM       (MHPRESP)/MHOCCUR {YN}       RPTESTCD {RPTESTCD}="Estimated Gestational Age" on mother's ID         RPORRES       RPORRES       RPORRESU {AGEU}



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#### CDISC's Participation: Annotated CRFs

	CORE CASE RECORD FORM
	DEMOGRAPHICS
	Clinical centre name: DM.SITEID Country: DM.COUNTRY {ISO-3166-1 Alpha-3}
	Enrolmentdate: [_D_][_D_]/[_M_][_M_]/[_2_][_0_][_Y_][_Y_] DM.RFICDTC
	Ethnic group <i>(check all that apply):</i> <u>Arab</u> Black East Asian South Asian West Asian Latin American White Aboriginal/First Nations Other: <u></u> Unknown <u>RACE {RACE}</u> <u>RACE1, RACE2, etc. {RACEC</u>
	Employed as a Healthcare Worker? DYES DODDN/A ERTESTCD {ERTESTCD} ERORRES
EK	Employed in a microbiology laboratory?   YES  NO KA ERTESTCD {ERTESTCD} ERORRES
SC	Sex at Birth:  Male Female Not specified SCTESTCD {SCTESTCD} SCORRES
	Estimated Age [][_]years OR [_][_]months AGE AGEU {AGEU}
/IH/RP	Pregnant?       YE\$       NO_       Unknown       N/A If YES: Gestational weeks assessment:       [][]] weeks         MHTERM       (MHPRESP)/MHOCCUR {YN}       RPTESTCD {RPTESTCD}="Estimated Gestational Age" on mother's
	RPORRES RPORRESU {AGEU}



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MODULE 1: MESENTATION ADMISSION CASE REPORT FORM GINGA INCUSION OFTEN petel a primei sui promite 1240 (Linksie: 015-010

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# CDISC's Participation: Mapping Spreadsheet

	В	C	D	E	F
	Domain	SDTM Variable	Value	CDASH Variable If Different	Notes
1				from SDTM	
8					
	DM	RACE		CRACE	While this item indicates that ethnic group is collected, the concept
					described most closely matches RACE in CDISC. The RACEs listed on the
					CRF do not correspond to CDISC RACE codelist. Capture in CRACE using
9					the RACEC CT and map into SDTM RACE as needed.
10	SUPP.DM	RACEOTH			
11					
	ER	ERTERM	EMPLOYED AS A HEALTHCARE		
12			WORKER		
13	ER	ERPRESP	Y		
14	ER	EROCCUR	Y; N; NA		
15	ER	ERCAT	COVID-19 RISK FACTOR		
16					
22	SC	SCTESTCD	SEXRBRTH		
23	SC	SCTEST	Sex Reported at Birth		
24	SC	SCORRES	M; F; U		"U" corresponds to "Not Specified"on the CRF
24					
25					
26	DM	AGE			
27	DM	AGEU	YEARS; MONTHS		
	REA	D ME Clinical Inc	lusion Criteria   Epi factors   Demogra	aphics Comorbidities Onset and	Admission SS at Hos (+)





#### Outcomes



#### Twelve annotated CRF pages

Mapping file with 950 variables, associated metadata, implementation recommendations and CDASH equivalent variables

Most concepts mapped with minimal adjustment A few concepts required non-standard variables or new modeling



CRFs and mapping file facilitate production of SDTM tables for combination with data from other sources

Perhaps future efforts can be designed to align a little more closely





#### Controlled Terminology for CDISC COVID-19 Guidance

Dr. Erin Muhlbradt, Clinical/Biomedical Information Specialist, MSC Inc. and NCI-EVS  $\cal{c}$ 

05.06.2020



#### Agenda

- 1. Publication timeline
- 2. COVID-19 (Package 41a) Publication Release (2020-05-08)
- 3. Questions

# Controlled Terminology COVID-19 Package 41a Publication

2020-05-08

#### **Controlled Terminology Publication Schedule**

Package Number	Team Cutoff (requests must be received at least two months before this date)	Public Review Start Date (1 wk from Team Cutoff)	Public Review Closed Date (4 wks)	Final Changes to NCI EVS (4 wks)	Publication Date (6 wks)		Codelists to be Included		
39	6/14/2019	6/21/2019	7/19/2019	8/16/2019	9/27/2019	Devices	ECG	General	Lab
39						Microbiology	Oncology	PK	Protocol Entities
39						SDTM Domain	SEND	SEND-AR	Unit
40	9/13/2019	9/20/2019	10/18/2019	11/8/2019	12/20/2019	ADaM	Define-XML	Device	General
40						CDISC Glossary	Lab	Microbiology	Oncology
40						PK	Protocol Entities	QRS	SDTM Doma
40						SEND	Spectype Speccond	Unit	
41	12/13/2019	12/20/2019	1/17/2020	2/14/2020	3/27/2020	ADaM	Define-XML	Device	ECG
41						General	Lab	Microbiology	Oncology
41						PK	Protocol Entities	QRS	SDTM Doma
41				Ń	ン	SEND	Spectype Speccond	Unit	
41a	4/20/2020	n/a	n/a	n/a	5/8/2020	CDASH	CDISC Glossary	General	Lab
41a					よう	Microbiology	PK	QRS	Spectype Speccond
41a						Unit			
42	3/13/2020	3/20/2020	4/17/2020	5/15/2020	6/26/2020	ADaM	Define-XML	Device	General
42						Lab	Microbiology	Oncology	PK
42						Protocol Entities	QRS	SEND	Unit
43	6/12/2020	6/19/2020	7/17/2020	8/14/2020	9/25/2020				
43									
43									
44	9/11/2020	9/18/2020	10/16/2020	11/13/2020	12/18/2020				

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4/21/2020

#### Agenda

- 1. Publication timeline
- 2. COVID-19 (Package 41a) Publication Release (2020-05-08)
- 3. Questions

- Updates to SDTM, SEND, CDASH, and CDISC Glossary Terminology
- CDISC CT Version Date: 2020-05-08
- Therapeutic Area Support:
  - QRS Terminology to support: COVID-19 Therapeutic Area User Guide.
  - SDTM/CDASH/Glossary Terminology to support: COVID-19 Therapeutic Area User Guide.



- General Terminology Team:
  - New Terms Added to Existing Codelists:
    - PROCEDUR; RETEST-CD; SCTEST-CD; TSPARM-CD; VSTEST-CD; RACEC
  - 4 New Codelists:
    - Health Care Encounters Dictionary Derived Term (HODECOD)
    - Mode of Subject Contact (CNTMODE)
    - COVID-19 Findings About Test Name (C19FAT)
    - COVID-19 Findings About Test Code (C19FATCD)
  - No changes to published terms



- CDISC Glossary Team
  - 8 additions to the CDISC Glossary:
    - epidemic
    - pandemic
    - endemic disease
    - immediately life-threatening disease or condition
    - medical countermeasure
    - morbidity rate
    - mortality rate
    - pre-approval access



- CDISC CDASH Terminology
  - New Terms Added to Existing Codelists: Race As Collected (RACEC)
  - To support the WHO ISARIC CRF annotation



- Laboratory Terminology Team
  - New Terms Added to Existing Codelists:
    - LBTEST-CD; METHOD; CLMETH; SPECTYPE
  - 4 changes to published terms:
    - FIO2/Fraction of Inspired Oxygen: Definition updated to better differentiate this term from FIO2/PAO2
    - PO2/Partial Pressure Oxygen: Synonyms added
- Units of Measure
  - 2 changes to published terms
    - mL/24h added as a synonym to C67410/mL/day in the UNIT and PKUNIT codelists – to support the WHO ISARIC CRF annotation



- Microbiology Terminology Team
  - New Terms Added to Existing Codelists:
    - MICROORG; MBTEST-CD

<u>IBTESTCD</u>	MBTEST
SARSCOV2	Severe Acute Resp Syndrome Coronavirus 2
SAR2IGG	SARS-CoV-2 IgG Antibody
SAR2IGM	SARS-CoV-2 IgM Antibody
SAR2IGA	SARS-CoV-2 IgA Antibody
SAR2IGGM	SARS-CoV-2 IgG/IgM Antibody
IERSRNA	MERS-CoV RNA
SAR2RNA	SARS-CoV-2 RNA
RSV	Respiratory Syncytial Virus
DENOVIR	Adenoviridae
CRONAVIR	Coronaviridae
IERSRNA	MERS-CoV RNA
SAR2RNA	SARS-CoV-2 RNA



- Microbiology Terminology Team
  - 2 changes to 1 existing term
    - MICROORG: CDISC Submission value of SARS CORONAVIRUS updated to SEVERE ACUTE RESPIRATORY SYNDROME-RELATED CORONAVIRUS. Also 'SARS-CoV' added as a synonym to this term.
  - New terms from P42 being published early with this package:
    - MICROORG: SEVERE ACUTE RESPIRATORY SYNDROME CORONAVIRUS 2
    - MBTEST/CD: Severe Acute Resp Syndrome Coronavirus 2/SARSCOV2



- New Codelists:
  - Clinical Classification (RS)
    - National Early Warning Score 2 (NEWS2)
    - Richmond Agitation-Sedation Scale (RASS)
    - Riker Sedation-Agitation Scale (SAS)



- SEND Terminology Team
  - New Terms Added to Existing Codelists:
    - LBTEST/CD; VSTEST/CD
  - 6 changes to existing terms -> LBTEST/CD, UNIT, PKUNIT as described previously

\*Because SEND makes use of some SDTM terminology, SEND Terminology will also be updated with this package release to reflect changes in shared codelists.





#### P41a Codetable Updates

- Updates with P41a
  - RACEC\_ETHNICC\_Mapping
  - SC\_Codetable\_Mapping
  - TS\_Codetable\_Mapping
  - VS\_Codetable\_Mapping
  - Unit-UCUM\_Codetable



No Additions or Changes being proposed by the following codelist-associated teams:

- ADaM Team
- Define-XML Team
- Medical Devices Team
- ECG Terminology Team
- Oncology Terminology Team
- CDISC PGx Terminology Team
- Protocol Entities Terminology Team



# If you are interested in contributing to the CDISC Terminology Initiative, please contact us...

Erin Muhlbradt, <u>muhlbradtee@mail.nih.gov</u> Dana Booth, <u>dbooth@cdisc.org</u>

> <u>CDISC New term request form:</u> <u>https://ncitermform.nci.nih.gov/ncitermform/?version=cdisc</u>



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#### Q&A




## How do we apply the guidelines if we are using SDTM 3.2 for a study?





Use of non-standard variables to flag COVID. What is recommendation: SDTM/ADaM? FDA preference (i.e. need for SDTM)?









Will there be a distinction between acquiring the illness and the restrictions on travel, i.e. for protocol deviations?

75



Will you have CRF examples in this TAUG? And did you map the CDC Covid Person Under Investigation CRF to CDASH and SDTM?







## When will the NEWS2 supplement be available?



77



What's the most suitable domain to map Antibody information between IS and MB?









NSV will be captured in SUPPxx domain and not in parent domain, correct?





Should we capture non-standard indicator variables onto Supplementary dataset or it will be added into main SDTM domains?









Diane - What was the rationale behind suggesting the use of a custom dataset VE (Visit Events)? The visit events are already mapped in the appropriate SV dataset. The additional findings about these visits (such as occurrence, contact mode) can perfectly be mapped into the FA (Findings About Events or Interventions) dataset.





@Diane: It seems to me that the custom Visit Events domain would be useful to have routinely, not just in disrupted studies. Will the SDTM team consider adding this to the SDTMIG?









Diane - Is there any guidance on reporting COVID-19 related deviations which are no longer a deviation following an emergency protocol amendment. These should also be reported to FDA.





At page #44 of presentation, why is 'sex at birth' annotated for SC domain instead of DM domain?









Do the Non std vars reside in parent domains? If yes, can you give an example? Thanks.





What was the rational behind mapping some information to NSV and others SUPPQUAL in the Interim User Guide for COVID-19?









Erin: Will Package 41a be a comprehensive, or incremental to just covered COVID-19 specific codelists & terms?



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China	29 May - 3 July	Weekly	Mandarin	

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Japan	28 May - 11 June	Weekly	Japanese	
China	29 May - 12 June	Weekly	Mandarin	

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Date	Webinar Title
2 JUN 2020	CDISC Library Virtual Workshop (Members-only!)
7 JUL 2020	Controlled Terminology Updates for Q3
13 OCT 2020	Controlled Terminology Updates for Q4





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**Presenters:** Saad Yousef, Senior Manager, Education Operations, CDISC







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Date and Time: TUE 2 JUN 2020 11:00 AM - 12:30 PM Eastern US Daylight Time

**Presenters:** Sam Hume, VP of Data Science, CDISC Anthony Chow, Director of Data Science, CDISC







Technology Digital Media Great Ideas!



## **Thank You!**

Questions, comments, concerns? Email bklinke@cdisc.org

