1	Sex differences in CSF biomarkers vary by Alzheimer's disease stage and APOE $\varepsilon$ 4
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45 Keywords: Alzheimer's Disease, CSF biomarkers, Tau, Aβ42, sex-differences, APOE, AD spectrum.

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#### 67 ABSTRACT

68 Objective: To evaluate sex-differences in cerebrospinal fluid (CSF) biomarkers, taking the potential
69 modifying role of clinical disease stage and APOEe4 genotype into account.

70 Method: We included participants (n=1801) with probable AD dementia (n=937), Mild Cognitive

71 Impairment (MCI; n=437) and Subjective Cognitive Decline (SCD; n=427). Main outcomes were CSF

amyloid  $\beta$ 1-42 (A $\beta$ 42), total tau (Tau) and tau phosphorylated at threonine 181 (pTau) levels. Age

73 corrected three-way interactions between sex, disease stage (i.e. syndrome diagnosis at baseline) and

APOEe4 were tested with linear regression analyses for each outcome measure. In case of significant

75 interactions (p<0.05), sex-differences were further evaluated by stratifying analyses for clinical

76 disease stage and APOEe4 genotype including age as a covariate. Covariates included age (model 1),

77 and additionally MMSE and educational level (model 2).

- **Results:** Three-way interactions were significant for Tau (p<0.001) and pTau (p<0.01), but not A $\beta$ 42.
- 79 In APOE carriers, women showed higher (p)Tau concentrations than men in SCD (Tau: $\beta \pm se=$

80  $0.25\pm0.08$ , p=0.002; pTau: $\beta\pm$ se= 0.16±0.06, p=0.009) and MCI (Tau: $\beta\pm$ se=0.29±0.07, p<0.001;

PTau: $\beta \pm se=0.21\pm0.06$ , *p*<0.001), but not AD dementia. In APOE non-carriers, women showed higher

82 (p)Tau concentrations in MCI (Tau: $\beta \pm se = 0.22 \pm 0.09$ , p=0.012; pTau: $\beta \pm se = 0.19 \pm 0.08$ , p=0.013) and

83 AD dementia (Tau:  $\beta \pm se = 0.20 \pm 0.07$ , p=0.006; pTau: $\beta \pm se = 0.14 \pm 0.06$ , p=0.014), but not in SCD.

84 **Conclusions:** Within APOEe4 carriers, sex-differences in CSF (p)Tau are more evident in early

disease stages, whereas for APOEe4 non carriers sex-differences are more evident in advanced disease

- stages. These findings suggest that the effect of APOE e4 on sex-differences in CSF biomarkers
- 87 depends on disease stage in AD.

### 88 Introduction

Alzheimer's Disease (AD) is a global health care challenge due to the rapidly growing disease 89 prevalence and the lack of preventive or curative treatment.<sup>1</sup> Therefore, knowledge regarding 90 the underlying pathophysiological process of AD that could potentially contribute to the 91 development of treatments is needed.<sup>2</sup> Current knowledge indicates that AD is characterized 92 by initial brain depositions of amyloid- $\beta$  (A $\beta$ ), followed by accumulation of neurofibrillary 93 tangles (NFT).<sup>3–5</sup> A growing body of literature is pointing towards sex-differences in AD 94 neuropathology, with women showing a higher NFT burden, while differences in A $\beta$  are less 95 apparent.<sup>6-10</sup> Additionally, there are indications that sex-differences in AD biomarkers are 96 modified by the presence of the Apolipoprotein (APOE) e4 allele, the major genetic risk 97 factor for sporadic AD.<sup>6,9</sup> In-vivo studies have shown that female APOEe4 allele carriers 98 have higher cerebrospinal fluid (CSF) total Tau and phosphorylated Tau (pTau) 99 concentrations than male APOEe4 carriers.<sup>6,9,10</sup> However, post-mortem studies do not show 100 this sex-specific association.<sup>6,7</sup> A possible explanation for these seemingly discrepant results 101 is that *in-vivo* studies mostly included pre-dementia subjects (i.e. normal cognition or Mild 102 Cognitive Impairment (MCI)), whereas post-mortem studies largely included end-stage 103 dementia patients. It could be hypothesized that the association between APOEe4 genotype 104 and sex-differences in AD neuropathology may be present in initial phases of the disease, but 105 diminishes as the disease progresses, and is no longer seen in end-stage AD dementia during 106 post-mortem examinations. Therefore, in the present study, we investigated whether sex-107 differences in CSF Aβ42, Tau and pTau are modified by APOEe4 genotype and clinical 108 109 disease stage.

## 111 Methods

112 Subjects

113	We selected 1801 subjects who visited our outpatient clinic between October 2000 and July
114	2015. Selection from the Amsterdam Dementia Cohort was based on a clinical diagnosis of
115	probable AD dementia ( $n=937$ ), MCI ( $n=437$ ) or Subjective Cognitive Decline (SCD; $n=427$ ,
116	and the availability of CSF biomarker results and APOE genotype. <sup>11,12</sup> There were no
117	exclusion criteria. All participants underwent a standardized dementia screening at baseline
118	that included physical and neurological examination, a neuropsychological test battery
119	including a Mini Mental State Examination (MMSE), Electroencephalogram (EEG),
120	Magnetic Resonance Imaging (MRI), and laboratory tests. Clinical diagnosis was given by
121	consensus in a multidisciplinary team according to international research and clinical criteria.
122	Subjects were labeled as having SCD when results of clinical examinations and test results
123	were normal (i.e. criteria for MCI or dementia were not fulfilled, and no psychiatric diagnosis
124	was given). MCI subjects were labeled according to the criteria by Petersen et al. and the
125	National Institute on Aging and Alzheimer's Association (NIA-AA) clinical criteria. <sup>13,14</sup> The
126	core clinical NIA-AA criteria were met for all probable AD patients. <sup>15,16</sup> Clinical diagnosis at
127	time of lumbar puncture, that is at baseline visit, was used to reflect clinical AD disease stage
128	(i.e. syndrome diagnosis). Sex was self-reported and defined as a biological characteristic that
129	discriminate women from men. <sup>17</sup> Educational levels were reported according to the Verhage
130	scoring system. <sup>18</sup>

131 Patient consents and Data availability statement

All subjects gave written informed consent and the study was approved by the local ethical
review board.<sup>11,12</sup> Anonymized data will be shared by request from any qualified investigator.

### 135 In-vivo markers of AD pathology

CSF AB42, Tau and pTau concentrations were used as *in-vivo* markers for the presence of AD 136 pathology. CSF samples were collected and processed according to international consensus 137 protocols as previously described.<sup>19,20</sup> Commercially available ELISAs were employed to 138 measure baseline A $\beta$ 42, Tau and pTau (Innotest  $\beta$ -amyloid(1-42), Innotest hTAU-Ag and 139 Innotest Phosphotau(181P); Fujirebio, Ghent, Belgium) concentrations. Intra- and inter-assay 140 variations for all analyses were below 3.2% and 10.9% respectively.<sup>21</sup> The team performing 141 the CSF analyses was not aware of the clinical diagnosis. To correct for the drift in CSF Aβ42 142 concentrations throughout the analysis-years we used adjusted A $\beta$ 42 concentrations.<sup>22,23</sup> Cut-143 offs to determine abnormality were <813 pg/ml for AB42<sup>23</sup> and >375 pg/ml for t-tau<sup>24</sup>. 144 MRI measurements 145 MRI measurements were acquired on 3T whole-body MR system (Discovery; GE Medical 146 Systems Milwaukee, WI, USA), using an eight-channel head coil at the Amsterdam UMC, 147 location VUmc. Medial temporal lobe atrophy (MTA) scores ranged from 0-4, and were rated 148 on coronal reconstructions of T1-weighted images.<sup>25</sup> Posterior cortical atrophy (PCA) and 149 global cortical atrophy (GCA) scores ranged from 0-3, and were rated on the combination of 150 T1-weighted and FLAIR sequences (PCA), or FLAIR sequences alone (GCA).<sup>26</sup> White matter 151 hyperintensities (WMH) were rated on FLAIR images using the Fazekas scale, with scores 152 ranging from 0-3.<sup>27</sup> The imaging took about 40 minutes in total. There was no intravenous 153 contrast administration. All scans were evaluated by an experienced neuroradiologist. 154 155

156 *Apolipoprotein E genotyping* 

- 157 DNA was isolated from 10 ml vacutainer tubes containing EDTA using the QIAamp DNA
- 158 blood isolation kit from Qiagen (Venlo, The Netherlands). Followed by genotype

159	determination	using the	LightCycler	ApoE mutation	Detection Kit	(Roche Di	agnostics,
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160 GmbH, Mannheim, Germany). Subjects with at least one APOEe4 allele were defined as

161 APOEe4 carriers, whereas no e4 allele defined subjects as non-carriers.

#### 162 *Statistical analysis*

Statistical analyses were completed using R studio (version: 3.3.2; "sincere Pumpkin Patch"). 163 Prior to performing statistical analyses. Tau and pTau were log transformed as they were not 164 normally distributed. Demographical and clinical data were compared between groups using 165 independent t-tests, chi-square tests and Mann-Whitney U tests as appropriate. To assess how 166 167 sex-differences in biomarkers depend on APOEe4 genotype and clinical disease stage, we used General Linear Models (GLM) with factors sex, APOEe4 genotype and clinical disease 168 stage, their 2-way interactions and 3-way interactions, and age was included as a covariate 169 170 (see table 3 for full models). In case of a significant interaction between sex, APOEe4 genotype and clinical disease stage, we performed GLM in CSF biomarker concentrations 171 stratified for APOE genotype and clinical disease stage as shown in figure 1. These analyses 172 included sex as a factor and age as covariate. and additionally MMSE and education (model 173  $\frac{2}{2}$ . We repeated analyses restricted to subjects with abnormal CSF AB42 concentrations (<813) 174 pg/ml) to study the specificity of the findings for the AD spectrum. Reported Effect Sizes 175 were calculated as the difference of the means of two groups divided by the weighted pooled 176 standard deviations of these groups according to Cohen's d statistics. We adjusted for multiple 177 testing by multiplying *p*-values with the number of tests (i.e. 12) according to the Bonferroni 178 method. In an additional set of sensitivity analyses, three-way interactions were repeated in 179 the total sample stratified by age (cut-off: median 67 years) to assess the effects of age. Three-180 181 way interactions included factors sex, clinical disease stage and APOEe4 genotype, their 2way interactions, and main factors. Separate models were run for CSF Tau and pTau. Finally, 182 we performed two additional analyses to test APOE genotype dose effects: 1) we repeated 183

184	analyses after excl	luding APOE e2e4 of	carriers (n=43), as the	e conveyed risk of APOE e2e4
	2	0		

185 carriers for AD is not fully known; 2) We evaluated the effect of e2 ((i.e. e2e2 = 4; e2e3 = 4)

186 111)) and e4 allele carriers (i.e. e3e4 =690; e4e4 = 298) against the e3 allele for sex effects on

187 CSF biomarkers. p < 0.05 was considered significant for main and interaction effects.

188

### 189 **Results**

- 190 Within SCD, the majority was male (61%), and females and males showed a similar
- 191 frequency of APOEe4 (F: 38% vs M: 37%, p > 0.05). Within MCI, the majority was male
- 192 (62%), females showed a higher frequency of APOEe4 than males (F: 65% vs M: 50%,
- 193 p=0.002), and females had less atrophy than males (table 1). Within AD dementia the
- 194 percentage of females and males was similar (52% vs 48%), as was the frequency of APOEe4
- 195 carriers between females and males (F: 67% vs. M: 68%, p>0.05). Females had less medial
- 196 temporal atrophy than males (F:1.32 vs M:1.52, p>0.01). Females and males did not differ in
- age within clinical disease stages. In MCI and AD dementia, males had higher educational
- 198 levels and higher MMSE scores compared to females. Subject characteristics of the CSF
- amyloid positive cohort was largely comparable to the total cohort (table 2).
- 200 For CSF Aβ42, age adjusted general linear models including sex, APOEe4 genotype,

201 diagnosis and all interactions showed main effects for diagnosis and APOEe4 genotype, but

202 not for sex. There were no significant interactions. For CSF Tau and pTau there was a three-

- 203 way interaction between sex, APOEe4 genotype, and clinical diagnosis (full models shown in
- table 3). Therefore, we further stratified these analyses for APOEe4 and clinical diagnosis asshown in figure 1.
- 206 Within APOEe4 carriers, females showed higher Tau and pTau concentrations than males in
- 207 SCD (Cohens *d* (95%CI): Tau= 0.52 (0.19-0.84),  $p_{adj}$ = .008; pTau=0.44 (0.11-0.77)  $p_{adj}$ =.05)

- and MCI (Cohens *d* (95%CI): Tau= 0.54 (0.28-0.80),  $p_{adj}$ =.0006; pTau=0.52 (0.26-0.77)
- 209  $p_{adj}=.001$ , but not in AD dementia (figure 1; table 2). Within APOEe4 non-carriers, females
- showed higher Tau and pTau concentrations than males in MCI (Cohens *d* (95%CI): Tau=
- 211 0.49 (0.17-0.80),  $p_{adj}=.02$ ; pTau=0.47 (0.16-0.78)  $p_{adj}=.04$ ) and AD dementia (Cohens d
- 212 (95%CI): Tau= 0.42 (0.19-0.65),  $p_{adj}$ =.006; pTau=0.38 (0.15-0.61)  $p_{adj}$ =.02), but not in SCD
- 213 (figure 1; table 2). When restricting analyses to individuals with abnormal CSF A $\beta$ 42, results
- for Tau and pTau were largely comparable with that of the total cohort, albeit significance
- 215 was overall somewhat attenuated for Tau and pTau, and was lost for Tau in APOE e4 carriers
- in the SCD stage and for pTau in non-carriers in the MCI stage due to a smaller effect size
- 217 (table 4).
- 218 In an additional analysis, we stratified for age and found a significant three way interaction
- 219 between sex, APOE genotype and diagnosis for CSF (p)Tau in older individuals (median
- [IQR]: 72.2 [69.5-76.0]), but not in younger individuals (median [IQR]: 61.3 [58.4-64.1])
- (full models shown in table 5). When we repeated the analyses in the sample excluding APOE
- e2e4 carriers (n=43), results remained essentially unchanged (full models shown in table 6).
- Finally, we studied dose effects for APOE genotypes and found that APOEe2 carriers
- behaved similar to APOEe3 carriers for all clinical disease stages (supplementary table 1).

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## 225 **Discussion**

226	Our findings suggest that APOE differentially affects sex-differences in CSF biomarkers
227	throughout the AD spectrum. Within APOEe4 carriers, females show higher Tau and pTau
228	concentrations in early disease stages (i.e. SCD and MCI) which equalized in the later
229	dementia stage. Within APOE non-carriers, we observed an opposite pattern, with females
230	showing higher Tau and pTau concentrations in later disease stages (i.e. AD dementia and
231	MCI ), but not in the early disease stage of SCD. We did not find sex-differences in $A\beta$
232	concentrations between females and males for any disease stage or APOEe4 genotype.
233	Although derived from cross-sectional data our findings suggest that within APOEe4 carriers
234	sex-differences in Tau and pTau become less evident in advanced disease stages, whereas for
235	APOEe4 non-carriers sex-differences in Tau and pTau become more evident in advanced
236	disease stages.
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249	It has been suggested that the sex-difference in APOEe4 carriers diminishes with increasing
250	age, as previous studies only found a sex-specific interaction with APOE e4 in younger
251	individuals (<75 years). <sup>6,32</sup> Seemingly in contrast to these previous studies, stratification for
252	age in our study revealed that results were largely attributable to older elderly individuals
253	(>67 years). However, closer inspection of the data in fact shows that our 'older' participants
254	fall within the same age range as the 'younger' participants in former studies (i.e. 65-75
255	years). <sup>6,32</sup> Another explanation for these age dependent sex-differences could be that younger
256	and older participants might have different underlying AD pathological mechanisms. For
257	instance, younger participants with a similar cognitive status as older individuals, may not
258	reflect an earlier phase of AD than older participants and vice versa. Therefore, it could be
259	possible that younger individuals might have more (unknown) genetic risk factors for AD,
260	which in turn influence Tau accumulation and sex-differences in Tau concentrations. <sup>33</sup> Taken
261	together, our results support the idea that the sex by APOEe4 interaction depends on age, and
262	we further show that this interaction effect depends also on clinical disease stage as well.
263	
264	In the latest framework for AD, Tau is considered a marker for neuronal injury <sup>5</sup> . As such,
265	higher Tau and pTau concentrations in female APOEe4 carriers may imply an initial steeper
266	pathological disease course, and more neurodegenerative change compared to male APOEe4
267	carriers. Increased neurodegenerative changes in female APOEe4 carriers was also implied by
268	
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269 270 271	previous studies who reported increased hypometabolism and (hippocampal) atrophy <sup>34</sup> , and a decreased hippocampal connectivity. <sup>35</sup> However, other studies have shown discordant findings. Previous population studies reported lower hippocampal volume in males compared to females <sup>29</sup> , and more rapid parahippocampal atrophy in amyloid positive males compared
269 270 271 272	previous studies who reported increased hypometabolism and (hippocampal) atrophy <sup>34</sup> , and a decreased hippocampal connectivity. <sup>35</sup> However, other studies have shown discordant findings. Previous population studies reported lower hippocampal volume in males compared to females <sup>29</sup> , and more rapid parahippocampal atrophy in amyloid positive males compared to amyloid negative females. <sup>36</sup> In the current study, we observed more atrophy in males than

274 levels we found in females. This suggests that discrepant findings between CSF Tau and

atrophy may reflect different pathological processes, and should perhaps not be used

276 interchangeably. Similar discrepant findings between tau and MRI have been reported

277 previously, which may depend on sex as well.<sup>37</sup> Future research combining *in-vivo* CSF

278 biomarker and MRI data with pathology data is needed to examine the relationship with each

279 other and neuropathology.

280

Possible explanatory biological mechanisms for higher levels of Tau and pTau in females 281 might be related to abrupt hormonal changes that have occurred in post-menopausal women.<sup>17</sup> 282 A drastic drop of estradiol levels in post-menopausal women, has shown to lead to an 283 increased activity of enzymes involved in Tau phosphorylation (GSK3-β and Protein Kinase 284 A), thus resulting in a higher pTau concentrations.<sup>38,39</sup> In addition, post-menopausal women 285 286 show indications of increased oxidative stress and mitochondrial dysfunction, which through cell death and apoptosis mechanisms may lead to increases in CSF Tau concentrations.<sup>5,39</sup> 287 APOE e4 has also been associated with higher levels of CSF Tau and NFT's,<sup>2</sup> and it was 288 shown to stimulate Tau phosphorylation as a result of impaired cholesterol exchange between 289 neuronal and non-neuronal cells.<sup>6,31,41,42</sup> Therefore, it is conceivable that the lack of 290 neuroprotective effects of estrogen together with the presence of APOE e4 might act 291 synergistically, leading to increased Tau concentrations in female APOE e4 carriers. Further 292 supporting this hypothesis is the observation that post-menopausal female APOE e4 carriers 293 on estrogen replacement therapy show more signs of neuroprotection compared to non-treated 294 female carriers. Another factor that may influence tau levels is cerebrovascular injury. <sup>43–45</sup> 295 Therefore, it could be that the sex-difference in CSF Tau concentrations is caused by a 296 difference in the amount cerebrovascular injury between both sexes. However, we did not 297 observe a difference between sexes in Fazekas scores, which are considered a marker of 298

299	cerebrovascular injury .46	Further (fundamental) research is needed to discover the true
300	underlying cause of the se	ex-differences seen in AD.

302	Other possible explanations for higher Tau and pTau concentrations in female APOEe4
303	carriers in the earlier stages of the disease spectrum, could be a difference in survival between
304	females and males. A faster disease progression in males or a higher mortality rate of the
305	"very sick" men dying from comorbidities at young ages could possibly cause an
306	overrepresentation of "healthier" males with lower levels of neuropathology (i.e. CSF Tau
307	concentrations) therefore making it seem as though women have higher Tau concentrations.
308	However, in our cohort this seems less likely since men had more atrophy in general.
309	
310	In our MMSE and education adjusted analyses, we found that females with similar MMSE
311	scores and educational level to men, had higher Tau and pTau levels. Moreover, despite
312	having higher Tau levels, and, possibly more neurodegeneration, females were given the same
313	clinical diagnosis as men. These findings seemingly imply that the females in our cohort have
314	more cognitive reserve than males. Cognitive reserve has been defined as the difference
315	between individuals in their ability to preserve cognitive function in the presence of
316	neuropathology. <sup>47</sup> Previous ADNI studies have shown that the female advantage in verbal
317	memory is maintained despite similar levels of temporal hypometabolism and moderate
318	hippocampal atrophy between females and males. <sup>48,49</sup> This would imply that women at first
319	better compensate for neuropathology and maintain cognitive function, thus have a greater
320	<del>cognitive reserve.</del>

- 321 In our study population, females and males showed similar A $\beta$ 42 concentrations within both
- 322 APOEe4 genotypes throughout clinical stages of AD. Our findings align with previous work,
- which suggest that sex-differences in AD pathology mainly occur in Tau and pTau

- 324 concentrations, downstream from amyloid deposition.<sup>6,7,9,10,29–31,50</sup> Interestingly, within the
- 325 current NIA-AA criteria<sup>5</sup> CSF total tau and pTau are seen as markers for separate pathological
- 326 mechanisms, where one represents neuronal injury (Tau) and the other is a specific marker for
- 327 Alzheimer's disease pathology (pTau). However, in our cohort Tau and pTau are highly
- 328 correlated (r= 0.93, p<0.001), and sex-differences in both APOEe4 carriers and non-carriers
- 329 were similar for Tau and pTau. This suggests that Tau and pTau to some extent reflect similar
- 330 or overlapping aspects of neuronal injury.

Among the limitations of our study is that to adequately assess the temporal order of AD

pathology markers, longitudinal data are needed. Moreover, our data are derived from a

- tertiary memory clinic cohort which consist of a relatively young population, and may result
- in a lack of generalizability of our findings. Among the strengths of this study is the use of a
- large well-defined cohort, and the careful clinical work-up that was used to diagnose all
- participants. Moreover, as our findings were largely replicable in an A $\beta$ 42 positive subset, we
- 337 were able to show that our findings were specific for the AD spectrum. Our data show that a
- 338 woman's brain can be more susceptible to Tau pathology depending on disease stage and
- 339 APOEe4 genotype. The effect sizes we found for (p)Tau concentrations between women and
- 340 men were moderate, and therefore not large enough for clinical use, for instance by
- 341 developing sex-specific cut-offs for Tau or pTau. However, a moderate difference in the
- 342 underlying pathology of AD between women and men is large enough to be taken into
- 343 consideration when developing disease modifying therapies.
- 344

In conclusion, within APOEe4 carriers sex-differences in Tau and pTau become less evident
in advanced disease stages, whereas in APOEe4 non-carriers sex-differences in Tau and pTau
become more evident in advanced disease stages. These findings largely remain for the
amyloid positive subgroup. Our findings imply a difference in neuropathological trajectories

- for women and men depending on APOEe4 genotype, and add to a growing body of evidence
- 350 of sex-differences in the underlying mechanism of AD.

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Tables

Table 1. Subj	ect characteristics for the total cohort, st	tratified per APOE e	e4 genotype and clinic	cal disease stage.			
		Total cohort		APOE e4 carrier		APOE e4 non-carrier	
		females males		females males		females males	
SCD	n, (%)	167 (39)	260 (61)	63 (40)	95 (60)	104 (39)	165 (61)
	Age, mean (SD), y	64.9 (6.5)	64.4 (6.2)	64.6 (6.2)	64.6 (6.4)	64.8 (6.7)	64.0 (6.1)
	MMSE, mean (SD)	28 (1.6)	28 (1.8)	28 (1.8)	28 (1.7)	28 (1.5)	28 (1.7)
	Education, mean (SD) <sup>a</sup>	5.3 (1.3)	5.4 (1.3)	5.4 (1.3)	5.5 (1.2)	5.3 (1.3)	5.5 (1.3)
	CSF Aβ42, pg/ml, mean (SD)	1032 (252)	1070 (263)	908 (245)	964 (265)	1114 (224)	1131 (232)
	Abnormal CSF Aβ42, n (% within sex) <sup>b</sup>	39 (23)	52 (19)	26 (41)	29 (31)	12 (12)	17 (11)
	CSF LN Tau, pg/ml, mean (SD)	5.7 (0.5)	5.6 (0.5)	5.9 (0.5)	5.6 (0.5)**	5.5 (0.5)	5.6 (0.5)
	Abnormal CSF Tau, n (% within sex) <sup>b</sup>	42 (24)	60 (22)	25 (40)	18 (19)*	15 (15)	33 (20)
	Fazekas, mean (SD) <sup>c</sup>	0.69 (0.76)	0.74 (0.74)	0.56 (0.82)	0.75 (0.82)	0.75 (0.74)	0.75 (0.75)
	MTA, mean (SD) <sup>c</sup>	0.35 (0.51)	0.38 (0.50)	0.33 (0.59)	0.45 (0.59)	0.36 (0.47)	0.34 (0.48)

	PCA, mean (SD) <sup>c</sup>	0.54 (0.70)	0.62 (0.68)	0.49 (0.79)	0.60 (0.76)	0.55 (0.67)	0.65 (0.68)
	GCA, mean (SD) <sup>c</sup>	0.38 (0.60)	0.47 (0.59)	0.32 (0.63)	0.49 (0.63)	0.41 (0.59)	0.49 (0.60)
MCI	n, (%)	168 (38)	269 (62)	109 (45)	134 (55)	59 (30)	135 (70)
	Age, mean (SD), y	68.5 (6.9)	68.3 (7.1)	68.1 (6.4)	67.7 (6.5)	68.9 (7.6)	68.1 (7.2)
	MMSE, mean (SD)	26 (2.4)	27 (2.4)*	26 (2.5)	27 (2.6)	26 (2.6)	27 (2.2)*
	Education, mean (SD) <sup>a</sup>	4.8 ( 1.3)	5.2 (1.4)*	4.9 (1.3)	5.3 (1.4)*	4.7 (1.4)	5.2 (1.4)*
	CSF Aβ42, pg/ml, mean (SD)	774 (247)	867 (290)***	688 (156)	757 (249)*	925 (311)	977 (283)
	Abnormal CSF Aβ42, n (% within sex) <sup>b</sup>	127 (72)	163 (53)***	89 (84)	94 (71)*	28 (48)	46 (34)
	CSF LN Tau, pg/ml, mean (SD)	6.2 (0.6)	5.9 (0.6)***	6.3 (0.5)	6.0 (0.6)***	6.0 (0.6)	5.7 (0.5)**
	Abnormal CSF Tau, n (% within sex) <sup>b</sup>	124 (70)	142 (46)***	82 (77)	79 (59)*	33 (57)	41 (30)**
	Fazekas, mean (SD) <sup>c</sup>	1.16 (0.98)	1.01 (0.93)	1.18 (0.99)	0.88 (0.93)*	1.12 (1.03)	1.15 (1.00)
	MTA, mean (SD) <sup>c</sup>	0.64 (0.85)	1.03 (0.81)***	0.62 (0.82)	0.94 (0.77)**	0.67 (0.95)	1.14 (0.94)**
	PCA, mean (SD) <sup>c</sup>	0.71 (0.74)	0.93 (0.70)**	0.73 (0.76)	0.92 (0.73)	0.68 (0.77)	0.93 (0.76)*
	GCA, mean (SD) <sup>c</sup>	0.68 (0.68)	0.90 (0.65)***	0.66 (0.68)	0.89 (0.64)**	0.72 (0.76)	0.93 (0.75)

AD dementia	n, (%)	488 (52)	449 (48)	325 (51)	307 (49)	163 (53)	142 (47)
	Age, mean (SD), y	67.4 (7.2)	67.3 (7.2)	66.7 (6.7)	67.5 (6.8)	67.7 (7.5)	67.0 (7.5)
	MMSE, mean (SD)	20 (4.8)	21 (5.1)***	20 (4.8)	21 (8.1)*	19 (5.0)	22 (4.7)***
	Education, mean (SD) <sup>a</sup>	4.6 (1.3)	5.0 (1.4)***	4.7 (1.2)	5.0 (1.4)**	4.5 (1.3)	5.1 (1.4)***
	CSF Aβ42, pg/ml, mean (SD)	663 (173)	649 (164)	638 (132)	626 (129)	709 (221)	705 (214)
	Abnormal CSF Aβ42, n (% within sex) <sup>b</sup>	460 (90)	447 (91)	296 (82)	286 (82)	130 (82)	112 (82)
	CSF LN Tau, pg/ml, mean (SD)	6.5 (0.5)	6.4 (0.5)***	6.5 (0.5)	6.4 (0.5)	6.5 (0.5)	6.2 (0.6)***
	Abnormal CSF Tau, n (% within sex) <sup>b</sup>	437 (85)	390 (79)*	268 (85)	248 (82)	135 (85)	103 (75)*
	Fazekas, mean (SD) <sup>c</sup>	1.0 (0.88)	0.95 (0.85)	1.04 (0.90)	0.94 (0.91)	0.95 (0.87)	0.95 (0.82)
	MTA, mean (SD) <sup>c</sup>	1.32 (0.90)	1.51 (0.88)**	1.32 (0.93)	1.53 (0.93)**	1.28 (0.90)	1.45 (0.85)
	PCA, mean (SD) <sup>c</sup>	1.26 (0.83)	1.33 (0.81)	1.17 (0.83)	1.32 (0.84)*	1.45 (0.87)	1.37 (0.83)
	GCA, mean (SD) <sup>c</sup>	1.11 (0.73)	1.20 (0.71)	1.03 (0.74)	1.21 (0.74)**	1.27 (0.78)	1.14 (0.72)

Table shows mean (SD), unless otherwise specified. Independent t-test. chi-square test and Mann-Whitney U test were applied where applicable. P < 0.05 is considered significant: \*

p<0.05, \*\* < 0.01, \*\*\* p<0.001. <sup>a</sup> Education according to Verhage score. <sup>b</sup> Cut-off for Aβ42: 813 pg/ml, and Tau: 375 pg/ml. <sup>c</sup> adjusted for age. Abbreviations: APOE, Apolipoprotein E4; SCD, Subjective Cognitive Decline; MCI, Mild Cognitive Impairment; AD, Alzheimer's Disease; MMSE, Mini Mental State Exam; CSF, Cerebrospinal Fluid; Aβ42, Amyloidβ 1-42; Tau, total tau, MTA, Medial-Temporal Atrophy; PCA, Posterior Cortical Atrophy; GCA, Global Cortical Atrophy. Table 2. Subject characteristics CSF Aβ42 positive subgroup stratified per APOE e4 genotype and clinical disease stage. APOE e4 carrier APOE e4 non-carrier Females Males Females Males SCD n, (%) 26 (47) 29 (53) 12 (41) 17 (59) Age, mean (SD), y 66.7 (5.7) 66.8 (6.1) 67 (8.1) 70 (7.5) MMSE, mean (SD) 28 (1.5) 28 (2.0) 29 (1.1) 29(0.8) Education, mean (SD)<sup>a</sup> 5.5 (1.1) 5.1 (1.3) 4.8 (1.4) 5.6 (1.4) CSF A $\beta$ 42, pg/ml, mean (SD) 688 (82) 657 (113) 711 (85) 687 (114) CSF LN Tau, pg/ml, mean (SD) 6.1 (0.5) 5.9 (0.6) 5.8 (0.8) 5.8 (0.7) CSF Tau positive, n (% within sex)<sup>b</sup> 14 (58) 10 (42) 5 (36) 9 (64) MCI n, (%) 89 (49) 94 (51) 28 (38) 46 (62) 68.5 (6.7) 70.8 (7.7) 69.5 (7.2) Age, mean (SD), y 68.7 (6.0) MMSE, mean (SD) 26 (2.5) 26 (2.5) 26 (2.9) 27 (2.5) Education, mean (SD)<sup>a</sup> 4.9 (1.4) 5.4 (1.4)\* 4.9 (1.3) 5.2 (1.6) CSF A $\beta$ 42, pg/ml, mean (SD) 638 (98) 629 (106) 686 (87) 659 (110) CSF LN Tau, pg/ml, mean (SD) 6.4 (0.4) 6.2 (0.5)\*\* 6.3 (0.6) 6.0 (0.6)\* CSF Tau positive, n (% within sex)<sup>b</sup> 74 (51) 72 (49) 22 (48) 24 (52) AD 296 (51) 112 (46) n, (%) 286 (49) 130 (54) 66.6 (6.7) 67.3 (6.7) 66.9 (7.6) 66.7 (7.3) Age, mean (SD), Y MMSE, mean (SD) 20 (4.9) 21 (5.0) 19 (5.0) 21 (4.6) \*\*\*

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	Education, mean (SD) <sup>a</sup>	4.8 (1.2)	5.0 (1.4)**	4.6 (1.3)	5.2 (1.4)**			
	CSF Aβ42, pg/ml, mean (SD)	618 (98)	609 (99)	624 (96)	620 (100)			
	CSF LN Tau, pg/ml, mean (SD)	6.5 (0.5)	6.4 (0.5)	6.6 (0.5)	6.3 (0.6)			
	CSF Tau positive, n (% within sex) <sup>b</sup>	256 (52)	238 (48)	122 (58)	90 (42)			
Subject char	acteristics for the amyloid positive subg	group shown per c	clinical disease sta	ge and APOE e4	genotype.			
Independent	t-test, chi-square test and Mann Whitne	ey U test were app	plied were applica	ble. <i>P</i> <0.05 is co	nsidered			
significant: * p<0.05, ** p<0.01, *** p<0.001. <sup>a</sup> Education according to Verhage score. <sup>b</sup> Cut-off Tau: 375 pg/ml.								
Abbreviations: APOE, Apolipoprotein E4; SCD, Subjective Cognitive Decline; MCI, Mild Cognitive Impairment; AD,								
Alzheimer's	Disease; MMSE, Mini Mental State Ex	kam; IQR, Interqu	artile Range.					

Table 3. Full models for sex, APOE e4 genotype and clinical disease stage per CSF biomarker.

	aβ42		Tau	d	pTau <sup>d</sup>	
	$\beta$ (se)	<i>p</i> -value	$\beta$ (se)	<i>p</i> -value	$\beta$ (se)	<i>p</i> -value
Sex: male <sup>a</sup>	14.5 (26.3)	0.58	0.04 (0.07)	0.58	0.02 (0.05)	0.68
Diagnosis: MCI <sup>b</sup>	-180.9(34.2)	<0.001	0.45 (0.09)	< 0.001	0.27 (0.07)	< 0.001
Diagnosis: AD <sup>b</sup>	-398.9(26.5)	< 0.001	0.91 (0.07)	< 0.001	0.60 (0.05)	<0.001
APOE: carrier <sup>c</sup>	-206.7(33.2)	<0.001	0.34 (0.09)	<0.001	0.24 (0.07)	<0.001
Sex: male* diagnosis: MCI <sup>ab</sup>	35.5 (41.8)	0.40	-0.32 (0.11)	0.004	-0.25 (0.08)	0.004
Sex: male* diagnosis: AD <sup>ab</sup>	-18.2 (35.8)	0.61	-0.27 (0.09)	0.003	-0.18 (0.07)	0.01
Sex: male* APOE: carrier <sup>ac</sup>	41.3 (42.8)	0.33	-0.31 (0.11)	0.006	-0.20 (0.09)	0.03
Diagnosis: MCI* APOE: carrier <sup>bc</sup>	-31.5 (47.4)	0.51	0.009 (0.12)	0.98	0.05 (0.10)	0.62
Diagnosis: AD* APOE: carrier <sup>bc</sup>	133.4 (38.9)	<0.001	-0.33 (0.10)	< 0.001	-0.22 (0.08)	0.004
Sex: male*diagnosis: MCI* APOE: carrier <sup>abc</sup>	-24.0 (60.2)	0.69	0.28 (0.15)	0.07	0.21 (0.12)	0.09
Sex: male*diagnosis: AD* APOE: carrier <sup>abc</sup>	-49.0 (52.0)	0.35	0.46 (0.13)	<0.001	0.33 (0.11)	0.002
We used Constal Linear Models (CLM) with fo	ators say ADOE	al construng	and aliniaal disaas	a stage their	2 way interaction	a and 2 may

We used General Linear Models (GLM) with factors sex, APOE e4 genotype and clinical disease stage, their 2-way interactions and 3-way interactions, and age was included as a covariate. This full model was run separately for each of the three biomarkers (i.e.  $a\beta42$ , Tau and pTau). <sup>a</sup> Reference is female sex. <sup>b</sup> Reference is SCD. <sup>c</sup> Reference is APOE non-carrier. p <0.05 is considered significant. <sup>d</sup> The natural logarithm of Tau and pTau concentrations are shown. Abbreviations: APOE, Apolipoprotein E4; SCD, Subjective Cognitive Decline; MCI, Mild Cognitive Impairment; AD, Alzheimer's Disease.

Table 4. Difference	Table 4. Differences in CSF Tau and pTau concentrations in men compared to women in the amyloid positive subgroup stratified per clinical diagnosis and APOE e4 genotype.									
			Tau			pTau				
		Adjusted mean difference $\beta(se)^a$	Effect size (95%CI)	<i>p</i> -value	Adjusted <i>p</i> -value	Adjusted mean difference $\beta$ (se) <sup>a</sup>	Effect size (95%CI)	<i>p</i> -value	Adjusted <i>p</i> -value	
APOE e4 carrier	SCD	-0.26 (0.14)	0.44 (-0.11-0.99)	0.07	0.84	-0.15 (0.11)	0.33 (-0.22–0.87)	0.18	2.16	
	MCI	-0.20 (0.07)	0.42 (0.12-0.71)	0.006	0.07	-0.14 (0.06)	0.37 (0.08– 0.67)	0.01	0.12	
	AD dementia	-0.07 (0.04)	0.13 (-0.03–0.29)	0.13	1.56	-0.03 (0.04)	0.06 (-0.10–0.23)	0.45	5.4	
APOE e4 non-carrier	SCD	-0.15 (0.27)	0.02 (-0.76–0.79)	0.59	7.08	-0.01 (0.22)	-0.13(-0.90-0.65)	0.95	11.4	
	MCI	-0.30 (0.15)	0.52 (0.04 - 1.0)	0.04	0.48	-0.24 (0.13)	0.46 (-0.03-0.94)	0.08	0.96	
	AD dementia	-0.26 (0.07)	0.24 (0.24–0.75)	<0.001	0.004	-0.18 (0.06)	0.44 (0.18-0.69)	0.002	0.02	

Differences in CSF Tau and pTau for women and men in the amyloid positive subgroup stratified per APOE e4 genotype and clinical diagnosis. Tau and pTau were log transformed to meet assumptions of normality. We performed GLM in CSF biomarker concentrations stratified for APOE genotype and clinical disease stage as shown in figure 1. These analyses included sex as a factor and age (model 1), and additionally MMSE and education (model 2) as covariate. Cohen's d statistics were used to calculate effect sizes: small=0.2, medium= 0.5, large= 0.8. *p* <0.05 is considered significant. <sup>a</sup> Reference is female sex. <sup>b</sup> Adjustments for age at time of lumbar puncture. Abbreviations: APOE, Apolipoprotein E4; SCD, Subjective Cognitive Decline; MCI, Mild Cognitive Impairment; AD, Alzheimer's Disease; MMSE, Mini Mental State Exam; 95% CI, 95% Confidence Interval.

Table 5. Full models for sex, APOE e4 genotype and clinical disease stage for (p)Tau stratified per age group.									
		<67	years		≥67 years				
	Ta	u <sup>d</sup>	pTa	u <sup>d</sup>	Ta	u <sup>d</sup>	рТа	u <sup>d</sup>	
	$\beta$ (se)	<i>p</i> -value							
Sex: male <sup>a</sup>	0.05 (0.08)	0.57	0.01 (0.07)	0.84	0.04 (0.11)	0.73	0.04 (0.09)	0.66	
Diagnosis: MCI <sup>b</sup>	0.40 (0.13)	0.002	0.30 (0.10)	0.005	0.37 (0.12)	0.003	0.24 (0.10)	0.02	
Diagnosis: AD <sup>b</sup>	1.05 (0.09)	<0.001	0.71 (0.07)	<0.001	0.74 (0.11)	<0.001	0.49 (0.09)	<0.001	
APOE: carrier <sup>c</sup>	0.20 (0.10)	0.05	0.13 (0.09)	0.12	0.57 (0.14)	<0.001	0.40 (0.11)	<0.001	
Sex: male* diagnosis: MCI <sup>a,b</sup>	-0.35 (0.16)	0.03	-0.32 (0.13)	0.01	-0.28 (0.16)	0.08	-0.22 (0.13)	0.09	
Sex: male* diagnosis: AD <sup>a,b</sup>	-0.29 (0.12)	0.02	-0.19 (0.09)	0.04	-0.26 (0.15)	0.07	-0.19 (0.12)	0.10	
Sex: male* APOE: carrier <sup>a,c</sup>	-0.20 (0.13)	0.13	-0.11 (0.11)	0.30	-0.39 (0.19)	0.04	-0.29 (0.15)	0.06	
Diagnosis: MCI* APOE: carrier <sup>b,c</sup>	0.28 (0.17)	0.10	0.21 (0.14)	0.13	-0.29 (0.18)	0.10	-0.14 (0.15)	0.33	

Diagnosis: AD* APOE: carrier <sup>b,c</sup>	-0.14 (0.13)	0.26	-0.12 (0.10)	0.25	-0.62 (0.16)	<0.001	-0.40 (0.13)	0.002
Sex: male*diagnosis: MCI* APOE: carrier <sup>a,b,c</sup>	0.03 (0.21)	0.88	0.08 (0.17)	0.64	0.46 (0.24)	0.05	0.34 (0.19)	0.08
Sex: male*diagnosis: AD* APOE: carrier <sup>a,b,c</sup>	0.22 (0.17)	0.18	0.17 (0.14)	0.20	0.69 (0.22)	0.001	0.50 (0.18)	0.004
Full models: We used General Linear Models (GLM)	with factors sex, AP	OE e4 genotype	e and clinical disease	e stage, their 2-	way interactions an	d 3-way interact	ions, and age was i	ncluded as a
covariate. This full model was run separately for each	of the three biomark	ters. <sup>a</sup> Referenc	e is female sex. <sup>b</sup> Re	ference is SCD	<sup>c</sup> Reference is APC	DE non-carrier. J	o <0.05 is considere	d significant. d
The natural logarithm of Tau and pTau concentrations	are shown. Abbrevi	ations: APOE,	Apolipoprotein E4;	SCD, Subjectiv	e Cognitive Decline	e; MCI, Mild Co	ognitive Impairment	t; AD,
Alzheimer's Disease.								

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	Total cohort excluding APOE e2e4 carriers				Cohort containing only APOE e2 allele carriers			
	Tai	1 <sup>d</sup>	pTau <sup>d</sup>		Tau <sup>d</sup>		pTau <sup>d</sup>	
	$\beta$ (se)	<i>p</i> -value	$\beta$ (se)	<i>p</i> -value	$\beta$ (se)	<i>p</i> -value	$\beta$ (se)	<i>p</i> -value
Sex: male <sup>a</sup>	0.04 (0.07)	0.58	0.02 (0.05)	0.76	0.09 (0.16)	0.59	0.09 (0.14)	0.54
Diagnosis: MCI <sup>b</sup>	0.45 (0.09)	< 0.001	0.31 (0.07)	< 0.001	0.37 (0.20)	0.06	0.27 (0.17)	0.11
Diagnosis: AD <sup>b</sup>	0.94 (0.09)	< 0.001	0.63 (0.05)	< 0.001	0.75 (0.18)	< 0.001	0.44 (0.15)	0.004
APOE: carrier <sup>c</sup>	0.39 (0.07)	< 0.001	0.27 (0.07)	<0.001	NA	NA	NA	NA
Sex: male* diagnosis: MCI <sup>a,b</sup>	-0.31 (0.11)	0.004	-0.24 (0.09)	0.004	-0.44 (0.25)	0.08	-0.49 (0.21)	0.02
Sex: male* diagnosis: AD <sup>a,b</sup>	-0.28 (0.09)	0.003	-0.19 (0.07)	0.01	-0.34 (0.25)	0.17	-0.24 (0.21)	0.26
Sex: male* APOE: carrier <sup>a,c</sup>	-0.36 (0.11)	0.002	-0.24 (0.09)	0.01	NA	NA	NA	NA
Diagnosis: MCI* APOE: carrier <sup>b,c</sup>	-0.05 (0.12)	0.71	0.02 (0.10)	0.85	NA	NA	NA	NA
Diagnosis: AD* APOE: carrier <sup>b,c</sup>	-0.38 (0.10)	<0.001	-0.26 (0.08)	0.002	NA	NA	NA	NA

Sex: male*diagnosis: MCI* APOE: carrier <sup>a,b,c</sup>	0.34 (0.16)	0.03	0.25 (0.13)	0.05	NA	NA	NA	NA
Sex: male*diagnosis: AD* APOE: carrier <sup>a,b,c</sup>	0.52 (0.14)	<0.001	0.28 (0.11)	<0.001	NA	NA	NA	NA
Full models: We used General Linear Models (GLM	(1) with factors sex	, APOE e4 genot	ype and clinical di	sease stage, their	2-way interaction	s and 3-way intera	actions, and age w	vas included as a
covariate. This full model was run separately for each	ch of the biomarke	ers. <sup>a</sup> Reference is	female sex. <sup>b</sup> Refe	rence is SCD. <sup>c</sup> R	eference is APOE	non-carrier. p <0	0.05 is considered	significant. <sup>d</sup>
The natural logarithm of Tau and pTau concentrations are shown. Abbreviations: APOE, Apolipoprotein E4; SCD, Subjective Cognitive Decline; MCI, Mild Cognitive Impairment; AD,								
Alzheimer's Disease; NA, Not Applicable.								

Name	Location	Role	Contribution
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Betty M Tijms, PhD	Alzheimer Center & Department of Neurology, Amsterdam UMC, VU University Medical Center, Amsterdam, the Netherlands	Author	Interpreted the data; revised the manuscript for intellectual content
Philip Scheltens, MD, PhD	Alzheimer Center & Department of Neurology, Amsterdam UMC, VU University Medical Center, Amsterdam, the Netherlands	Author	Interpreted the data; revised the manuscript for intellectual content
Frederik Barkhof, MD, PhD	<ol> <li>Department of Radiology and Nuclear Medicine, AmsterdamUMC Location VUmc, Amsterdam, the Netherlands</li> <li>Institutes of Neurology and Healthcare Engineering, UCL, London, UK</li> </ol>	Author	Interpreted the data; revised the manuscript for intellectual content
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