

HL7 Clinical Genomics Weekly Call - Jan 3, 2017

Attendees

1. Bob Milius - NMDP - bmilius@nmdp.org
2. JD Nolen - Cerner - [johndavid.nolen@cerner.com](mailto: johndavid.nolen@cerner.com)
3. Joel Schneider - NMDP - [jschneid@nmdp.org](mailto: jschneid@nmdp.org)
4. Bob Freimuth - Mayo Clinic - Happy New Year (It can't possibly be any more weird than 2016)
5. Kevin Power - Cerner - [kpower@cerner.com](mailto: kpower@cerner.com)
6. Jeremy Warner - Vanderbilt - [jeremy.warner@vanderbilt.edu](mailto: jeremy.warner@vanderbilt.edu)
7. Xin Liu - BCH - [xinliu215@gmail.com](mailto: xinliu215@gmail.com)
8. Shannon Lu - NLM - [shennon.lu@nih.gov](mailto: shennon.lu@nih.gov)
9. Clem McDonald - NLM - [clemmcdonald@mail.nih.gov](mailto: clemmcdonald@mail.nih.gov)
10. David Kreda - HMS - [david.kreda@gmail.com](mailto: david.kreda@gmail.com)
11. Joseph Kane - Epic - [jkane@epic.com](mailto: jkane@epic.com)
12. Perry Mar - Partners HealthCare System - [pmar@partners.org](mailto: pmar@partners.org)
13. Joe Quinn - Optum - [joseph.quinn@optum.com](mailto: joseph.quinn@optum.com)
14. Jonathan Holt - SeqTechDx - [jholt@seqtechdx.com](mailto: jholt@seqtechdx.com)
15. Andrea Pitkus - IMO - [-apitkus@imo-online.com](mailto: -apitkus@imo-online.com)

Discussion

- **Brief reports from external efforts (discussion only if needed)**
 - GA4GH
 -
 - National Academies
 - No updates - JD Nolen
 - Clingen/Clinvar
 - No updates due to holidays (Bob F)
 - GA4GH Variant Modeling Collaborative (VMC)
 -
 - NHGRI
 -
- **Upcoming calls through next WGM in San Antonio**
 - Jan 3 (today) - V2/LRI summary reconciliation spreadsheet distribution; WGM prep
 - Jan 10 - V2/LRI block vote on subgroup recommendations; WGM prep
- **FHIR - update (if any)**
 - *nothing new*
- **V2/LRI update**
 - summary uploaded to Clinical Genomics document center:
 - <http://www.hl7.org/documentcenter/public/wg/clingenomics/2017%201%2003-%20HL7%20V2%20CG%20IG%20September%20-%20Merged%20Reconciliation%20Comments.xls>
 - **LRI** - Ballot Close Date: Monday, January 9, 2017
- **WGM preparation**
 - Agenda
 - http://wiki.hl7.org/index.php?title=File:HL7_WGM_Jan2017_-_Clinical_Genomics_Agenda.docx
 - Working session: discussion captured in agenda document

- Tue Q2:
 - Domain Analysis Model (DAM) update
 - Next iteration of DAM?
 - Workflow issues?
 - Clem – need to clarify serological and where it’s discussed
 - David - may need to expand scope?
 - Clem – don’t start with new, but just tune this one
 - Tue Q4:
 - V2
 - More ballot reconciliation
 - Discussion of Need to able to have revision of genetic results
 - Do we need a Next version LRI/V2?
 - Wed Q3:
 - Information Modeling – overview of work;
 - Update group with subgroup work – model
 - Talk about PSS for IM
 - VMC – variant model collaborative review
 - Decision Making Process
 - <https://docs.google.com/document/d/18ZxNAjMukUKXxbNPRtRdjytMCvnRns4srIDe0EBs0FI/edit>
 - short url = <http://bit.ly/2ikjXiV>
 - SWOT
 - https://docs.google.com/document/d/1zFUzRYLfCmrnThBU8xXVS_JiScDACBi13tzFJep751k/edit
 - short url = <http://bit.ly/2iklUw0>
- Other
 - Co-chair elections
 - We’ll be electing **TWO** co-chairs at the WGM in San Antonio. If you aren't able to attend, but still want to vote, you can email your absentee vote to **nominations@HL7.org NO LATER than 12:00 p.m. ET on Wednesday, January 11, 2017.**
 - Per Section 09.02.06.03 of the GOM, *absentee ballots will only be accepted from current individual members and the designated voting representatives of current organizational or Affiliate members who are subscribers of a Work Group’s primary list server but who are not able to attend the WGM. Write-in candidates will be accepted at the meeting*
- Chat
 - none