QIBA FDG-PET Biomarker Committee Call

06 January 2017 at 9 AM CT Draft Call Summary

In attendance: RSNA

Uma Ranjan, MSc, PhD

Scott Wollenweber, PhD (co-chair)Jeff Kolthammer, PhDAnne Smith, PhDJoe KoudelikOrest Boyko, MD, PhDMartin Lodge, PhDMitsuaki Tatsumi, MDJulie LisieckiJanice Campbell, PhDNancy Obuchowski, PhDTimothy Turkington, PhDHoward Higley, PhDAmy Perkins, PhDRichard Wahl, MD

Moderator: Dr. Wollenweber

Discussion Topics included:

Paul Kinahan, PhD

RSNA 2016 (QIBA Breakout Session) Discussion Overview

- Profile status and next steps beyond the Technically Confirmed phase
- Extension of the claims scope under consideration to address mixed reviews regarding asymmetrical repeatability

Jeffrey Yap, PhD

- Test-retest groundwork studies may be necessary to update the Profile
- Other issues discussed included
 - Point spread function
 - Lean body mass
 - Time of flight
 - Asymmetric claim structure and achieving consensus on this topic
 - Next steps to achieve Profile Clinical Confirmation via crowd-sourcing (i.e. multiple sites to submit testretest data)
 - O How the Profile will address treatment response criteria needs additional discussion

Profile Status

- The Profile is currently at the <u>Technically Confirmed</u> stage and Dr. Kinahan has posted the <u>most recent version of</u> <u>the Profile</u> (v1.13) to the QIBA wiki
- Moving to the next phase will include the use of test-retest studies and Dr. Subramaniam's guidance will be needed
- Complications that may cause skewing of the data are related to asymmetric limits and differences in the ways that test-retest studies are analyzed
- These issues will need to be resolved by the group in order to have consensus moving forward
- Feedback from test sites would be ideal; however, determining logistics may prove difficult, as a coordinating site manager will be needed to address Technologist questions while implementing the Profile
- Possibly utilizing a divide-and-conquer approach by scanner expert may be helpful; however, this remains a challenge

Checklist Update

- The final version of the checklist has been included as an Appendix for the Profile
- The checklist has been broken down into two checklists for more tailored use, posted to the QIBA wiki:
 - A site checklist
 - A scanner-specific checklist for vendors
 - o Comments or edits may be sent to: timothy.turkington@duke.edu and kinahan@uw.edu

Action items for next call (February 3rd):

- All are asked to review the <u>most recent version of the Profile</u> posted on the QIBA wiki prior to the next call
- If available, Dr. Subramaniam to provide details regarding the test-retest studies and suggestions on moving the Profile toward the Clinically Confirmed stage

Notes from RSNA QIBA FDG-PET/CT Biomarker Committee Breakout session

Date: 30 Nov 2016

Reviewed on Biomarker Committee WebEx 2017-01-06

Provided by: Paul Kinahan, PhD

Summary of topics raised in discussion:

- 1. We should review asymmetric claim at next BC call
 - What are implications for response criteria (i.e. PERCIST)?
- 2. We should review checklists on next call
 - Tim to lead
 - Should we ask technologists to review?
 - Good opportunity to refresh everyone on current Profile
- 3. We should synchronize profile versions
 - (not sure what this means, perhaps w.r.t. with the standard QIBA form?)
- 4. How do we get to clinical confirmation?
 - There was agreement this is good, but not clear if it is feasible
 - can also discuss a crowd-sourced trial for clinical confirmation
 - Leveraging iCROs and or moonshot to add re-test studies
 - see funding and/or support from ECOG-ACRIN
- 5. Claim should be reassessed for LBM, PSF and TOF
 - Could start with IQ phantom data from harmony project comparing w/ w/o PSF TOF
- 6. Other tracers: FLT and Ga68-PSMA.
- 7. What about PET/MR?
- 8. Note that Technically Confirmed Profile (version 1.13) is on Wiki site http://qibawiki.rsna.org/index.php/Profiles

Nuclear Medicine WebEx Schedule:

Jan 13: ad hoc Amyloid PET Physics call

Jan 20: Amyloid BC

Jan 27: NM Leadership / TBD

Feb 03: FDG-PET BC