

QIBA Perfusion, Diffusion, and Flow (PDF-MRI) Biomarker Committee: 2017 Overview



Michael A. Boss¹, John Kirsch², Mark A. Rosen³, Amita Shukla-Dave⁴, Thomas L. Chenevert⁵, Dariya Malyarenko⁵, Hendrik Laue⁶, Caroline Chung⁷, Krishna Nayak⁸, Mark Shiroishi⁸, Kirt Gill⁸, Lillian Lai⁸, James Provenzale⁹, Walter Schneider¹⁰, Ona Wu², Bradley Erickson¹¹, Xavier Golay¹², Rik Achten¹³, Matthias Günther⁶, Nancy Obuchowski¹⁴, Alexander R. Guimaraes¹⁵, Daniel Sullivan⁸, and Edward F. Jackson¹⁶

¹National Institute of Standards and Technology, ²Massachusetts General Hospital, ³University of Pennsylvania, ⁴Memorial Sloan-Kettering, ⁵University of Michigan, ⁶Fraunhofer-MEVIS, ⁷MD Anderson Cancer Center, ⁸University of Southern California, ⁹Duke University, ¹⁰University of Pittsburgh, ¹¹Mayo Clinic, ¹²University College London, ¹³Ghent University, ¹⁴Cleveland Clinic, ¹⁵Oregon Health & Science University, ¹⁶University of Wisconsin

SUMMARY AND GOALS OF THE PDF-MRI BIOMARKER COMMITTEE

The RSNA QIBA Perfusion, Diffusion, and Flow MRI Biomarker Committee (PDF-MRI BC) is composed of scientists representing imaging device manufacturers, image analysis laboratories, biopharmaceutical industry, academia, government research organizations, imaging core labs, and professional societies. The goal of the PDF-MRI BC is to define technical performance standards (QIBA Profiles) for data acquisition, data processing, and quality control procedures that enable consistent and reliable quantitative imaging biomarkers for assessment of physiologic measures related to perfusion, diffusion, and/or blood flow in normal and abnormal tissues. There are now five Profiles in development, with the addition of ASL in 2017, each with a corresponding task force. 2017 also saw the release of the DWI Profile for public comment.

translational research, and pharmaceutical studies continues to grow. Thus, there appears to be a promising future of these techniques for both clinical research and in routine clinical practice, particularly in the era of precision medicine. However, in order to fulfill this promise, it is essential that common quantitative endpoints are used and that results are reproducible and unbiased across imaging platforms, clinical sites, and time.

The efforts of the PDF-MRI BC are motivated by the emergence of perfusion/diffusion/flow quantitative imaging biomarkers as a means of diagnosing pathologies, staging disease, and evaluating responsiveness to therapy. Despite variance in imaging techniques, parameter choices, vendor specifications, and analytic methods, the application of these physiologic measures in clinical medicine,

Summary of PDF-MRI Biomarker Committee Goals
 To develop consensus technical performance standards (QIBA Profiles), based on existing literature and and groundwork projects, regarding the appropriate data acquisition, data processing, and quality control procedures necessary to provide reproducible quantitative MR imaging biomarker measures of normal and diseased tissues.

ORGANIZATIONAL UPDATES, PROFILE AND CONFORMANCE PROGRESS

PDF-MRI

Co-Chairs: Dan Barboriak, Michael Boss, John Kirsch

DCE Task Force

Co-Chairs: Caroline Chung, Hendrik Laue

The main clinical impact DCE is the identification and staging of various tumors. As a quantitative biomarker, K^{trans} is of special interest in treatment monitoring of antiangiogenic therapies.

The DCE TF developed preliminary claims for K^{trans} for two different tumor types:
Claim 2a: A measured change of K^{trans} in brain tumors (GBM) of 21.33 % or larger indicates that a true change has occurred with 95 % confidence.
Claim 2b: A measured change of K^{trans} in prostate tumors of 50.7 % or larger indicates that a true change has occurred with 95 % confidence.
 The claim definitions for prostate and brain are based on test-retest data from Alonsi¹ et al., 2010 and Jackson² et al., 2003, respectively. The claims will be updated with new, soon-to-be-published test-retest data.

¹ Jackson, A., Jayson, G. C., Li, K., Zhu, X. P., Checkley, D. R., Teiser, J. J., L., & Wadden, J. C. (2003). Reproducibility of quantitative dynamic contrast-enhanced MRI in newly presenting glioma. British Journal of Radiology, 76(903), 153–162. <http://dx.doi.org/10.1259/bjr/70652746>
² Alonsi, R., Taylor, N. J., Stirling, J. J., d'Arcy, J. A., Collins, D. J., Saunders, M. I., et al. (2010). Reproducibility and correlation between quantitative and semiquantitative dynamic and intrinsic susceptibility-weighted MRI parameters in the benign and malignant human prostate. Journal of Magnetic Resonance Imaging - JMRI, 32(1), 155–164. <http://doi.org/10.1002/jmri.22215>

Brain: Comprehensive Search Results: 2080 papers Papers reviewed for profile content: 653 papers	Head and Neck: Comprehensive Search Results: 672 papers Papers reviewed for profile content: 127 papers	Prostate: Comprehensive Search Results: 395 papers Papers reviewed for profile content: 202 papers
Outcome measures K^{trans} , V_e , V_p , K_{ep} , K_i , AUGC, AUC, CBV , CBF , MTT, Max SI, max SI (ratelapse)	Outcome measures K^{trans} , V_e , V_p , K_{ep} , K_i , AUGC, AUC, CBV , CBF , MTT, Max SI, max SI (ratelapse), peak height, time to peak, relative max enhancement, relative washout ratio, vascular permeability, leakage space, contrast index, tumor blood volume, tumor blood flow, VWF	Outcome measures K^{trans} , V_e , V_p , K_{ep} , K_i , AUGC, AUC, CBV , CBF , MTT, Max SI, max SI (ratelapse), wash-in/wash-out
Field strength 0.5T: 2 (on MS); 1.5T: 29; 3T: 21	Field strength 1.5T: 55; 3T: 47; 4.7T: 2; 8.4T: 1	Field strength 1.5T: 79; 3T: 94; 1.5T + 3.0T mixed: 11
Acquisition parameters for brain tumors TR: 3.8-11.9ms (majority 3.8-6ms); TE: 1.4-2 ms FA: 8-30° Temporal resolution: 1-60 s (majority 1.25-6.5 s) Slice thickness: 2.5-6 mm Overall scan time: 1.5 – 12 min (majority 2.8-6.5 min)	Acquisition parameters for H&N tumors TR: 2.3-20 ms for GRE (majority 2.3-9 ms); TE: 0.9-4.2 ms for GRE Temporal resolution: 0.5-10 s (majority between 0.5-10 s) Slice thickness: 1-10 mm Overall scan time: < 1-10 min (majority 1.5-6 min)	Acquisition parameters for prostate tumors TR: 3.8-11.9 (majority between 3.8-6ms) TE: 1.4-2 ms FA: 5-70° Temporal resolution: 1-90 s (majority 1.25-6.5 s) Slice thickness: 2.5-6 mm Overall scan time: 1.5 – 12 min (majority 2.8-6.5 min)

The DCE task force has recently completed extensive literature searches in three main body sites, informing our profile with regards to scan parameters and reproducibility metrics.

ASL Task Force

Co-Chairs: Xavier Golay (WG1), Rik Achten (WG2), Matthias Günther (WG3)

The QIBA Steering Committee approved the formation of the ASL task force, in conjunction with ESR's EIBALL. This task force has organized 3 working groups: WG1 focuses on statistics and technical definition of the claims; WG2 on the clinical aspects of the claims; WG3 addresses phantom usage and sequence parameters.

DTI Task Force

Co-Chairs: Jim Provenzale, Walt Schneider

The DTI TF is in the process of drafting its Profile. The TF has conducted an initial literature review regarding the reproducibility of several different anisotropic diffusion metrics, including fractional anisotropy and mean diffusivity. In parallel, the TF has scanned a phantom with anisotropic components consisting of hollow polypropylene fibers with inner and outer diameters on the scale of micrometers to determine optimal scan protocols and reproducibility of metrics with existing clinical protocols (see groundwork project).

DSC Task Force

Co-Chairs: Brad Erickson, Ona Wu

The clinical application of the DSC profile is characterization of brain tumors. The TF has recently completed two ground work projects to develop a physical phantom and DRO generation tool. A literature search of relevant articles yielded 2472 articles in 2015; an additional 562 articles have been added since. The draft profile is presently being updated to the latest QIBA profile template.

DWI Task Force

Co-Chairs: Michael Boss, Tom Chenevert

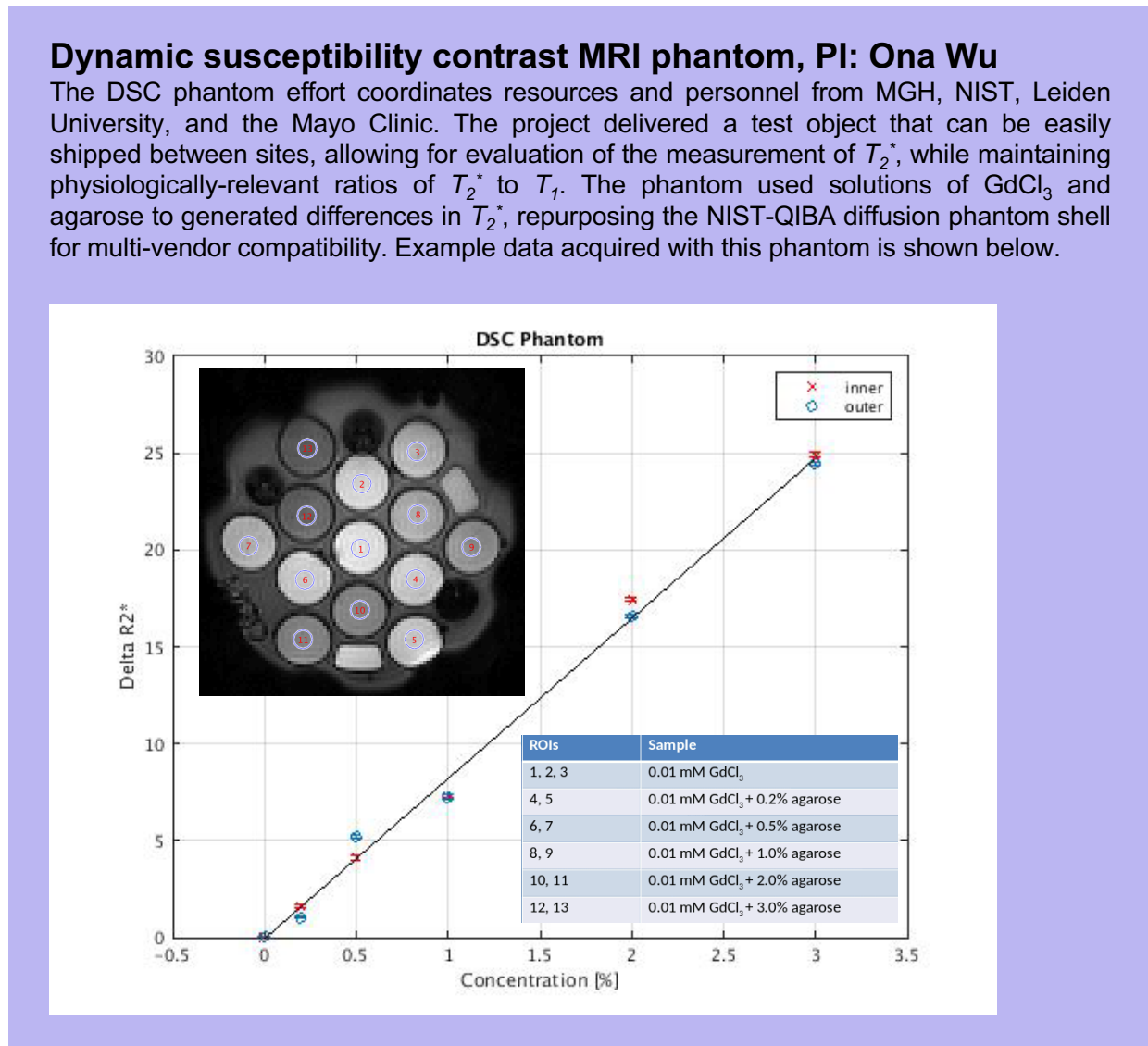
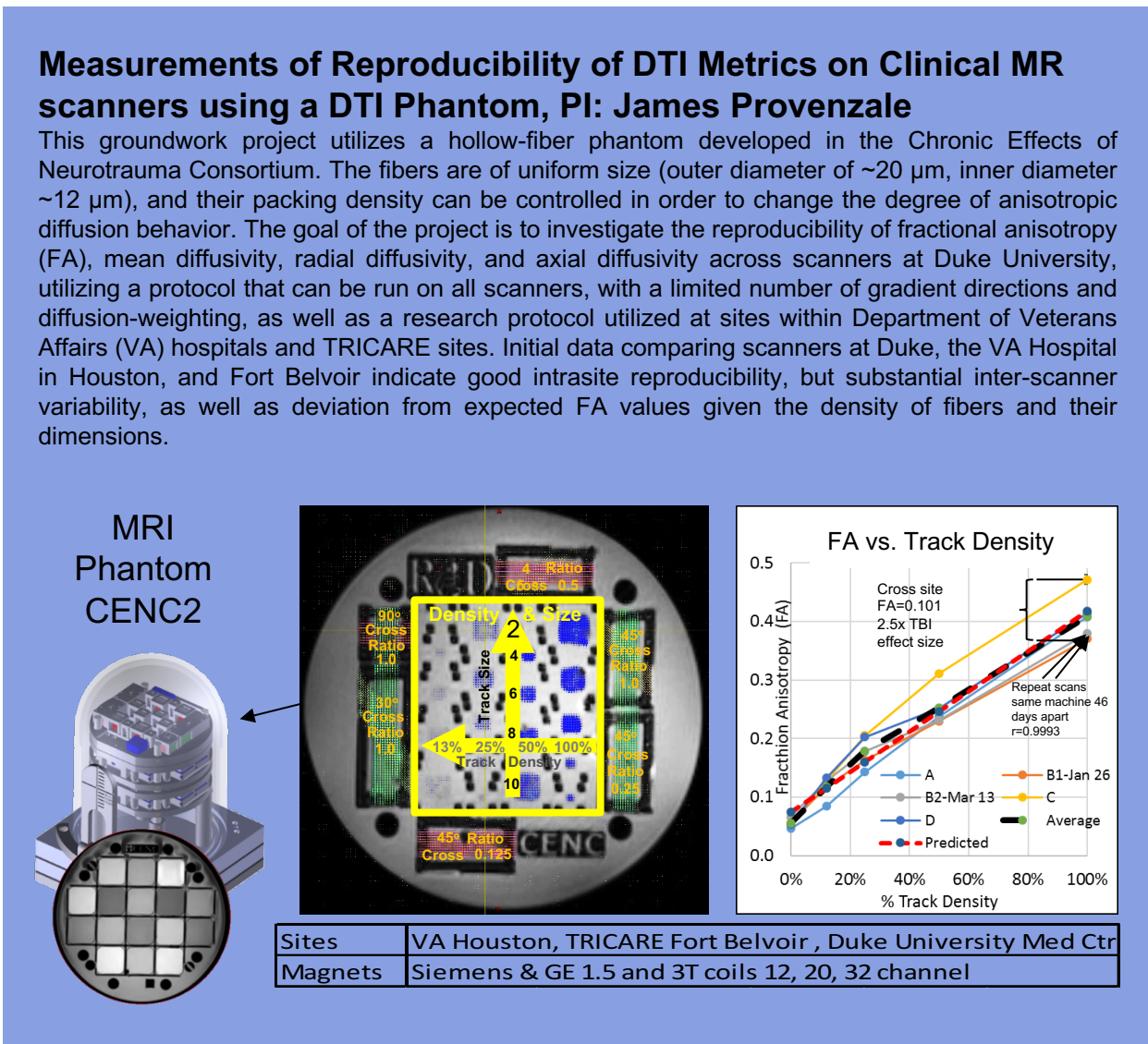
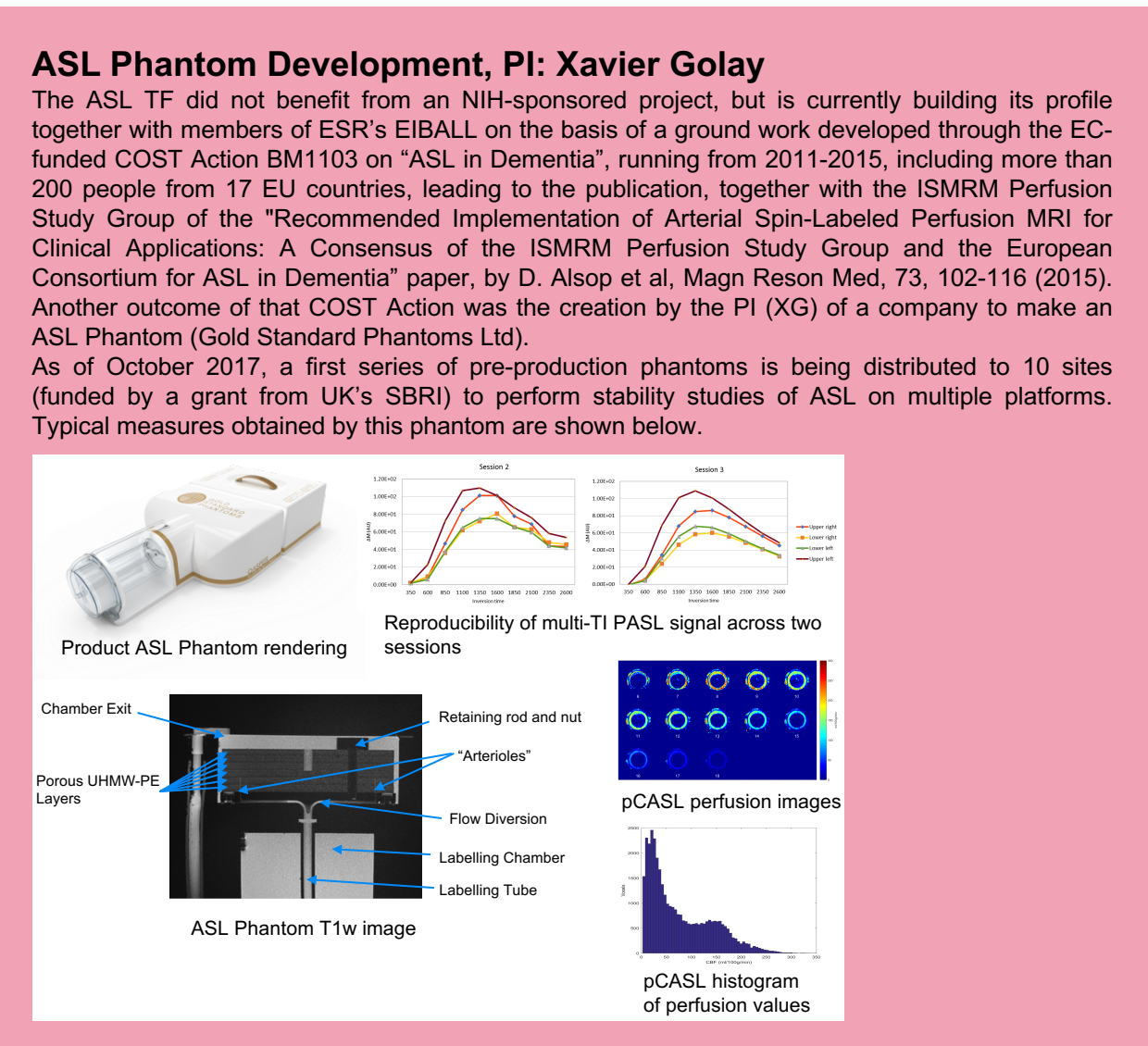
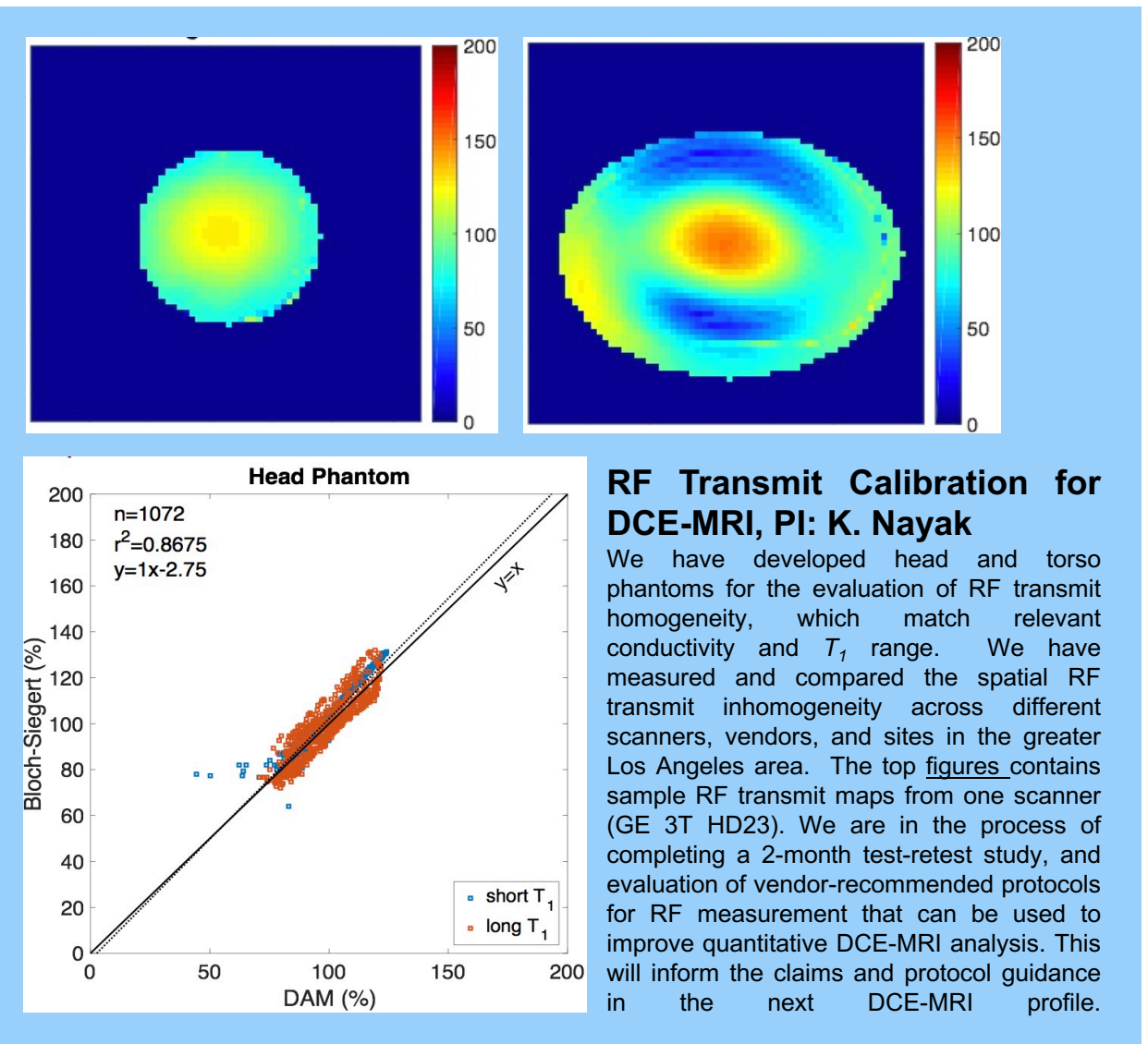
The DWI TF successfully released v1.0 of its profile for Public Comment, and began work on a checklist to facilitate profile use. The claims for prostate are shown below:
Claim 3a: A measured change in the ADC of a prostate lesion of 47% or larger indicates that a true change has occurred with 95 % confidence.
Claim 3b: A 95% CI for the true change in ADC of a prostate lesion is given below, where Y_1 and Y_2 are the ADC measurements at the two time points:
 $(Y_2 - Y_1) \pm 1.96 \times \sqrt{(Y_1 \times 0.17)^2 + (Y_2 \times 0.17)^2}$
 The TF received about 100 comments, and created a spreadsheet to track user-submitted proposals and the final TF-approved resolution; examples are shown below.

Name	New Line	Old Line	Section	Issue	Proposal	Priority	Resolution
Anon	896	344	3.6.2	stated TE does not match PI-RADS	TE = 90 ms	H	ADTLCDM: Change to "Acceptable <= 90ms" (will match PI-RADS)
Anon	896	344	3.6.2	stated maximal b-value does not match PI-RADS	maximal b-value of at least 1,400 (stated in PI-RADS in numerous locations); 500-1,000 is not sufficient	H	ADTLCDM: Leave unchanged to agree with primary pubs and emphasis on ADC (not DWI)

GROUNDWORK PROJECTS

Groundwork projects are specific investigational activities, often funded by the National Institute of Biomedical Imaging and Bioengineering (NIBIB), to aid in the implementation of PDF-MRI BC areas of investigation. Groundwork projects seek to provide material resources (e.g., phantoms, digital reference objects, software) to aid investigators seeking to obtain reproducible quantitative imaging biomarkers, or to provide data from field testing of the robustness of quantitative imaging biomarkers when applied to data acquisition (phantom or human data) in a multi-

institutional framework. Groundwork projects are not themselves designed to test specific aspects of biological or pharmaceutical phenomena reflected by the quantitative imaging biomarkers. Rather, they enable development of tangible products to aid in Profile development or revision, to allow demonstration of conformance with such Profiles, and to demonstrate robustness of the quantitative imaging biomarkers(s) in practice. Recently groundwork projects in PDF include:



A web-based tool for creating DSC Digital Reference Objects, PI: Brad Erickson
 On the QIBA website (<http://www.rsna.org/QIDW/>)

QIDW Data Inventory

- COPD/Asthma Phantom
- HEI CT Digital Reference Object
- FMRB Digital Reference Object
- USC-SIEMENS Digital Phantoms
- DWI Phantom
- QIBA DCE-MRI DRO
- QIBA DCE-MRI WG

QIDW Tools

- DSC DRO Modeling Website

The DSC DRO website will create digital simulations of DSC perfusion acquisitions. The user may select 1 of 3 models, and select values for the many different acquisition parameters and assumptions about the imaged tissue. The website will then create a 4D DICOM image that the user can download that simulates those acquisition conditions.

One of 3 models available: The BNI DRO image generator

PUBLICATIONS

1. EM Palacios, AJ Martin, MA Boss, et al. *Towards Precision and Reproducibility of Diffusion Tensor Imaging: A Multicenter Diffusion Phantom and Traveling Volunteer Study*, AJNR 38:537-545; <https://doi.org/10.3174/ajnr.A5025>
2. SJ Hectors, M Wagner, I Corcuera-Solano, et al. *Comparison Between 3-Scan Trace and Diagonal Body Diffusion-Weighted Imaging Acquisitions: A Phantom and Volunteer Study*, Tomography 2:411-420; <https://dx.doi.org/10.18383/tom.2016.00229>
3. D Malyarenko et al. *Toward uniform implementation of parametric map DICOM in multi-site quantitative diffusion imaging studies*, J Med Imag 6:011006; <https://doi.org/10.1117/1.JMI.5.1.011006>
4. DC Newitt, D Malyarenko, TL Chenevert, et al. *Multi-site concordance of apparent diffusion coefficient measurements across the NCI Quantitative Imaging Network*, J Med Imag 5:011003; <https://doi.org/10.1117/1.JMI.5.1.011003>

QIBA Projects and activities have been funded in whole or in part with Federal funds from the National Institute of Biomedical Imaging and Bioengineering, National Institutes of Health, Department of Health and Human Services, under Contract No. **HHSN268201500021C**

