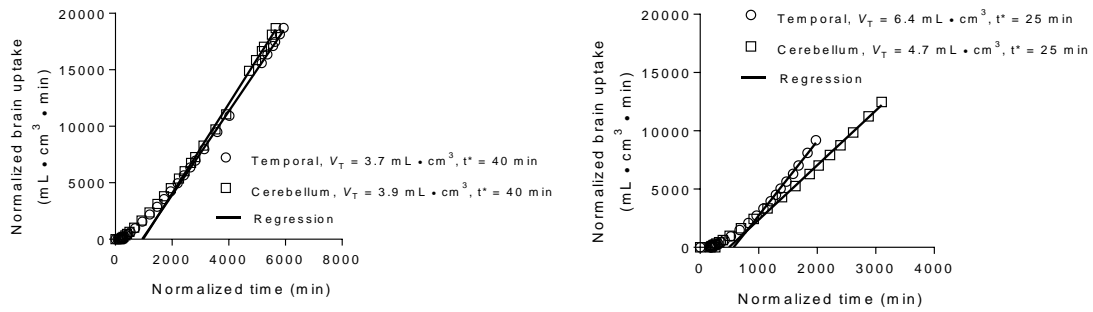
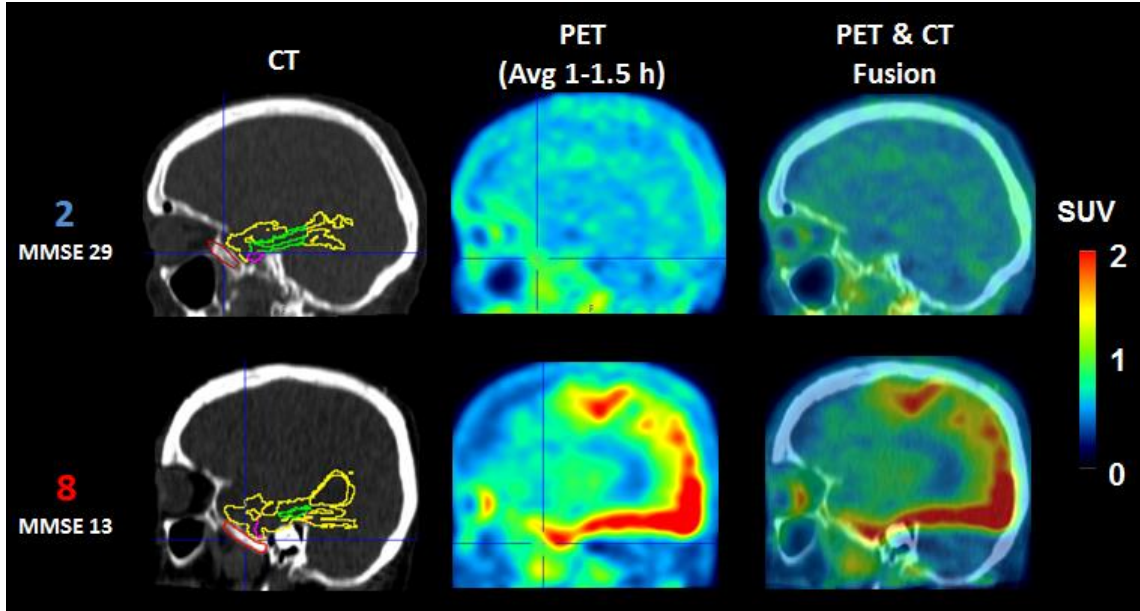


**SUPPLEMENTAL FIGURE 1.** Comparison of total  $V_T$  in various brain regions between cognitively HE subjects (1, 2, and 3) and AD subjects (6, 7, and 10) who received arterial sampling.  $V_T$  values were calculated using unconstrained 2-tissue compartmental model (2-TCM).

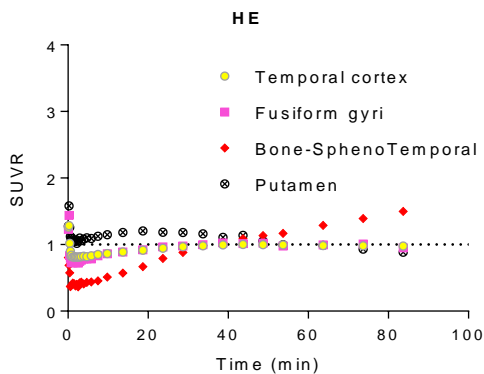


**SUPPLEMENTAL FIGURE 2.** Regional Logan plots (open symbols) and linear regressions (solid lines) derived from invasive Logan plot in a representative HE (Subject 3 – left plot) and AD (Subject 10 – right plot). Normalized time on x-axis is derived by the equation:  $\int_0^T C_p(t)dt/C_t(T)$ . Normalized brain uptake on y-axis is derived by the equation:  $\int_0^T C_t(t)dt/C_t(T)$ .

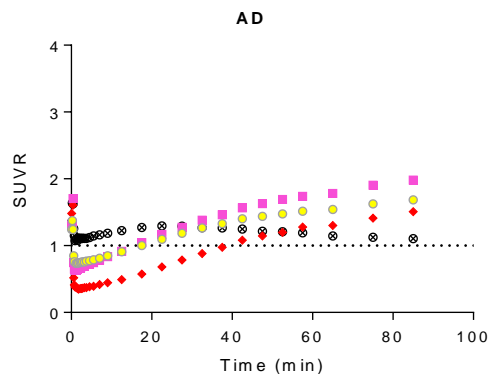
A



B



C



**SUPPLEMENTAL FIGURE 3.** Effect of bone uptake due to potential defluorination on brain cortical uptake. (A) Sagittal cross-sectional images (CT, PET, and PET and CT fusion) of a representative HE (2) subject and AD (8) subject showing atlas-derived regional VOI overlays (temporal cortex in yellow, fusiform gyri in pink, parahippocampal gyri in green) and manually drawn speno-temporal bone VOIs (in red). PET images are averaged between 60-90 min and are color scaled as SUV between 0 and 2. (B) Regional (Brain cortices,

adjacent bone, and putamen) SUVR time course in HE subjects with cerebellar cortex as a reference. Curves represent a mean of 4 subjects. (C) Regional (Brain cortices, adjacent bone, putamen) SUVR time course in AD subjects with cerebellar cortex as reference. Curves represent a mean of 6 subjects.

**SUPPLEMENTAL TABLE 1: CSF, Amyloid PET, and MRI characteristics in AD cohort**

Subject ID	Clinical diagnosis	CSF			Amyloid PET	MRI			
		A $\beta$ 42 (>853 pg/mL)	Total tau (<400 pg/mL)	<sup>181</sup> phosphotau (<65 pg/mL)	(SUVR <1.29)	MTA R	MTA L	ARWMC	GRE Hyperintensities
5	Dementia	439	163	29.4	N/A	2	2	3	8
6	Dementia	N/A	N/A	N/A	N/A	0	1	1	1
7	MCI	707	496	66	1.73	3	3	0	0
8	Dementia	409	1042	164	N/A	3	3	1	0
9	Dementia	N/A	N/A	N/A	N/A	2	3	0	0
10	MCI	782	1336	165	1.71	2	3	0	0

ARWMC = Age-related white matter change (based on reference 1), CSF = Cerebrospinal fluid (cutoff criteria based on reference 2), GRE = Gradient recalled echo, MCI = Mild cognitive impairment, MTA R/L= Medial temporal lobe atrophy score (based on reference 3) right/left, N/A = not applicable, SUVR = Regional standard uptake value ratio. The values between brackets indicate the normal range. The amyloid PET was performed using <sup>18</sup>F-florbetaben.

**SUPPLEMENTAL TABLE 2:** Rate constants and total distribution volumes from two-tissue compartment model in HE subjects

Brain region	Two-tissue compartmental rate constants								Two-tissue $V_T$ (mL • cm <sup>-3</sup> )			
	$K_1$ (mL • cm <sup>-3</sup> • min <sup>-1</sup> )		$k_2$ (min <sup>-1</sup> )		$k_3$ (min <sup>-1</sup> )		$k_4$ (min <sup>-1</sup> )					
Temporal	0.326	(1.2%)	0.135	(2.1%)	0.012	(6.5%)	0.012	(8.4%)	5.2	±	1.3	(2.2%)
Hippocampus	0.283	(1.5%)	0.115	(3%)	0.012	(11.3%)	0.017	(12.3%)	4.4	±	0.9	(2.2%)
Amygdala	0.286	(2.5%)	0.108	(4.6%)	0.008	(18.2%)	0.013	(23.4%)	4.2	±	0.9	(4%)
Caudate	0.274	(1.6%)	0.120	(3.1%)	0.012	(14.1%)	0.016	(12.1%)	3.7	±	0.8	(2%)
Putamen	0.417	(1.3%)	0.124	(2.9%)	0.010	(17.2%)	0.023	(14%)	4.8	±	1.2	(1.4%)
Cerebellar cortex	0.349	(1.2%)	0.143	(2%)	0.010	(6%)	0.010	(9.3%)	5.8	±	1.7	(2.8%)

Rate constants are presented as median values and  $V_T$  values as mean ± SD from 3 subjects. For each brain region, median standard errors are listed in parentheses and are expressed as % of the variable itself.

**SUPPLEMENTAL TABLE 3:** Rate constants and total distribution volumes from two-tissue compartment model in AD subjects

Brain region	Two-tissue compartmental rate constants				Two-tissue $V_T$ (mL • cm <sup>-3</sup> )
	$K_1$ (mL • cm <sup>-3</sup> • min <sup>-1</sup> )	$k_2$ (min <sup>-1</sup> )	$k_3$ (min <sup>-1</sup> )	$k_4$ (min <sup>-1</sup> )	
Temporal	0.336 (1.1%)	0.144 (2.2%)	0.022 (4.1%)	0.015 (5.2%)	5.9 ± 0.9 (2%)
Hippocampus	0.267 (1.8%)	0.117 (4.5%)	0.033 (8.2%)	0.018 (8.4%)	6.0 ± 0.3 (2.9%)
Amygdala	0.234 (2.3%)	0.103 (5.9%)	0.035 (9.1%)	0.011 (13.9%)	9.8 ± 1.5 (6.6%)
Caudate	0.233 (2.5%)	0.141 (5.3%)	0.021 (13.2%)	0.023 (10.5%)	3.1 ± 0.5 (2.1%)
Putamen	0.470 (1.2%)	0.145 (2.8%)	0.021 (9.2%)	0.034 (6.7%)	5.3 ± 1.0 (1%)
Cerebellar cortex	0.335 (0.9%)	0.145 (1.7%)	0.017 (4.6%)	0.016 (6.4%)	5.4 ± 1.3 (1.7%)

Rate constants are presented as median values and  $V_T$  values as mean ± SD from three subjects. For each brain region, median standard errors are listed in parentheses and are expressed as % of the variable itself.

**SUPPLEMENTAL TABLE 4:** SUVRs and DVRs from different models in HE subjects

<b>Brain region</b>	<b>SUVR<sub>60-90 min</sub></b>	<b>DVR<sub>2-TCM</sub></b>	<b>DVR<sub>Logan Plot</sub></b>	<b>DVR<sub>LoganRefTissue</sub></b>
Temporal	0.98 ± 0.07	0.88 ± 0.08	0.92 ± 0.06	0.94 ± 0.04
Hippocampus	0.93 ± 0.10	0.76 ± 0.12	0.82 ± 0.09	0.91 ± 0.07
Amygdala	0.84 ± 0.11	0.70 ± 0.08	0.75 ± 0.09	0.84 ± 0.07
Caudate	0.79 ± 0.06	0.67 ± 0.10	0.70 ± 0.07	0.79 ± 0.04
Putamen	0.91 ± 0.04	0.84 ± 0.11	0.89 ± 0.06	1.01 ± 0.06

DVR<sub>2TCM</sub> and DVR<sub>Logan Plot</sub> values are Mean ± SD from n = 3 HE subjects, whereas SUVR<sub>60-90 min</sub> and DVR<sub>LoganRefTissue</sub> are from n = 4

HE subjects



**SUPPLEMENTAL TABLE 5: SUVRs and DVRs from different models in AD subjects**

<b>Brain region</b>	<b>SUVR<sub>60-90 min</sub></b>	<b>DVR<sub>2TCM</sub></b>	<b>DVR<sub>Logan Plot</sub></b>	<b>DVR<sub>LoganRefTissue</sub></b>
Temporal	1.64 ± 0.72	1.12 ± 0.21	1.12 ± 0.22	1.39 ± 0.57
Hippocampus	1.37 ± 0.25	1.16 ± 0.05	1.20 ± 0.13	0.99 ± 0.50
Amygdala	1.67 ± 0.40	1.61 ± 0.14	1.73 ± 0.15	1.20 ± 0.60
Caudate	0.71 ± 0.16	0.62 ± 0.06	0.65 ± 0.08	0.68 ± 0.10
Putamen	1.15 ± 0.35	0.97 ± 0.12	1.03 ± 0.08	1.19 ± 0.20

DVR<sub>2TCM</sub> and DVR<sub>Logan Plot</sub> values are Mean ± SD from n = 3 AD subjects, whereas SUVR<sub>60-90 min</sub> and DVR<sub>LoganRefTissue</sub> are from n = 6 AD subjects

**SUPPLEMENTAL TABLE 6: Correlations between SUVR and DVRs across subjects**

Subject	SUVR <sub>60-90min</sub> vs. DVR <sub>2-TCM</sub>			SUVR <sub>60-90min</sub> vs. DVR <sub>Logan Plot</sub>			SUVR <sub>60-90min</sub> vs. DVR <sub>Logan Ref.Tissue</sub>		
	R <sup>2</sup>	Slope	Intercept	R <sup>2</sup>	Slope	Intercept	R <sup>2</sup>	Slope	Intercept
1	0.61	0.64	0.39	0.78	0.8	0.21	0.88	1.18	-0.17
2	0.54	0.80	0.29	0.65	1.01	0.06	0.5	1.02	0
3	0.77	0.99	0.05	0.78	0.86	0.17	0.63	1.07	-0.08
4	N/A	N/A	N/A	N/A	N/A	N/A	0.31	0.9	0.14
5	N/A	N/A	N/A	N/A	N/A	N/A	0.85	1.17	-0.01
6	0.94	1.14	-0.09	0.95	1.13	-0.12	0.83	1.47	-0.44
7	0.96	1.06	-0.09	0.94	1	0	0.79	1.36	-0.29
8	N/A	N/A	N/A	N/A	N/A	N/A	0.98	1.08	0.14
9	N/A	N/A	N/A	N/A	N/A	N/A	0.98	1.38	-0.22
10	0.96	1.13	-0.09	0.96	1.11	-0.08	0.9	1.51	-0.45

N/A: Not applicable due to no arterial sampling.

## SUPPLEMENTAL REFERENCES

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