

Investigating demyelination, iron accumulation, and synaptic loss in Alzheimer's disease using multimodal imaging techniques

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Promoters:

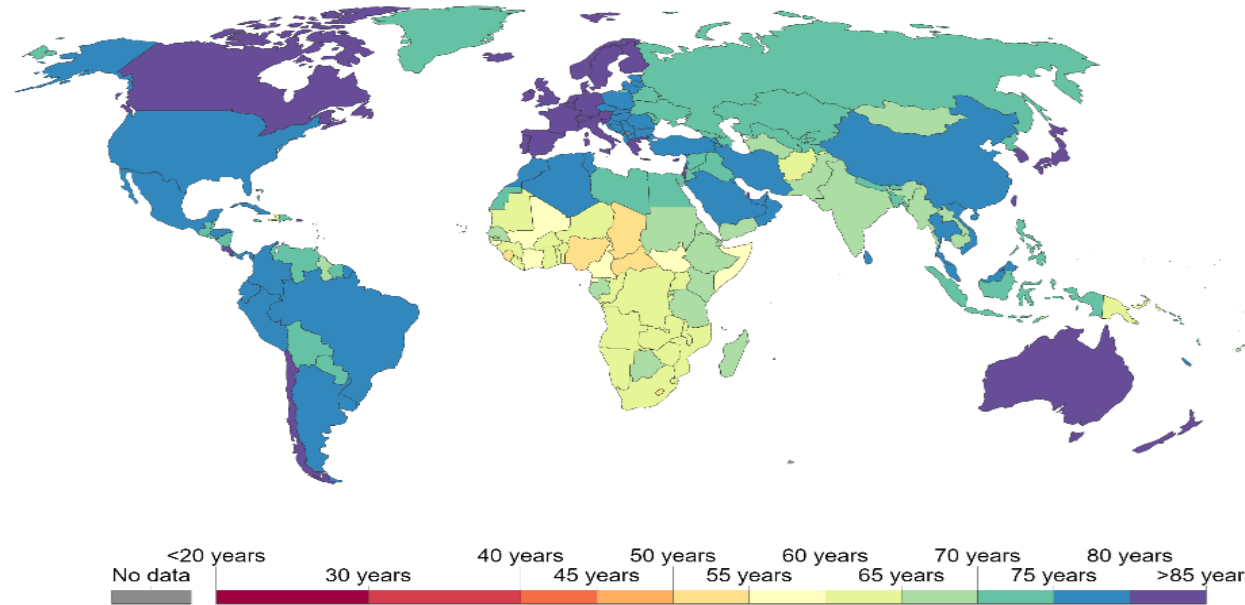
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Life expectancy, 2019

Our World
in Data

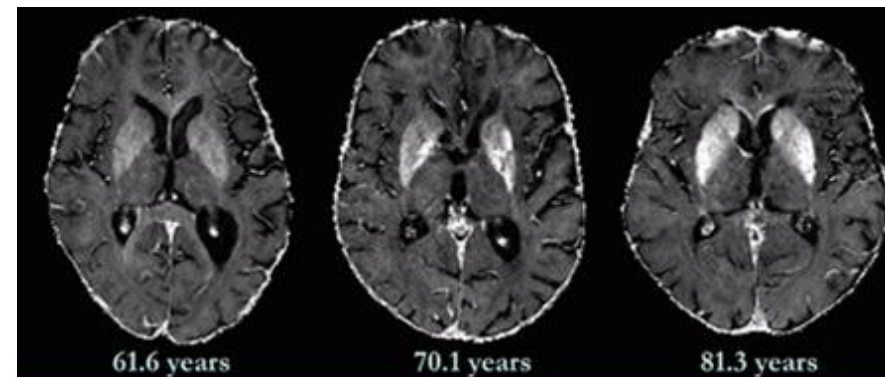
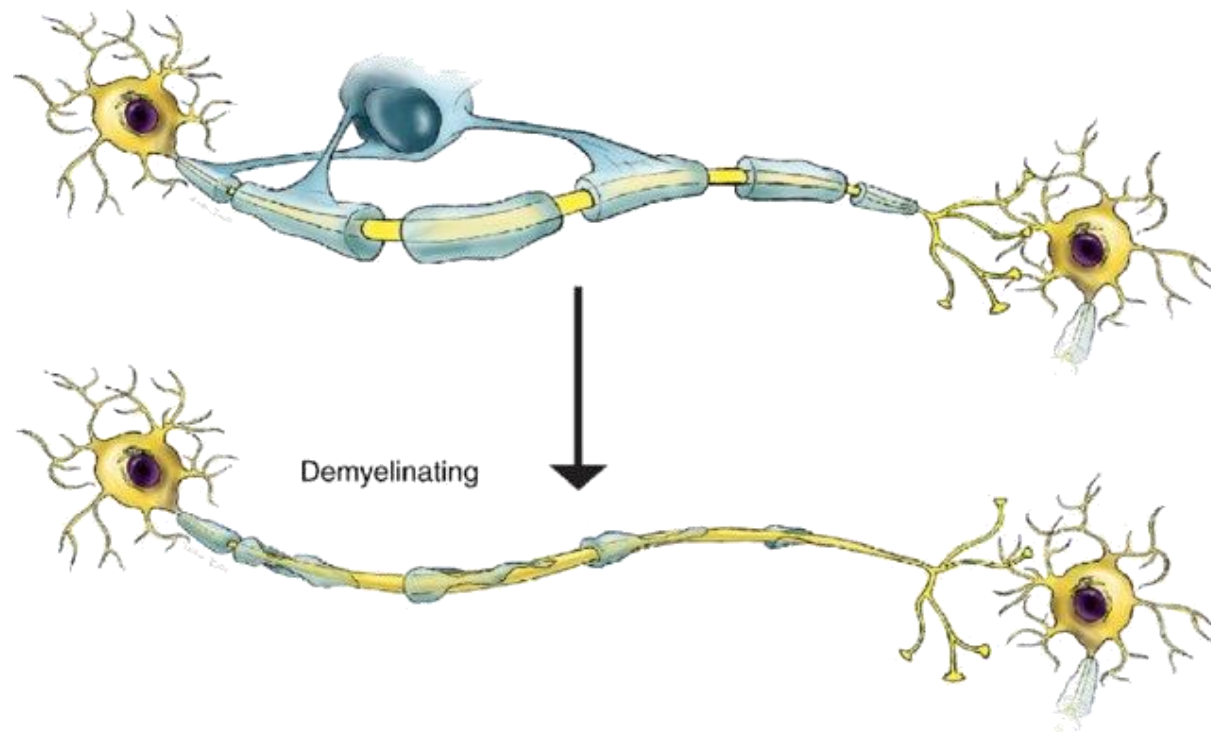
Source: Riley (2005), Clivio (2015), and UN Population Division (2019). OurWorld
Life expectancy at birth, the average number of years a newborn would live if the pattern
remained throughout its life.

The total number of people with **AD dementia** in 2050 is projected to be **13.8 million**, with **7.0 million aged 85 years or older** in the US.

By 2050, the number of people aged 65 and older with **Alzheimer's dementia** is projected to reach 12.7 million. (ONLY IN the US)

57 million people were living with dementia globally in **2019**, a number expected to rise to **153 million by 2050**.

- Hebert, L. E, et. Al., 2013 *Neurology*. <https://doi.org/10.1212/WNL.0b013e31828726f5>
- 2022 Alzheimer's Disease Facts and Figures, Special Report More Than Normal Aging: Understanding Mild Cognitive Impairment
- Alzheimer's disease international, 2022





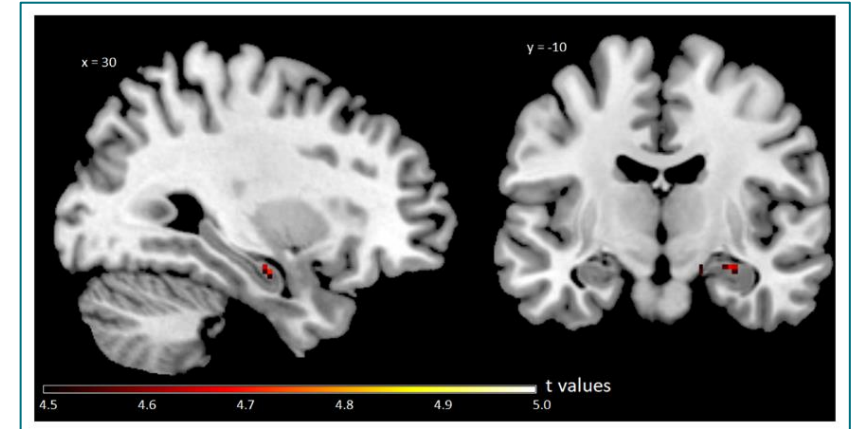
European Journal of Nuclear Medicine and Molecular Imaging (2020) 47:390–402
<https://doi.org/10.1007/s00259-019-04461-x>

ORIGINAL ARTICLE

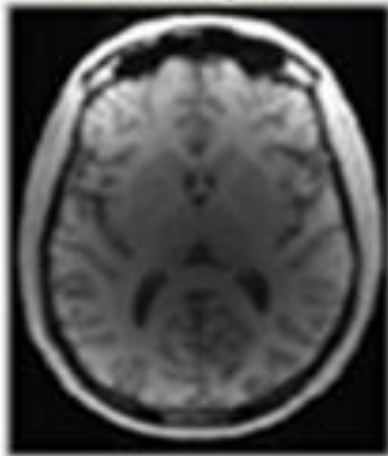


In vivo imaging of synaptic loss in Alzheimer's disease with [18F] UCB-H positron emission tomography

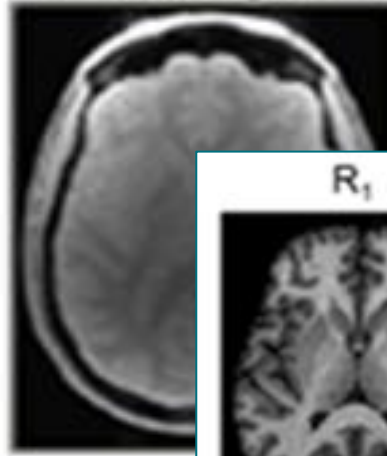
Christine Bastin¹  · Mohamed Ali Bahri¹ · François Meyer¹ · Marine Manard¹ · Emma Delhaye¹ · Alain Plenevaux¹ · Guillaume Becker¹ · Alain Seret¹ · Christine Mella¹ · Fabrice Giacomelli¹ · Christian Degueldre¹ · Evelyne Balteau¹ · André Luxen¹ · Eric Salmon¹



T1 Weighted



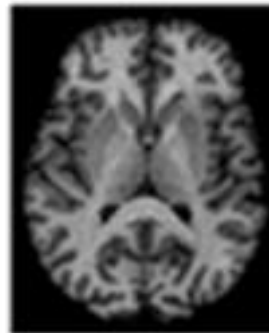
PD Weighted



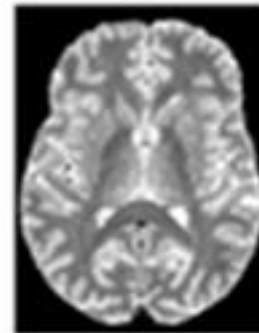
MT Weighted



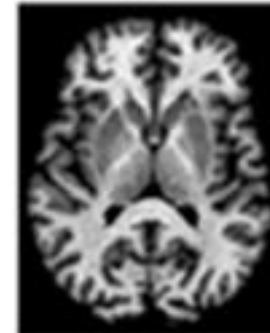
R_1



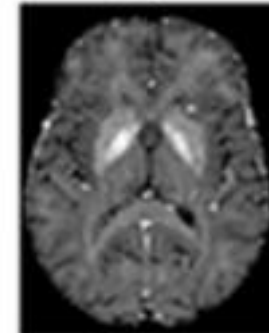
PD*



MT



R_2^*





Characterize patients from healthy controls using different image-derived biomarkers for AD

Myelination

Iron level

Synaptic density

MTsat

R2*

PET_vt

Myelination

Iron level

Synaptic density

Investigate the association of image-derived markers in a multivariate model

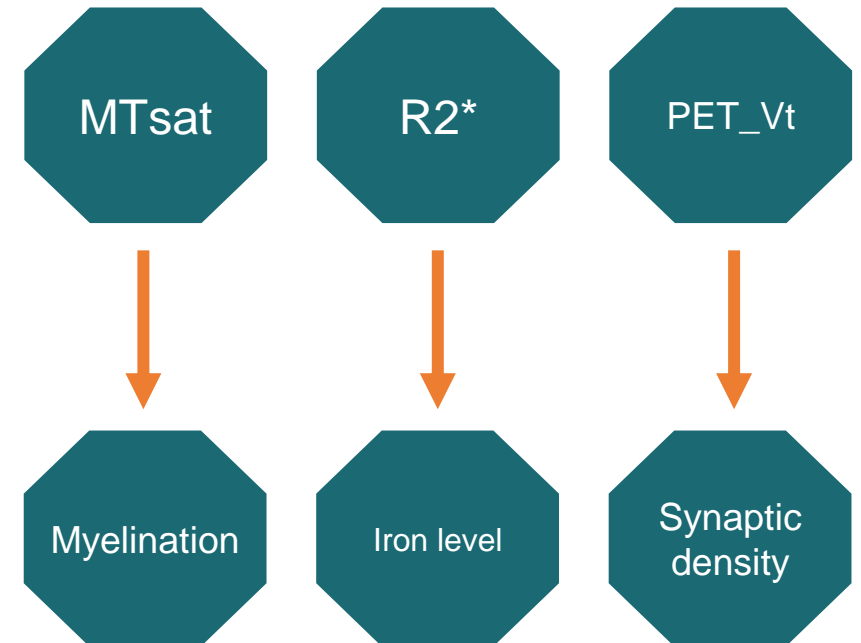
Iron level

Myelination

Synaptic density

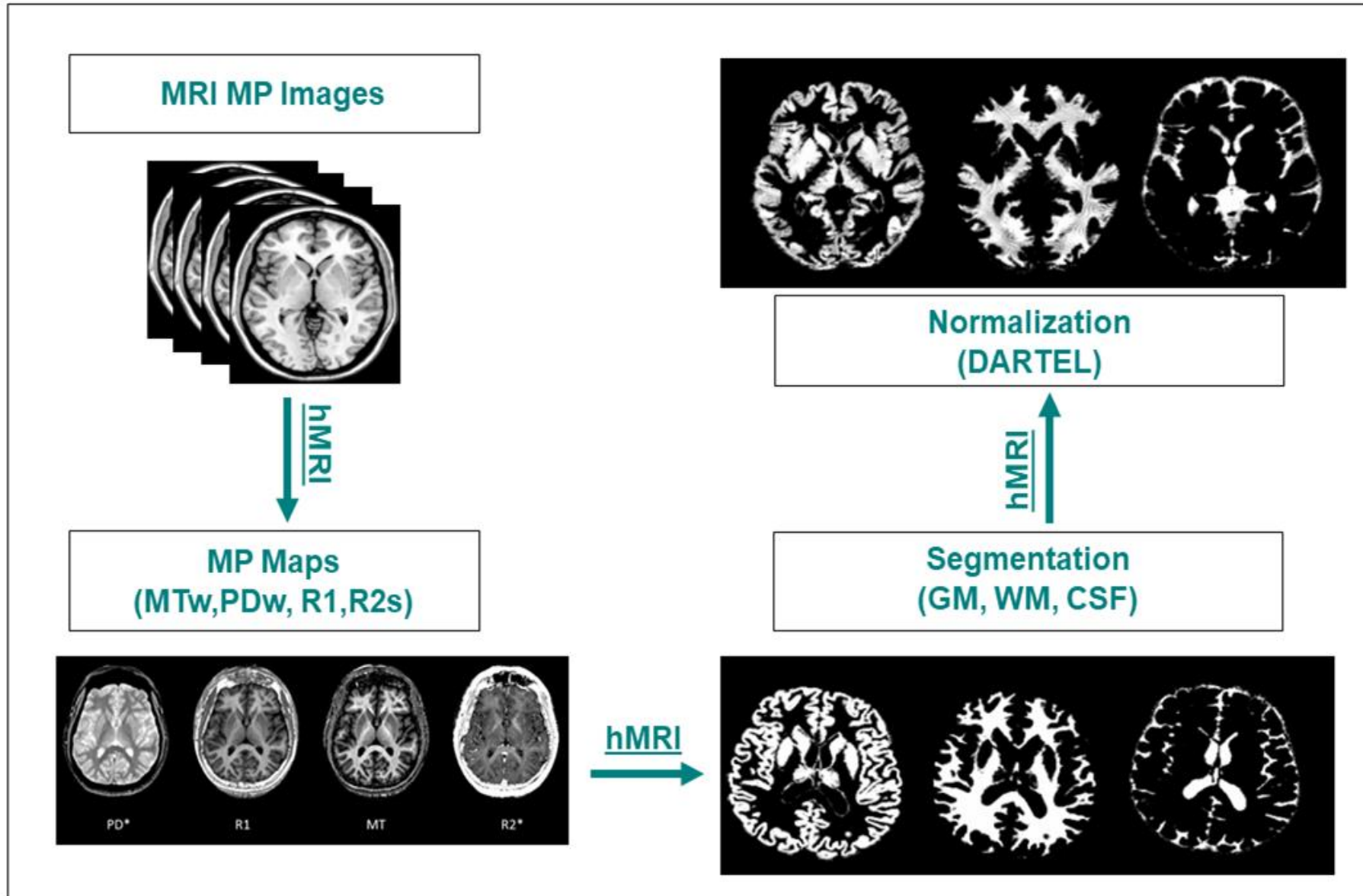


Data sets	# Subjects	Gender	Age Range
Alzheimer's disease	24 (11 males - 13 females)		60-80
Healthy control	19 (9 males -10 females)		



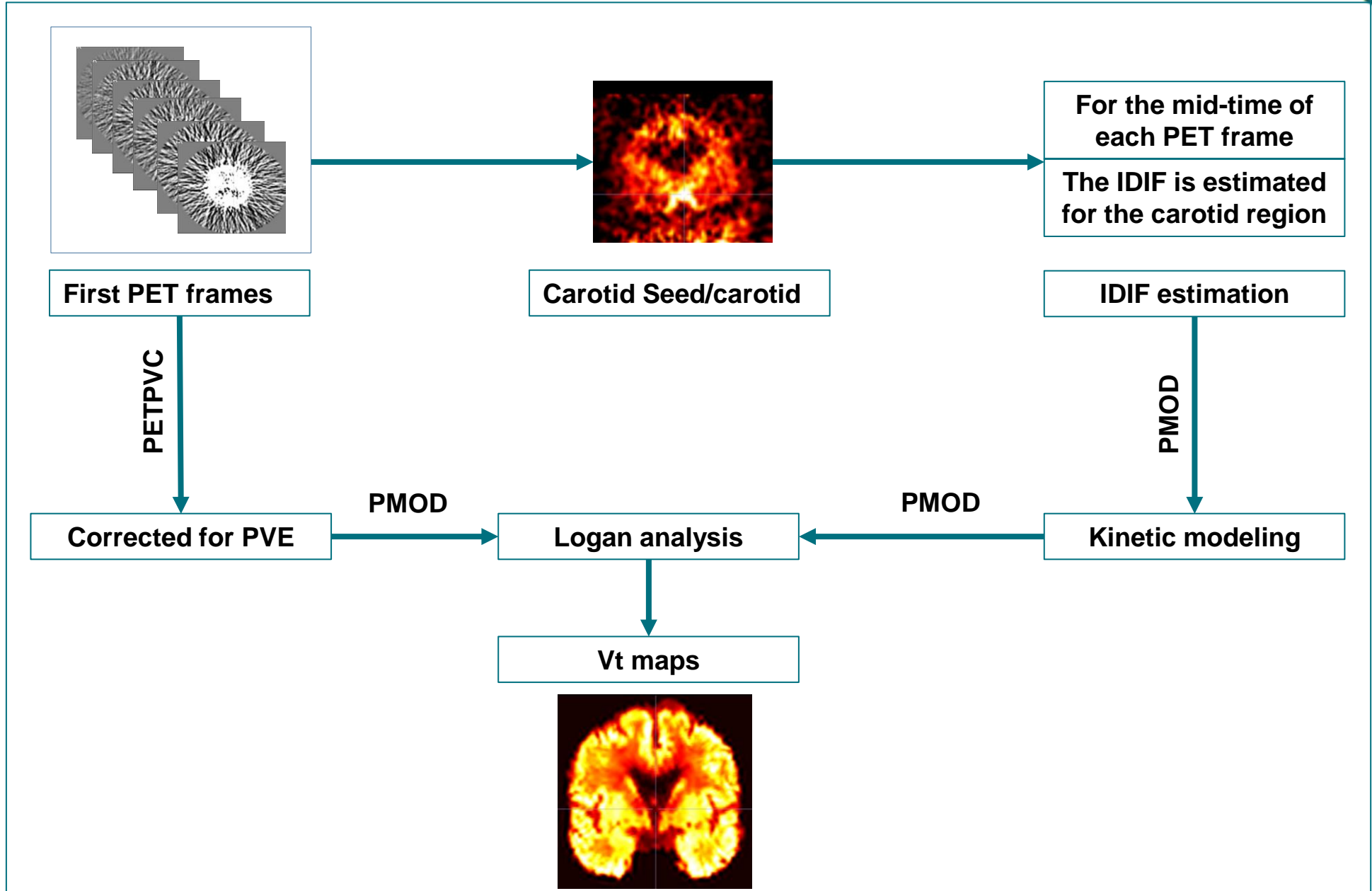


Quantitative maps creation





Vt maps creation





All maps are in the MNI space...

Good enough for the univariate analysis

Each modality represents a different property and unit!

The data is not comparable, and we cannot use them in a multivariate analysis

Z-transformation



$$Y = X\beta + \varepsilon$$

Univariate GLM:

$$X = \begin{bmatrix} AD_m & 0 & age & sex \\ 0 & HC_m & age & sex \end{bmatrix}_{43 \times 4} \quad \beta = \begin{bmatrix} \mu_{11} \\ \mu_{12} \\ \mu_{21} \\ \mu_{22} \end{bmatrix}$$

Test the difference between groups using one modality

$$H_0: C\beta = 0$$

$$H_1: C\beta \neq 0$$

Between Subject contrast

$$C = [1 \quad -1 \quad 0 \quad 0]$$

Multivariate GLM:

$$X = \begin{bmatrix} AD_{MT} & 0 & age & sex \\ AD_{R2S} & 0 & age & sex \\ AD_{PET} & 0 & age & sex \\ 0 & HC_{MT} & age & sex \\ 0 & HC_{R2S} & age & sex \\ 0 & HC_{PET} & age & sex \end{bmatrix}_{129 \times 4} \quad B = \begin{bmatrix} \mu_{111} & \mu_{112} \\ \mu_{121} & \mu_{122} \\ \mu_{211} & \mu_{212} \\ \mu_{221} & \mu_{222} \end{bmatrix}$$

MSPM

Test the difference between groups using all modalities at once

$$H_0: CBL = 0$$

$$H_1: CBL \neq 0$$

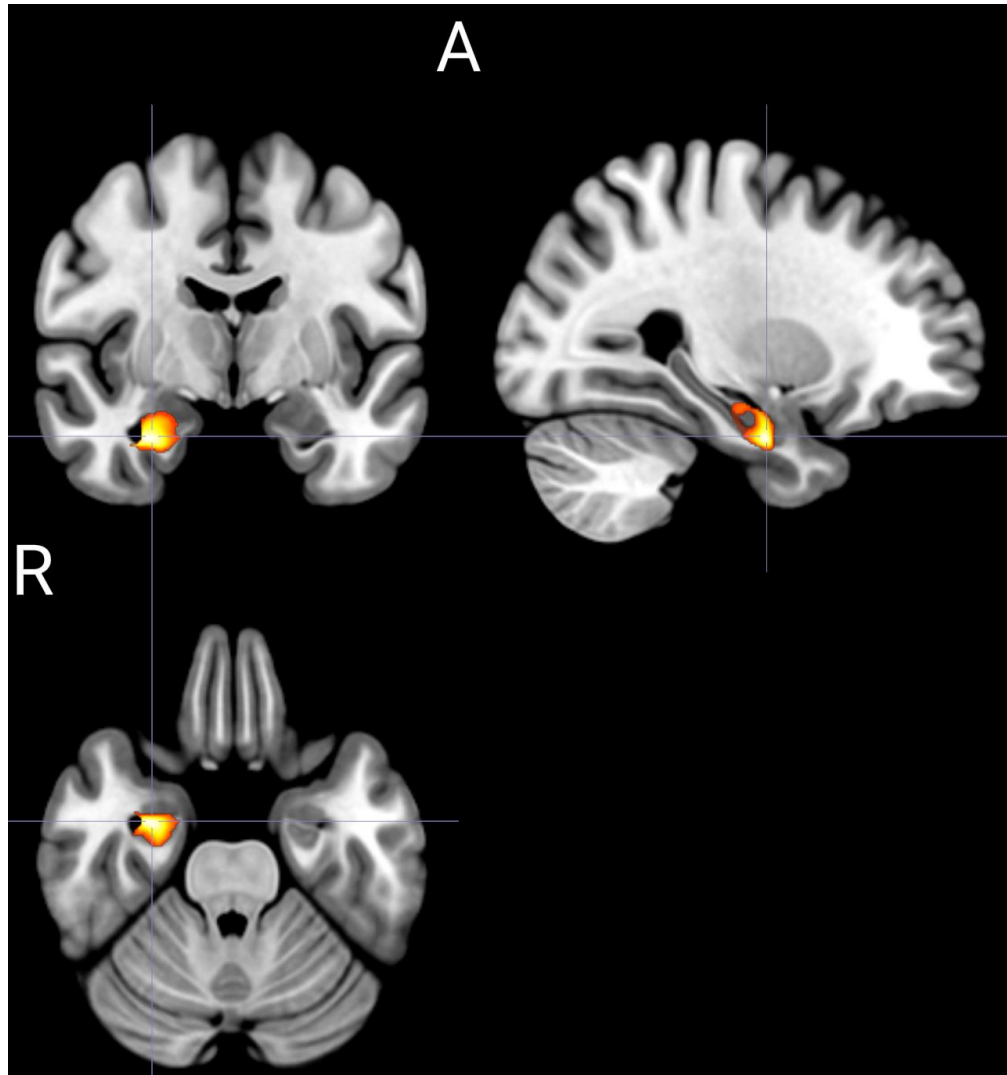
Within Subject parameter

$$L = \begin{bmatrix} 1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \end{bmatrix}_{3 \times 3}$$

Between Subject contrast

$$C = [1 \quad -1 \quad 0 \quad 0]$$

* Here "m" represents different modalities


Statistics: p -values adjusted for search volume

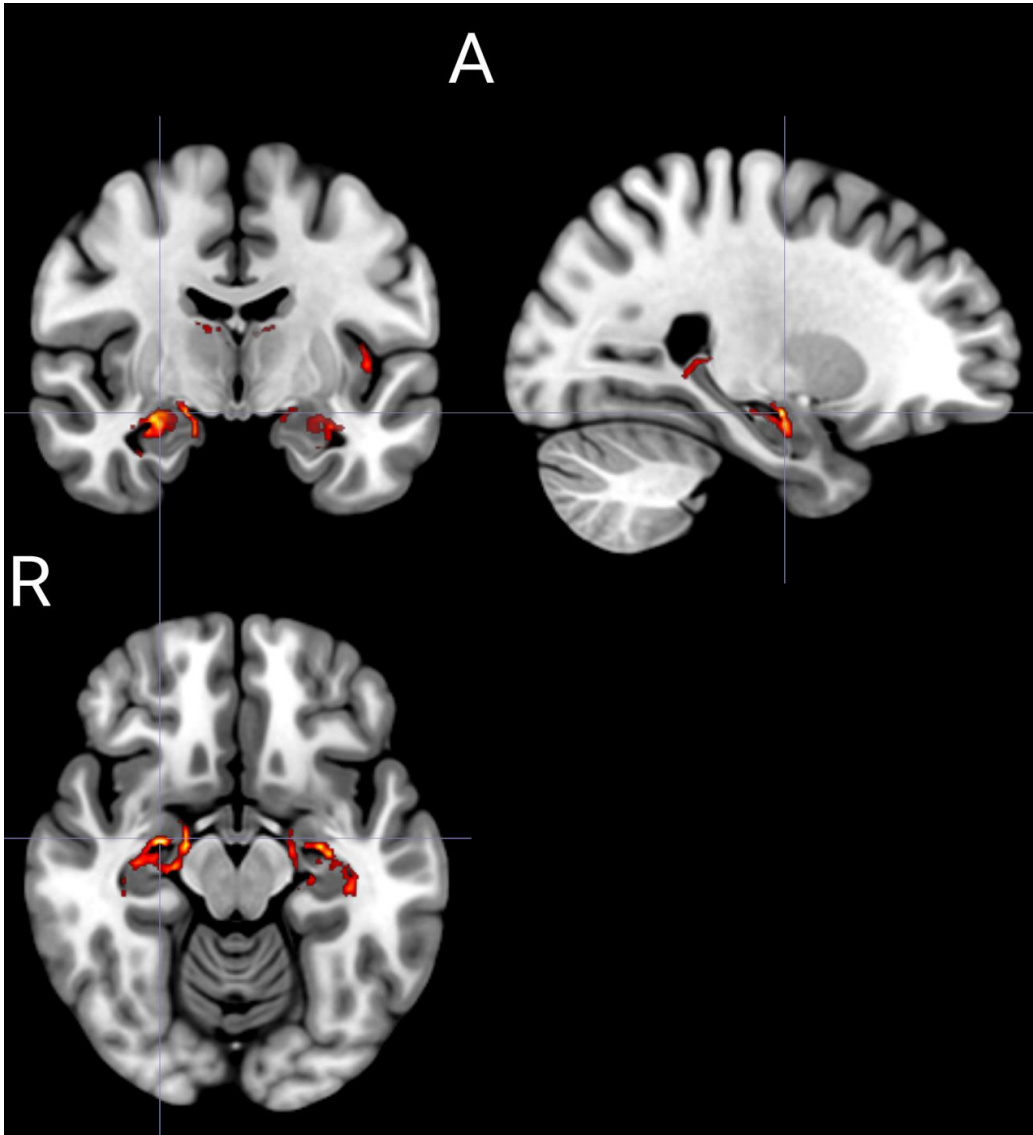
set-level		cluster-level				peak-level					mm mm mm		
p	c	$p_{\text{FWE-corr}}$	$q_{\text{FDR-corr}}$	k_E	p_{uncorr}	$p_{\text{FWE-corr}}$	$q_{\text{FDR-corr}}$	F	(Z_E)	p_{uncorr}			
0.845	3	0.001	0.001	1993	0.000	0.201	0.186	28.25	4.43	0.000	31	-7	-27
						0.520	0.307	23.72	4.12	0.000	37	-21	-12
						0.782	0.307	21.04	3.91	0.000	20	-11	-20
		0.972	0.940	131	0.203	0.604	0.307	22.86	4.06	0.000	-66	-49	-15
		0.636	0.486	315	0.057	0.997	0.752	16.28	3.48	0.000	-24	-10	-25
						0.999	0.816	15.48	3.40	0.000	-19	-6	-35

table shows 3 local maxima more than 8.0mm apart

Height threshold: $F = 12.66$, $p = 0.001$ (1.000)
 Extent threshold: $k = 100$ voxels, $p = 0.264$ (0.991)
 Expected voxels per cluster, $\langle k \rangle = 86.513$
 Expected number of clusters, $\langle c \rangle = 4.67$
 FWEp: 34.309, FDRp: Inf, FWEc: 1993, FDRc: 1993

Degrees of freedom = [1.0, 39.0]
 FWHM = 11.0 9.9 10.1 mm mm mm; 11.0 9.9 10.1 {voxels}
 Volume: 912858 = 912858 voxels = 689.3 resels
 Voxel size: 1.0 1.0 1.0 mm mm mm; (resel = 1098.45 voxels)

Table of results - AD vs HC - ($P < 0.001$)


Statistics: p -values adjusted for search volume

set-level		cluster-level				peak-level					mm mm mm		
p	c	$p_{\text{FWE-corr}}$	$q_{\text{FDR-corr}}$	k_E	p_{uncorr}	$p_{\text{FWE-corr}}$	$q_{\text{FDR-corr}}$	F	(Z_E)	p_{uncorr}			
0.000	6	0.000	0.012	8	0.002	0.001	0.108	64.16	6.01	0.000	25	-10	-14
		0.006	0.170	2	0.085	0.001	0.108	64.07	6.01	0.000	14	-10	-21
		0.001	0.032	5	0.011	0.001	0.108	63.74	6.00	0.000	18	-11	-13
		0.015	0.211	1	0.211	0.010	0.355	54.13	5.67	0.000	40	49	22
		0.015	0.211	1	0.211	0.025	0.568	50.10	5.52	0.000	-29	-14	-15
		0.015	0.211	1	0.211	0.026	0.568	49.87	5.51	0.000	-27	-12	-15

table shows 3 local maxima more than 8.0mm apart

Height threshold: $F = 47.00$, $p = 0.000$ (0.050)

Extent threshold: $k = 0$ voxels

Expected voxels per cluster, $\langle k \rangle = 0.686$

Expected number of clusters, $\langle c \rangle = 0.07$

FWEp: 47.001, FDRp: Inf, FWEc: 1, FDRc: 5

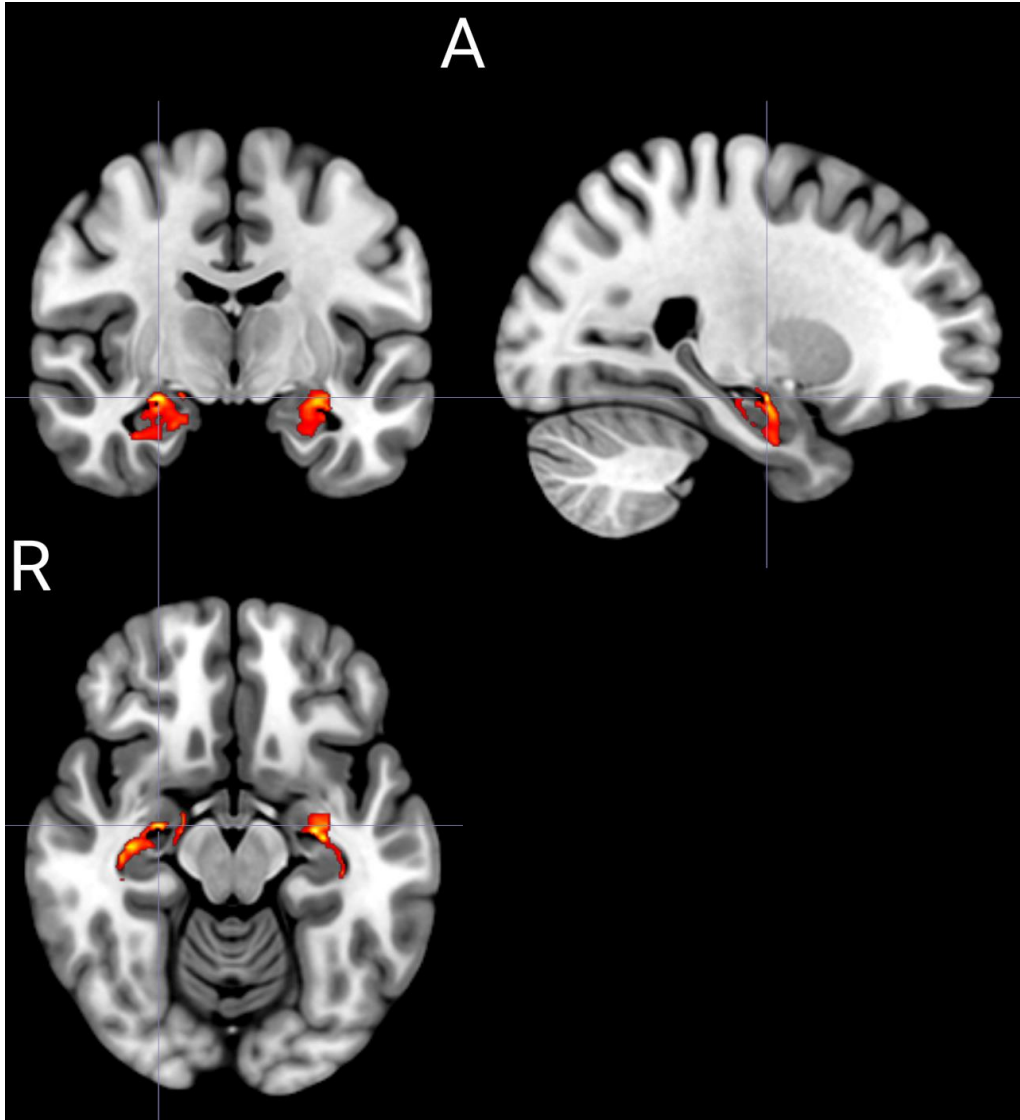
Degrees of freedom = [1.0, 39.0]

FWHM = 4.0 3.9 4.1 mm mm mm; 4.0 3.9 4.1 {voxels}

Volume: 1477777 = 1477777 voxels = 21987.7 resels

Voxel size: 1.0 1.0 1.0 mm mm mm; (resel = 63.34 voxels)

Table of results - AD vs HC - ($P < 0.05$ FWE)


STATISTICS: p -values adjusted for search volume

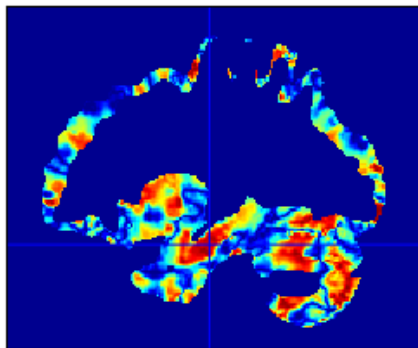
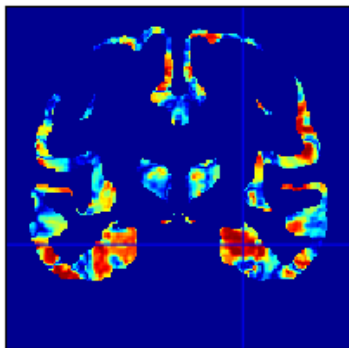
set-level		cluster-level				peak-level					mm mm mm		
p	c	$p_{\text{FWE-corr}}$	$q_{\text{FDR-corr}}$	k_E	p_{uncorr}	$p_{\text{FWE-corr}}$	$q_{\text{FDR-corr}}$	F	(Z_E)	p_{uncorr}			
0.000	10	0.022	0.810	8	0.430	0.005	0.438	21.11	5.37	0.000	25	-10	-14
		0.018	0.810	11	0.353	0.005	0.438	21.00	5.36	0.000	-27	-12	-15
		0.027	0.810	5	0.540	0.026	0.998	17.65	5.00	0.000	35	-18	-13
		0.041	0.810	1	0.810	0.028	0.998	17.51	4.99	0.000	40	49	22
		0.041	0.810	1	0.810	0.037	0.998	16.99	4.92	0.000	37	52	25
		0.041	0.810	1	0.810	0.040	0.998	16.80	4.90	0.000	35	-16	-15
		0.041	0.810	1	0.810	0.045	0.998	16.59	4.88	0.000	-62	-44	35
		0.041	0.810	1	0.810	0.046	0.998	16.55	4.87	0.000	30	-7	-22
		0.041	0.810	1	0.810	0.047	0.998	16.51	4.87	0.000	8	34	-27
		0.041	0.810	1	0.810	0.050	0.998	16.40	4.85	0.000	38	-25	-12

table shows 3 local maxima more than 8.0mm apart

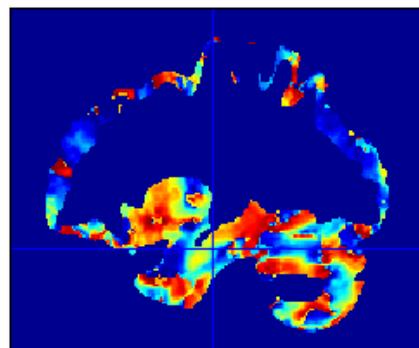
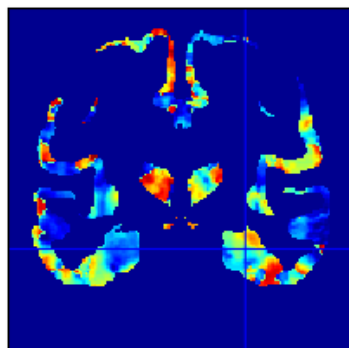
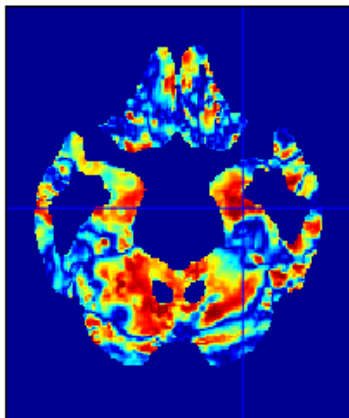
Height threshold: $F = 16.40$, $p = 0.000$ (0.050)
 Extent threshold: $k = 0$ voxels
 Expected voxels per cluster, $\langle k \rangle = 13.738$
 Expected number of clusters, $\langle c \rangle = 0.05$
 FWEp: 16.397, FDRp: Inf, FWEc: 1, FDRc: Inf

Degrees of freedom = [3.0, 37.0]
 FWHM = 11.0 9.9 10.1 mm mm mm; 11.0 9.9 10.1 {voxels}
 Volume: 816797 = 816797 voxels = 689.3 resels
 Voxel size: 1.0 1.0 1.0 mm mm mm; (resel = 1098.45 voxels)

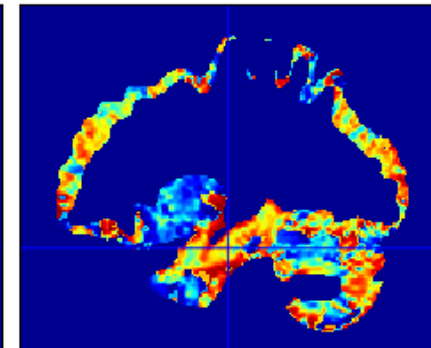
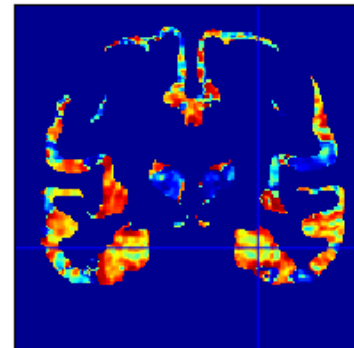
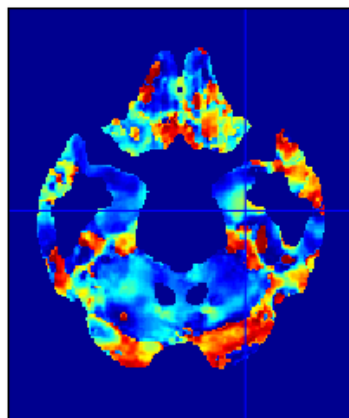
Table of results - AD vs HC - ($P < 0.05$ FWE)

 $Y = 0.921536$

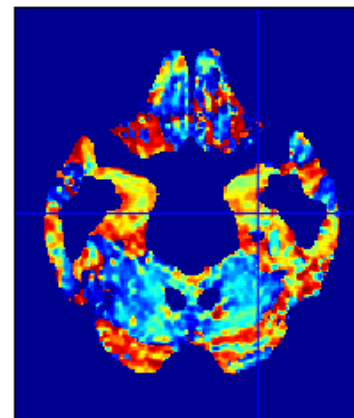
MTsat

 $Y = -0.116768$

R2*

 $Y = 0.362678$

PET_Vt





Take home message:

- Quantitative maps can be used to assess AD in its early stages.
- Alteration in **myelin content** and **synaptic density** in early AD occurs in the hippocampus.
- Accounting for the interactions between myelin, iron, and SV2A protein we can get a more accurate characterization of the difference between the groups of AD and HC.
- **Multivariate GLM model** is preferable to multiple univariate models to test for AD, which relies on a cascade of processes.



THANK YOU!



<https://hdl.handle.net/2268/296010>





Mtsat(P<0.001 FWE uncorrected at voxel level analysis)

Cluster number	Coordinate at peak	F-Value at peak	Cluster P-value*	Cluster size	Brain region
1	(28 -7 -27)	26.58	0.002	1993	Right hippocampus

PET_vt (P<0.05 FWE corrected at voxel level analysis)

1	(25,-10,-14)	64.16	0.000	8	Right hippocampus
2	(14,-10,-21)	64.07	0.006	2	Right hippocampus
3	(18,-11,-13)	63.74	0.001	5	Right amygdala

Multimodal (P<0.05 FWE corrected at voxel level analysis)

1	(25,-10,-14)	21.11	0.022	8	Right hippocampus
2	(-27,-12,-15)	50.10	0.018	11	Left hippocampus
3	(35,-18,-13)	34.02	0.027	5	Right hippocampus



MSPM toolbox:

Example :

- Three modalities
- 43 subjects
- Two groups

Selected 3/[1-...] files. (Added 1/1 file.)
 E:\UCBH\univariate_analysis\z_MT\SPM.mat
 E:\UCBH\univariate_analysis\z_R2s\SPM.mat
 E:\UCBH\univariate_analysis\z_PET\SPM.mat

Panel: Select L contrasts...
 Listbox
 New contrast

Panel: Select c contrasts...
 Listbox
 New contrast
 Done

$$X = \begin{bmatrix} AD_{MT} & AD_{MT} \\ AD_{R2s} & AD_{R2s} \\ AD_{PET} & AD_{PET} \\ HC_{MT} & HC_{MT} \\ HC_{R2s} & HC_{R2s} \\ HC_{PET} & HC_{PET} \end{bmatrix}_{43 \times 2}$$

$$L = \begin{bmatrix} 1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \end{bmatrix}_{3 \times 3}$$

$$C = [0 \ 1]_{2 \times 1}$$