As of 11/28/18, the protocols are finalized here:
https://markvcid.partners.org/imaging-protocols

Please complete a full scan of a human subject using the complete set of core scan types that your site will be performing during the UH3 phase as soon as practicable and upload the anonymized data using Globus. If possible, please scan a patient, which will better help judge the quality of the images; if not a healthy control is acceptable.

To ensure the MarkVCID Data Management system can seamlessly handle data from your site, we request that each site send the coordinating center MRI data following the procedures on https://markvcid.partners.org/imaging-data-collection

This data should be human data, acquired using the protocols for the approved biomarkers that your site will be collecting in the course of the UH3 phase. We also ask this data be anonymized using the DICOM Data Anonymization SOP and sent to us though Globus (see SOP).

Note that you will **not** be registering the patient nor the data for this test case on the website, so you will not have a patient or data ID for these data. Please follow the directory structure and naming convention in the SOP documents with the exception of:

<site>-testdata.zip (e.g., mgh-testdata.zip) for the name of the zip archive and "<site>-patientID" (e.g. mgh-patientID) as the top-level directory name.

(The lower directory levels are named by the scan type using the abbreviations given in the DICOM Data Anonymization SOP.)

The test/exercise allows you to practice the anonymization and data-upload processes with actual data and allows the MarkVCID Data Management team to identify any issues with this process and our handling of data that is specific to your site.

Questions/Comments?
Contact: helmer@nmr.mgh.harvard.edu