

## HL7 Clinical Genomics Weekly Call - Nov 22, 2016

### Attendees

1. Bob Milius - NMDP - [bmilius@nmdp.org](mailto:bmilius@nmdp.org) (presiding co-chair)
2. Jonathan Holt - SeqTechDx - [jholt@seqtechdx.com](mailto:jholt@seqtechdx.com)
3. Kevin Power - Cerner - [kpower@cerner.com](mailto:kpower@cerner.com)
4. Joel Schneider - NMDP - [jschneid@nmdp.org](mailto:jschneid@nmdp.org)
5. Tianlong Chen - USTC - [wiwjp619@gmail.com](mailto:wiwjp619@gmail.com)
6. Joseph Kane - Epic - [jkane@epic.com](mailto:jkane@epic.com)
7. Amnon Shabo (Shvo) - Philips - [amnon.shvo@gmail.com](mailto:amnon.shvo@gmail.com)
8. Yi Wang - USTC - [panzer.wy@gmail.com](mailto:panzer.wy@gmail.com)
9. Xiao Luo - USTC - [l.xander.233@gmail.com](mailto:l.xander.233@gmail.com)
10. Clem McDonald - NLM -
11. Perry Mar - Partners HealthCare System - [pmar@partners.org](mailto:pmar@partners.org)
12. Xin Liu - BCH - [xinliu215@gmail.com](mailto:xinliu215@gmail.com)
13. David Kreda - [david.kreda@gmail.com](mailto:david.kreda@gmail.com)
14. Andrea Pitkus - IMO [apitkus@imo-online.com](mailto:apitkus@imo-online.com)
15. Deven Atnoor - [datnoor@gmail.com](mailto:datnoor@gmail.com)
16. Eric Whitebay
17. Joseph Kane

### Discussion

- Minutes approval
  - [http://wiki.hl7.org/index.php?title=File:HL7\\_CG\\_20161115.pdf](http://wiki.hl7.org/index.php?title=File:HL7_CG_20161115.pdf)
  - motion - Bob, second David K
  - discussion - no
  - abstains - Joseph, Kevin Power,
  - nays - none
  - yeas - rest
- Brief reports (skip this week)
  - In the future, email individuals to update this section ahead of time
  - only bring up for discussion if there are any questions about this.
- V2 "Lite" has become preparation for LRI V2 ballot preparation
  - LRI PSS update
    - Bob M shared the new LRI PSS by OO
    - summarized here:  
<http://www.hl7.org/Special/committees/clingenomics/projects.cfm?action=edit&ProjectNumber=1294>
  - David - two types of specifications that we CG and HL7 is actively involved in: V2, FHIR,
    - also CDA, V3 message Pedigree
    - Brett - FHIR->V3->FHIR?
  - David - is managing V2 as complicated as FHIR?
    - Bob M - it's different, FHIR has more technical specifications associated with it, V2 is a Implementation Guide, a document to read.
  - Bob M - presented slides with ideas on how to formalize process going forward

- [http://www.hl7.org/documentcenter/public/wg/clingenomics/HL7CG\\_proposed\\_process.pdf](http://www.hl7.org/documentcenter/public/wg/clingenomics/HL7CG_proposed_process.pdf)
  - David - good starting point, but we need more detail; V2 and FHIR take up most of our time; need clear understanding of what happens at the group-wide calls
  - Jon - need to look at high level as well
    - Bob M - IM group looking at high level concepts
    - David - how has IM affected FHIR and V2? it hasn't
    - Joel - IM is not only harmonizing HL7, but also other external groups.
    - Perry - IM very productive, issues come up and we're educating other on issues and dealing with them
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  - FHIR Ballot reconciliation update
    - FHIR reconciliation spreadsheet
      - Short url - <http://bit.ly/2gaglig>
      - Long url - <https://docs.google.com/spreadsheets/d/17fqBTWkrlvPysZXlhlhLX4proKpv95inFQGVNO54axk/edit#gid=0>
    - FHIR Subgroup Meeting minutes:
      - Short url - <http://bit.ly/2gvRSbq>
      - Long url: [https://docs.google.com/document/d/1FGCQRtxJKyHhnC1uB\\_t4sJZ9yXbLMGOqPXHPr5tSLLQ/edit?pref=2&pli=1#heading=h.ntntuef6iwzkDAM](https://docs.google.com/document/d/1FGCQRtxJKyHhnC1uB_t4sJZ9yXbLMGOqPXHPr5tSLLQ/edit?pref=2&pli=1#heading=h.ntntuef6iwzkDAM)
    - David gave update: voted to accept current disposition
      - Perry - germline question partly resolved
      - Amnon's comment - voted non-persuasive
        - Bob - agreed out of scope for this STU3, but could be an active development option for STU4 - have side by side options and do a bake-off to determine what STU4 should be
      - David K - Amnon needs an opportunity to present his ideas for future development
        - Jon - motions that Amnon given a full hour at a Tues call to discuss his FHIR comment, date to be determined, but soon.
          - David K - move it past next week to give time to prepare, including questions to Amnon to address
          - motion - Jon; David - second
          - abstain - none
          - nay - none
          - yea - motion carries
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- Clem - thought some of his own comments weren't yet addressed
- DAM
  - <https://docs.google.com/document/d/1BrpqPbgUCvtrPvRdf-oikK0pHiKUxo5QdZahxX9Z7dU/edit>
  - Closed for comments/questions, but Bob F had a lot of questions and want to open it up for him to add input.
  - Added after the meeting, notes by David Kreda:
  - **DAM Update**
    - *CGWG Group Edits – Comments and Mark-Ups closed*
    - *Target for final vote-ready version: Tuesday, December 6, pre-meeting*
      - *Suggest a 20-minute slot at top of that call [if call not devoted to Amon's Presentation] to walkthrough key changes due to group input*
    - *1 week to read and vote, so decision by Tuesday, December 13*
    - *Materials will include:*
      - *Google doc - with mark-ups and comments – so people can see changes (exclusive of formatting, typos, doc naming, table of contents, etc.)*
      - *Word doc (and/or PDF) would be clean = The vote candidate*
    - *Should emphasize this DAM is about “Clinical Sequencing” – not “Omics”*
    - ***Proposed to allow, if CGWG group was amenable, that Bob Freimuth would be asked to provide answers to his own DAM comments in the next week such that and DAM editors might yet be able to review and, if sensible to them, meld them into the document.***
    - *RESTATING: Would expect a “by poll” vote up or down week of December 13*
      - *If YES gets published PDF on site, disseminated*
      - *If NO, figure out what went wrong!*
- Decision Making Processes
  - [http://www.hl7.org/documentcenter/public/wg/clingenomics/HL7\\_WG\\_DMP\\_v3.0\\_CG\\_v2\\_09172014%20final.doc](http://www.hl7.org/documentcenter/public/wg/clingenomics/HL7_WG_DMP_v3.0_CG_v2_09172014%20final.doc)
- Other
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