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Longitudinal Changes in Anger, Anxiety, and Fatigue are Associated with Cerebrospinal Fluid Biomarkers of Alzheimer's disease

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Abstract

Alzheimer's disease (AD) studies in cognitively-normal (CN) older adults aged 65 suggest depression is associated with molecular biomarkers (imaging and cerebrospinal fluid [CSF]). This study used linear mixed models (covariance pattern model) to assess whether baseline CSF biomarkers ($A\beta_{42}/A\beta_{40}$, t-Tau/ $A\beta_{42}$, p-Tau/ $A\beta_{42}$) predicted changes in non-depressed mood states in CN older adults (N=248), with an average of three follow-up years. Participants with higher levels of CSF biomarkers developed more anger, anxiety, and fatigue over time compared to those with more normal levels. Non-depressed mood states in preclinical AD may be a prodrome for neuropsychiatric symptoms in symptomatic AD.

Keywords

Cerebrospinal Fluid; Alzheimer disease; Biomarkers; Older adults; Anger; Anxiety

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INTRODUCTION

Symptomatic Alzheimer's disease (AD) is often accompanied by depression, neuropsychiatric symptoms (NPS) and/or mild behavioral impairment (MBI) in later stages [1–3]. Conversely, depressive symptoms and a depression diagnosis have independently been associated as a prodrome and increased risk factor for both all-cause dementia, and AD specifically [4, 5]. Depression has been identified as a modifiable risk factor for reducing dementia prevalence [6]. In a similar vein, older adults (age 65 and older) who developed NPS (e.g., nighttime behaviors, irritability) experienced a faster change progression to symptomatic AD [7, 8]. Given the complex interactions and difficulty disentangling causality, emerging studies using molecular biomarkers have begun assessing the associations between underlying pathophysiology and early changes in mood [9].

Cross-sectional and longitudinal studies of amyloid imaging using Positron Emission Tomography (PET) suggest that more abnormal levels of amyloid are associated with depressive symptoms [10–12]. One study investigated and found that PET-tau (not amyloid imaging) was associated with a greater risk of depression diagnosis [13]. In AD, depression as a mood disorder has many distinguishing characteristics that may impact the results, including symptoms vs. diagnosis, active vs. remote, late-life onset vs. lifelong depression, and antidepressant use. Yet, there are limited studies that investigate whether mood states beyond depression are associated with molecular AD biomarkers. A prior study examining cerebrospinal fluid (CSF) and PET-amyloid biomarkers found that older adults with more change in confusion, anxiety, in addition to depression and NPS over one-year had more elevated levels of CSF and PET biomarkers [9]. However, it is unclear if changes in the non-depressed mood states are sustained over time given the limited follow-up interval. This study builds upon our past research to examine associations between non-depressed, nonbehavioral mood states assessed longitudinally and CSF biomarkers in a cognitively-normal sample of older adults.

METHODS

Sample

Participants were enrolled in studies conducted at the Knight Alzheimer's Disease Research Center at Washington University in St. Louis. Cognitive normality was defined by a 0 on the Clinical Dementia Rating (CDR®) [14]. Participants completed an annual clinical and neurological exam, demographics, mood, a health history, and neuropsychological assessments. Biomarker testing (CSF) was obtained every two to three years. Participants were included if they had a CDR® of 0 at baseline assessment and did not progress upon subsequent follow-up. Additionally, CSF data within two years of the baseline mood was used. Mood follow-up data was used until March 1, 2020 to exclude any potential confounding effects of the pandemic on mood. This study was approved by the Institutional Review Board of Washington University in Saint Louis, and all participants provided signed informed consent.

CSF

Following an overnight fast, a trained neurologist used a 22-gauge Sprotte spinal needle to collect 20–30 mL of CSF via standard lumbar puncture into a polypropylene tube at 8:00 A.M. as previously described [15–17]. CSF analytes including amyloid-beta₄₀ (A β_{40}), amyloid-beta₄₂ (A β_{42}) total tau (t-Tau), and phosphorylated tau₁₈₁ (p-Tau₁₈₁) were measured using an automated electrochemiluminescence immunoassay (Lumipulse G1200, Fujirebio). CSF biomarker ratios (A β_{42} /A β_{40} , t-Tau/A β_{42} , p-Tau/A β_{42}) were used to demarcate preclinical AD given their robustness, sensitivity, and specificity in predicting progression from cognitive-normality to symptomatic AD [17–20]. CSF amyloid cutoffs have high concordance with amyloid PET [21]. Participants were classified as preclinical AD positive if A β_{42} /A β_{40} ratio 0.0673, t-Tau/A β_{42} ratio 0.488, and p-Tau/A β_{42} 0.0649 based on established cut-offs [22].

Mood States

The Profile of Mood States—Short Form (POMS-SF) is a self-report, 30-item assessment of six mood states (positive and negative) and provides a total mood disturbance (TMD) score [23]. The six subscales detect presence and severity of anxiety, depression, anger, vigor, fatigue, and confusion (range 30–80). Higher scores on the vigor subscale indicates positive mood while the remaining subscales indicate more negative mood. The TMD score is summation of anxiety, depression, anger, fatigue, and confusion subscales, minus the vigor subscale. TMD scale ranges from –20 to 100, where higher scores indicate greater mood disturbance.

Statistical Analysis

Descriptive statistics summarized key demographics variables. Longitudinal analyses assumed a linear relationship between POMS subscales and CSF biomarkers within a two-year window of the index date/baseline POMS measurement. A covariance pattern model (linear mixed model) was used to predict mean POMS subscales based on CSF biomarkers. CSF biomarkers were dichotomized using aforementioned cutoffs to reflect negative (without preclinical AD) and positive (with preclinical AD) biomarker values. This model assumed a common variance-covariance correlation structure among the repeatedly measured POMS subscales over time from the same participant. Appropriate correlation matrix was selected based on model fit statistic Akaike Information Criterion (AIC). Analyses adjusted for age, gender, and education. Least squares means of POMS subscales for CSF groups were predicted from the covariance pattern model. Regression diagnostics such as residual plots were used to check the assumptions of the linear mixed models. Parallel analyses were conducted substituting the POMS and subscales with the Neuropsychiatric Inventory Questionnaire (NPIQ) and Geriatric Depression Scale (GDS) using the same biomarker models and demographic adjustments. All the statistical analyses were two-tailed at a significance level of 0.05 and performed with SAS 9.4 (SAS Institute, Cary, NC).

RESULTS

Participants were on average older, more educated with at least a bachelors' level of education, has a similar gender distribution, and were a majority of non-Hispanic white (Table 1). There were no impairments in cognitive functioning as reflected by the Mini-Mental State Examination (MMSE), and only nine percent had a depression diagnosis at baseline assessment. This ongoing study included data from seven years of follow-up POMS data (mean: 3.1 years; range 1–7 years). Approximately one third of the sample had at least one copy of the Apolipoprotein E ε 4 allele. Across the three biomarker ratios: A $\beta_{42}/A\beta_{40}$ (38.3%), t-Tau/A β_{42} (33.9%), p-Tau/A β_{42} (31.9%), roughly a third of the sample were classified as having preclinical AD. On average, the baseline scores on the POMS score centered around the nadir for the five negative moods states suggesting little symptoms and higher on the vigor subscale indicating optimal energy.

In the longitudinal analyses using the covariance pattern models, three of the subscales of the POMS revealed statistically significant differences between older adults with and without preclinical AD (Table 2; Figure 1). In the anger subscale, participants who were classified as preclinical AD by CSF p-Tau/A β_{42} and A $\beta_{42}/A\beta_{40}$ had slightly higher scores compared to those with more normal biomarker levels. Based on p-Tau/A β_{42} , older adults with more abnormal levels had higher scores on the tension/anxiety and fatigue subscale, respectively. There were some marginally significant results between the tension/anxiety subscale and A $\beta_{42}/A\beta_{40}$ (*p*=0.062) and t-Tau/A β_{42} (*p*=0.051). Similarly, the fatigue subscale was marginally significant with t-Tau/A β_{42} (*p*=0.063). There were no significant group differences between the CSF biomarkers on the POMS depression subscale or the TMD. We re-ran the models substituting biomarker group with APOE4 status (+/-) and did not find any group differences with APOE4 across the six mood states or TMD. Analyses conducted with biomarker models(p-Tau/A β_{42} , A $\beta_{42}/A\beta_{40}$, Tau/A β_{42}) and the NPIQ and GDS, independently were not statistically significant.

DISCUSSION

Understanding the association between mood states, NPS, and biomarkers, in addition to their changes can provide valuable insight into the trajectory of AD. In this longitudinal study with seven years of follow-up data in cognitively-normal older adults, participants with higher levels of CSF biomarkers developed more anxiety, fatigue, and anger over time compared to those with more normal biomarker levels. There were no statistically significant differences in depression mood state, overall mood disturbances, the NPIQ or GDS between both groups.

Anger was consistently higher across CSF p-Tau/A β_{42} and A β_{42} /A β_{40} biomarker groups with more elevated levels. In a study published three decades ago, anger was found to be more prevalent in patients with probable AD and associated with increased cognitive loss as measured by the MMSE, however, depression was not associated with cognitive changes [24]. A recent systematic review found only a handful of biomarker studies examining associations between agitation and aggression which identified positive correlations between CSF A β_{42} , p-Tau, and t-Tau [25]. The cross-sectional nature of the small number of

biomarker studies was a key limitation identified by the authors. Our longitudinal study followed a cognitively-normal cohort in their natural trajectory of mood changes and found that greater amyloid and tau pathology predicted more change in anger. One potential explanation may be that increased anger in the preclinical stage of AD may be a prodrome for agitation and aggression in NPS or MBI as a premonitory stage [3], when a person progresses to symptomatic AD.

A higher score on the tension/anxiety subscale on the POMS was associated with preclinical AD based on the p-Tau/A β_{42} and Tau/A β_{42} biomarkers (A β_{42} /A β_{40} was marginal; p=0.06). Prior studies examining anxiety and AD have found inconsistent results given the shared symptomology with depression [26–28]. A study using the Alzheimer's Disease Neuroimaging Initiative (ADNI) cohort found that anxiety symptoms in participants with amnestic mild cognitive impairment predicted progression to symptomatic AD, independent of depression [29]. Anxiety was assessed using a single item (including severity rating) on the NPIQ and changes in structural volumetric MRI biomarkers (e.g. intracranial, hippocampal, amygdala) was included in the models with the ADNI cohort. A recent study using the Harvard Aging Brain Study (HABS) cohort [11] found that greater cerebral amyloid (via imaging) was associated with higher anxious-depressive scores based on the GDS symptoms in a cognitively-normal sample followed up to five years. The results from our study extend the findings from the ADNI and HABS cohorts by parceling out depression, using a cognitively-normal cohort, and with a longer follow-up period. Finally, greater fatigue was associated with higher levels of p-Tau/A β_{42} and was marginally associated with t-Tau/A β_{42} (p=0.06). Fatigue, a self-report of tiredness resulting from mental or physical exertion and increased vulnerability to stressors, increases with biological age and can have a negative impact on overall health and well-being [30]. A recent study of Multidomain Alzheimer Preventive Trial cohort found chronic fatigue was associated with greater white matter hyperintensities in a cognitive normal cohort. However, there are limited AD studies examining the relationship between fatigue and amyloid, tau, and neurodegeneration biomarkers.

The extant literature examined mood, largely depression and preclinical AD and consistently finds a moderate association with depression symptoms and diagnosis with imaging biomarkers (amyloid and tau tracers) [10, 12, 13]. A key difference between prior research and this study is the use of the POMS survey rather than the GDS or the NPIQ and preclinical AD based on CSF biomarkers. In our prior study [9], we examined differences in the POMS and CSF biomarkers with one year of follow-up data. We found a group difference between t-Tau/A β_{42} but not p-Tau/A β_{42} (A β_{42} /A β_{40} was not available at the time). In this study, while t-Tau/A β_{42} did not reach statistical significance, it was marginal (*p*=0.06) providing consistent results with our prior work but also elucidating new relationships with non-depressed mood states and AD biomarkers.

MBI is conceptualized as the late-life onset of behaviors (low motivation, affect dysregulation, impulse dyscontrol, social inappropriateness, abnormal thought content), which increase the conversion risk from mild cognitive impairment to dementia and may presage transition from cognitive-normality to preclinical stage[31, 32]. MBI is also associated with biomarkers— PET amyloid and plasma neurofilament light, a marker of

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axonal damage signifying neurodegeneration[33, 34]. Future studies could examine the associations between MBI (behavior), mood states, and AD biomarkers. Cognitive reserve may also impact emergence of negative mood states and moderate the relationship with conversion to prodromal AD and should be examined to assess mediating or moderating roles [35].

There are some limitations despite the longitudinal follow-up period and modest sample size with biomarkers. Obtaining CSF biomarkers is specialized, costly, burdensome, and not widely available. The slightly skewed means on the POMS TMD and subscales may reflect the overall high physical and mental health of the participants in this cohort. This was also reflected in the low prevalence of depression in the sample. Pharmacological therapy including specific medication classes (e.g., antidepressants) can influence the presence and magnitude of behavior and mood states. We did not have data on current medication available to examine and adjust in our analyses. Our sample included a large proportion of non-Hispanic whites, who were well educated, most lacking significant physical disabilities, psychiatric, or neurologic conditions/diagnosis, which may not be representative of the general population. As a result, findings may not be easily generalizable to the larger population.

In sum, the findings from this study of mood states suggest that longitudinal changes in anger, tension/anxiety, and fatigue were associated with baseline preclinical AD biomarkers. These changes in non-depressed mood states may be a prodrome of future NPS or indicative of MBI as an older adult progresses to symptomatic AD. Assessment of various mood states in the preclinical and symptomatic stages of AD may provide more information for clinical diagnostics and to understand the trajectory on how emotive and cognitive changes co-occur to impact function.

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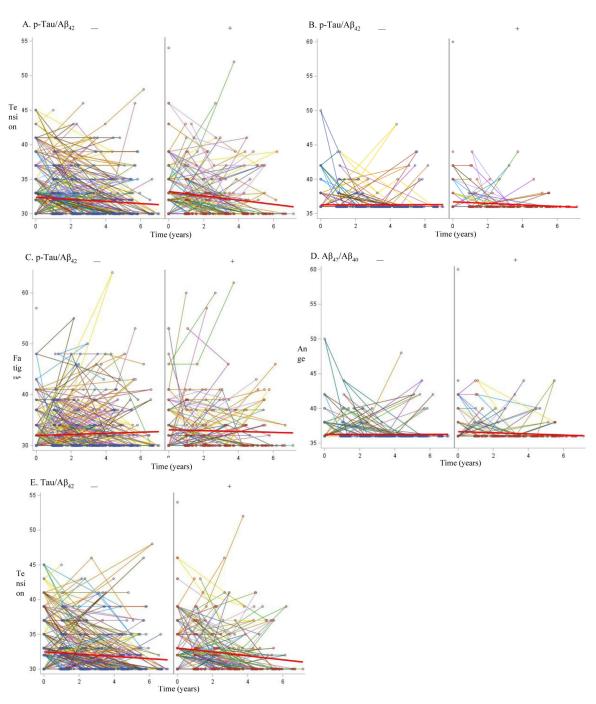
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Spaghetti plots show the changes in POMS mood subscales for tension/anxiety, anger, and fatigue for $A\beta_{42}/A\beta_{40}$, t-Tau/A β_{42} , p-Tau/A β_{42} , with preclinical AD – and + groups to the left and right, respectively. Individual lines represent longitudinal data for a single participant. POMS = Profile of Mood States.

Table 1.

Baseline demographics (N=248)*

Age (years)	72.78±4.80				
Education (years)	16.33	3±2.45			
Women, N (%)	127 (5	51.21%)			
Race, Caucasian, N (%)	220 (8	38.71%)			
APOE4, N	82 (3	3.47%)			
MMSE (out of 30)	29.28	8±1.00			
Depression, N (%)	23 (9.27%)				
GDS (out of 15)	$0.69{\pm}1.76$				
NPIQ (out of 36)	0.92±1.36				
Follow-up time (years)	3.12±2.31				
POMS					
TMD	-4.74 ± 8.61				
Tension/Anxiety	32.83	3±3.87			
Depression	32.52±2.29				
Anger	36.49±2.11				
Vigor	58.91±9.74				
Fatigue	32.36±4.48				
Confusion	38.08±3.78				
Biomarker status (%)	+	-			
$A\beta_{42}/A\beta_{40}$	95 (38.31%)	153 (61.69%)			
t-Tau/A β_{42}	84 (33.87%)	164 (66.13%)			
$p\text{-Tau}/A\beta_{42}$	79 (31.85%)	169 (68.15%)			

Abbreviations: APOE4 = Apolipoprotein E e4 allele; MMSE = Mini-Mental State Examination; POMS = Profile of Mood States; TMD = Total Mood Disturbance; GDS = Geriatric Depression Scale; NPIQ = Neuropsychiatric Inventory Questionnaire

*Mean \pm Standard Deviation or count (percentage)

Table 2.

Group means of mood subscales across CSF ratio biomarkers

POMS	p-Tau/Aβ ₄₂					
	+		_		р	
	LSM	95% CI	LSM	95% CI		
TMD	-4.06	(-5.80, -2.33)	-5.58	(-6.77, -4.39)	0.159	
Anxiety	33.09	(32.39, 33.79)	31.97	(31.49, 32.45)	0.010	
Depression	32.94	(32.48, 33.41)	32.46	(32.14, 32.78)	0.095	
Anger	36.67	(36.39, 36.97)	36.25	(36.06, 36.45)	0.017	
Vigor	59.78	(57.94, 61.61)	58.75	(57.50, 60.00)	0.367	
Fatigue	33.21	(32.31, 34.09)	32.08	(31.47, 32.69)	0.042	
Confusion	38.41	(37.74, 39.09)	37.71	(37.25, 38.17)	0.091	
GDS	0.999	(0.785, 1.121)	1.023	(0.710, 1.336)	0.902	
NPIQ	0.770	(0.573, 0.966)	0.852	(0.562, 1.143)	0.644	
	$A\beta_{42}/A\beta_{40}$					
		+		-	р	
	LSM	95% CI	LSM	95% CI		
TMD	-4.57	(-6.16, -2.98)	-5.42	(-6.68, -4.17)	0.412	
Anxiety	32.81	(32.17, 33.46)	32.03	(31.52, 32.54)	0.062	
Depression	32.79	(32.36, 33.22)	32.51	(32.17, 32.84)	0.308	
Anger	36.64	(36.38, 36.90)	36.23	(36.03, 36.44)	0.015	
Vigor	59.72	(58.04, 61.40)	58.68	(57.36, 60.00)	0.341	
Fatigue	32.79	(31.97, 33.561	32.23	(31.58, 32.87)	0.266	
Confusion	38.21	(37.59, 38.83)	37.76	(37.27, 38.25)	0.263	
GDS	1.005	(0.778, 1.231)	1.010	(0.725, 1.296)	0.974	
NPIQ	0.777	(0.570, 0.985)	0.826	(0.561, 1.091)	0.779	
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	LSM	95% CI	LSM	95% CI		
TMD	-4.06	(-5.73, -2.38)	-5.64	(-6.85, -4.43)	0.132	
Anxiety	32.88	(32.20, 33.56)	32.04	(31.55, 32.53)	0.050	
Depression	32.96	(32.51, 33.41)	32.44	(32.11, 32.76)	0.066	
Anger	36.58	(36.31, 36.86)	36.29	(36.09, 36.49)	0.093	
Vigor	59.21	(57.44, 60.97)	59.01	(57.73, 60.29)	0.859	
Fatigue	33.10	(32.24, 33.95)	32.09	(31.47, 32.71)	0.063	
Confusion	38.29	(37.64, 38.94)	37.75	(37.28, 38.22)	0.189	
GDS	1.006	(0.789, 1.225)	1.007	(0.789, 1.309)	0.996	
NPIQ	0.786	(0.585, 0.986)	0.815	(0.537, 1.093)	0.8692	

Abbreviations: POMS = Profile of Mood States; LSM = Least Squares Means; TMD = Total Mood Disturbance; GDS = Geriatric Depression Scale; NPIQ = Neuropsychiatric Inventory Questionnaire

Bold = <.05; *Italics* = marginally significant; 95% CI = 95% Confidence Interval;

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