

## Supplementary Materials

### Amyloid PET imaging

All subjects also underwent [<sup>11</sup>C]Pittsburgh Compound B (PiB) PET imaging over 60 min, as previously described [29,30]. An average measure of global amyloid- $\beta$  burden was estimated in a large neocortical aggregate of frontal, lateral temporoparietal and retrosplenial (FLR) regions, based on distribution volume ratios (DVR) computed using the reference Logan graphical method, with the CerGM as reference region ( $t^* = 40$  min). Subjects with PiB DVR values in the FLR region greater than 1.13 were considered as PiB-positive. The 1.13 threshold was determined by applying gaussian mixture model to all available [<sup>11</sup>C]PiB DVR data ( $n = 268$ ) collected on the same scanner as this study.

### Spectral analysis

Spectral analysis is a data-driven method for evaluating tracer kinetics without assumptions on kinetic compartment configuration. In spectral analysis, tracer activity ( $C_T(t)$ ) in the tissue at time  $t$  is described as the convolution of the tracer activity in the plasma ( $C_P(t)$ ) and the impulse response function (IRF( $t$ )):

$$C_T(t) = C_P(t) \otimes \text{IRF}(t) \quad (1)$$

IRF( $t$ ) contains information about the kinetics of the tissue in response to tracer delivery and can be solved as an analytical sum of distinct exponential terms:

$$\text{IRF}(t) = \sum_{j=0}^M \alpha_j e^{-\beta_j t} \quad (2)$$

Where  $\beta_j$  ( $\beta_0 = 0, \beta_j \geq 0$ , unit  $\text{min}^{-1}$ ) are the frequency of the spectral components and  $\alpha_j$  (unit  $\text{mL cm}^{-3} \text{min}^{-1}$ ) are the corresponding amplitudes.  $M + 1$  is the maximum number of terms in the model. The values of  $\beta_j$  are predefined to cover a range of spectral frequencies from:

- Very large  $\beta_j$ : high frequency spectral components, which correspond to blood volume fraction
- Very small  $\beta_j$ : low frequency spectral components, which often correspond to irreversible trapping in the tissue compartments
- Intermediate  $\beta_j$ : spectral components in equilibrium, which indicate the number of identifiable reversible tissue compartments

After fixing the values of  $\beta_j$  to cover an appropriate spectral range, the  $\alpha_j$  values were estimated from  $C_T(t)$  and  $C_P(t)$  by using non-negative least square (NNLS) algorithm. Several macro-parameters can then be derived from the kinetic spectrum:

- Influx rate constant:  $K_{1\_SA} = \sum_{j=0}^M \alpha_j$ , unit:  $\text{mL cm}^{-3} \text{min}^{-1}$  (3)

- Volume of distribution  $V_{T\_SA} = \sum_{j=0}^M \alpha_j / \beta_j$ , unit:  $\text{mL cm}^{-3}$  (4)

In this study, spectral analysis was implemented using a pre-defined grid of 100 components for  $\beta_j$ , ranging from  $10^{-5}$  to  $1 \text{ min}^{-1}$  on a logarithmic scale.

### Effect size calculation (Hedge's g coefficients)

Due to the small sample size, effect sizes were estimated using Hedge's g coefficients using the formula:

$$g = J(df) \times \frac{(\bar{x} - \bar{y})}{s}$$

where  $df$  is the degree of freedom,  $\bar{x}$  and  $\bar{y}$  are the mean values for each group or time point,  $s$  is the pooled standard deviation and  $J(df)$  is the bias correction term. The pooled standard deviation and the bias correction term are defined as follows:

$$s = \sqrt{\frac{(n_1 - 1)S_1^2 + (n_2 - 1)S_2^2}{n_1 + n_2 - 2}}$$

where  $n_1$  and  $n_2$  are the sample size of each group or time point.

$$J(df) = \frac{(\Gamma(\frac{df}{2}))}{\sqrt{df/2\Gamma(\frac{df-1}{2})}}$$

where  $\Gamma$  is the gamma function.

The 95% confidence interval for the  $g$  coefficients was estimated using bootstrapping (1000 iterations).

**Supplementary Table 1.** Published [<sup>18</sup>F]MK-6240 human PET imaging studies: Imaging parameters and reference regions.

Author, year	Journal †	Tomograph	FWHM *	Acquisition duration	SUVR **	Reference region
Pascoal 2018	Alzheimer's Research and Therapy	Siemens HRRT	8mm after smoothing	Up to 180 min	Varied, 90-110 min chosen	Cerebellar gray matter
Bethhauser 2019, 2020	J Nucl Med	Siemens ECAT Exact HR+	5 mm	Up to 120 min	70-90 min	Inferior cerebellar gray matter
Lohith 2019	J Nucl Med	Siemens Hirez Biograph	5 mm	0-90 min, break, 120-150 min	60-90 min	Cerebellar gray matter
Guehl 2019	EJNMMI	1. Siemens ECAT Exact HR+ 2. GE Discovery MI	1. 5mm 2. 5mm	Up to 135 min	1. 70-90 min 2. 90-120 min	Cerebellar gray matter
Salinas 2020 ^	JCBFM	Siemens ECAT Exact HR+	5 mm	Up to 150 min	90-120 min	Inferior cerebellar gray matter
Pascoal 2020	Brain	Siemens HRRT	8mm after smoothing	90-110 min	90-110 min	Inferior cerebellum
Pascoal 2021	Nature Medicine	Siemens HRRT	8 mm after smoothing	90-110 min	90-110 min	Cerebellar Crus I gray matter
Therriault 2021 ^^	Neurology	Siemens HRRT	8 mm after smoothing	90-110 min	90-110 min	Inferior cerebellar gray matter
Gogola 2021	J Nucl Med	Siemens ECAT Exact HR+	5 mm	70-90 min	70-90 min	Cerebellar gray matter
Smith 2021	NeuroImage: Clinical	Siemens ECAT Exact HR+	5 mm	70-90 min	70-90 min	Inferior cerebellar gray matter
Mertens 2022 ^^	JCBFM	GE Signa PET/MR	4 mm	90-120 min	90-120 min	Cerebellar gray matter

† J Nucl Med: Journal of Nuclear Medicine; EJNMMI: European Journal of Nuclear Medicine and Molecular Imaging; JCBFM: Journal of Cerebral Blood Flow and Metabolism. \* FWHM = full width at half maximum. \*\*SUVR = standardized uptake value ratio. ^ Salinas et.al., 2020: Only example here to apply a 5 mm cerebellar erosion from the outermost voxels. ^^ Examples here to apply region- and voxel- level partial volume correction on the PET data.

**Supplementary Table 2.** Comparative size (in voxels) of the candidate reference regions and extracerebral masks in the PET space, for a representative subject.

	<b>CerGM</b>	<b>CerGM<sub>3mm</sub></b>	<b>InfCer</b>	<b>WM</b>	<b>WM<sub>4mm</sub></b>	<b>Pons</b>	<b>Extracerebral</b>
<b>ROI size (number of voxels)</b>	22566	6556	10174	88295	15310	3014	121544

**Supplementary Table 3.** Baseline SUV<sub>90-110</sub>, longitudinal changes of SUV<sub>90-110</sub> in candidate reference regions and extracerebral signal at 6-month and 1-year follow-ups.

		Baseline SUV <sub>90-110</sub> (g/mL) (n=32: 24 CN, 8 AD/MCI)	ΔSUV <sub>90-110</sub> at 6-month (n=25: 20 CN, 5 AD/MCI)	ΔSUV <sub>90-110</sub> at 1-year (n=16: 11 CN, 5 AD/MCI)
<b>CerGM</b>	<b>CN</b>	0.62±0.12	0.04±0.07	0.04±0.13
	<b>AD/MCI</b>	0.59±0.17	(-0.09, 0.18)	(-0.15, 0.26)
<b>CerGM<sub>3mm</sub></b>	<b>CN</b>	0.53±0.13	0.05±0.06	0.04±0.11
	<b>AD/MCI</b>	0.51±0.17	(-0.10, 0.16)	(-0.10, 0.26)
<b>PVC CerGM</b>	<b>CN</b>	0.54±0.20	0.03±0.14	-0.002±0.17
	<b>AD/MCI</b>	0.62±0.26	(-0.32, 0.25)	(-0.22, 0.36)
<b>Inferior CerGM</b>	<b>CN</b>	0.67±0.13	0.04±0.08	0.04±0.13
	<b>AD/MCI</b>	0.59±0.17	(-0.12, 0.22)	(-0.10, 0.24)
<b>WM</b>	<b>CN</b>	0.44±0.09*	0.05±0.06	0.02±0.11
	<b>AD/MCI</b>	0.75±0.20*	(-0.05, 0.16)	(-0.24, 0.21)
<b>WM<sub>4mm</sub></b>	<b>CN</b>	0.42±0.11	0.05±0.05	0.02±0.10
	<b>AD/MCI</b>	0.51±0.15	(-0.04, 0.17)	(-0.25, 0.20)
<b>PVC WM</b>	<b>CN</b>	0.34±0.14	0.02±0.09	-0.01±0.12
	<b>AD/MCI</b>	0.56±0.10	(-0.22, 0.20)	(-0.28, 0.19)
<b>Pons</b>	<b>CN</b>	0.29±0.07	0.02±0.05	-0.002±0.07
	<b>AD/MCI</b>	0.32±0.09	(-0.06, 0.08)	(-0.16, 0.13)
<b>PVC Pons</b>	<b>CN</b>	0.18±0.09	0.02±0.06	-0.01±0.09
	<b>AD/MCI</b>	0.27±0.09	(-0.11, 0.13)	(-0.19, 0.16)
<b>Extracerebral</b>	<b>CN</b>	0.79±0.15	0.01±0.12	0.0002±0.16
	<b>AD/MCI</b>	0.74±0.17	(-0.22, 0.39)	(-0.20, 0.38)

\* = significantly different between the CN and AD/MCI groups. Baseline SUV<sub>90-110</sub> are reported as mean±std and longitudinal changes are reported as mean±std (min, max).

**Supplementary Table 4.** Target region distribution volume ratio (DVR) and SUVR<sub>90-110</sub> values (entorhinal, amygdala, inferior temporal and precuneus) for each candidate reference region in the CN and AD/MCI groups. Target region DVR were calculated using two-tissue compartmental model (2TCM), Logan graphical analysis, reference tissue methods (multilinear reference tissue modeling (MRTM2) and reference Logan graphical analysis) and SUVR using the 90-110 min of dynamic data.

	CerGM		CerGM3mm		InfCer		WM		WM <sub>4mm</sub>		Pons	
	CN	AD/MCI	CN	AD/MCI	CN	AD/MCI	CN	AD/MCI	CN	AD/MCI	CN	AD/MCI
<b>2TCM DVR</b>												
Entorhinal	1.1±0.21	1.57±0.44	1.25±0.33	1.74±0.52	1.01±0.15	1.55±0.42	1.56±0.4	1.76±0.35	1.86±0.61	2.32±0.7	2±0.52	2.5±0.89
Amygdala	0.79±0.22	1.35±0.57	0.89±0.25	1.49±0.63	0.74±0.24	1.33±0.57	1.1±0.23	1.51±0.6	1.31±0.4	1.99±0.9	1.4±0.31	2.17±1.11
Inferior Temporal	1.27±0.28	2.06±1.08	1.45±0.43	2.29±1.28	1.16±0.21	2.02±1.04	1.81±0.51	2.26±0.99	2.15±0.77	3.05±1.71	2.31±0.68	3.32±1.99
Precuneus	0.92±0.1	1.56±0.97	1.04±0.13	1.75±1.15	0.86±0.13	1.53±0.93	1.3±0.16	1.71±0.91	1.54±0.26	2.34±1.54	1.67±0.25	2.54±1.76
<b>Logan DVR</b>												
Entorhinal	1.02±0.08	1.53±0.44	1.18±0.15	1.67±0.66	0.91±0.11	1.51±0.41	1.61±0.33	1.69±0.4	1.84±0.44	2.31±0.93	2.03±0.42	2.51±1.04
Amygdala	0.75±0.21	1.27±0.5	0.87±0.25	1.37±0.6	0.68±0.21	1.26±0.53	1.16±0.3	1.38±0.44	1.33±0.37	1.89±0.85	1.46±0.39	2.07±0.98
Inferior Temporal	1.3±0.28	1.93±0.94	1.53±0.49	2.13±1.28	1.14±0.16	1.9±0.9	2.08±0.69	2.1±0.89	2.38±0.85	2.96±1.81	2.63±0.91	3.23±2
Precuneus	0.89±0.11	1.47±0.84	1.02±0.13	1.64±1.14	0.79±0.16	1.45±0.79	1.38±0.19	1.61±0.82	1.58±0.29	2.28±1.61	1.75±0.29	2.48±1.77
<b>MRTM2 DVR</b>												
Entorhinal	1.04±0.12	1.75±0.63	1.13±0.18	1.93±0.82	0.99±0.1	1.77±0.64	1.41±0.23	1.48±0.22	1.58±0.32	2.44±0.84	1.83±0.49	2.75±1.06
Amygdala	0.81±0.11	1.39±0.49	0.86±0.12	1.52±0.65	0.78±0.12	1.41±0.5	1.07±0.13	1.18±0.2	1.19±0.16	1.87±0.64	1.31±0.22	2.16±0.79
Inferior Temporal	1.16±0.15	2.3±1.08	1.28±0.25	2.58±1.41	1.1±0.11	2.33±1.1	1.6±0.32	1.87±0.47	1.8±0.44	3.12±1.4	2.12±0.69	3.74±1.89
Precuneus	0.93±0.08	2.2±1.45	0.99±0.1	2.52±1.86	0.9±0.08	2.24±1.48	1.23±0.11	1.71±0.65	1.38±0.17	2.86±1.78	1.52±0.24	3.63±2.5
<b>Reference Logan DVR</b>												
Entorhinal	1.04±0.12	1.7±0.57	1.13±0.17	1.84±0.71	0.99±0.1	1.73±0.58	1.34±0.21	1.43±0.21	1.46±0.28	2.17±0.65	1.75±0.42	2.5±0.85
Amygdala	0.82±0.12	1.38±0.47	0.87±0.13	1.49±0.6	0.79±0.12	1.4±0.49	1.05±0.12	1.16±0.18	1.15±0.15	1.72±0.47	1.3±0.2	2.02±0.67
Inferior Temporal	1.15±0.15	2.22±1	1.28±0.25	2.44±1.24	1.1±0.12	2.25±1.02	1.5±0.28	1.79±0.44	1.63±0.37	2.83±1.18	2.04±0.62	3.31±1.52
Precuneus	0.93±0.08	2.1±1.3	0.99±0.1	2.33±1.58	0.9±0.08	2.14±1.33	1.2±0.11	1.64±0.6	1.32±0.17	2.67±1.48	1.5±0.24	3.17±1.97
<b>SUVR<sub>90-110</sub></b>												
Entorhinal	1.14±0.16	2.12±0.9	1.34±0.25	2.57±1.4	1.06±0.14	2.17±0.95	1.58±0.28	1.6±0.26	1.73±0.38	2.44±0.84	2.41±0.58	3.84±1.52
Amygdala	0.8±0.21	1.64±0.72	0.94±0.23	1.98±1.1	0.75±0.22	1.68±0.77	1.1±0.19	1.23±0.25	1.19±0.23	1.87±0.64	1.66±0.31	2.97±1.21
Inferior Temporal	1.25±0.18	2.71±1.43	1.47±0.29	3.31±2.11	1.16±0.15	2.78±1.49	1.74±0.37	1.98±0.5	1.92±0.5	3.11±1.4	2.66±0.71	4.94±2.46
Precuneus	0.89±0.14	2.51±1.87	1.04±0.17	3.13±2.68	0.83±0.15	2.58±1.95	1.23±0.14	1.75±0.74	1.34±0.23	2.86±1.77	1.87±0.36	4.58±3.18

**Supplementary Table 5.** Cross-sectional effect size (Hedge's  $g$  coefficients) for differentiating target region  $SUVR_{90-110}$  between diagnostic and  $A\beta$  status groups using the candidate reference regions. Values are expressed as mean (95% confidence interval estimated using bootstrap).

<b>AD/MCI (n=8) and CN (n=24)</b>						
	<b>CerGM</b>	<b>CerGM<sub>3mm</sub></b>	<b>InfCer</b>	<b>WM</b>	<b>WM<sub>4mm</sub></b>	<b>Pons</b>
<b>Entorhinal</b>	1.37 (0.24,2.45)	1.47 (0.3,2.58)	1.1 (0.04,2.09)	-0.04 (-0.82,0.74)	1 (0.02,1.93)	1.1 (0.08,2.07)
<b>Amygdala</b>	1.48 (0.32,2.59)	1.55 (0.36,2.67)	1.21 (0.13,2.25)	0.58 (-0.26,1.41)	1.29 (0.22,2.31)	1.33 (0.23,2.38)
<b>Inferior Temporal</b>	1.29 (0.17,2.35)	1.36 (0.22,2.45)	1.09 (0.04,2.09)	0.45 (-0.4,1.28)	1.04 (0.04,2)	1.11 (0.07,2.1)
<b>Precuneus</b>	1.1 (0.04,2.12)	1.14 (0.06,2.16)	0.99 (-0.05,1.97)	0.86 (-0.13,1.81)	1.06 (0.02,2.06)	1.06 (0.01,2.06)
<b><math>A\beta+</math> (n=18) and <math>A\beta-</math> (n=14)</b>						
	<b>CerGM</b>	<b>CerGM<sub>3mm</sub></b>	<b>InfCer</b>	<b>WM</b>	<b>WM<sub>4mm</sub></b>	<b>Pons</b>
<b>Entorhinal</b>	0.89 (0.17,1.59)	0.78 (0.08,1.47)	0.97 (0.24,1.68)	0.18 (-0.52,0.87)	0.48 (-0.2,1.14)	0.37 (-0.3,1.04)
<b>Amygdala</b>	1.17 (0.4,1.91)	1.19 (0.42,1.94)	1.03 (0.29,1.75)	1.3 (0.53,2.05)	1.11 (0.38,1.82)	0.81 (0.1,1.5)
<b>Inferior Temporal</b>	0.78 (0.07,1.46)	0.84 (0.13,1.53)	0.71 (0.01,1.39)	0.19 (-0.5,0.88)	0.44 (-0.22,1.1)	0.4 (-0.27,1.06)
<b>Precuneus</b>	0.68 (-0.01,1.35)	0.72 (0.02,1.39)	0.64 (-0.05,1.31)	0.45 (-0.22,1.11)	0.55 (-0.11,1.2)	0.5 (-0.17,1.15)

**Supplementary Table 6.** Longitudinal effect size (Hedge's  $g$  coefficients) for detecting changes over a one-year period in target region  $SUVR_{90-110}$  in AD/MCI and in  $A\beta+$  individuals using the candidate reference regions. Values are expressed as mean (95% confidence interval estimated using bootstrap).

	<b>AD/MCI (n=5)</b>					
	<b>CerGM</b>	<b>CerGM<sub>3mm</sub></b>	<b>InfCer</b>	<b>WM</b>	<b>WM<sub>4mm</sub></b>	<b>Pons</b>
<b>Entorhinal</b>	-0.12 (-0.35,0.18)	-0.11 (-0.41,0.2)	-0.13 (-0.57,0.11)	-0.28 (-2.01,0.37)	0.01 (-1.36,0.55)	0.12 (-0.99,0.66)
<b>Amygdala</b>	-0.03 (-0.22,0.17)	-0.05 (-0.44,0.39)	-0.06 (-0.68,0.14)	0.03 (-0.18,0.14)	0.12 (-0.06,0.82)	0.2 (-0.06,1.12)
<b>Inferior Temporal</b>	0 (-0.19,0.56)	-0.02 (-0.26,0.42)	-0.02 (-0.23,1.13)	0.14 (-1.01,0.64)	0.14 (-0.28,0.71)	0.2 (-0.18,0.85)
<b>Precuneus</b>	-0.05 (-0.16,0.17)	-0.07 (-0.28,0.07)	-0.06 (-0.2,0.17)	-0.05 (-0.34,0.26)	-0.02 (-0.44,0.26)	0.03 (-0.27,0.29)
	<b>A<math>\beta</math>+ (n=8)</b>					
	<b>CerGM</b>	<b>CerGM<sub>3mm</sub></b>	<b>InfCer</b>	<b>WM</b>	<b>WM<sub>4mm</sub></b>	<b>Pons</b>
<b>Entorhinal</b>	-0.08 (-0.31,0.11)	-0.09 (-0.37,0.15)	-0.09 (-0.35,0.13)	-0.1 (-0.77,0.51)	0.06 (-0.32,0.47)	0.12 (-0.23,0.53)
<b>Amygdala</b>	0 (-0.11,0.12)	-0.02 (-0.15,0.09)	-0.02 (-0.17,0.13)	0.12 (-0.17,0.47)	0.16 (-0.16,0.56)	0.19 (-0.05,0.51)
<b>Inferior Temporal</b>	-0.02 (-0.28,0.23)	-0.04 (-0.36,0.26)	-0.03 (-0.32,0.25)	-0.01 (-0.55,0.52)	0.08 (-0.26,0.47)	0.13 (-0.25,0.57)
<b>Precuneus</b>	-0.04 (-0.16,0.07)	-0.06 (-0.25,0.11)	-0.04 (-0.19,0.09)	-0.05 (-0.28,0.17)	-0.01 (-0.19,0.18)	0.04 (-0.18,0.29)



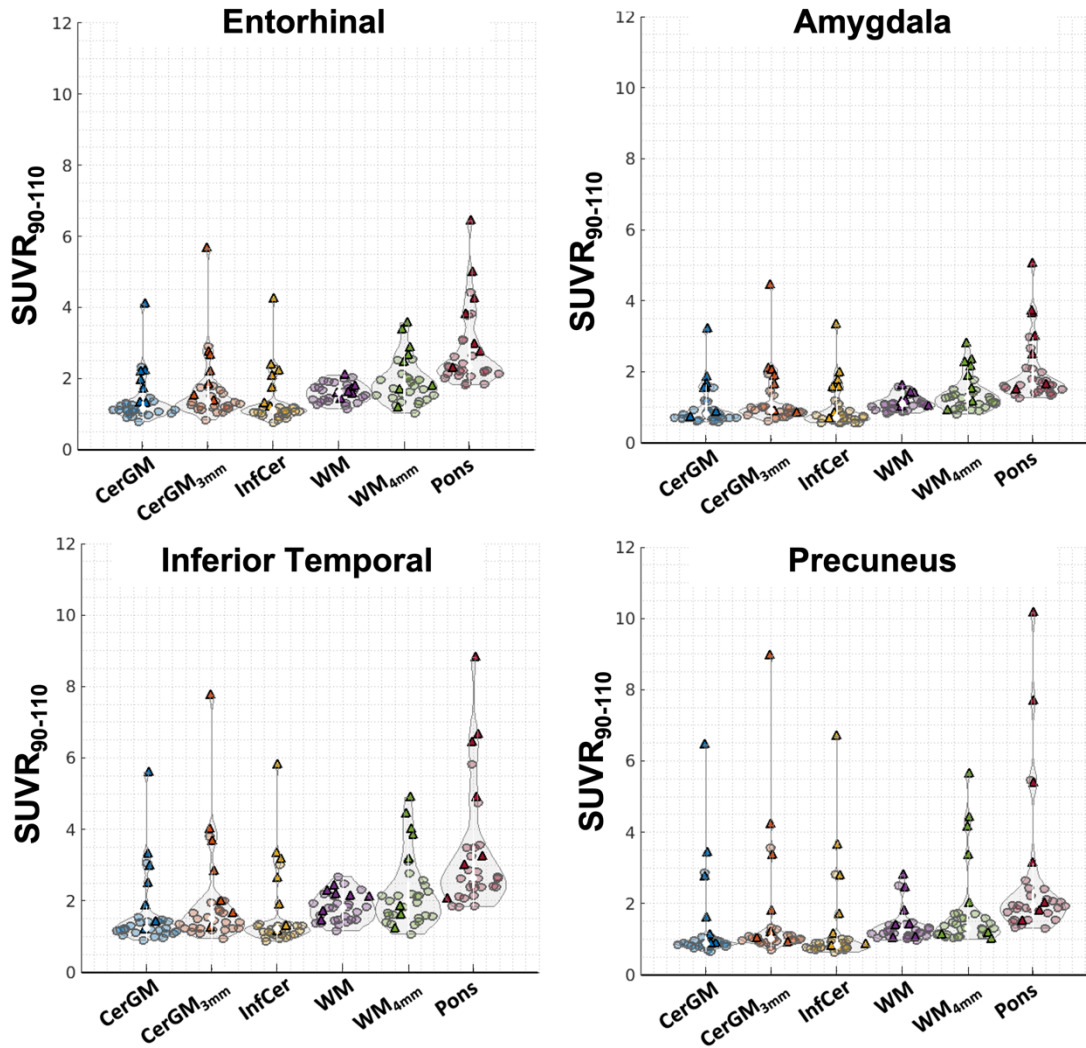
**Supplementary Table 6.** Extracerebral [<sup>18</sup>F]MK-6240 uptake (SUVR<sub>90-110</sub>) at baseline and follow-ups for the high and low extracerebral uptake groups. The SUVR<sub>90-110</sub> values are expressed as mean ± standard deviation.

	<b>High extracerebral (SUVR &gt;1.39)</b>		<b>Low extracerebral (SUVR ≤1.39)</b>	
	<b>n</b>	SUVR <sub>90-110</sub>	<b>n</b>	SUVR <sub>90-110</sub>
<b>Baseline (32)</b>	16 (12 CN, 4 AD/MCI)	2.01±0.70	16 (12 CN, 4 AD/MCI)	1.14±0.14
<b>6 months (26)</b>	12 (8 CN, 4 AD/MCI)	2.04±0.47	13 (12 CN, 1 AD/MCI)	1.07±0.20
<b>1 year (16)</b>	7 (3 CN, 4 AD/MCI)	2.04±0.62	9 (8 CN, 1 AD/MCI)	1.09±0.15

**Supplementary Figure 1.** Target region  $SUVR_{90-110}$  obtained using different reference regions for the CN and AD/MCI subjects.

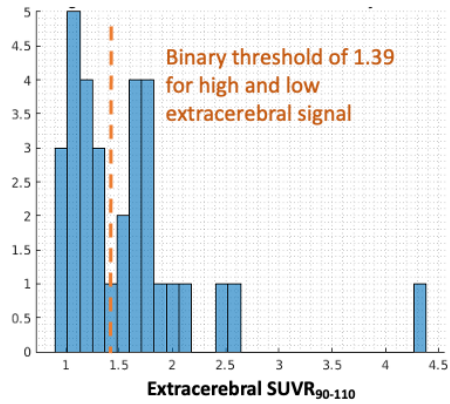
**Target region  $SUVR_{90-110}$  in CN and AD/MCI subjects**

○ CN (n=24)    △ AD/MCI (n=8)



**Supplementary Figure 2.** Extracerebral SUVR<sub>90-110</sub> threshold. **a)** Histogram of extracerebral SUVR<sub>90-110</sub> identified a binary threshold of 1.39, **b)** which separated all subjects into high-extracerebral-signal and low-extracerebral-signal groups. Eroded CerGM by 3mm was used as the reference region.

**(a) Histogram of extracerebral SUVR<sub>90-110</sub>**



**(b) Extracerebral SUVR<sub>90-110</sub>**

