**Orders & Observations Conference Call**

**14 December 2016**

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**WebURL:** [**https://join.me/vernetzt.us**](https://join.me/vernetzt.us)

**Attendees:**

|  |  |  |
| --- | --- | --- |
|  | Name | Organization |
| 1 | Andrea Pitkus | IMO |
| 2 | Kathy Walsh | LabCorp |
| 3 | Carolyn Knapik | CAP |
| 4 | Riki Merrick | Vernetzt, LLC / APHL |
| 5 | JD Nolen | Cerner |
| 6 | Raj Dash | Duke |
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 Regrets: Rob Hausam

**Co-Chair**: Riki Merrick

**Scribe:** Riki Merrick

Agenda/Minutes:

1. Agenda Review
2. Call next week – will have it – even if we don’t have quorum, we will continue and complete the spreadsheet review and edit the document – then we will bring to OO for review and vote
3. Approve minutes – defer until we have completed the spreadsheet
	1. from May 6, 2016: : <http://wiki.hl7.org/index.php?title=File:SPM_Minutes_20160506_ConCall.docx>
	2. from May 27, 2016:<http://wiki.hl7.org/index.php?title=File:SPM_Minutes_20160527_ConCall.docx>
	3. from June 3, 2016: <http://wiki.hl7.org/index.php?title=File:SPM_Minutes_20160603_ConCall.docx>
	4. from June 10, 2016: <http://wiki.hl7.org/index.php?title=File:SPM_Minutes_20160610_ConCall.docx>
	5. from June 17, 2016: <http://wiki.hl7.org/index.php?title=File:SPM_Minutes_20160617_ConCall.docx>
	6. from June 24, 2016: <http://wiki.hl7.org/index.php?title=File:SPM_Minutes_20160624_ConCall.docx>
	7. NO CALL ON July 1, 2016
	8. From July 8, 2016: <http://wiki.hl7.org/index.php?title=File:SPM_Minutes_20160708_ConCall.docx>
	9. From July 15, 2016: <http://wiki.hl7.org/index.php?title=File:SPM_Minutes_20160715_ConCall.docx>
	10. From July 22, 2016: <http://wiki.hl7.org/index.php?title=File:SPM_Minutes_20160722_ConCall.docx>
	11. From July 29, 2016: <http://wiki.hl7.org/index.php?title=File:SPM_Minutes_20160729_ConCall.docx>
	12. From August 5, 2016: <http://wiki.hl7.org/index.php?title=File:SPM_Minutes_20160805_ConCall.docx>
	13. From August 12, 2016 – NO Quorum
	14. From August 19, 2016 – <http://wiki.hl7.org/index.php?title=File:SPM_Minutes_20160819_ConCall.doc>
	15. From August 26, 2016 - <http://wiki.hl7.org/index.php?title=File:SPM_Minutes_20160826_ConCall.doc>
	16. From September 2, 2016 – NO Quorum
	17. From September 9, 2016 – No call
	18. From September 16, 2016 - <http://wiki.hl7.org/index.php?title=File:SPM_Minutes_20160916_ConCall.docx>
	19. From October 14, 2016 – NO Quorum
	20. From October 28, 2016 – NO Quorum
	21. From November 2, 2016 - <http://wiki.hl7.org/index.php?title=File:SPM_Minutes_20161102_ConCall.docx>
	22. From November 9, 2016 - NO Quorum
	23. From November 169, 2016 - NO Quorum
4. Compare Specimen DAM to biologic specimen model in BRIDG - map specification source is BRIDG and the map to is specimen DAM

Spreadsheet

EA file is here:

<http://www.hl7.org/documentcenter/public/wg/orders/V3_DAM_Specimen_R1_2014May-BRIDG_Discussions.zip>

The spreadsheet: <http://www.hl7.org/documentcenter/public/wg/orders/Copy%20of%20BRIDG%20to%20Specimen%20DAM%20Mapping%20Spreadsheet%20as%20of%20Nov%202%202016.xlsx>

Ballot document:

<http://www.hl7.org/documentcenter/public/wg/orders/V3_DAM_Specimen_R2_INFORM_2017JAN_20161102.docx>

Also Raj sent this file:



We have these items left in these classes:

* SpecimenProcessingProtocol
	+ Add referencedProtocolIdentifier to specimenCollectionProcedure as O 0..1
	+ Add referencedProtocolIdentifier to specimenProcessingActivity as O 0..1
	+ For biospecimen best practices generally recommends referencing SOP
	+ On last call we discussed that we also want to document when there was deviation from the referenced protocol, what it was and possibly why – will need to find the elements for that:
		- * referencedProtocolDeviationComment as text, O 0..\* - this could hole the reason for the deviation type or any other information not captured by referencedProtocolDeviationType - TO ADD TO SPREADSHEET to both specimenCollectionProcedure and specimenProcessingActivity
		- do we need to document the reason for the specific comment it belongs
			* may be start just with comment – since free text, it could include the reason as well for starters
			* over time we can add a coded element for referencedProtocolDeviationType as code O 0..\* - TO ADD TO SPREADSHEET to both specimenCollectionProcedure and specimenProcessingActivity
			* Raj can request the top 10 reasons seen today to create starter set
* Subject – with related considerations for attributes in Material, Person, Non-human-living entity, Biologic entity
	+ - we decided to keep ONLY the ID for linkage, so that all information related to subject is in one place – no need for reconciliation between 2 locations etc
		- capture the type of specimen (environmental, human, animal, food, etc) using the specimen.ClassCode aas that may be quite important to know without linking to the subject
		- keep SubjectCharacteristicsAtTimeOfCollection that would allow additional elements to be recorded, should the link need to be broken for de-identification
			* Do we need to remove the “atTimeOfCollection” specifier, so we can track subjectCharacteristics at different times?
			* Can that include a picture for forensic purposes? Should be allowed – is normal observation, so don’t see why not support for image
			* Do we need specimenCharacteristics attribute to describe:
				+ taking images of the culture plate daily during specimenProcessingActivity?
				+ would that not be a ProcessingActivity in its own right and the resulting image is a derived specimen?
				+ it would be more like an analysis or observation on the specimen – we will need to add a reference from specimen to separate class of testing / observation so the specimen can be linked that way

prefer to capture the image

amount of extracted DNA etc that way

lab test result

* + - * + do we need a specimenProcessingActiivityIdentifier to reference specimen events?
			* How would you document that you dropped a specimen?
				+ Assume specimenProcessingActivity per SOP #x and then record the DeviationType and a comment?
* Specimen
	+ Accession number – is used very variable across organizations, or even inside organizations, in BRIDG is defined as the goupID for the same specimen collection – would that be the specimenCollectionProcedureIdentifier? No could use multiple collection methods during same accession – is more often linking to the ordered tests for the visit – so may be create this element: specimen.RelatedOrderIdentifier as Attribute, with datatype string, cardinality 0..\*, definition: The alphanumeric sequence that defines the specimen that are collected to fulfill a specific order = mark as approved
	+ Specimen.Condition – should be able to record more than one condition, so change cardinality to 0..\* and add the following text: For example hemolyzed, thawed, lipemic etc This is specifically allowed to repeat, in case more than one condition needs to be captured. = mark as approved

Call adjourned 5:01 PM ET

* StorageEquipment Parameter = Kathy working on this one – she said descriptions all there, so need to check if anything left
* SpecimenProcessingActivity
* SpecimenCollectionProcedure
* Performer
* Specimen
* SpecimenProcessingActivity
* SpecimenCollectionProcedure
* Performer
* Datatype change for identifier (add type code into the datatype)

Some items from prior minutes, so check on these:

PerformedAdministrativeActivityVarianceReasonCode – proposed source change to add to SpecimenProcessingActivity and

specimenCollectionProcedure, IF we have a protocolReference (or SOP reference) there as well – yes will add that

PerformedAdministrativeActivityVarianceTypeCode - – proposed source change to add to SpecimenProcessingActivity and

specimenCollectionProcedure, IF we have a protocolReference (or SOP reference) there as well

TargetAnatomicSiteConditionCode – do we need this?

Subject.performingSpecimen – how should we represent when tissue from a tissue bank is a specimen – we think it would still fall under person, only de-identified, because currently our material is ONLY from non-living subject

Subject.performingSpecimen: find out if we need this

Specimen.containingSpecimenCollectionGroup – and related attributes

1. Resources:
	1. Link to BRIDG model: <http://bridgmodel.nci.nih.gov/files/BRIDG_Model_4.0_html/index.htm> - chose VIEW:BSP - biospecimen
	2. Link to Specimen DAM: <http://wiki.hl7.org/index.php?title=Specimen> – scroll to bottom for image