

## Flortaucipir (AV-1451) processing methods

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### Summary

ADNI flortaucipir regional summary data are updated regularly and uploaded to LONI by our group. Our image analysis pipeline includes the flortaucipir scan and an MPRAGE for each subject that is usually acquired at the same visit as the flortaucipir image. This MPRAGE is segmented and parcellated with Freesurfer (version 5.3.0) to define a variety of regions of interest in each subject's native space. We then coregister each flortaucipir scan to its corresponding MPRAGE and calculate mean flortaucipir uptake within each Freesurfer-defined region. Mean regional uptake can be calculated across several regions of interest (e.g. Braak stage composite regions – see below) and divided by a reference region (cerebellar GM or hemispheric WM) to generate flortaucipir SUVRs.

#### **Are the flortaucipir data in our dataset already intensity normalized?**

Please note that the Stage 3 flortaucipir images as well as the Stage 4, fully pre-processed flortaucipir images (“AV1451 Coreg, Avg, Std Img and Vox Siz, Uniform Resolution”) are SUVR images that have been **approximately** intensity normalized using an atlas-space cerebellar grey matter region defined by Bob Koeppel during his pre-processing procedures (<http://adni.loni.usc.edu/methods/pet-analysis-method/pet-analysis/#pet-pre-processing-container>). These procedures include defining an atlas-space cereb GM region using a coregistered FDG or MPRAGE scan and reverse normalizing this region back onto the native space flortaucipir image. Because this initial intensity normalization carries with it some noise associated with the warping procedures, we defined native-space reference regions (as well as regions of interest) more precisely using Freesurfer. We then replace (e.g. divide out) the initial intensity normalization carried out by Bob Koeppel with a subsequent intensity normalization using our Freesurfer-defined / native space reference regions as described below. Note that the Freesurfer-defined cortical SUVRs listed in our dataset include *only* Bob Koeppel's cerebellar GM intensity normalization, so in order to generate SUVRs that take advantage of our Freesurfer-based reference regions, you have to divide a region of interest SUVR mean (e.g. Braak12) by one of the reference regions we provide in our dataset (e.g. inferior cerebellar grey matter).

### Method

#### ***Acquisition of flortaucipir and MRI image data from LONI***

We download flortaucipir data from LONI in the most fully pre-processed format (series description in LONI Advanced Search: “AV1451 Coreg, Avg, Std Img and Vox Siz, Uniform Resolution”). Each subject's pre-processed flortaucipir image is coregistered using SPM to that subject's MRI image (series description: ADNI 1 scans \*N3;\* and ADNI GO/2 scans \*N3\*”) that was closest in time to the flortaucipir scan. Typically the MRI and PET images are within 3 months, but when a concurrent MRI is not available we use an MRI scan acquired at another visit.

### ***Calculation of flortaucipir SUVR***

We have investigated a number of strategies for quantifying and staging tau using flortaucipir [1-4]. This ADNI UC Berkeley flortaucipir dataset includes a broad set of regional flortaucipir means and their corresponding Freesurfer-defined volumes (in mm<sup>3</sup>). This set includes cortical and subcortical regions of interest and reference regions such as cerebellar grey matter and eroded hemispheric WM. Additionally, we approximate uptake in the anatomical Braak stages [5] by calculating volume-weighted means of groups of FreeSurfer-defined regions, specified in the “Braak ROIs” section.

As described in the box above, flortaucipir SUVRs can be calculated by dividing a region of interest (with or without an adjustment for regional volume) by a reference region.

### ***Flortaucipir Partial Volume Correction***

We also provide a separate dataset with flortaucipir data corrected for partial volume effects using the Geometric Transfer Matrix (GTM) approach [6] as implemented for flortaucipir by Suzanne Baker [1, 2]. The GTM approach we are currently using models all FreeSurfer-defined ROIs (see list below) as well as regions in which off-target binding is common (e.g. choroid plexus) in order to reduce contamination from these regions into neighboring regions of interest.

In order to reduce the influence of off-target flortaucipir binding that has been observed in the dorsal cerebellum, we defined an **inferior cerebellar GM reference region** using the SUIT template [7] (<http://www.diedrichsenlab.org/imaging/suit.htm>) and reverse-normalized this region back to each subject's native space as described in Baker et al. NeuroImage 2017[2].

In our flortaucipir PVC and nonPVC datasets, we use the individual Freesurfer-defined SUVRs and volumes to calculate weighted averages of the following composite regions (Braak I/II, Braak III/IV, Braak V/VI) that approximate the spread of tau as depicted by Braak and Braak [5] and described in Scholl et al. [4] and Maass et al [3].

We recommend normalizing either composite (e.g. Braak) or individual PVC ROI values by a PVCed reference region (e.g. inferior cerebellar grey matter) to ensure standardized units.

### ***Freesurfer-defined region codes for Braak ROIs***

#### **Braak 1 and 2 composite region (Braak12):**

##### **Braak 1**

1006 L\_entorhinal  
2006 R\_entorhinal

##### **Braak 2**

17 L\_hippocampus  
53 R\_hippocampus

#### **Braak 3 and 4 composite region (Braak34):**



**Braak 3**

1016 L\_parahippocampal  
1007 L\_fusiform  
1013 L\_lingual  
18 L\_amygdala  
2016 R\_parahippocampal  
2007 R\_fusiform  
2013 R\_lingual  
54 R\_amygdala

**Braak 4**

1015 L\_middletemporal  
1002 L\_caudantcing  
1026 L\_rostantcing  
1023 L\_postcing  
1010 L\_isthmusing  
1035 L\_insula  
1009 L\_inferiortemporal  
1033 L\_temppole  
2015 R\_middletemporal  
2002 R\_caudantcing  
2026 R\_rostantcing  
2023 R\_postcing  
2010 R\_isthmusing  
2035 R\_insula  
2009 R\_inferiortemporal  
2033 R\_temppole

**Braak 5 and 6 composite region (Braak56):**

**Braak 5**

1028 L\_superior\_frontal  
1012 L\_lateral\_orbitofrontal  
1014 L\_medial\_orbitofrontal  
1032 L\_frontal\_pole  
1003 L\_caudal\_middle\_frontal  
1027 L\_rostral\_middle\_frontal  
1018 L\_pars\_opercularis  
1019 L\_pars\_orbitalis  
1020 L\_pars\_triangularis  
1011 L\_lateraloccipital  
1031 L\_parietalsupramarginal  
1008 L\_parietalinferior  
1030 L\_superiortemporal  
1029 L\_parietalsuperior  
1025 L\_precuneus  
1001 L\_bankSuperiorTemporalSulcus  
1034 L\_tranvtemp  
2028 R\_superior\_frontal



2012 R\_lateral\_orbitofrontal  
2014 R\_medial\_orbitofrontal  
2032 R\_frontal\_pole  
2003 R\_caudal\_middle\_frontal  
2027 R\_rostral\_middle\_frontal  
2018 R\_pars\_opercularis  
2019 R\_pars\_orbitalis  
2020 R\_pars\_triangularis  
2011 R\_lateraloccipital  
2031 R\_parietalsupramarginal  
2008 R\_parietalinferior  
2030 R\_superiortemporal  
2029 R\_parietalsuperior  
2025 R\_precuneus  
2001 R\_bankSuperiorTemporalSulcus  
2034 R\_tranvtemp

#### **Braak 6**

1021 L\_pericalcarine  
1022 L\_postcentral  
1005 L\_cuneus  
1024 L\_precentral  
1017 L\_paracentral  
2021 R\_pericalcarine  
2022 R\_postcentral  
2005 R\_cuneus  
2024 R\_precentral  
2017 R\_paracentral

#### ***PVC input regions***

#### **All Braak regions listed above**

#### **Other non-Braak-related regions used as PVC input**

Choroid plexus: 31, 63  
28 Left-VentralDC  
30 Left-vessel  
60 Right-VentralDC  
62 Right-vessel  
77 WM-hypointensities  
80 non-WM-hypointensities  
85 Optic-Chiasm  
1000 ctx-lh-unknown  
1004 ctx-lh-corpuscallosum  
2000 ctx-rh-unknown  
2004 ctx-rh-corpuscallosum



**Not included in PVC model (set to zero). Note that bone, soft tissue, and CSF outside the brain are omitted and are all implicitly set to zero [2]**

- 4 Left-Lateral-Ventricle
- 5 Left-Inf-Lat-Vent
- 14 3rd-Ventricle
- 15 4th-Ventricle
- 24 CSF
- 43 Right-Lateral-Ventricle
- 44 Right-Inf-Lat-Vent
- 72 5th-Ventricle

#### **Inferior Cerebellar Gray Matter definition**

- 8 Left-Cerebellum-Cortex
- 47 Right-Cerebellum-Cortex

#### **SUIT ROI numbers used for Inferior Cerebellar Gray definition [7]**

Inferior cerebellar inclusion mask: SUIT codes 6, 8-28, 33, 34

Superior cerebellar exclusion mask (bilateral lobules I-VI): SUIT codes 1-5, 7

## **References**

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