HL7 Clinical Genomics Weekly Call - Oct 18, 2016

Attendees

- 1. Bob Milius NMDP bmilius@nmdp.org
- 2. Kevin Power Cerner kpower@cerner.com
- 3. Tianlong Chen USTC wiwjp619@gmail.com
- 4. Brett johnson bch icanbrj@gmail.com
- 5. JD Nolen Cerner jondavid.nolen@cerner.com
- 6. Yi Wang USTC panzer.wy@gmail.com
- 7. Amnon Shabo (Shvo) Philips amnon.shvo@gmail.com
- 8. Jonathan Holt SeqTechDx iholt@seqtechdx.com
- 9. Mollie Ullman-Cullere OM1 mollie.ullmancullere@gmail.com
- 10. Joseph Kane Epic jkane@epic.com
- 11. Bob Freimuth Mayo Clinic freimuth.robert@mayo.edu
- 12. Bret Heale Intermountain Healthcare bheale@gmail.com
- 13. Clem McDonald National Library of Medicine -
- 14. Andrea Pitkus Intelligent Medical Objects apitkus@e-imo-online.com
- 15. Perry Mar Partners HealthCare System pmar@partners.org
- 16. Terry McDonnell Syapse terrym@syapse.com

Discussion

- Brief reports
 - ClinGen/ClinVar
 - Bob F continued work on developing examples for allele interpretation
 - o GA4GH
 - Oct 16-18 in Vancouver
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 - National Academies
 - JD nothing new
 - Other
 - nothing new
- Interim Co-chair
 - https://docs.google.com/forms/d/1t0GAOqtd8JkOShywD9JvaOihQr3VInn5Ui8OpXF54JI
 - 2016-10-19: Co-Chair call for nominations opens
 - 2016-11-18: Co-Chair Nominations Close at 5:00 pm Eastern
 - 2016-12-09: Co-Chair election statements due to HQ

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- Ballot Reconciliation updates
 - Ballot Reconciliation Howto
 - http://wiki.hl7.org/index.php?title=Reconciliation HowTo
 - o FHIR
 - Gil traveling (GA4GH & ASHG)
 - o V2 Lite
 - discussions with Kevin re complexity

- maybe combine sections
- another issue is there anyone who will test/adopt?
 - Mollie may be necessary for to ballot again
 - need two orgs to adopt for normative ballot
 - Joseph Kane Epic very interested in the standard, but have not prioritized so can't commit in the near term; strong interest
 - Kevin Cerner similar answer; would like to find a client interested; will
 check; not the one to make the decision but can start the conversation
 - Mollie reach out to Don Rule?
 - Clem may talk to Kevin/Joseph separately this week re complexity
 - Jon should get a lab to generate the data for Cerner/Epic to store
 - Mollie Don Rule would be perfect for that
 - o Clem Cerner/Epic probably knows labs
 - Kevin that's where the clients would come into play; would like to get them involved
 - Brett are we coordinating these standards efforts with labs in the sense that we are offering all these solutions to vendors (V2, V3, FHIR, CDA)?

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- Can we add that solution here? Base64 JSON payload was FHIR in V2 message?
- Clem tried to include FHIR discussion in V2 Lite IG but was told that it didn't belong there
- Andrea from a Lab perspective, FHIR not mature enough for labs; has to be CLIA compliant
- Jon in San Antonio ~2 yrs ago someone GeneDx was at the meetings; we should reach out them his name was Christopher Lauricella [mailto:chrisgc@genedx.com]
- Also Marc Bellar from Vanderbilt created a very mature pharmacogenomics solution using HL7 V2
- Clem issues about multiplicity of databases
 - cosmic three different databases
 - ncbi parallel but not clear
 - bob m to send Clem the imgt/hla database

Other

- Bob M terminology services
 - http://hl7.org/fhir/2016Sep/terminologies-systems.html
 - Jon need to identify what the different services do
 - Bret if we have a terminology service then we bind to the service
 - o do we want to constrain?
- Bret how will it be developed and maintain who will pay it?

Recorded Chat

FHIR compliant terminology server