SCAN PET Protocol

Largely based on ADNI – meant for compatibility **Scanner qualification Acquisition Protocols** Data uploads **Quality Control Pre-processing Data analysis**

Scanner Qualification

If a scanner is qualified for ADNI, no need for re-qualification

All scanners not previously qualified will need to follow the standard ADNI protocol: Imaging of Hoffman brain phantom (supplied by SCAN) with F18 radioactivity x 2 uploaded for QC

As in ADNI, a single scanner must be used for all imaging in SCAN

PET Acquisition Protocols

	PIB	10-15 mCi	40-60 or 50-70 min frames
Amyloid	Florbetapir	10 mCi	50-70 or 50-60 min frames
	Florbetaben	8 mCi	90-110 min frames
	Flortaucipir	10 mCi	80-100 min frames
Tau	MK6240	5-10 mCi	70-90 or (90-110 possible) min frames
	PI2620	5 mCi	AD: 45-75 min frames Non AD: 0-60 or 30-60
	Other tau tracers	Pending	Pending
	FDG	5 mCi	30-60 min frames

Protocol Modifications/Deviation

Centers wishing to collect novel frames or dynamic data are free to do so, but will be required to upload the standard frames

Scans that do not comply with protocol will not be QC'd, curated, or counted towards center numbers

Imaging protocols for new tracers will be defined by consensus and data

Data Uploads

Image data will be managed by NACC and LONI together

Uploads of images to NACC/LONI

All scans accompanied by an "information sheet" listing acquisition conditions/parameters

Quality Control/Pre-Processing University of Michigan – Bob Koeppe (ADNI Methods)

<u>QC</u>

Subject motion

Check Field of View

Artifacts

Check data against headers and protocol

Pre-processing

Coregister and average frames that pass QC

Standard voxel size/grid

Reorientation/intensity normalization

Smoothing on a per-scanner basis to 8 mm isotropic resolution

Data Analysis

Contemporaneous MRI is desired but not required We will use MRI-free processing pipelines

All numerical data returned to NACC where they can be linked to participant

All QC'd and processed scans to NACC/LONI

Data Analysis: PET

Amyloid PET Regional FreeSurfer based SUVRs Centiloid Conversion of all data Amyloid positivity based on accepted thresholds

<u>Tau PET</u>

Regional FreeSurfer based SUVRs Aggregate summary ROIs (Braak stages, meta-ROI)

<u>FDG-PET (Depending on need)</u> FreeSurfer based SUVRs Atlas-based spatial normalization with calculation of SUVRs in meta-ROI