



UNIVERSITÀ
DI TORINO

Dipartimento
Neuroscienze
"Rita Levi Montalcini"



IronAD project

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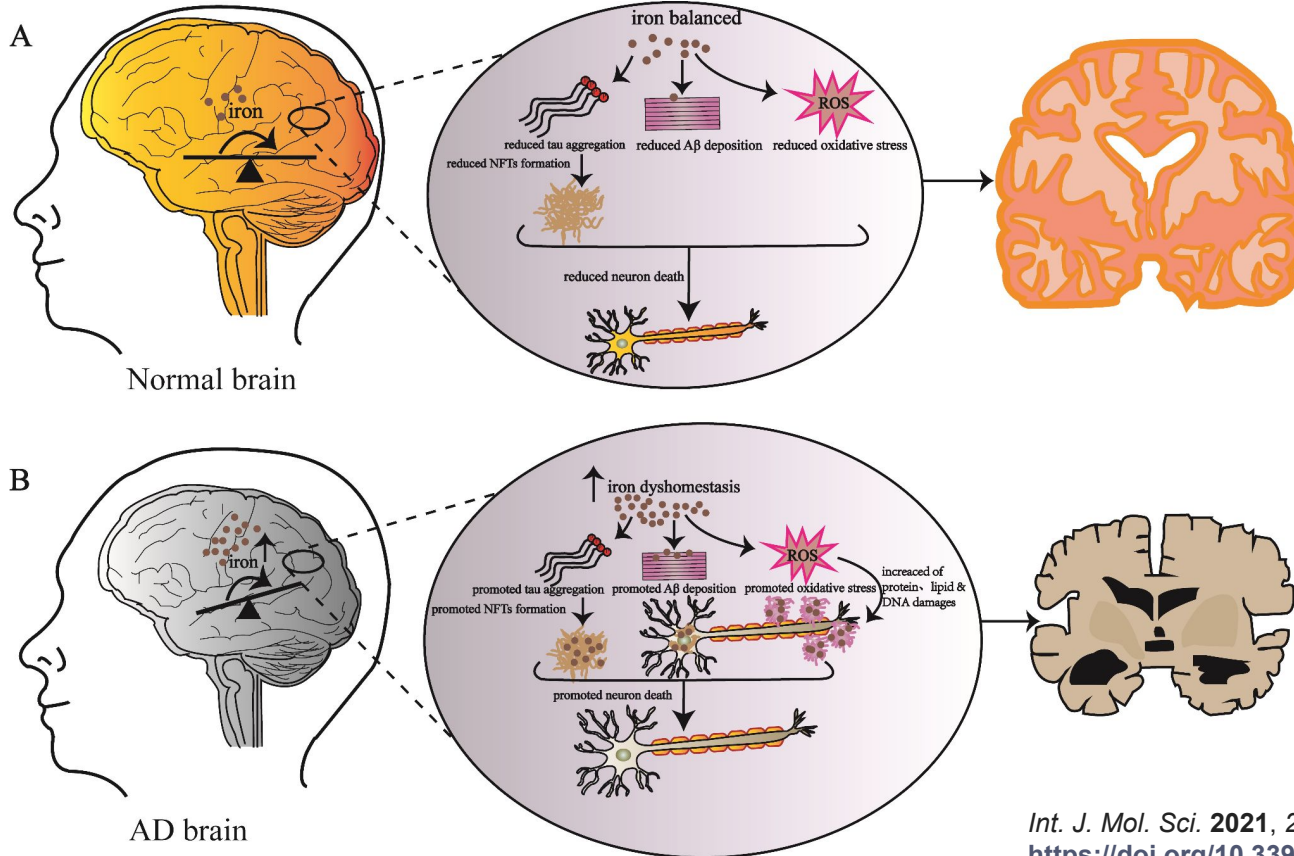
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CN1-Spoke 8 – In silico Medicine & Omics data
WP8.2 Digital Twins & in-silico trials

Outline

- Background
- Aims
- Datasets
- Tools for Neuroimaging processing
- Preliminary results
- Goals to be achieved by the end of the project.

Background: iron homeostasis



Background: iron homeostasis

Iron Dyshomeostasis and Ferroptosis: A New Alzheimer's Disease Hypothesis?

Feixue Wang^{1,2}, Jiandong Wang^{1,2}, Ying Shen^{1,2}, Hao Li², Wolf-Dieter Rausch⁴ and Xiaobo Huang^{1,2*}

NeuroImage 244 (2021) 118584



Contents lists available at ScienceDirect

NeuroImage

journal homepage: www.elsevier.com/locate/neuroimage



The effect of beta-amyloid and tau protein aggregations on magnetic susceptibility of anterior hippocampal laminae in Alzheimer's diseases

Text

Zhiyong Zhao³, Lei Zhang^{3,c}, Qingqing Wen³, Wanrong Luo³, Weihao Zheng³, Tingting Liu³, Yi Zhang³, Keqing Zhu^{3,c,*}, Dan Wu^{3,*}



Machine Learning Profiling of Alzheimer's Disease Patients Based on Current Cerebrospinal Fluid Markers and Iron Content in Biofluids

Eleonora Ficiarà^{1*}, Silvia Boschi^{1,2}, Shoeb Ansari¹, Federico D'Agata¹, Ornella Abollino³, Paola Caroppo⁴, Giuseppe Di Fede⁴, Antonio Indaco⁴, Innocenzo Rainero¹ and Caterina Guiot¹



Iron homeostasis can be disturbed by:

- genetic factors
- environmental factors
- aging



iron metabolism diseases, including many neurodegenerative diseases such as **Alzheimer's disease (AD)**.

Aims

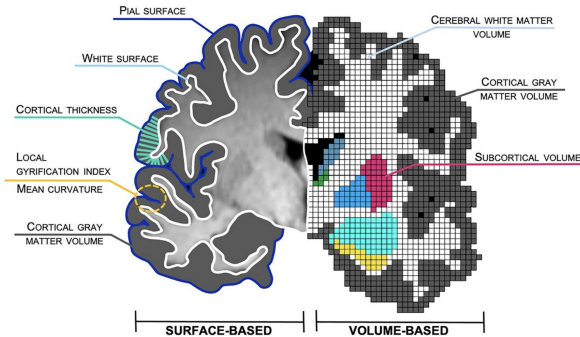
1. *Evaluate the relationship between the presence/accumulation of iron and morphological abnormalities in healthy subjects and those with AD, in order to use this characteristic as a potential biomarker for the early diagnosis of AD.*
2. *Analysis of candidate genes of interest for AD.*

Aims

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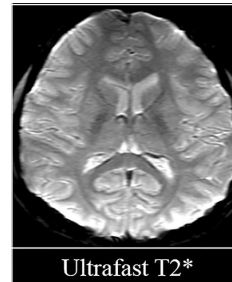
Methods

Atrophy assessed by VBM and SBM



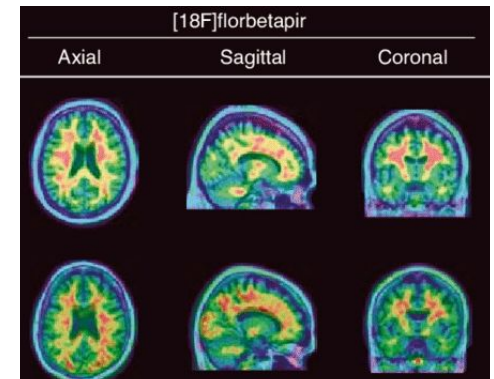
Backhausen, L.L., *Neuropsychol Rev* 32, 400–418 (2022).
<https://doi.org/10.1007/s11065-021-09496-2>

Iron concentration assessed by T2* MRI



<https://andreaforneris.com/web/protocollo-encefalo-standard-in-rm-in-un-minuto/>

Metabolism analysis assessed by PET



<https://www.ncbi.nlm.nih.gov/pmc/articles/PMC3468774/>

Aims

1. *Evaluate the relationship between the presence/accumulation of iron and morphological abnormalities in healthy subjects and those with AD, in order to use this characteristic as a potential biomarker for the early diagnosis of AD.*
2. *Analysis of candidate genes of interest for AD.*

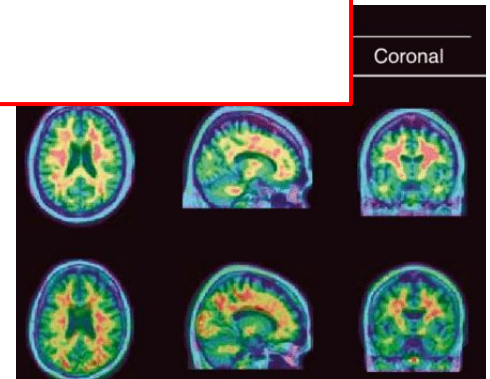
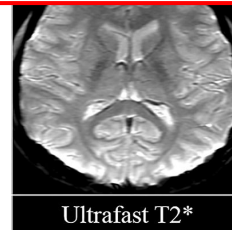
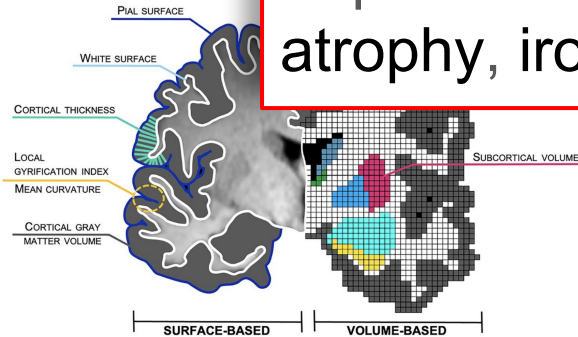
Methods

Atrophy assessed by

VBM and

Metabolism analysis by PET

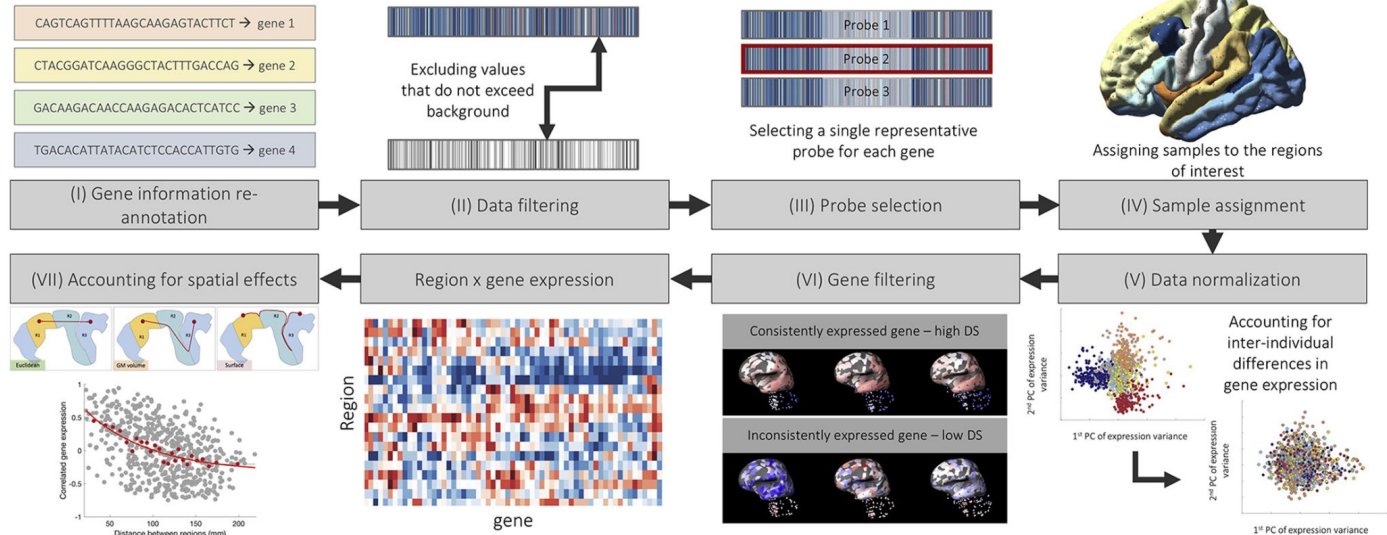
Expected results: overlapping patterns between atrophy, iron concentration and PET.



Aims

1. Evaluate the relationship between the presence/accumulation of iron and morphological abnormalities in healthy subjects and those with AD, in order to use this characteristic as a potential biomarker for the early diagnosis of AD.
2. Analysis of candidate genes of interest for AD.

Methods



abagen

IMAGES DATASETS



6000 subjects

ADNI is a longitudinal multicenter study designed to develop clinical, imaging, genetic, and biochemical biomarkers for the early detection and tracking of Alzheimer's disease (AD).

<https://adni.loni.usc.edu/>



3045 subjects

The Australian Imaging, Biomarker and Lifestyle (**AIBL**) Flagship Study of Ageing is an ongoing observational cohort study helping researchers unlock new insights into the onset and progression of Alzheimer's disease.

<https://aibl.org.au/>



2600 subjects

OASIS-3 is a longitudinal multimodal neuroimaging, clinical, cognitive, and biomarker dataset for normal aging and Alzheimer's Disease.

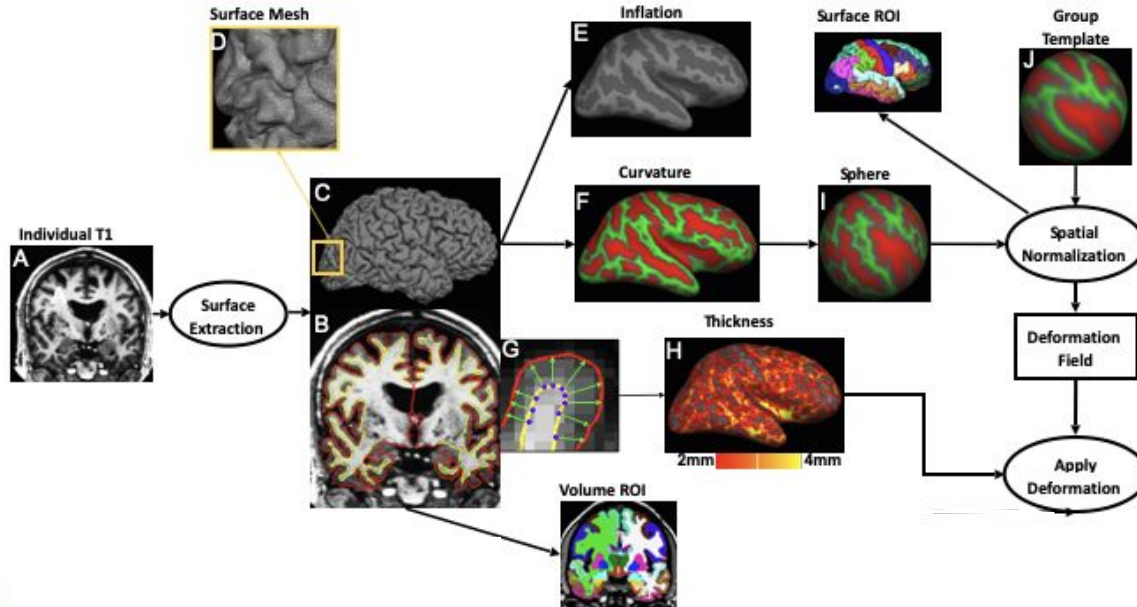
<https://www.oasis-brains.org/>

TOOLS

Softwares

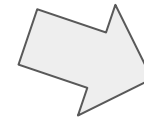
FreeSurfer

FreeSurfer Analysis Pipeline Overview



Freesurfer pipeline (T1w):

1. Data conversion from DICOM to Nii.gz
2. T1W1 input
3. Skull stripping
4. Registration (spatial normalization)
4. Intensity normalization
5. Volumetric labeling
6. Tissue segmentation
7. Surface atlas registration
8. Surface Extraction



Singularity
container



FreeSurfer Analysis Pipeline Overview

Surface Mesh Inflation Surface ROI Group

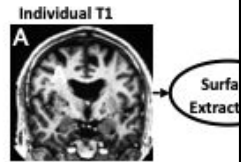
Computational Requirements

Runtime -> 12 hours / subject

Can be run with 8GB memory

Each subject output folder is ~95 MB

10k subjects -> ~1 year!!



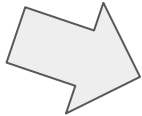
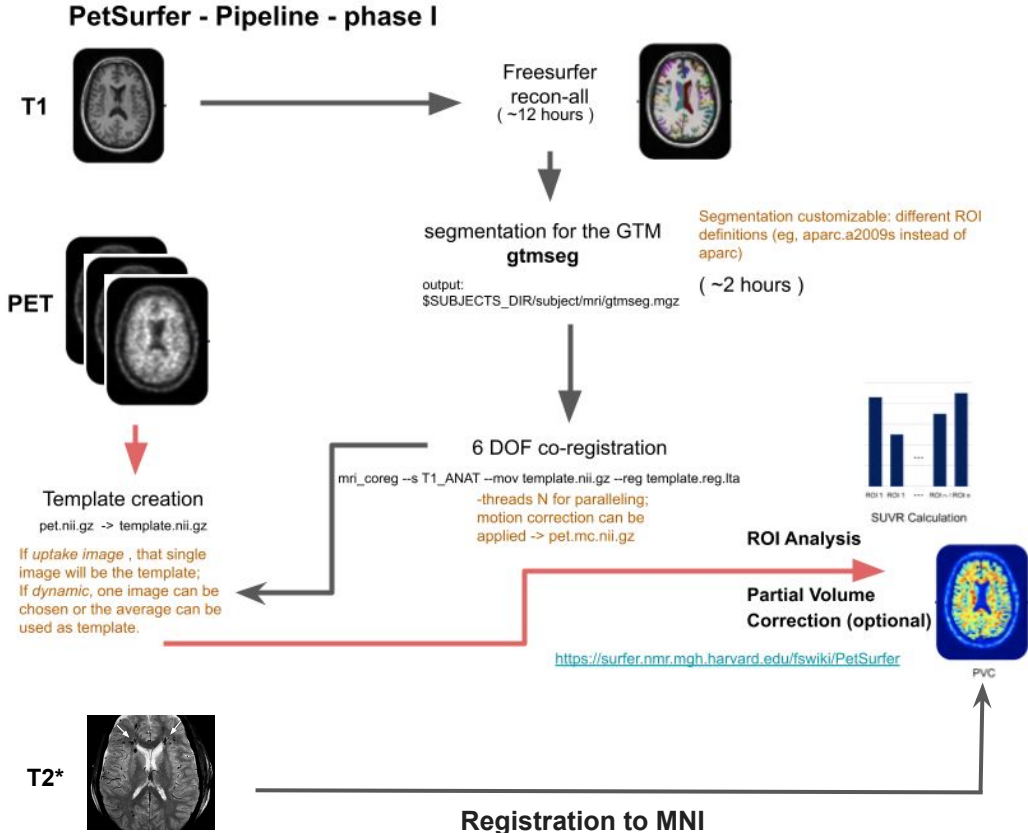
FreeSurfer pipeline (T1w):

- Conversion from DICOM to Nii.gz
- Input
- Stripping
- Registration (spatial normalization)
- Intensity normalization
- Metric labeling
- Segmentation
- Template atlas registration
- Surface Extraction



PetSurfer

1. Template creation
2. Partial volume correction
3. Motion correction across PET frames
4. Co-registration PET images with T1
5. Intensity normalization



Singularity container



PET and T2* pipeline: workflow showing the preprocessing steps for the PET data analysis, using the PetSurfer pipeline (<https://surfer.nmr.mgh.harvard.edu/fswiki/PetSurfer>). The output will be superimposed to the T2* maps to detect the iron deposition related to metabolic abnormalities.

PRELIMINARY RESULTS

- ❑ 206 CN / 160 AD including T1w, T2* e β amiloide PET (AV45) from ADNI
- ❑ Building up T1 e PET pipeline
- ❑ Building up VM on Cloud CINECA
- ❑ Running the T1 pipeline using FreeSurfer on Cloud CINECA

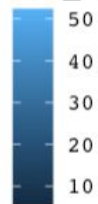
ADNI: 160 AD, 206 CN

Volumes: AD < CN

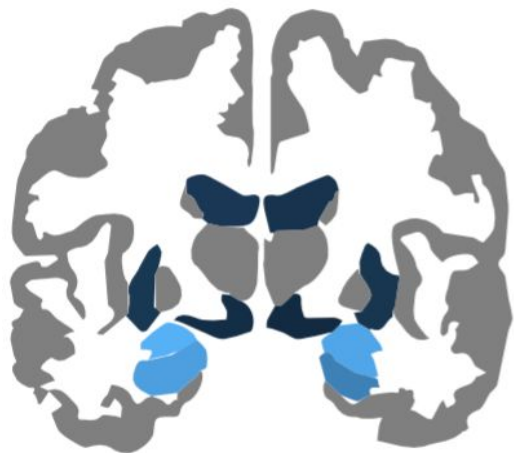
(corrected for age, sex, TIV)

$-\log_{10}(pval)$

log_val



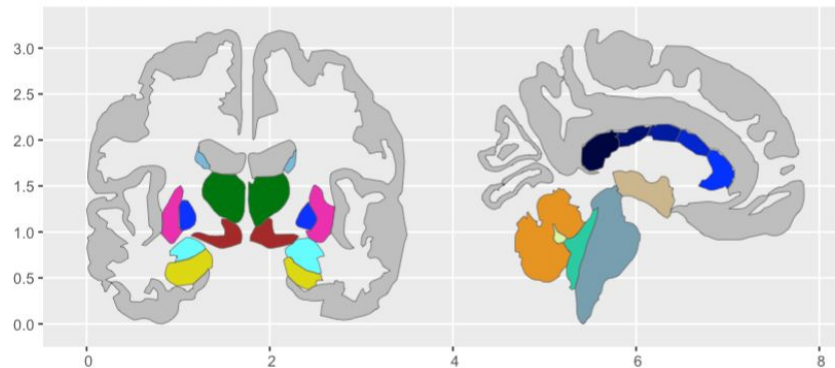
(bonferroni corrected)



coronal

Atlas for ROI nomenclature

aseg subcortical atlas



- 3rd ventricle
- 4th ventricle
- amygdala
- brain stem
- caudate
- CC anterior
- CC central
- CC mid anterior
- CC mid posterior
- CC posterior
- cerebellum cortex
- cerebellum white matter
- hippocampus
- pallidum
- putamen
- thalamus proper
- ventral DC

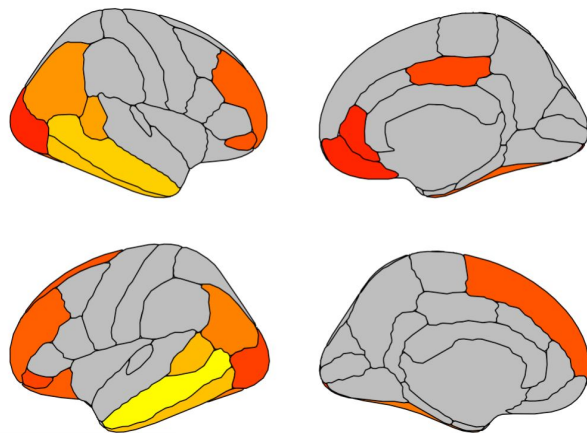
Cortical surface area: AD < CN

(corrected for age, sex, TIV)

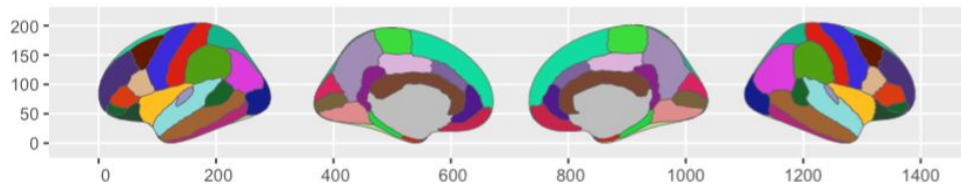
$-\log_{10}(pval)$



(bonferroni corrected)



dk cortical atlas



Achievements by the ended of the project and UPDATES

- ❑ PET pipeline to run on CINECA
- ❑ T2* - PET β amiloide comparison
- ❑ Apply predictive models (ML) using longitudinal data

- ❑ RAC approved 06/21/2024:
 - ❑ 100.000 core-hours on Leonardo DCGP and 20 TB of storage

Thank you!

ASSESSING IRON OVERLOAD AND
ABNORMAL MORPHOLOGIES IN
NORMAL VS AD BRAIN
BASED ON COMPARISON BETWEEN
PET & RMI IMAGING BASED ON
AVAILABLE LARGE DATASETS

FOCUS:

- ABETA PLAQUES ARE DETECTABLE BY PET AND ARE EXPECTED TO BE SUPERPOSED TO IRON DEPOSITION SITES
- MRI IMAGES ARE MORE EASILY AVAILABLE AND SOME SPECIFIC SEQUENCES ARE EXPECTED TO BE SENSITIVE FOR THE MAGNETIC PROPERTIES OF SMALL IRON DEPOSITS
- MORPHOLOGICAL DIFFERENCES IN BRAIN REGIONS, E.G. HIPPOCAMPAL SHRINKING AND/OR ABNORMAL SULCAL FEATURES DETECTABLE FROM MRI IN CONTROL AND AD PATIENTS

MRI-pet tools: t1, t2, pet ab-tau

