

# Alzheimer disease brain atrophy subtypes are associated with cognition and rate of decline

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## ABSTRACT

**Objective:** To test the hypothesis that cortical and hippocampal volumes, measured in vivo from volumetric MRI (vMRI) scans, could be used to identify variant subtypes of Alzheimer disease (AD) and to prospectively predict the rate of clinical decline.

**Methods:** Amyloid-positive participants with AD from the Alzheimer's Disease Neuroimaging Initiative (ADNI) 1 and ADNI2 with baseline MRI scans ( $n = 229$ ) and 2-year clinical follow-up ( $n = 100$ ) were included. AD subtypes (hippocampal sparing [HpSp<sub>MRI</sub>], limbic predominant [LP<sub>MRI</sub>], typical AD [tAD<sub>MRI</sub>]) were defined according to an algorithm analogous to one recently proposed for tau neuropathology. Relationships between baseline hippocampal volume to cortical volume ratio (HV:CTV) and clinical variables were examined by both continuous regression and categorical models.

**Results:** When participants were divided categorically, the HpSp<sub>MRI</sub> group showed significantly more AD-like hypometabolism on <sup>18</sup>F-fluorodeoxyglucose-PET ( $p < 0.05$ ) and poorer baseline executive function ( $p < 0.001$ ). Other baseline clinical measures did not differ across the 3 groups. Participants with HpSp<sub>MRI</sub> also showed faster subsequent clinical decline than participants with LP<sub>MRI</sub> on the Alzheimer's Disease Assessment Scale, 13-Item Subscale (ADAS-Cog<sub>13</sub>), Mini-Mental State Examination (MMSE), and Functional Assessment Questionnaire (all  $p < 0.05$ ) and tAD<sub>MRI</sub> on the MMSE and Clinical Dementia Rating Sum of Boxes (CDR-SB) (both  $p < 0.05$ ). Finally, a larger HV:CTV was associated with poorer baseline executive function and a faster slope of decline in CDR-SB, MMSE, and ADAS-Cog<sub>13</sub> score ( $p < 0.05$ ). These associations were driven mostly by the amount of cortical rather than hippocampal atrophy.

**Conclusions:** AD subtypes with phenotypes consistent with those observed with tau neuropathology can be identified in vivo with vMRI. An increased HV:CTV ratio was predictive of faster clinical decline in participants with AD who were clinically indistinguishable at baseline except for a greater dysexecutive presentation. *Neurology*® 2017;89:2176-2186

## GLOSSARY

**A $\beta$**  =  $\beta$ -amyloid; **AD** = Alzheimer disease; **ADAS-Cog<sub>13</sub>** = Alzheimer's Disease Assessment Scale, 13-Item Subscale; **ADNI** = Alzheimer's Disease Neuroimaging Initiative; **CDR-SB** = Clinical Dementia Rating Sum of Boxes; **CTV** = cortical total volume; **EOAD** = early-onset Alzheimer disease; **FAQ** = Functional Assessment Questionnaire; **FDG** = <sup>18</sup>F-fluorodeoxyglucose; **GMV** = gray matter volume; **HCI** = hypometabolic convergence index; **HpSp** = hippocampal sparing; **HV** = hippocampal volume; **LOAD** = late-onset Alzheimer disease; **LP** = limbic predominant; **MMSE** = Mini-Mental State Examination; **NFT** = neurofibrillary tangle; **tAD** = typical Alzheimer disease; **TMT** = Trail Making Test; **vMRI** = volumetric MRI.

When tracked longitudinally with cognitive or functional instruments, people with Alzheimer disease (AD) exhibit varying rates of clinical decline. Emerging evidence links this heterogeneity to differences in the underlying biomarker and neuropathology profiles. Recent neuropathology studies have sought to formalize one aspect of this variability by defining AD subtypes on the basis of the different relative densities of pathologic tau deposits in cortical and hippocampal regions of participants with equivalently staged AD.<sup>1,2</sup> These categorical Murray-Dickson subtypes, called hippocampal sparing (HpSp), typical AD (tAD), and limbic predominant (LP),

Supplemental data  
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Data used in preparation of this article were obtained from the Alzheimer's Disease Neuroimaging Initiative (ADNI) database ([adni.loni.usc.edu](http://adni.loni.usc.edu)). As such, the investigators within the ADNI contributed to the design and implementation of ADNI and/or provided data but did not participate in analysis or writing of this report. A complete listing of ADNI investigators can be found in the coinvestigators list at [Neurology.org](http://Neurology.org).

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were associated with differences in age at diagnosis and death, clinical presentation, and rate of antemortem clinical progression, with individuals with the HpSp variant being younger, more commonly showing an atypical clinical presentation, and declining faster.<sup>1,2</sup> In vivo brain atrophy measurements have shown varying anatomic patterns and degree of atrophy across participants, with increased cortical atrophy in a subgroup associated with more executive dysfunction reminiscent of the HpSp subtype.<sup>3–5</sup> Moreover, a within-participant comparison demonstrated that the ratio of cortical to hippocampal volumes (HVs) from antemortem volumetric MRI (vMRI) correlates with the postmortem tau neuropathologic variant.<sup>6</sup>

The goal of this study was to test the hypothesis that measures of regional cortical and HV, measured in vivo from vMRI, can be used to define disease subtypes with phenotypes consistent with those based on tau neuropathology and that these features would prospectively predict differential clinical presentations and rates of clinical decline in participants with AD, explaining part of the variability in symptomatology and progression.

**METHODS Participant sample.** Data used in the preparation of this article were obtained from the Alzheimer's Disease Neuroimaging Initiative (ADNI) database (<http://adni.loni.usc.edu>). Appendix e-1 and table e-1 at [Neurology.org](http://www.adni-info.org), <http://www.adni-info.org>, <http://adni.loni.usc.edu>, and previous reports<sup>7–20</sup> give more information.

**Standard protocol approvals, registrations, and patient consents.** Written informed consent was obtained according to the Declaration of Helsinki, and procedures were approved by site-specific Institutional Review Boards for the Protection of Human Subjects.

We selected amyloid-positive participants with AD from ADNI, diagnosed as previously described (<http://www.adni-info.org>), with baseline 3-dimensional T1 magnetization-prepared rapid acquisition gradient echo vMRI scans ( $n = 229$ ). Amyloid positivity was defined as having a CSF  $\beta$ -amyloid<sub>1–42</sub> ( $A\beta_{1–42}$ )  $< 192$  pg/mL on the University of Pennsylvania assay.<sup>21</sup> If CSF was not available, a <sup>18</sup>F-florbetapir-PET cortical standardized uptake value ratio  $> 1.11$  based on the University of California, Berkeley quantification was used.<sup>22</sup> This cohort was used to define the AD subtypes and to assess baseline demographics, age at onset, memory, and executive function.<sup>23,24</sup> In addition, the mean hypometabolic convergence index (HCI),<sup>25</sup> a measure of the severity of an AD-like hypometabolism pattern on <sup>18</sup>F-fluorodeoxyglucose (FDG)-PET provided in the ADNI database, was assessed.

To characterize longitudinal changes, we also evaluated a sub-cohort of participants with AD with 2-year clinical follow-up scores on the Mini-Mental State Examination (MMSE), Alzheimer's Disease Assessment Scale, 13-Item Subscale (ADAS-

Cog<sub>13</sub>), Clinical Dementia Rating Sum of Boxes (CDR-SB), executive and memory composite, and Functional Assessment Questionnaire (FAQ). These tests were administered as previously described (<http://www.adni-info.org>). The rate of change in these scales was estimated as the slope of change from baseline to the 2-year visit, including all intermediate visits. Longitudinal analyses were completed only for those who had complete data (at baseline and 6, 12, and 24 months), including  $n = 100$  for CDR-SB and FAQ,  $n = 99$  for MMSE and memory composite,  $n = 97$  for executive function composite, and  $n = 88$  participants for the ADAS-Cog<sub>13</sub>. Those excluded from the longitudinal analysis for missing data ( $n = 119$ ) were not different from those included except they had a higher baseline ADAS-Cog<sub>13</sub> and a shorter disease duration and were more likely not to be non-Hispanic white ( $p < 0.05$ , data not shown).

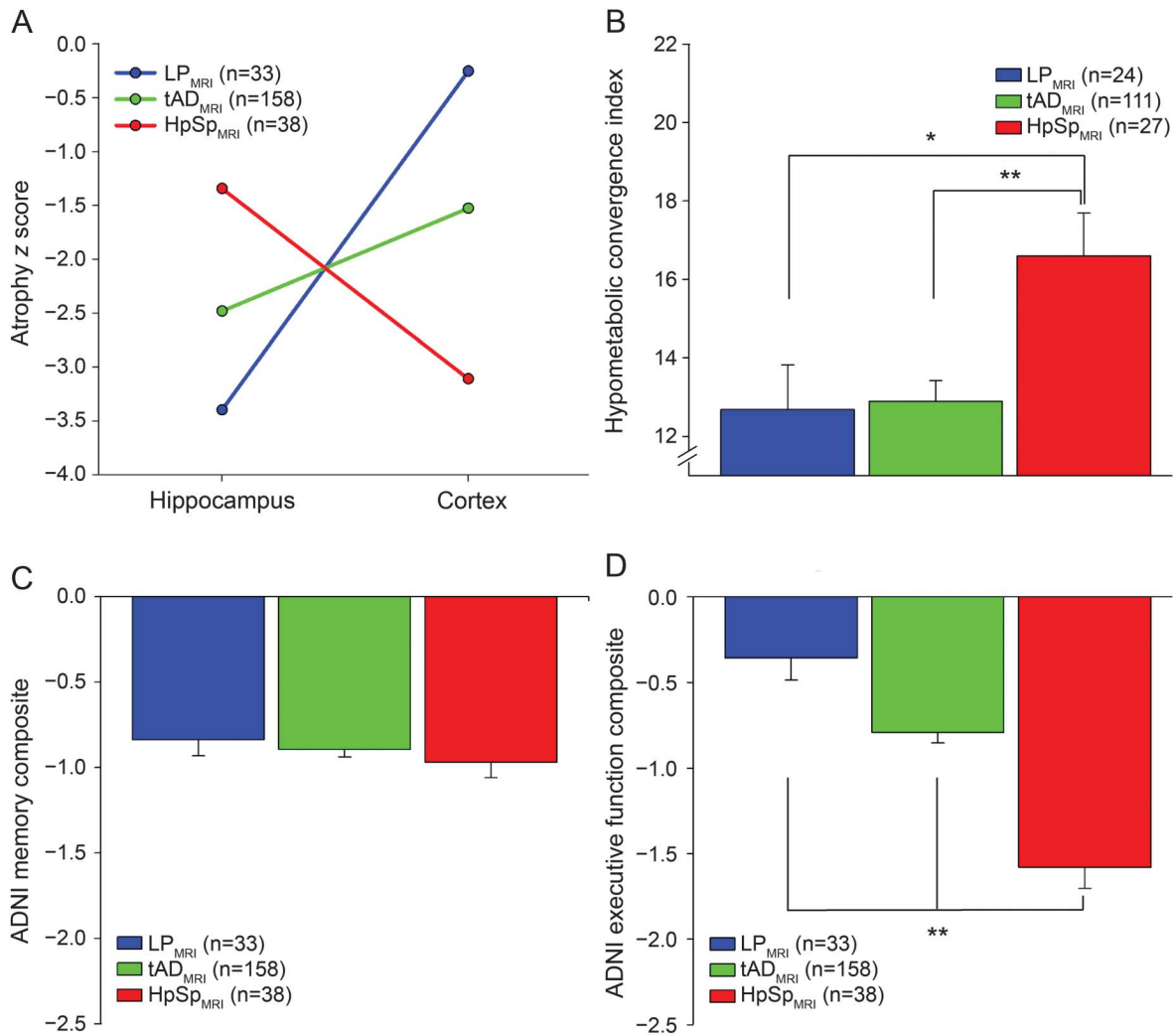
**CSF measures.** CSF amyloid and tau analytes were collected and processed as described<sup>15</sup> and downloaded from the Laboratory of Neuro Imaging site (<http://adni.loni.usc.edu>).

**vMRI analysis and endpoints.** Volumetric measures were calculated from the 3-dimensional T1 images with FreeSurfer (version 5.1). Specifically, left and right gray matter volumes (GMVs) from lateral frontal (caudal and rostral midfrontal, pars opercularis, pars triangularis), superior temporal, and lateral parietal (inferior parietal, superior parietal, supramarginal) cortices in both cerebral hemispheres were summed to provide a measure of bilateral cortical total volume (CTV)<sup>1,6</sup> (appendix e-1, *Freesurfer Regions*). HVs were also summed to create a bilateral total. Both the CTV and HV measures were preadjusted for the effects of intracranial volume, scanner strength (1.5T vs 3T), age, and sex with  $\beta$  coefficients estimated with a regression model estimated on all amyloid-negative, stable, cognitively normal controls from ADNI (see appendix e-1, *Pre-adjustment Formula for Volumetric Measures*). The residual values for CTV and HV were then used to calculate the HV:CTV ratio (see appendix e-1, equation 3).

**Definition of AD subtypes.** In the original presentation of the subtype algorithm, the HpSp, tAD, and LP subtypes were defined with a 2-step procedure based on the neurofibrillary tangle (NFT) counts in the hippocampus and cortical regions.<sup>1</sup> In our study, the HV:CTV ratio was first split (stage 1) at the 25th and 75th percentiles. Participants with HV:CTV ratios below the 25th percentile were provisionally designated as having LP<sub>MRI</sub> (HV:CTV ratio  $\leq 0.0408$ ,  $n = 57$ ); those with HV:CTV ratios above the 75th percentile were provisionally designated as having HpSp<sub>MRI</sub> (HV:CTV ratio  $\geq 0.0501$ ,  $n = 57$ ); and the remainder were considered to have tAD<sub>MRI</sub> ( $n = 115$ ). In a second step (stage 2), only participants with HpSp<sub>MRI</sub> whose HV was greater than the median adjusted HV (median = 5,726.20 cm<sup>3</sup>) and CTV was less than the median adjusted CTV (median = 128,916.82 cm<sup>3</sup>) were considered as definitively having HpSp<sub>MRI</sub> ( $n = 33$ ). Furthermore, only participants with LP<sub>MRI</sub> whose CTV was greater than the median adjusted CTV and HV was less than the median adjusted HV were retained as having LP<sub>MRI</sub> ( $n = 38$ ). The remainder of participants were reclassified as tAD<sub>MRI</sub> ( $n = 158$ ). Note that in our study, designations are reversed to reflect that MRI volumes decrease with disease severity, in contrast to counts of NFT pathology, which increase. Figure 1A displays the relative hippocampal to cortical atrophy in the 3 subtypes.

Because a single-step procedure would be logistically simpler to operationalize and because in a direct comparison of antemortem MRI to postmortem pathologic subtypes the simple ratio of

**Figure 1** Difference in baseline memory, executive function, and HCl between baseline atrophy subtypes



(A) Relative atrophy in the hippocampus and cortex is represented as a z score relative to the amyloid-negative cognitively normal population from the Alzheimer’s Disease Neuroimaging Initiative (ADNI). The limbic predominant (LP<sub>MRI</sub>; blue) subtype shows substantial hippocampal atrophy and limited cortical atrophy. The hippocampal sparing (HpSp<sub>MRI</sub>; red) subtype shows the opposite pattern, with greater cortical atrophy than hippocampal atrophy. The typical Alzheimer disease presentation (tAD<sub>MRI</sub>; green) shows nearly equal relative atrophy in both the hippocampus and cortex. (B) A significant difference by subtype is also observed for the <sup>18</sup>F-fluorodeoxyglucose hypometabolic convergence index (HCl), a measure of hypometabolism in typical AD cortical regions ( $p = 0.008$ ). In particular, the HpSp<sub>MRI</sub> subtype shows the greatest level of hypometabolism on this measure relative to the other subtypes (LP, tAD,  $p < 0.05$ ). (C) Finally, the atrophy subtypes (LP<sub>MRI</sub>, tAD<sub>MRI</sub>, and HpSp<sub>MRI</sub>) do not show significant differences in memory performance at baseline ( $p > 0.05$ ). (D) However, the HpSp<sub>MRI</sub> subtype is associated with significantly reduced baseline executive function relative to the tAD<sub>MRI</sub> or LP<sub>MRI</sub> subtype ( $p < 0.001$ ). \* $p < 0.05$ , \*\* $p < 0.01$ .

hippocampal-to-cortical GMW was found to significantly predict the postmortem neuropathologic tau subtype,<sup>6</sup> we also assessed the subgroups obtained after stage 1 only of the algorithm.

**Statistical analyses.** Relationships between subtype categories and age at onset, continuous demographic variables, and baseline clinical, cognitive, CSF A $\beta_{1-42}$ , total tau, and phosphorylated tau<sub>181</sub> and FDG HCl variables were assessed with a 1-way analysis of covariance. The relationship between subtype category and 2-year change on clinical and cognitive measures was assessed with a repeated-measures analysis of covariance with correction for sphericity. The relationships of subtype category and categorical demographic and genetic variables were evaluated with a  $\chi^2$  test. Finally, the linear relationships between baseline HV, CTV, or HV:CTV as continuous variables and slope of the 2-year change in clinical and cognitive measures were assessed with a stepwise linear regression model. Significant associations

between HV:CTV and clinical and cognitive measures were then evaluated with a partial Pearson correlation. Age, sex, and years of education were included in all models when appropriate. A threshold of  $\alpha = 0.05$  was used for statistical significance, and post hoc pairwise comparisons between categories were corrected for multiple comparisons with Bonferroni adjustment. SPSS version 24.0 (SPSS Inc, Chicago, IL) was used for all statistical analyses.

**RESULTS Demographics.** The characteristics of the sample by atrophy subtypes are summarized in table 1. The prevalence of the atypical subtypes was 31% (14.4% for HpSp<sub>MRI</sub> and 16.6% for LP<sub>MRI</sub>) after stage 2 of the algorithm. People assigned to the HpSp<sub>MRI</sub> subtype were on average younger than

**Table 1** Baseline and clinical change characteristics of overall sample and categorical Murray-Dickson subtypes determined from MRI volumetry (mean ± SD)

	Baseline sample (n = 229)					Longitudinal sample (n = 100)				
	LP <sub>MRI</sub>	tAD <sub>MRI</sub>	HpSp <sub>MRI</sub>	p Value	Pairwise (p < 0.05)	LP <sub>MRI</sub>	tAD <sub>MRI</sub>	HpSp <sub>MRI</sub>	p Value	Pairwise (p < 0.05)
n	33	158	38	NA	14	67	19	NA		
Female, %	30.3	47.5	34.2	0.097	NA	28.6	50.7	36.8	NS	NA
Age, y	75.5 ± 7.2	74.3 ± 7.7	71.1 ± 9.2	0.043	None	73.9 ± 7.3	75.3 ± 7.6	69.7 ± 8.4	0.023	tAD <sub>MRI</sub> > HpSp <sub>MRI</sub>
Education, y	15.3 ± 2.2	15.3 ± 3.1	16.4 ± 2.6	NS	None	15.5 ± 2.3	14.7 ± 3.3	16.8 ± 2.6	0.034	HpSp <sub>MRI</sub> > tAD <sub>MRI</sub>
Non-Hispanic white, %	93.9	90.5	86.8	NS	NA	100	95.5	94.7	NS	NA
Age at onset, y <sup>a</sup>	72.9 ± 7.7	72.0 ± 8.0	69.2 ± 9.9	NS	None	70.0 ± 7.9	72.3 ± 8.1	67.2 ± 9.2	0.062	NA
EOAD (<65 y), % <sup>a</sup>	9.4	18.4	36.8	0.01	NA	15.4	17.9	42.1	0.067	NA
Time since onset, y <sup>a</sup>	3.0 ± 2.6	2.8 ± 2.7	2.3 ± 2.1	NS	None	4.2 ± 3.1	3.6 ± 2.8	2.9 ± 2.4	NS	None
APOE ε4+, % <sup>b</sup>	81.8	76.9	52.8	0.006	NA	92.9	85.1	52.6	0.003	NA
% MAPT H1/H1 <sup>c,d</sup>	61.3	70.3	57.6	NS	NA	50.0	75.4	63.1	NS	NA
Follow-up time, y						2.1 ± 0.1	2.1 ± 0.1	2.2 ± 0.1	NS	NA
CSF Aβ <sub>1-42</sub> , pg/mL <sup>e,f</sup>	126.5 ± 26.6	131.4 ± 20.8	128.7 ± 23.0	NS	NA	121.0 ± 28.6	131.2 ± 20.8	134.6 ± 21.5	NS	NA
CSF total tau, pg/mL <sup>g,h</sup>	129.6 ± 51.7	131.0 ± 62.0	130.8 ± 67.8	NS	NA	123.4 ± 44.6	127.4 ± 56.7	129.1 ± 66.8	NS	NA
CSF p-tau <sub>181</sub> , pg/mL <sup>e,f</sup>	56.3 ± 35.5	54.0 ± 31.7	47.8 ± 17.7	NS	NA	50.5 ± 22.6	51.6 ± 30.5	43.3 ± 17.1	NS	NA
Mean global cortical <sup>18</sup> F-florbetapir SUVR <sup>i,j</sup>	1.4 ± 0.2	1.4 ± 0.2	1.4 ± 0.2	NS	NA	1.3 ± 0.1	1.5 ± 0.1	1.3 ± 0.4	0.014	tAD <sub>MRI</sub> > LP <sub>MRI</sub>
CDR-SB score	4.4 ± 1.9	4.5 ± 1.6	4.2 ± 1.3	NS	None	4.2 ± 1.5	4.3 ± 1.6	4.1 ± 1.4	NS	None
Δ CDR-SB score						1.4 ± 1.5	1.4 ± 1.1	2.5 ± 1.6	0.010	HpSp <sub>MRI</sub> > tAD <sub>MRI</sub>
FAQ score <sup>k</sup>	13.0 ± 6.8	13.0 ± 6.8	12.9 ± 6.7	NS	None	13.9 ± 7.0	12.1 ± 6.8	10.9 ± 6.7	NS	None
Δ FAQ score						2.2 ± 2.3	3.6 ± 2.4	4.9 ± 2.5	0.012	HpSp <sub>MRI</sub> > LP <sub>MRI</sub>
MMSE score	23.4 ± 1.6	23.2 ± 2.1	23.0 ± 2.2	NS	None	23.7 ± 1.5	23.2 ± 2.0	23.2 ± 1.6	NS	None
Δ MMSE score <sup>l</sup>						-1.8 ± 3.0	-1.8 ± 2.0	-3.9 ± 2.5	0.003	LP <sub>MRI</sub> , tAD <sub>MRI</sub> > HpSp <sub>MRI</sub>
ADAS-Cog <sub>13</sub> score <sup>m,n</sup>	29.3 ± 6.2	30.5 ± 8.3	31.6 ± 9.9	NS	None	26.8 ± 6.1	28.8 ± 6.7	30.4 ± 9.1	NS	None
Δ ADAS-Cog <sub>13</sub> score <sup>o</sup>						2.2 ± 3.3	4.7 ± 3.9	7.2 ± 3.9	0.004	HpSp <sub>MRI</sub> > LP <sub>MRI</sub>
Memory composite score	-0.8 ± 0.4	-0.9 ± 0.5	-1.0 ± 0.7	NS	None	-0.8 ± 0.3	-0.8 ± 0.5	-0.9 ± 0.7	NS	None

Continued

Table 1 Continued

	Baseline sample (n = 229)					Longitudinal sample (n = 100)				
	LP <sub>MRI</sub>	tAD <sub>MRI</sub>	HpSp <sub>MRI</sub>	p Value	Pairwise (p < 0.05)	LP <sub>MRI</sub>	tAD <sub>MRI</sub>	HpSp <sub>MRI</sub>	p Value	Pairwise (p < 0.05)
$\Delta$ Memory composite score <sup>p</sup>						-0.1 ± 0.2	-0.2 ± 0.2	-0.3 ± 0.2	0.041	None
Executive function composite score	-0.4 ± 0.8	-0.8 ± 0.8	-1.6 ± 0.7	<0.001	LP <sub>MRI</sub> > tAD <sub>MRI</sub> > HpSp <sub>MRI</sub>	-0.3 ± 1.0	-0.8 ± 0.7	-1.5 ± 0.7	<0.001	LP <sub>MRI</sub> , tAD <sub>MRI</sub> > HpSp <sub>MRI</sub>
$\Delta$ Executive function composite score <sup>q</sup>						-0.2 ± 0.3	-0.3 ± 0.3	-0.3 ± 0.3	NS	None

Abbreviations: A $\beta$  =  $\beta$ -amyloid; ADAS-Cog<sub>13</sub> = Alzheimer's Disease Assessment Scale-13-Item Subscale; APOE = apolipoprotein E; CDR-SB = Clinical Dementia Rating Scale-Sum of Boxes; EOAD = early-onset Alzheimer disease (age <65 years); FAQ = Functional Assessment Questionnaire; HpSp<sub>MRI</sub> = hippocampal sparing; LP<sub>MRI</sub> = limbic predominant; MAPT = microtubule associated protein tau; MMSE = Mini-Mental State Examination; NA = not available; p-tau181 = tau phosphorylated at threonine 181; SUVR = standardized uptake value ratio (normalized to whole cerebellum); tAD<sub>MRI</sub> = typical AD.

For p values: female, non-Hispanic white, APOE  $\epsilon$ 4+, and MAPT haplotype based on  $\chi^2$  test; all others based on analysis of covariance.

<sup>a</sup>One participant missing age at onset, percent EOAD, and years since onset from both cross-sectional (CS) and longitudinal analysis (1 LP<sub>MRI</sub>).

<sup>b</sup>Four participants missing APOE genotype from CS analysis (2 tAD<sub>MRI</sub>, 2 HpSp<sub>MRI</sub>).

<sup>c</sup>Seventeen participants missing MAPT genotype from CS analysis (1 LP<sub>MRI</sub>, 10 tAD<sub>MRI</sub>, 5 HpSp<sub>MRI</sub>).

<sup>d</sup>Two participants missing MAPT genotype from longitudinal analysis (2 tAD<sub>MRI</sub>).

<sup>e</sup>Fifteen participants missing CSF A $\beta$ <sub>1-42</sub> and p-tau from CS analysis (2 LP<sub>MRI</sub>, 9 tAD<sub>MRI</sub>, 4 HpSp<sub>MRI</sub>).

<sup>f</sup>One participant missing CSF A $\beta$ <sub>1-42</sub> and p-tau from longitudinal analysis (1 tAD<sub>MRI</sub>).

<sup>g</sup>Twenty-two participants missing CSF total tau from CS analysis (5 LP<sub>MRI</sub>, 13 tAD<sub>MRI</sub>, 4 HpSp<sub>MRI</sub>).

<sup>h</sup>Three participants missing CSF total tau from longitudinal analysis (1 LP<sub>MRI</sub>, 2 tAD<sub>MRI</sub>).

<sup>i</sup>Ninety-five participants missing <sup>18</sup>F-florbetapir SUVR from CS analysis (10 LP<sub>MRI</sub>, 63 tAD<sub>MRI</sub>, 12 HpSp<sub>MRI</sub>).

<sup>j</sup>Seventy-four participants missing <sup>18</sup>F-florbetapir SUVR from longitudinal analysis (8 LPMRI, 49 tADMRI, 17 HpSpMRI).

<sup>k</sup>One participant missing baseline FAQ from CS analysis (1 LPMRI).

<sup>l</sup>One participant missing  $\Delta$  MMSE from longitudinal analysis (1 HpSpMRI).

<sup>m</sup>Four participants missing baseline ADAS-Cog<sub>13</sub> from CS analysis (3 tADMRI, 1 HpSpMRI).

<sup>n</sup>One participant missing baseline ADAS-Cog<sub>13</sub> from longitudinal analysis (1 tADMRI).

<sup>o</sup>Seven participants missing  $\Delta$  ADAS-Cog<sub>13</sub> from longitudinal analysis (2 LPMRI, 2 tADMRI, 3 HpSpMRI).

<sup>p</sup>One participant missing  $\Delta$  memory composite from longitudinal analysis (1 LPMRI).

<sup>q</sup>Three participants missing  $\Delta$  executive function composite from longitudinal analysis (1 tADMRI, 2 HpSpMRI).

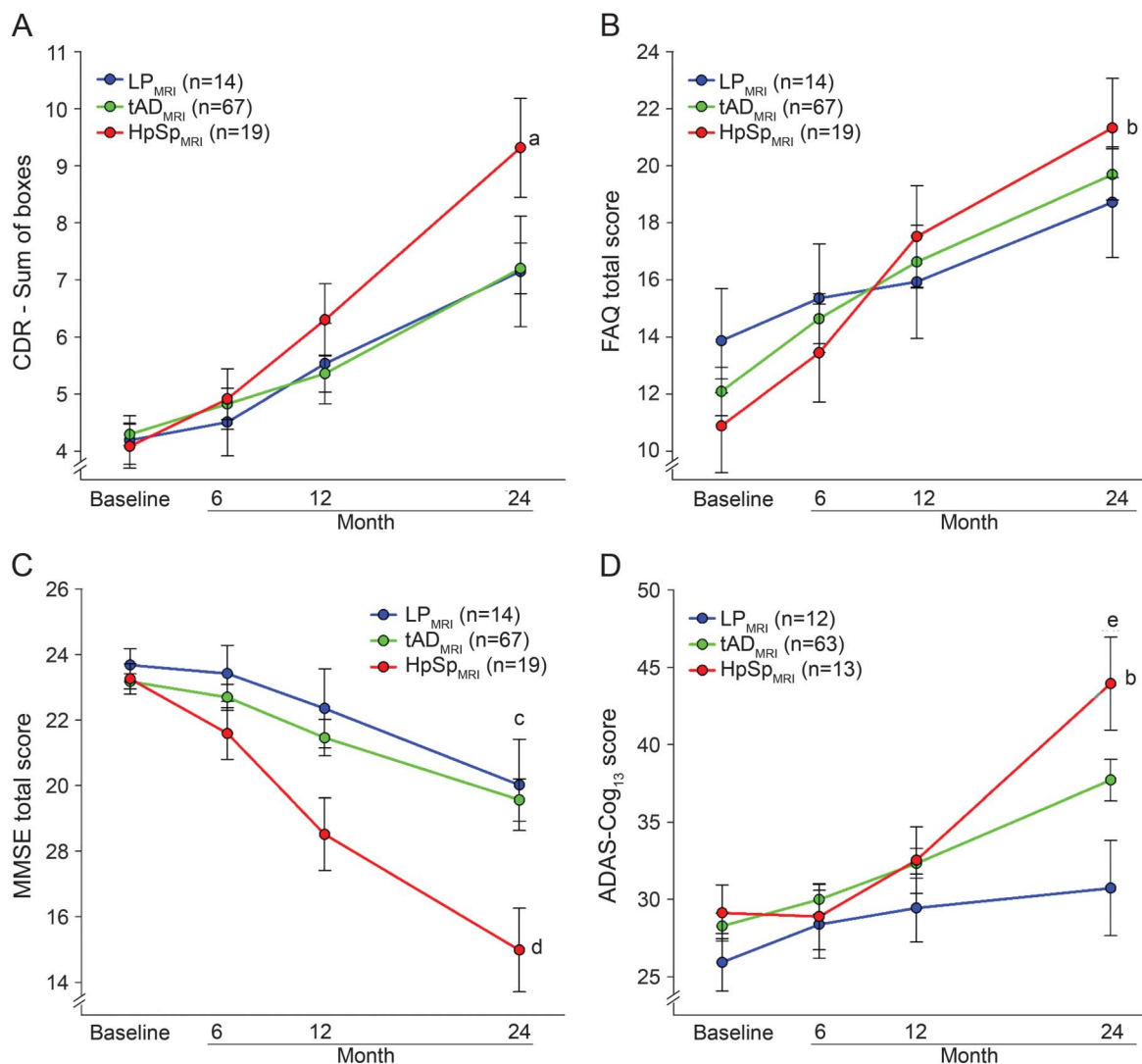
those in both the  $tAD_{MRI}$  and  $LP_{MRI}$  subtypes. The prevalence of  $APOE \epsilon 4$  was significantly lower in the  $HpSp_{MRI}$  subtype, while  $MAPT$  H1/H1 haplotype prevalence was not different between subtypes. Finally, the  $HpSp_{MRI}$  subtype tended to have an earlier age at onset, but this difference was not significant. However, the percentage of individuals classified as having early-onset AD (EOAD; onset before 65 years of age) was different across the groups in the cross-sectional sample, with the  $HpSp_{MRI}$  group showing the highest percentage of individuals with EOAD (table 1).

**Cross-sectional analyses.** Baseline CSF measures of  $A\beta_{1-42}$ , total tau, and phosphorylated tau<sub>181</sub> were not different between groups (table 1). Furthermore,

no differences between subtypes were observed at baseline in the MMSE, ADAS-Cog<sub>13</sub>, CDR-SB, FAQ, or memory composite score (table 1 and figure 2C), but the  $HpSp_{MRI}$  subtype scored worse on the executive function composite (table 1 and figure 2D). The FDG HCI, an index of AD-like hypometabolism, was different between atrophy subtypes, with  $HpSp_{MRI}$  showing a greater hypometabolic pattern (figure 2B).

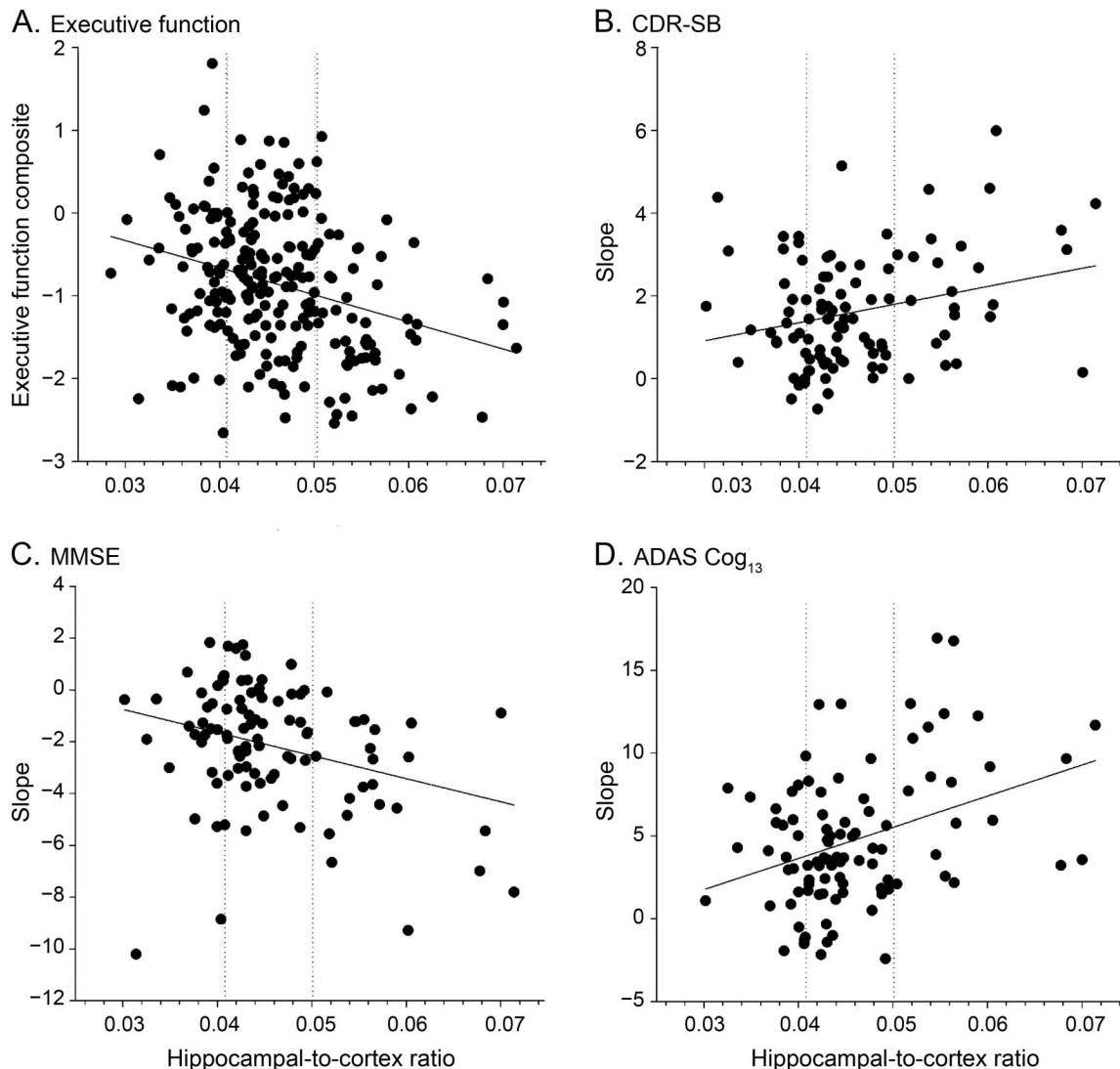
**Longitudinal analyses.** Mean follow-up time was not different across groups (table 1). Subtype category was associated with 2-year decline on the CDR-SB, FAQ, ADAS-Cog<sub>13</sub>, and MMSE (table 1 and figure 3) but not in the memory or executive function composite scores. In post hoc comparisons, the

**Figure 2** Relationship between baseline atrophy subtype and subsequent 2-year change in clinical and cognitive measures



Two-year changes in clinical and cognitive measures differed significantly between the atrophy subtypes (limbic predominant [ $LP_{MRI}$ ], typical AD [ $tAD_{MRI}$ ], and hippocampal sparing [ $HpSp_{MRI}$ ]). Participants with  $HpSp$  decline more quickly than those with the other subtypes, including a significantly greater rate of increasing (A) clinical dementia severity (Clinical Dementia Rating [CDR] Sum of Boxes,  $p = 0.013$ ) and (B) functional impairment (Functional Assessment Questionnaire [FAQ],  $p = 0.020$ ), as well as (C) faster cognitive decline on the Mini-Mental State Examination (MMSE) ( $p = 0.002$ ) and (D) Alzheimer's Disease Assessment Scale, 13-Item Subscale (ADAS-Cog<sub>13</sub>,  $p < 0.001$ ). a =  $\Delta HpSp > \Delta tAD$ ,  $p < 0.05$ ; b =  $\Delta HpSp > \Delta LP$ ,  $p < 0.05$ ; c =  $LP, tAD > HpSp$ ,  $p < 0.05$ ; d =  $\Delta tAD, \Delta LP > \Delta HpSp$ ,  $p < 0.05$ ; and e =  $HpSp > LP$ ,  $p < 0.01$ .

**Figure 3** Relationship between HV:CTV ratio and baseline and 2-year change in clinical and cognitive measures



Significant linear relationships between the hippocampal volume to cortical volume (HV:CTV) ratio and baseline executive function and 2-year change in clinical and cognitive measures were observed. (A) Specifically, baseline HV:CTV ratio was significantly associated with baseline executive function ( $r_p = -0.294, p < 0.001$ ). Baseline HV:CTV ratio was associated with (B) a faster 2-year increase in Clinical Dementia Rating Sum of Boxes (CDR-SB,  $r_p = 0.283, p = 0.003$ ) score, (C) a faster decline in Mini-Mental State Examination (MMSE,  $r_p = -0.303, p = 0.001$ ) score, and (D) a faster increase in Alzheimer's Disease Assessment Scale, 13-Item Subscale (ADAS-Cog<sub>13</sub>) score ( $r_p = 0.370, p < 0.001$ ). Dotted lines represent the atrophy subtype cutoffs for LP<sub>MRI</sub> vs tAD<sub>MRI</sub> (lower bound) and HpSp<sub>MRI</sub> vs tAD<sub>MRI</sub> (higher bound).

HpSp<sub>MRI</sub> group declined more rapidly than the LP<sub>MRI</sub> group on the FAQ, ADAS-Cog<sub>13</sub>, and MMSE and more rapidly than the tAD<sub>MRI</sub> group on the CDR-SB and MMSE ( $p < 0.05$ ).

We repeated all analyses using stage 1 definitions of the classification algorithm. The results were very similar to those with stage 2 classification (table e-1).

**Regression analyses.** Finally, we examined associations between HV:CTV and clinical phenotypes. At baseline, HV:CTV was not associated with any of the global scales (MMSE, ADAS-Cog<sub>13</sub>, CDR-SB, or FAQ) or the memory composite score but was associated with the executive function composite score, for which a higher HV:CTV ratio (reflecting

increased cortical relative to hippocampal atrophy) was associated with poorer executive function (figure 3A and table 2).

The HV:CTV ratio was also associated with 2-year change on the CDR-SB, MMSE, and ADAS-Cog<sub>13</sub>, with higher HV:CTV ratio associated with a faster rate of decline (figure 3, B–D, and table 2). Two-year change in the FAQ was not associated with either HV:CTV or any demographic variable, while 2-year change in the memory and executive composite scores was associated only with age and sex, respectively (table 2).

When HV and CTV were entered into the models as additional independent predictors, each was each independently associated with baseline CDR-SB,

**Table 2** Associations between continuous atrophy metrics and clinical and cognitive performance

	R <sup>2</sup>	Variable 1			Variable 2			Variable 3		
		Name	β	p Value	Name	β	p Value	Name	β	p Value
Including demographic variables and HV:CTV ratio only										
Baseline CDR-SB score	NA	None								
Baseline FAQ score	NA	None								
Baseline ADAS-Cog <sub>13</sub> score	NA	None								
Baseline MMSE score	NA	None								
Baseline memory score	NA	None								
Baseline executive function score	0.102	HV:CTV	-33.704	<0.001	Educ	0.035	0.047	NA		
Slope of Δ in CDR-SB score	0.080	HV:CTV	46.351	0.004	NA			NA		
Slope of Δ in FAQ score	NA	None								
Slope of Δ in ADAS-Cog <sub>13</sub> score	0.195	HV:CTV	164.998	0.002	Age	-0.124	0.013	NA		
Slope of Δ in MMSE score	0.092	HV:CTV	-89.170	0.002	NA			NA		
Slope of Δ in memory score	0.166	Age	0.011	<0.001	NA			NA		
Slope of Δ in executive function score	0.042	Sex	-0.121	0.044	NA			NA		
Including demographic variables, HV, CTV, and HV:CTV ratio										
Baseline CDR-SB score	0.069	HV	-3.9 × 10 <sup>-4</sup>	0.003	CTV	-1.7 × 10 <sup>-5</sup>	0.036	NA		
Baseline FAQ score	0.038	CTV	-1.0 × 10 <sup>-4</sup>	0.003	NA			NA		
Baseline ADAS-Cog <sub>13</sub> score	0.193	CTV	-2.8 × 10 <sup>-4</sup>	<0.001	Age	0.251	0.001	HV	-0.002	0.005
Baseline MMSE score	0.102	CTV	5.0 × 10 <sup>-5</sup>	<0.001	Age	-0.059	0.001	HV	-3.6 × 10 <sup>-4</sup>	0.029
Baseline memory score	0.122	CTV	2.0 × 10 <sup>-5</sup>	<0.001	HV:CTV	16.147	0.002	Age	-0.014	0.003
Baseline executive function score	0.296	CTV	3.9 × 10 <sup>-5</sup>	<0.001	Age	-0.025	<0.001	Educ	0.033	0.039
Slope of Δ in CDR-SB score	0.080	HV:CTV	46.351	0.004	NA			NA		
Slope of Δ in FAQ score	NA	None								
Slope of Δ in ADAS-Cog <sub>13</sub> score	0.244	CTV	-1.7 × 10 <sup>-4</sup>	<0.001	HV	0.001	0.041	NA		
Slope of Δ in MMSE score	0.181	CTV	8.6 × 10 <sup>-5</sup>	<0.001	NA			NA		
Slope of Δ in memory score	0.166	Age	0.011	<0.001	NA			NA		
Slope of Δ in executive function score	0.041	Sex	-0.121	0.044	NA			NA		

Abbreviations: ADAS-Cog<sub>13</sub> = Alzheimer's Disease Assessment Scale-13-Item Subscale; CDR-SB = Clinical Dementia Rating Scale-Sum of Boxes; CTV = cortical volume; Educ = education; FAQ = Functional Assessment Questionnaire; HV = hippocampal volume; MMSE = Mini-Mental State Examination; NA = not available.

ADAS-Cog<sub>13</sub>, and MMSE scores, with decreased volumes associated with an increased CDR-SB and ADAS-Cog<sub>13</sub> and decreased MMSE scores (table 2). CTV alone was independently associated with the baseline FAQ score and the executive function composite (and in this case HV:CTV was no longer significant), with decreased CTV associated with increased FAQ and reduced executive function score (table 2). Both CTV and the HV:CTV ratio were independently associated with the baseline memory composite, with a lower CTV and lower HV:CTV ratio associated with poorer memory (table 2). In the assessment of 2-year change and with HV and CTV included in the model, the HV:CTV ratio remained significantly independently associated with increasing clinical dementia severity (CDR-SB), with an increased HV:CTV ratio associated with a faster

increase in CDR-SB score (table 2). Two-year change in ADAS-Cog<sub>13</sub> score was independently associated with both CTV and HV (table 2), while change in MMSE score was associated only with CTV (table 2). Similar to the findings in the models including only HV:CTV, the slope of change in FAQ score was not associated with any atrophy or demographic variable and change in the memory and executive function composite scores was associated only with age and sex, respectively (table 2).

**DISCUSSION** We found that subtypes of AD consistent with those identified with postmortem NFT counts<sup>1</sup> could be identified in vivo from vMRI in cases with relatively mild dementia (mean MMSE [SD] score 23.2 [2.0], range 19–27). Specifically, an analog of the Murray-Dickson algorithm,



applied to hippocampal and cortical GMV to define HpSp<sub>MRI</sub>, tAD<sub>MRI</sub>, and LP<sub>MRI</sub> subtypes, yielded clinical phenotypes consistent with those reported in the autopsy study.<sup>1</sup> The HpSp<sub>MRI</sub> group was younger and declined more rapidly than both the tAD<sub>MRI</sub> and LP<sub>MRI</sub> groups on measures of global cognition despite comparable cognition at baseline. Moreover, the HpSp<sub>MRI</sub> subtype performed more poorly on a composite measure of executive function. When modeled as continuous variables, smaller CTV relative to HV was predictive of decreased baseline executive function and more 2-year clinical decline. When HV and CTV were modeled independently, CTV emerged as the main driver of the baseline performance and differential rates of decline across the cohort, although the ratio was independently predictive of 2-year change in dementia severity. Overall, given that the patterns of AD subtypes and associated clinical phenotypes were similar between those defined with atrophy measures from MRI and those defined with postmortem NFT counts, these findings suggest a localized association between the amount of tau pathology and the loss in GM consistent with a previous report.<sup>6</sup> Future studies with tau PET will help to further elucidate this relationship.

Unlike at baseline, subtype did not affect 2-year decline in the executive function composite score. However, this finding may be due to a floor effect. Major components of the executive function score are Trail Making Test (TMT) A and B, which have maximal scores for noncompletion (150 seconds for TMT A, 300 seconds for TMT B). Thus, if an individual could not complete the TMT at baseline or at follow-up, decline in executive function could not be captured.

The atrophy signature and cognitive profiles associated with the different subtypes identified in the present study are similar to those associated with sporadic EOAD. Specifically, increased cortical atrophy, especially in lateral and medial parietal areas, and a higher prevalence of atypical (dysexecutive, visuospatial) cognitive presentations have been reported in EOAD, in contrast to atrophy predominantly in the hippocampus and an amnesic cognitive profile in late-onset AD (LOAD).<sup>26–28</sup> Thus, the HpSp<sub>MRI</sub> subtype shows features similar to EOAD, whereas LOAD features are more similar to those of LP<sub>MRI</sub>. The fact that the subtype (or the continuous GMV) remained significantly associated with clinical presentation only when age was included in the statistical model and was a stronger predictor than age itself suggests that while an EOAD/LOAD age cut point provides a simple diagnostic rule, the clinical profile and trajectory are driven by the different underlying patterns of neurodegeneration, which may provide a more biologically driven basis for segregating

patients with AD into subtypes. Whereas a typical AD sequence of atrophy, similar to the stereotypical progression of tau pathology,<sup>29,30</sup> would show hippocampal atrophy preceding a more widespread decrease in cortical GMV, the HpSp<sub>MRI</sub> group appears to show the reversed sequence, with cortical atrophy preceding that of the hippocampus. The presence of distinct atrophy patterns in mild cognitive impairment and AD and the profiles of the subtypes identified in the present study are also consistent with data-driven cluster analyses, which identified differential brain atrophy patterns that were dominated either by medial temporal atrophy or by widespread cortical atrophy.<sup>3,31–33</sup>

The prevalence of the atypical Murray-Dickson subtypes found in the present study (14.4% HpSp<sub>MRI</sub>, 16.6% LP<sub>MRI</sub>) was comparable to that found in the original, substantially larger, autopsy study.<sup>1</sup> Subtypes defined solely after stage 1 of the Murray-Dickson algorithm (i.e., defined solely on the basis of the HV:CTV ratio) exhibited phenotypic relationships very similar to those obtained after stage 2. In particular, the HpSp<sub>MRI</sub> group (stage 1) progressed more rapidly and performed worse on executive relative to memory tasks. This finding is also consistent with the direct comparison of ante-mortem MRI to pathologic subtypes determined postmortem, in which the simple HV:GMV ratio (i.e., corresponding to step 1 of the algorithm) was found to significantly predict the postmortem neuropathologic tau subtype.<sup>6</sup>

To avoid selection bias, we calculated the subtype cutoffs from baseline data independently of whether the participants had follow-up data. If the subtypes were calculated just on the subset of participants who had 2-year follow-up data on all scales, the distribution of participants across the 3 subtypes was maintained and the findings were not substantially altered (only 4 participants showed different subtype categorization in stage 1 or 2). Thus, the cutoff values to determine subtype in the present study appear to be fairly consistent within the study population, supporting the presence of phenotypic differences within the AD cohort.

One drawback of the Murray-Dickson algorithm is that it requires subdividing a cohort of patients (when applied to vMRI) on the basis of the distribution of their hippocampal and cortical GMV and their ratio. Thus, this technique is not per se directly applicable prospectively to individual participants. However, the quartile and median values reported in the present study may provide suitable cut points for a decision tree to assign a subtype prospectively to new participants with mild AD with vMRI scans processed with the same processing pipeline and segmentation software. This hypothesis remains to be determined with replication in independent samples.

A few other limitations of the present study exist. Although AD pathology likely develops in preclinical and prodromal stages over many years, we focused only on patients with clinical AD in this study. Future studies in prodromal populations (mild cognitive impairment, particularly amnesic vs nonamnesic), as well as preclinical AD, are warranted. Furthermore, the ADNI study recruits from primarily academic medical institutions and may not be reflective of the broader AD community. In addition, the study has age (55–90 years only) and severity (mild AD or less) inclusion criteria and does not include atypical presentations of AD. However, the fact that we saw differences by atrophy subtype despite the relatively strict enrollment criteria suggests that these effects are robust and generalizable. Future studies in a broader AD population would help to better characterize these differential atrophy profiles.

AD subtypes based on brain atrophy defined with an algorithm originally derived from postmortem NFT counts identified participants with varying clinical profiles, genetic background, and differential rates of cognitive decline, consistent with those observed in the original autopsy study. In particular, patients with the HpSP<sub>MRI</sub> subtype, reflecting increased cortical rather than hippocampal atrophy, were generally younger, were less likely to be *APOE* ε4 positive, and had both a more dysexecutive cognitive profile and a more rapid rate of clinical decline. The rate of cognitive decline was driven primarily by cortical GMV loss. The ability to distinguish these subtypes and to determine neurodegenerative predictors of decline with in vivo imaging methods enables clinical trajectories to be predicted more accurately in living patients and points to the utility of considering atrophy patterns beyond the hippocampus in the assessment of patients with AD.

#### AUTHOR CONTRIBUTIONS

Dr. Shannon Risacher completed the design, conceptualization, and execution of the study, performed analysis and interpretation of the data, and was responsible for the drafting and revision of the manuscript. Mr. Wesley Anderson and Mr. Peter Castelluccio were involved in the analysis of the data contained in this manuscript and revision of the manuscript. Dr. Sergey Shcherbinin was involved in the interpretation of the data contained in this manuscript and revision of the manuscript. Dr. Andrew Saykin and Dr. Adam Schwarz were involved with the design, conceptualization, and execution of the study, interpretation of the data, and revision of the manuscript.

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