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Circulating Ethanolamine Plasmalogen Indices in Alzheimer's Disease: Relation to Diagnosis, Cognition, and CSF Tau

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Abstract:	<p>Introduction:Altered lipid metabolism is implicated in Alzheimer's disease (AD) but the mechanisms remain obscure. Aging-related declines in plasmalogens may increase AD risk by reducing plasmalogen availability. Methods:We measured 4 ethanolamine plasmalogens (PlsEtns) and 4 phosphatidylethanolamines (PtdEtns) from AD Neuroimaging Initiative (ADNI; n=1547 serum) and University of Pennsylvania (UPenn; n=112 serum) cohorts, and derived indices reflecting PlsEtn/PtdEtn metabolism: PL-PX (PlsEtns), PL/PE (PlsEtn/PtdEtn), and PBV (composite index). We tested associations with diagnosis, cognition, and CSF biomarkers. Results:Results revealed significant negative relationships in ADNI: AD vs. CN and LMCI vs. CN with PL-PX and PBV. In UPenn, AD vs. CN was associated negatively with PL/PE and PBV. In ADNI, cognition was negatively associated with plasmalogen indices (PL-PX and PBV). In both cohorts total-tau and t-tau/Aβ1-42 was negatively associated with plasmalogen indices. Discussion:These data indicate an association of decreased PlsEtns with AD, cognition, and CSF tau. Future studies are needed to define mechanistic relationships.</p>

Circulating Ethanolamine Plasmalogen Indices in Alzheimer's Disease: Relation to Diagnosis, Cognition, and CSF Tau

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6 *Data used in preparation of this article were obtained from the Alzheimer’s Disease
7 Neuroimaging Initiative (ADNI) database (adni.loni.usc.edu). As such, the investigators within
8 the ADNI contributed to the design and implementation of ADNI and/or provided data but did
9 not participate in analysis or writing of this report. A complete listing of ADNI investigators can
10 be found at:

11 http://adni.loni.usc.edu/wpcontent/uploads/how_to_apply/ADNI_Acknowledgement_List.pdf
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15 **Abbreviations:** AD: Alzheimer’s disease; ADAS-Cog13: 13-item Alzheimer’s Disease
16 Assessment Scale-Cognitive subscale; ADMC: Alzheimer’s Disease Metabolomics Consortium;
17 ADNI: Alzheimer’s Disease Neuroimaging Initiative; CN: Cognitively normal older control;
18 APOE: apolipoprotein E; APP: amyloid precursor protein; CNS: central nervous system; CSF:
19 cerebrospinal fluid; DHA: docosahexaenoic acid ; EMCI: Early mild cognitive impairment;
20 FDR: False discovery rate; LCFA: long-chain fatty acids; LMCI: Late mild cognitive
21 impairment; MCI: Mild cognitive impairment; MMSE: Mini-Mental State Exam; PDI:
22 Phenomenome Discoveries, Inc.; PBV: plasmalogen biosynthesis value; PlsEtns: ethanolamine
23 plasmalogens; PtdEtns: phosphatidylethanolamines; SMC: significant memory concerns; VL-
24 PUFA: very long chain polyunsaturated fatty acid
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28 **Key words:** Dementia, mild cognitive impairment, phosphatidylethanolamines,
29 plasmenylethanolamines, metabolomics, lipidomics, case-control study, mass spectrometry
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4 **Abstract**
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6 Introduction: Altered lipid metabolism is implicated in Alzheimer's disease (AD), but the
7 mechanisms remain obscure. Aging-related declines in circulating plasmalogens containing
8 omega-3 fatty acids may increase AD risk by reducing plasmalogen availability.
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11 Methods: We measured four ethanolamine plasmalogens (PlsEtns) and four closely-related
12 phosphatidylethanolamines (PtdEtns) from the AD Neuroimaging Initiative (ADNI; n=1547
13 serum) and University of Pennsylvania (UPenn; n=112 serum) cohorts, and derived indices
14 reflecting PlsEtn and PtdEtn metabolism: PL-PX (PlsEtns), PL/PE (PlsEtn/PtdEtn ratios), and
15 PBV (plasmalogen biosynthesis value; a composite index). We tested associations with baseline
16 diagnosis, cognition, and cerebrospinal fluid (CSF) AD biomarkers.
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19 Results: Results revealed statistically significant negative relationships in ADNI between AD
20 vs. CN with PL-PX ($p=0.007$) and PBV ($p=0.005$), LMCI vs. CN with PL-PX ($p=2.89 \times 10^{-5}$) and
21 PBV ($p=1.99 \times 10^{-4}$), and AD vs. LMCI with PL/PE ($p=1.85 \times 10^{-4}$). In the UPenn cohort, AD vs.
22 CN diagnosis associated negatively with PL/PE ($p=0.0191$) and PBV ($p=0.0296$).
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25 In ADNI, cognition was negatively associated with plasmalogen indices, including
26 ADAS-Cog13 (PL-PX: $p=3.24 \times 10^{-6}$; PBV: $p=6.92 \times 10^{-5}$) and MMSE (PL-PX: $p=1.28 \times 10^{-9}$; PBV:
27 $p=6.50 \times 10^{-9}$). In the UPenn, there was a trend towards a similar relationship of MMSE with
28 PL/PE ($p=0.0949$).
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31 In ADNI, CSF total-tau was negatively associated with PL-PX ($p=5.55 \times 10^{-6}$) and PBV
32 ($p=7.77 \times 10^{-6}$). Additionally, CSF t-tau/ $A\beta_{1-42}$ ratio was negatively associated with these same
33 indices (PL-PX, $p=2.73 \times 10^{-6}$; PBV, $p=4.39 \times 10^{-6}$). In the UPenn, PL/PE was negatively
34 associated with CSF total-tau ($p=0.031$) and t-tau/ $A\beta_{1-42}$ ($p=0.021$). CSF $A\beta_{1-42}$ was not
35 significant associated with any of these indices in either cohort.
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39 Discussion: These data extend previous studies by showing an association of decreased
40 plasmalogen indices with AD, MCI, cognition, and CSF tau. Future studies are needed to better
41 define mechanistic relationships, and to test the effects of interventions designed to replete serum
42 plasmalogens.
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1. Introduction

Alzheimer's disease (AD) is a major public health problem [1]. While much is known about its natural history, epidemiologically-identified risk factors, and characteristic neurodegenerative changes, there are currently no available treatments to modify the underlying disease process or prevent cognitive or functional decline. Moreover, relatively little is known regarding etiologic mechanisms that initiate or promote neurodegenerative processes in sporadic AD. Biomarkers that identify AD in preclinical stages may aid in elucidating these mechanisms. Currently, most accepted biomarkers relate to neuropathologic features of AD, e.g., cerebrospinal fluid (CSF) levels of tau and amyloid- β and neuroimaging of brain atrophy and amyloid or tau deposits. Blood-based biomarkers are potentially appealing due to lower participant burden. Serum- and plasma-based biomarkers have been used in other studies to predict cognitive declines, dementia diagnosis, and progression to AD [2-6].

Altered lipid metabolism is implicated in sporadic AD risk [7, 8]; however, the precise mechanisms accounting for findings from observational studies remains to be fully elucidated. Although cholesterol and its content in circulating lipoproteins are perhaps the most well-studied lipids in AD [8], recent data implicate other classes such as glycerophospholipids [2, 9-14]. Plasmalogens represent a subclass of glycerophospholipids carrying a vinyl ether linkage at the sn1 carbon of the glycerol backbone, vs. the more familiar fatty acyl ester linkage [15] (see **Supplemental Figure 1**). Plasmalogens are integral membrane components that appear to play important roles relevant to AD pathophysiology, including vesicle fusion [16, 17] necessary for synaptic neurotransmitter release [18], modulation of membrane fluidity and microdomain dynamics [19], membrane antioxidant functions [20, 21], and neuroprotection [22]. Plasmalogens can exert reciprocal effects with cholesterol on membrane fluidity and lipid microdomain composition, promoting the breakdown of amyloid precursor protein (APP) by alpha-secretase to non-amyloidogenic peptide products. This enhancement of alpha-secretase-mediated APP breakdown may reduce the production of beta-secretase-derived peptides and thus to the formation of the amyloid plaques characteristic of AD neuropathology [23].

The initial steps of endogenous plasmalogen synthesis occur in peroxisomes, notably in the liver [15, 24]. Peroxisomes also perform enzymatic functions necessary for synthesis of long-chain fatty acids (LCFA) and very long-chain fatty acids (VLCFA) such as docosahexaenoic acid (DHA, C22:6n-3), particularly beta-oxidation and side chain elongation steps. Plasmalogens

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4 are incorporated into circulating lipoproteins [25], from which they can be transported to other
5 organs including the brain [26, 27]. Serum plasmalogen levels decline with increasing age [28,
6 29] and previous studies show decreased levels of serum and postmortem brain plasmalogens in
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8 AD [9, 10, 13, 29-34].
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11 Ratios of metabolites measured at a given time point are frequently informative regarding
12 the homeostasis of metabolites and metabolic flux. Thus, these ratios can be used to identify
13 metabolic defects and to elucidate etiopathogenesis. One example from disorders of peroxisome
14 biogenesis is the use of very long chain fatty acid ratios in the diagnosis of Zellweger Spectrum
15 Disorders where elevation of C24/C22 and C26/C22 ratios are indicative of enzymatic defects in
16 the beta-oxidation of very long chain fatty acids in the peroxisomes [35]. Zellweger syndrome is
17 a peroxisomal biogenesis disorder with multi-enzyme defects in the peroxisomal beta-oxidation
18 pathway [35].
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26 Another commonly used lipid ratio is that of omega-6/omega-3 fatty acids which reflects the
27 relative activity of competing metabolic pathways in omega-6 and omega-3 fatty acid synthesis,
28 dietary intake and rate of utilization [36]. Plasmalogens act as reservoirs of polyunsaturated fatty
29 acid and they are selectively enriched in certain PUFA side chains [37]. Thus, fatty acid
30 composition in plasmalogens bring valuable insights not only about plasmalogen biosynthesis
31 but also about homeostasis of fatty acids and remodeling of plasmalogens.
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37 Ratios among plasmalogen species (PL/PL) having substrate-product relationships, such as
38 those described below, reflect the relative abundance of DHA, EPA (omega-3) and adrenic acid
39 (omega-6) fatty acid side chains in plasmalogens and represent the results of plasmalogen
40 remodeling, fatty acid biosynthesis/dietary intake, and peroxisomal beta-oxidation. Plasmalogen
41 and phosphatide levels are homeostatically regulated, such that a reduction in the circulating
42 levels of one subclass (e.g., plasmalogens) is often accompanied by a compensatory increase in
43 the other (e.g., phosphatides) [38]. Therefore, ratios of PL to PE reflect this equilibrium and
44 represent the capacity of the body to synthesize plasmalogens. In addition, the calculation of
45 composite indices from two or more ratios may result in reduced biological variability and thus
46 show greater comparability across various cohorts. Certain indices constructed in this fashion
47 were found to be associated with cognition in previous studies [10].
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57 We report here data obtained using a focused lipidomics platform to measure four
58 ethanolamine plasmalogens and four closely-related phosphatidylethanolamines in baseline
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4 serum specimens from the Alzheimer’s Disease Neuroimaging Initiative (ADNI)-1, -GO, and -2
5 cohorts, and in plasma samples from subjects at the Perelman School of Medicine, University of
6 Pennsylvania (UPenn). We selected these species based on previous work in several disparate
7 populations as described in [10]. We then calculated ratios of among omega-3 and omega-6-
8 bearing plasmalogens (PL/PL) and of plasmalogens to phosphatides (PL/PE), and constructed
9 composite indices from these ratios based on previous work [10]. Algorithm-based in vitro
10 diagnostic multivariate index assays are used in increasing frequency to identify the risks of
11 complex chronic diseases. Our overarching goal was to create an in vitro diagnostic multivariate
12 index that can be used to assess the risk of AD in pre-clinical stages. The use of these ratios
13 helped create a single index that can be tested in future longitudinal studies for early detection of
14 AD.
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17 We further note that this study was not intended to be an untargeted lipidomic study aimed
18 at generation of hypotheses, but rather was a focused analysis of selected members of certain
19 physiologically important lipid subclasses known to be associated with AD and cognitive
20 dysfunction. The molecules of interest have been well researched and published in peer-reviewed
21 journals [9, 10, 13, 29-34]. Preliminary results on the particular metabolite ratios mentioned here
22 have been presented in Alzheimer’s Association Conferences from years 2014-2016 and
23 published as abstracts in Alzheimer’s and Dementia [39, 40], as well as in a paper based on some
24 of these data that was recently published [10]. These metabolites and their ratios were selected
25 iteratively from comparisons of data across several large lipidomic datasets in previous studies
26 cited earlier. The objective of this study was to confirm these findings in well characterized
27 cohorts, such as ADNI and UPenn.
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29 We sought to test the following hypotheses:

- 30 1. Are indices of plasmalogen and related phosphatide metabolism associated with (a)
31 diagnosis or (b) cognitive function at the time of blood sampling?
- 32 2. Are indices of plasmalogen and related phosphatide metabolism associated with CSF
33 biomarkers of AD pathophysiology, i.e., amyloid-beta and tau?
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35 We present here findings showing a reduction in indices of plasmalogen remodeling that may
36 also reflect altered biosynthesis and peroxisomally-derived plasmalogen beta-oxidation
37 associated with AD.
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4 **2. Methods**
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7 **2.1. Study cohorts and samples:**
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10 **2.1.1. Alzheimer’s Disease Neuroimaging Initiative (ADNI)**
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12 Data were obtained from the Alzheimer’s Disease Neuroimaging Initiative (ADNI)
13 database (www.adni.loni.usc.edu). In this study, we utilized serum samples obtained at the
14 baseline study visit in the ADNI-1, -GO, and -2 cohorts (see **Supplemental Text 1** for a
15 summary of the study design and inclusion criteria for these cohorts). For additional
16 information, see www.adni-info.org. Metabolomics data and results have been made accessible
17 through the AMP-AD Knowledge Portal (<https://ampadportal.org>). The AMP-AD Knowledge
18 Portal is the distribution site for data, analysis results, analytical methodology and research tools
19 generated by the AMP-AD Target Discovery and Preclinical Validation Consortium and multiple
20 Consortia and research programs supported by the National Institute on Aging. Information on
21 data availability and accessibility is available in the **Data Availability** section.
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32 **2.1.2. University of Pennsylvania Alzheimer’s Disease Clinical Center (ADCC)**
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34 We selected subjects from the Integrated Neurodegenerative Diseases Database at the
35 Perelman School of Medicine, University of Pennsylvania for studies of plasma and
36 cerebrospinal fluid (CSF) metabolomics (see **Supplemental Text 1** for selection criteria). All
37 subjects gave informed consent for participation in research protocols approved by the Penn
38 Institutional Review Board for the collection of clinical and genetic data and biofluid samples. A
39 total of 127 samples were selected, representing 112 unique subjects; 12 subjects underwent
40 plasma sampling at two different times, and one subject was sampled on three occasions. In all
41 cases, the earliest plasma sample was used for the analyses described here.
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49 **2.2. Procedures (blood draw, sample storage/handling)**
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52 Blood samples were collected in the morning after an overnight fast (water was permitted)
53 prior to collection. Serum was obtained from blood drawn into 10-mL plain red top plastic tubes,
54 allowed to clot for 30 minutes at room temperature in a vertical position, and centrifuged at
55 1,500 relative centrifugal force (rcf) for 15 minutes. Serum was collected into polypropylene
56 tubes, frozen, and shipped on dry ice to the ADNI Biorepository. Serum samples were
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4 subsequently thawed and divided into 0.5-mL aliquots at the ADNI Biorepository, and frozen at -
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6 80C until retrieved for analysis.

7 8 **2.3. APOE Genotype**

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10 *APOE* genotyping was performed at the time of participant enrollment. The two SNPs
11 (rs429358, rs7412) that define the epsilon 2, 3, and 4 alleles, were genotyped using DNA
12 extracted by Cogenics from a 3-mL aliquot of EDTA blood.
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15 16 **2.4. Cognitive Function**

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18 Global cognition was measured using the modified Alzheimer Disease Assessment Scale
19 13-item cognitive subscale (ADAS-Cog13) where higher scores indicate worse cognition and the
20 Mini-Mental State Exam (MMSE) where higher scores indicate better cognition. ADAS-Cog13
21 and MMSE scores were available in the ADNI cohort, whereas MMSE scores were only
22 available in the UPenn cohort.
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25 26 **2.5. Biochemical analysis**

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28 We measured four plasmenylethanolamine (PlsEtn) and four phosphatidylethanolamine
29 (PtdEtn) glycerophospholipid species with the following sn-1/sn-2 compositions in 20-uL
30 aliquots of serum or EDTA plasma: PlsEtn 16:0/18:2, 18:0/20:5, 16:0/22:4, 16:0/22:6 and PtdEtn
31 16:0/18:3, 18:0/20:5, 18:0/22:4, 16:0/22:6 by a stable isotope dilution, flow-injection analysis
32 (FIA)-mass spectrographic (MS) method on an Ionics 3Q tandem mass spectrometer operating in
33 the negative ionization atmospheric pressure chemical ionization (APCI) mode. Methods are
34 summarized in **Supplemental Text 2** and described elsewhere [9, 13]. Presumed structures of
35 phospholipids corresponding to the specific mass/charge (m/z) ratio peaks are shown in
36 **Supplemental Figure 1**. Due to the specific hypotheses being tested, the data reported here will
37 concern ratios and indices derived from three of the above plasmalogen species (PlsEtn
38 18:0/20:5, 16:0/22:4, 16:0/22:6) and one phosphatidylethanolamine species (PtdEtn 16:0/22:6).
39 These ratios and indices have previously been shown to be useful composite measures of
40 plasmalogen physiology in other cohorts [10].
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2.6. Statistical analysis

2.6.1. Calculation of phosphoethanolamine ratios and indices and rationale:

Each bivariate ratio was scaled to the full dataset for each of the cohorts examined (i.e., ADNI and UPenn) by log transformation of the ratio, followed by normalization to the mean and standard deviation of the values for the cohort. ADNI values underwent \log_2 transformation before normalization; UPenn data were \log_{10} transformed. Composite scores were then created from these scaled values for each sample to assess:

- 1) peroxisomal plasmalogen (PL) biosynthesis and/or remodeling (PL-PX),
- 2) relative abundance of PLs vs. PEs (PL/PE); and
- 3) overall plasmalogen biosynthesis value (PBV).

These indices were calculated as follows and are summarized in **Supplemental Table 1**:

- A) **PL-PX**: Mean of PL226/PL224, PL205/PL224, and PL205/PL226;
provides an index of biosynthesis and/or remodeling of ethanolamine plasmalogens based on ratios of concentration of specific sn2 isoforms measured (i.e., those containing adrenic acid or 22:4; eicosapentaenoic acid, EPA or 20:5; and docosahexaenoic acid or 22:6)
- B) **PL/PE**: Mean of PL226/PE226 and PL205/PE226;
provides an index of the ratio of omega-3 fatty acid-containing plasmalogens to the DHA-containing phosphatide PtdEtn 16:0/22:6.
- C) **Overall PBV**: Mean of the following five ratios:
PL226/PE226, PL205/PE226 (constituents of index B, above),
PL226/PL224, PL205/PL224, and PL205/PL226 (constituents of index A, above);
provides an overall composite index of plasmalogen biosynthesis and/or remodeling.

2.7. Statistical models

Statistical analyses were conducted using R version 3.2.4 and MATLAB R2016b. Data were analyzed according to previously published methods [41]. The statistical pipeline steps applied to these data are summarized in **Supplemental Figure 2**.

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4 Phosphoethanolamines were measured in 829 samples from 809 subjects within the
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6 ADNI-1 cohort and in 905 samples from 873 subjects in the ADNI-GO/2 cohort. Biological
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8 replicates were averaged for 20 subjects in ADNI-1 and 17 subjects in ADNI-GO/2. From the
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10 original set of 1734 samples, we removed 104 who were not fasting, two missing BMI, and one
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12 missing sex. An additional 26 subjects were removed after determining they were multivariate
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14 outliers based on squared Mahalanobis distances significant at the upper 1-tailed 0.05 level of
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16 confidence adjusted for multiple comparisons by false discovery rate (FDR) correction. This
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18 resulted in an analysis dataset containing 1545 subjects, five bivariate ratios, and three composite
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20 indices.

21 The Kruskal-Wallis one-way ANOVA on ranks for continuous covariates and the
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23 Pearson chi-square test for categorical covariates were used to check for differences among
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25 covariate information collected at baseline grouped by baseline diagnosis (CN vs. LMCI vs.
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27 AD).

28 To assess the association of metabolic features and the odds of being in one subgroup
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30 relative to being in another subgroup for all pairs of subgroups (i.e. MCI versus CN, AD versus
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32 CN, and AD versus MCI) binary logistic regression models were used. As EMCI and SMC were
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34 not represented in ADNI-1, we only examined comparisons among AD, (L)MCI, and CN
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36 subjects in the logistic regression analyses. The MCI subjects in ADNI-1 were combined with
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38 the LMCI cohort in ADNI-2, given the comparability of criteria for selecting these groups [42].
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40 In the UPenn cohort, due to the relatively small sample of MCI patients, we only tested the
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42 association of AD vs. CN with the three composite phosphoethanolamine indices.

43 To test the association of each metabolic feature with the outcome of interest in both
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45 ADNI and UPenn cohorts, a separate linear regression (ADAS-Cog13, MMSE, and CSF t-tau,
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47 $A\beta_{1-42}$, and t-tau/ $A\beta_{1-42}$ ratio) was performed. ADAS-Cog13 and MMSE were square-root
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49 transformed and CSF t-tau/ $A\beta_{1-42}$ was \log_{10} -transformed prior to analysis. All models included
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51 age, sex, and *APOE* $\epsilon 4$ genotype carrier status as covariates to control for the confounding
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53 effects (*APOE* $\epsilon 4$ genotype carrier status was not included in the analyses of the UPenn cohort
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55 due to the number of subjects who had not been genotyped for *APOE*). Moreover, education in
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57 years was controlled for in the analyses modeling cognition.
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4 Since it is known that the genetic variance in the FADS1/FADS2 gene cluster can affect
5 the blood levels of EPA/DHA containing lipids [43], we conducted a candidate gene association
6 analysis for relationships between plasmalogen ratios and indices with genetic variants in this
7 gene cluster. In the ADNI cohort, to control for the confounding effect of medication and dietary
8 supplements on metabolic levels, we constructed additional models in which each ratio and index
9 was regressed on 42 medication and dietary supplement categories using previously published
10 methods [41]. **Supplemental Figure 3** depicts the percentage of ADNI subjects in each
11 diagnostic group who were recorded as taking a member of each of these medication classes at
12 the baseline study visit. Medication classes were backward-selected via Bayesian information
13 criteria to select an optimal combination of medications for minimizing confounding while
14 limiting model complexity. For this analysis, we removed acetylcholinesterase inhibitors and
15 NMDA receptor antagonist drugs from consideration, in order to avoid sampling bias due to the
16 extreme degree of heterogeneity among diagnostic groups in the frequency of their use (i.e., few
17 if any CN subjects were taking these drugs, while most AD subjects report taking them). The
18 residuals from the regression of metabolic levels on medications and supplements were kept to
19 create a separate analysis dataset. We did not conduct this medication-adjusted analysis on the
20 UPenn cohort due to the size of this sample, which limited our power to conduct a meaningful
21 analysis.

3. Results

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Characteristics of the combined ADNI-1/-GO/-2 participants at the time of baseline serum collection are depicted in **Table 1A**. Characteristics of the University of Pennsylvania cohort are presented in **Table 1B**.

The means and standard deviations of the five ratios and three composite indices in the ADNI diagnostic groups AD, LMCI, and CN are shown in **Supplemental Table 2**.

3.1. Phosphoethanolamine lipid indices by diagnosis

3.1.1. AD vs. CN

The results of phosphoethanolamine lipid ratios and indices in cross-sectional analysis, unadjusted for medications, in the ADNI cohort are shown in **Table 2A**. Of the composite indices examined using pairwise-by-diagnosis logistic regression models, the PL-PX (Odds

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4 Ratio (OR) = 0.784, $p = 0.007$) and overall PBV (OR= 0.775, $p = 0.005$) showed a significant
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6 negative relationship with the likelihood of AD vs. CN diagnosis that persisted after FDR
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8 correction. This result indicates that reduced values of these indices of plasmalogen metabolism
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10 were associated with an increased likelihood of having AD vs. CN. The other composite index,
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12 PL/PE, did not show significant relationships with AD vs. CN diagnosis (**Table 2A**).

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14 When all medications except those specific to AD (i.e., anticholinesterases and
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16 memantine) were included in the models (see **Supplemental Table 3** for the medication classes
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18 retained in these models due to significant contributions to the analysis of the three composite
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20 measures), none of the associations of plasmalogen ratios or indices with the AD vs. CN
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22 diagnosis was significant (all p -values > 0.1). See **Supplemental Table 4** for a summary of these
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24 results.

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26 In the UPenn ADCC cohort the results of phosphoethanolamine lipid ratios and indices
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28 are shown in **Table 2B**. As observed in ADNI, in the UPenn ADCC cohort we observed a
29
30 statistically significant association of AD vs. CN with PBV (OR= 0.695, $p = 0.0296$). We also
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32 observed a significant association of AD vs. CN with PL/PE (OR=0.630, $p = 0.0191$). The PL-
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34 PX index did not show a significant association with AD vs. CN.

3.1.2. LMCI vs CN

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39 The results of cross-sectional logistic regression analysis of phosphoethanolamine lipid
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41 ratios and indices for association with baseline diagnoses of LMCI vs. CN in ADNI are shown in
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43 **Table 2A** (unadjusted for medications) and **Supplemental Table 2** (adjusted for all but AD-
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45 specific medications). In the non-medication adjusted analysis, as in the AD vs. CN analysis,
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47 both the PL-PX composite index (OR=0.719, $p = 2.89 \times 10^{-5}$, $q = 1.73 \times 10^{-4}$) and the PBV
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49 composite (OR=0.745, $p = 1.99 \times 10^{-4}$, $q = 7.97 \times 10^{-4}$) showed significant associations with the
50
51 LMCI vs. CN comparison (**Table 2A**).

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53 When medications except those specific to AD were included in the model, the
54
55 associations remained statistically significant for the PL-PX composite (OR=0.767, $p = 0.0019$)
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57 and the PBV composite (OR=0.802, $p=0.0096$) with the LMCI vs. CN comparison.
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3.1.3. AD vs LMCI

As shown in **Table 2A**, the unadjusted logistic regression analysis in the ADNI cohort for phosphoethanolamine lipid ratios and indices showed a negative association between specific composite indices and AD vs. LMCI, indicating that lower values of these ratios and indices were associated with a higher likelihood of AD vs. LMCI. In this analysis, the PL/PE (OR= 0.757, $p = 6.15 \times 10^{-4}$), showed a negative association with AD vs. LMCI. PL-PX and PBV did not show significant associations with AD vs. LMCI.

After adjusting for all non-AD-specific medications in ADNI, as shown in **Supplemental Table 4**, the logistic regression analyses on these ratios and indices showed that the association of PL/PE with the likelihood of AD vs. LMCI remained significant (OR=0.73, $p = 1.85 \times 10^{-4}$).

3.2. Cognitive Function

The results of cross-sectional linear regression analysis of baseline ADAS-Cog13 score in the ADNI cohort with phospholipid ratios and indices are shown in **Table 2A** (unadjusted analyses) and **Supplemental Table 4** (analyses adjusted for all non-AD-specific medications). In the medication-unadjusted analysis (**Table 2A**), statistically significant relationships were seen between baseline ADAS-Cog13 score and PL-PX ($\beta = -0.1349$, $p = 3.24 \times 10^{-6}$) and overall PBV ($\beta = -0.1302$, $p = 6.92 \times 10^{-5}$). Since higher values of ADAS-Cog13 indicate greater degrees of cognitive impairment, the negative sign of the coefficient indicates that lower values of these indices are associated with greater cognitive impairment. The other index, PL/PE, did not show a significant relationship with ADAS-Cog13. When adjusted for all non-AD-specific medications, the above associations remained statistically significant. Thus, the composite indices PL-PX ($\beta = -0.0996$, $p = 0.0018$), and overall PBV ($\beta = -0.0938$, $p = 0.0031$) remained significant after non-AD-specific medications adjustment. The association of the composite measure PL/PE with ADAS-Cog13 remained non-significant.

In ADNI, baseline MMSE showed a significant association with the same two indices as for ADAS-Cog13. Thus, the PL-PX composite ($\beta = 0.0397$, $p = 1.28 \times 10^{-9}$) and the PBV composite ($\beta = 0.0379$, $p = 6.50 \times 10^{-9}$) were significantly and positively associated with MMSE. Since lower values of MMSE indicate greater degrees of cognitive impairment, the positive values of these coefficients indicate that lower values of PL-PX and PBV are associated with lower MMSE scores and thus greater cognitive impairment. After adjustment for all non-AD-

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4 specific medications, the above associations remained statistically significant (**Supplemental**
5 **Table 4**). Thus, the PL-PX ($\beta = 0.0343$, $p = 1.71 \times 10^{-6}$) and PBV composite ($\beta = 0.0318$, $p =$
6 8.44×10^{-6}) showed significant positive associations with baseline MMSE following adjustment
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8 for these medications.
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12 In the UPenn ADCC cohort, MMSE showed a trend towards an association with PL/PE (β
13 $= 1.123$, $p = 0.0949$; **Table 2B**). There were no other significant findings with MMSE.
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16 17 **3.3. CSF $A\beta_{1-42}$, Total tau, and Total tau/ $A\beta_{1-42}$ ratio**

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20 In cross-sectional analyses without adjustment for medications in the ADNI, CSF t-tau
21 showed significant negative correlations with the PL-PX ($\beta = -7.193$, $p = 5.55 \times 10^{-6}$) and PBV
22 composite ($\beta = -6.994$, $p = 7.77 \times 10^{-6}$) indices. Similarly, the \log_{10} CSF t-tau/ $A\beta_{1-42}$ showed
23 significant negative correlations with these same 2 indices (PL-PX, $\beta = -0.0422$, $p = 2.73 \times 10^{-6}$;
24 PBV, $\beta = -0.0408$, $p = 4.39 \times 10^{-6}$). CSF $A\beta_{1-42}$ did not show any significant relationships (**Table**
25 **3A**).
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31 After adjusting for all non-AD-specific medications in ADNI, the negative correlations of
32 CSF-t-tau with PL-PX ($\beta = -6.654$, $p = 1.13 \times 10^{-4}$) and with PBV ($\beta = -6.332$, $p = 1.89 \times 10^{-4}$),
33 and of the \log_{10} CSF t-tau/ $A\beta_{1-42}$ ratio with PL-PX ($\beta = -0.0396$, $p = 5.24 \times 10^{-5}$) and PBV ($\beta = -$
34 0.0374 , $p = 1.05 \times 10^{-4}$) remained statistically significant (**Supplemental Table 5**).
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38 In the UPenn ADCC cohort, the PL/PE index showed a significant negative relationship
39 with both CSF t-tau ($\beta = -12.231$, $p=0.0305$) and CSF t-tau/ $A\beta_{1-42}$ ($\beta = -0.0973$, $p = 0.0206$), but
40 not with CSF $A\beta_{1-42}$ (**Table 3B**). The PBV and PL-PX indices did not show statistically
41 significant relationships with any CSF measures (**Table 3B**).
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46 **3.4. Association of Plasmalogen Ratios and Indices with Variants in the**

47 ***FADS1/FADS2 Gene Cluster***

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50 A gene-based association analysis of the *FADS1/FADS2* gene cluster with plasmalogen
51 ratios and indices reported in this manuscript showed that *FADS2* was associated with the
52 PL226_224 ratio ($p=0.0348$; **Supplemental Figure 4**) in the ADNI-1/-GO/-2 cohort after
53 adjusting for multiple comparisons using permutation. A SNP (rs97384) in *FADS2* was most
54 significantly associated with PL226_224 ($p=0.0016$; **Supplemental Figure 4** – left figure lower
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4 left), where subjects with the minor allele (T) of SNP rs97384 had high levels of PL226_224
5 compared to subjects without the minor allele (central figure at the bottom of **Supplemental**
6 **Figure 4**). An expression quantitative trait locus (eQTL) analysis of SNP rs97384 with *FADS2*
7 gene expression in the temporal cortex from cognitively normal subjects showed that *FADS2*
8 was significantly highly expressed in subjects with the minor allele (T) of SNP rs97384 (lower
9 right figure in **Supplemental Figure 4**).
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18 **4. Discussion**

19 Age-related declines in circulating plasmalogens may increase the risk for AD by reducing
20 plasmalogen availability to the central nervous system [9, 10, 13, 14, 29, 39, 40]. Using a
21 focused lipidomics approach we measured four ethanolamine plasmalogens (PlsEtns) and four
22 closely-related phosphatidylethanolamines (PtdEtns) and derived indices from a subset of these
23 phosphoethanolamines reflecting PlsEtn and PtdEtn metabolism to explore the association of
24 these indices with baseline diagnosis, cognitive functioning, and CSF biomarkers of AD.
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30 In ADNI, without correcting for concomitant medications, we observed statistically
31 significant associations between AD diagnosis at the baseline visit and several serum indices of
32 ethanolamine plasmalogen biosynthesis, peroxisomal beta-oxidation, and phosphoethanolamine
33 remodeling. Most notably, baseline serum PL-PX and overall PBV were strongly associated with
34 baseline AD vs. CN and LMCI vs. CN, while baseline serum PL/PE did not show such an
35 association (**Table 2A**). The associations of the PL-PX and PBV indices with AD vs. LMCI
36 were not statistically significant; on the other hand, the PL/PE index showed statistically
37 significant relationships with AD vs. LMCI diagnosis (**Table 2A**).
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45 In the UPenn ADCC, we observed an association of AD vs. CN with PBV, replicating the
46 finding in ADNI, and also observed an association with the PL/PE index. The significant
47 association between AD vs. CN and PBV in both cohorts is consistent with findings from the
48 Religious Orders Study [9, 10, 13, 14, 29, 39, 40]/Memory and Aging Project cohort [10], and
49 provides further support for the utility of this measure as an index of plasmalogen biosynthesis
50 and remodeling that is applicable across cohorts, while individual analyte values may show more
51 variability among various subject populations.
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58 In ADNI, using baseline ADAS-Cog13 as a measure of global cognition, we also
59 observed statistically significant negative associations with the same two indices, PL-PX and
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4 overall PBV, as seen in comparisons of AD vs. CN and LMCI vs. CN, indicating an association
5 of reduced values of these plasmalogen metabolic indices with greater cognitive impairment.
6 However, we saw no evidence of an association between the baseline PL/PE index and baseline
7 ADAS-Cog13 (**Table 2A**).

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11 In the UPenn ADCC, we observed trends towards associations of impaired cognition as
12 measured by MMSE with PBV and PL/PE (**Table 2B**), although these did not reach statistical
13 significance. The lack of statistically significant effects in the UPenn cohort may be attributable
14 to its limited sample size (n=112 vs. 1547 for ADNI).

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18 In ADNI, we observed statistically significant negative associations between CSF t-tau
19 and of the CSF t-tau/A β_{1-42} ratio with PL-PX and PBV, i.e., the same indices associated with the
20 AD vs. CN and LMCI vs. CN and with the cognitive scores (**Table 3A**). On the other hand, we
21 did not see an association of CSF A β_{1-42} alone with any of these indices. The negative
22 association of CSF t-tau and its ratio with A β_{1-42} with these indices suggest that altered
23 plasmalogen biosynthesis and/or remodeling may associate more closely with the development
24 of tau than of amyloid pathology. Longitudinal studies including concomitant measurements of
25 plasmalogen indices, biomarkers of amyloid and tau pathology, and cognitive function are
26 needed to address this hypothesis.

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35 After adjustment for concomitant medication classes (other than anticholinesterases and
36 memantine) in ADNI, the association of the plasmalogen indices with the AD vs. CN was no
37 longer significant; however, those with LMCI vs. CN and AD vs. LMCI remained statistically
38 (**Supplemental Table 4**) In addition, the linear regression models showed persistence of
39 statistical significance of the associations of plasmalogen indices with cognition (**Supplemental**
40 **Table 4**). Likewise, the associations of CSF tau and the CSF t-tau/A β_{1-42} ratio with PL-PX and
41 PBV remained significant (**Supplemental Table 5**).

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48 These results are in broad agreement with previous studies of blood-based levels of
49 ethanolamine plasmalogens and related lipids in AD [9, 10, 13, 29-31], and suggest that reduced
50 indices of plasmalogen biosynthesis and/or remodeling of VL-PUFA-containing plasmalogens is
51 associated with an increased risk of AD. In addition, significant associations of PL-PX and PBV
52 with LMCI vs. CN indicates reduced plasmalogen indices occur at the MCI stage, and thus may
53 be associated with cognitive decline prior to the development of dementia. Finally, the
54 association of the PL/PE index, which is calculated from ratios of omega-3 fatty-acid-bearing
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4 plasmalogens to corresponding phosphatides, with AD vs. LMCI diagnosis may indicate that the
5 development of dementia is characterized in part by a specific compensatory failure of
6 peroxisomal mechanisms required to sustain plasmalogen biosynthesis in relation to that of
7 corresponding phosphatidylethanolamines which do not require peroxisomes for biosynthesis.
8 Increased remodeling by phospholipases [44], and degradation through lysoplasmalogenases [45]
9 and oxidative pathways [46] might contribute to relative reductions in plasmalogen levels. Fatty
10 acids made available by plasmalogen degradation might be directed towards PE synthesis [38].
11 These three phenomena, namely decreased biosynthesis, decreased remodeling/degradation, and
12 a putative increase in PE synthesis could contribute, individually or collectively, to reduction of
13 the PL/PE ratio. Since our observations were conducted at a single time point, we are unable to
14 distinguish between these possibilities. Studies examining the kinetics of plasmalogen
15 biosynthesis and metabolism in AD and related conditions are needed to better understand the
16 mechanisms underlying these findings. Attendant changes in signaling pathways, cholesterol
17 metabolism, and neurons might also contribute to pathogenesis of AD.

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19 Because genetic variance in the *FADS1/FADS2* gene cluster can affect the blood levels of
20 EPA/DHA containing lipids (cf. [43]), we conducted a candidate gene association analysis for
21 relationships between plasmalogen ratios and indices with genetic variants in this gene cluster in
22 the ADNI cohort. We found a significant gene-based association between *FADS2* and the
23 PL226_224 ratio ($p=0.0348$; **Supplemental Figure 4**) in the ADNI-1/-GO/-2 cohort after
24 adjusting for multiple comparisons using permutation. Five SNPs in this region were
25 significantly and independently associated with the PL226_224 ratio, most notably rs97384.
26 Individuals homozygous for the minor allele (T) had higher values of this ratio than those
27 without the minor allele ($p=0.0016$; see middle figure in **Supplemental Figure 4**); conversely,
28 an eQTL analysis showed that individuals homozygous for the major allele (C) had lower levels
29 of *FADS2* gene expression in the temporal cortex of cognitively normal subjects than those with
30 at least one minor allele ($p=1.8 \times 10^{-5}$, lower right figure in **Supplemental Figure 4**). These
31 findings suggest that *FADS2* genotype may influence the relative abundance of omega-3 (DHA)-
32 to omega-6 (adrenic acid)-containing plasmalogens both in circulating blood and in the brain.
33 Further studies are needed to assess the implications of this effect for AD, particularly with
34 regard to the balance of pro- and anti-inflammatory fatty acid-containing plasmalogens in brain.

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4 In contrast to most previous studies of plasmalogens and other phospholipids in AD, we
5 focused on examining indices calculated from specific, preselected ratios of plasmalogens, both
6 to each other and to comparable phosphatidylethanolamine species. The finding of significant
7 alterations of these ratios and indices in relation to cognition and diagnoses appears to be
8 informative in this setting. It will be important to determine if these ratios and indices are
9 similarly informative in other studies of AD and cognitive impairment conducted in other
10 populations and by different laboratories.
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19 **5. Conclusions**

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21 Further studies are needed to elucidate the specific mechanisms accounting for these
22 findings, including measurement of additional lipids representing upstream and downstream
23 pathways and enzyme activities (e.g., phospholipases and oxidative stress). In particular, the
24 steady-state measurements conducted here at a single time point do not allow discrimination
25 between decreased biosynthesis vs. increased degradation of plasmalogen species in AD.
26 Studies of the kinetics of plasmalogen biosynthesis may help distinguish between these
27 possibilities. Also, longitudinal measurement of these lipids will likely be informative regarding
28 the relative time courses of changes in serum lipids and of clinical, other biochemical, and
29 imaging outcomes in MCI and AD.
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38 These data further suggest the hypothesis that interventions that enhance plasmalogen
39 biosynthesis and availability to the central nervous system may mitigate cognitive decline and/or
40 brain atrophy in those at risk for AD. Future research will also address the potential relationships
41 of plasmalogen ratios and indices to genetic variation, neuroimaging measures, and other
42 characteristics of AD-type dementias.
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6. Data availability

Metabolomics datasets used in the current analyses for the ADNI-1 and ADNI-GO/-2 cohorts are available via the Accelerating Medicines Partnership-Alzheimer's Disease (AMP-AD) Knowledge Portal and can be accessed at <http://dx.doi.org/10.7303/syn5592519> (ADNI-1) and <http://dx.doi.org/10.7303/syn9705278> (ADNI-GO/-2). The full complement of clinical and demographic data for the ADNI cohorts are hosted on the LONI data sharing platform and can be requested at <http://adni.loni.usc.edu/data-samples/access-data/>.

The Alzheimer's Disease Metabolomics Consortium (ADMC): A complete listing of ADMC investigators can be found at: <https://sites.duke.edu/adnimetab/who-we-are/>

The Alzheimer's Disease Neuroimaging Initiative (ADNI): Data used in the preparation of this article were obtained from the ADNI database (<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 at: http://adni.loni.usc.edu/wp-content/uploads/how_to_apply/ADNI_Acknowledgement_List.pdf.

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59 **DISCLOSURES**

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5
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7
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25
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27
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29
30 sale of Avid to Eli Lilly as a co-inventor on imaging-related patents submitted by the University
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4 **Figure 1. Primary Lipid Pathways for Ethanolamine Plasmalogen and**
5 **Phosphatidylethanolamine Biosynthesis and Remodeling**
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10 *Legend.* The initial steps in plasmalogen biosynthesis, and the beta-oxidation of VL-PUFA, take
11 place within peroxisomes; later reactions occur in the endoplasmic reticulum. Fundamental lipid
12 species potentially available through dietary sources are metabolized in the endoplasmic
13 reticulum and are incorporated into plasmalogens in the peroxisome. The direction of alteration
14 in constituent lipid species and processes in AD based on the data reported here are indicated by
15 red arrows. Red hammerhead symbols indicate putative enzymatic steps proposed to be inhibited
16 in Alzheimer’s disease.
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Table 1A: Baseline demographic characteristics of the ADNI-1, -GO, -2 cohorts

Characteristic	N	Overall group (N=1545)	CN (n=364)	SMC (n=98)	EMCI (n=282)	LMCI (n=497)	AD (n=304)	p-value
Age yr (SD)	1545	73.67 (7.2)	74.63 (5.7)	72.24 (5.6)	71.14 (7.5)	74.09 (7.5)	74.65 (7.8)	<0.001
Male, % (#)	1545	54 (840)	48 (176)	42 (41)	55 (154)	61 (304)	54 (165)	<0.001
Education yr (SD)	1545	15.91 (2.8)	16.30 (2.77)	16.73 (2.5)	15.98 (2.6)	15.86 (2.9)	15.20 (3.0)	<0.001
APOE4 +, % (#)	1545	47 (726)	28 (103)	33 (32)	43 (120)	54 (270)	66 (201)	<0.001
ADAS-Cog13, (SD)	1534	16.82 (9.6)	9.18 (4.3)	8.82 (4.0)	12.57 (5.3)	18.70 (6.6)	29.74 (8.1)	<0.001
MMSE (SD)	1545	27.17 (2.6)	29.08 (1.1)	29.00 (1.2)	28.33 (1.5)	27.16 (1.8)	23.23 (2.0)	<0.001
CSF T-tau (SD)	1111	90.10 (53.5)	68.71 (22.3)	64.91 (31.3)	75.82 (47.3)	101.18 (55.8)	125.77 (59.9)	<0.001
CSF Abeta ₁₋₄₂ (SD)	1127	174.50 (53.8)	200.72 (51.6)	202.06 (48.5)	184.05 (50.9)	162.8 (51.4)	139.91 (39.5)	<0.001
T-tau/Abeta ₁₋₄₂	1111	0.62 (0.50)	0.39 (0.27)	0.35 (0.23)	0.39 (0.47)	0.72 (0.52)	0.97 (0.54)	<0.001
Log T-tau/Abeta ₁₋₄₂ (SD)	1111	-0.33 (0.33)	-0.49 (0.25)	-0.52 (0.24)	-0.43 (0.32)	-0.25 (0.32)	-0.08 (0.26)	<0.001

Table 1B: Baseline demographic characteristics of the University of Pennsylvania cohort

Characteristic	N	Overall group (N=112)	CN (n=51)	MCI (n=18)	AD (n=43)	p-value
Age yr (SD)	112	69.9 (9.2)	68.1 (9.8)	70.9 (5.9)	71.5 (9.3)	0.186
Male, % (#)	112	47 (53)	41 (21)	78 (14)	42 (18)	0.019
Education yr (SD)	112	16.0 (2.9)	16.3 (2.8)	15.7 (3.2)	15.7 (2.9)	>0.05
APOE4 +, % (#)	82	37.8 (31)	26.0 (13)	36.4 (4)	66.7 (14)	0.006
MMSE (SD)	109	27.72 (5.3)	29.16 (1.11)	27.35 (1.46)	20.98 (5.85)	<0.001
CSF T-tau (SD)	112	74.1 (51.2)	52.2 (19.0)	66.3 (45.7)	103.3 (64.6)	<0.001
CSF Abeta ₁₋₄₂ (SD)	112	232.9 (86.6)	281.3 (77.8)	224.0 (89.8)	179.2 (59.3)	<0.001
T-tau/Abeta ₁₋₄₂	112	0.39 (0.37)	0.20 (0.12)	0.39 (0.40)	0.62 (0.43)	<0.001
Log T-tau/Abeta ₁₋₄₂ (SD)	112	-0.54 (0.34)	-0.74 (0.22)	-0.57 (0.37)	-0.29 (0.28)	<0.001

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4 *Note for the UPenn Cohort: APOE genotype available on 50 CN, 11 MCI, 21 AD. MMSE
5 missing on 1 CN, 1 MCI, and 1 AD subject
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8 Abbreviations: AD: Alzheimer's Disease; ADAS-Cog13: Alzheimer's Disease Assessment
9 Scale, Cognitive Subscale; APOE: Apolipoprotein ϵ 4; CN: Cognitively Normal; CSF:
10 Cerebrospinal Fluid; EMCI: Early Mild Cognitive Impairment; MCI: Mild Cognitive
11 Impairment; MMSE: Mini-Mental State Exam; LMCI: Late MCI; SMC: Subjective Memory
12 Concerns; T-tau: Total tau
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Table 2A. Serum Plasmalogen/Phosphatide Indices in ADNI-1/GO/2 Cohort: Associations with Diagnosis and Cognition

Index/ratio	Logistic Regression Analyses									Linear Regression Analyses					
	AD vs. CN			LMCI vs. CN			AD vs. LMCI			ADAS-Cog13			MMSE		
	OR (95% CI)	p- value	q- value	OR (95% CI)	p- value	q- value	OR (95% CI)	p- value	q- value	β (95% CI)	p- value	q- value	β (95% CI)	p- value	q- value
PBV	0.775 (0.648,0.925)	0.005	0.030	0.745 (0.638,0.869)	1.99E-04	7.97E-04	0.999 (0.859,1.162)	0.994	0.994	-0.130 (-0.187,-0.074)	6.92E-06	2.77E-05	0.038 (0.025,0.051)	6.50E-09	2.60E-08
PL-PX	0.784 (0.656,0.935)	0.007	0.030	0.719 (0.615,0.838)	2.89E-05	1.73E-04	1.051 (0.904,1.222)	0.517	0.564	-0.135 (-0.192,-0.078)	3.24E-06	1.94E-05	0.040 (0.027,0.052)	1.28E-09	1.25E-08
PL/PE	0.894 (0.745,1.072)	0.228	0.304	1.141 (0.983,1.328)	0.085	0.145	0.757 (0.644,0.887)	6.15E-04	2.46E-03	0.0120 (-0.047,0.070)	0.696	0.759	-0.003 (-0.016,0.010)	0.646	0.887
PL205_224	0.787 (0.658,0.937)	0.008	0.030	0.843 (0.728,0.976)	0.023	0.056	0.901 (0.775,1.047)	0.176	0.235	-0.077 (-0.133,-0.020)	0.008	0.023	0.032 (0.020,0.045)	5.99E-07	1.80E-06
PL205_226	0.902 (0.754,1.078)	0.257	0.308	1.163 (1.000,1.355)	0.051	0.102	0.718 (0.609,0.844)	6.90E-05	4.14E-04	0.059 (0.002,0.117)	0.044	0.105	0.003 (-0.010,0.016)	0.677	0.887
PL226_224	0.804 (0.673,0.959)	0.015	0.046	0.684 (0.583,0.801)	2.70E-06	3.24E-05	1.133 (0.972,1.321)	0.110	0.165	-0.148 (-0.205,-0.092)	2.95E-07	3.54E-06	0.039 (0.026,0.052)	2.08E-09	1.25E-08
PL226_PE226	0.928 (0.774,1.111)	0.418	0.456	1.116 (0.961,1.299)	0.152	0.203	0.822 (0.702,0.961)	0.014	0.024	-0.002 (-0.060,0.056)	0.951	0.951	-0.005 (-0.018,0.008)	0.448	0.767
PL205_PE226	0.887 (0.740,1.061)	0.190	0.285	1.191 (1.025,1.386)	0.023	0.056	0.692 (0.586,0.815)	1.17E-05	1.41E-04	0.042 (-0.015,0.099)	0.150	0.225	-0.001 (-0.014,0.012)	0.891	0.934

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Table 2B. Serum Plasmalogen/Phosphatide Indices in U. Penn. Cohort: Associations with Diagnosis and Cognition

Index/Ratio	AD vs. CN		MMSE	
	OR (95% CI)	p-value	β (95% CI)	p-value
PBV	0.695 (0.039, 1.399)	0.0433	0.948 (-0.808, 2.705)	0.286
PL-PX	0.293 (-0.236, 0.846)	0.285	0.190 (-1.276, 1.655)	0.797
PL/PE	0.630 (0.121, 1.193)	0.02	1.123 (-0.195, 2.442)	0.094

P-values were estimated from logistic (diagnosis) or linear (cognitive function) regression
Models as described in Methods. All models were adjusted for age, sex, and APOE e4 status (note: analyses with UPenn cohort did not adjust for APOE due to number of subjects not genotyped). ADAS-Cog13 and MMSE models also adjusted for years of education. Above results were not adjusted for concomitant medications.
Abbreviations: AD-Alzheimer's Disease; ADAS-Cog13-Alzheimer Disease Assessment Scale 13-item cognitive subscale; CN-Cognitively normal; LMCI-Late Mild Cognitive Impairment; MMSE- Mini-Mental State Exam.

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Table 3A. Serum Plasmalogen/Phosphatide Indices in ADNI-1/GO/2 Cohort: Associations with CSF AD Biomarkers

Ratio/Index	CSF Total tau (Ttau)			CSF Aβ1-42			Log10(CSF Ttau/CSF Aβ1-42)		
	β (95% CI)	P value	Q value	β (95% CI)	P value	Q value	β (95% CI)	P value	Q value
PBV	-6.994 (-10.049,-3.94)	7.77E-06	3.11E-05	1.970 (-0.868,4.807)	0.173	0.694	-0.041 (-0.058,-0.023)	4.39E-06	1.76E-05
PL-PX	-7.193 (-10.285,-4.101)	5.55E-06	3.11E-05	2.048 (-0.826,4.922)	0.162	0.694	-0.042 (-0.06,-0.025)	2.73E-06	1.64E-05
PL/PE	-0.038 (-3.205,3.13)	0.981	0.981	0.518 (-2.391,3.428)	0.727	0.988	0.000 (-0.018,0.018)	0.9664	0.9664
PL205_224	-5.11 (-8.096,-2.123)	8.15E-04	2.44E-03	1.649 (-1.11,4.407)	0.241	0.724	-0.029 (-0.046,-0.012)	0.0009	0.0026
PL205_226	0.589 (-2.507,3.685)	0.709	0.945	0.137 (-2.711,2.984)	0.925	0.988	0.006 (-0.012,0.023)	0.5157	0.9664
PL226_224	-7.543 (-10.653,-4.432)	2.21E-06	2.65E-05	2.150 (-0.745,5.045)	0.145	0.694	-0.045 (-0.062,-0.027)	7.38E-07	8.86E-06
PL226_PE226	-0.238 (-3.409,2.932)	0.883	0.963	0.581 (-2.329,3.492)	0.695	0.988	-0.002 (-0.02,0.016)	0.8097	0.9664
PL205_PE226	0.302 (-2.81,3.414)	0.849	0.963	0.428 (-2.434,3.289)	0.769	0.988	0.003 (-0.015,0.021)	0.7322	0.9664

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Table 3B. Serum Plasmalogen/Phosphatide Indices in UPenn Cohort: Associations with CSF AD Biomarkers

Ratio/Index	CSF Total tau (Ttau)		CSF A β 1-42		log10(CSF Ttau/A β 1-42)	
	β (95% CI)	P value	β (95% CI)	P value	β (95% CI)	P value
PBV	-6.444 (-19.997,7.110)	0.348	-4.118 (-27.553,19.318)	0.728	-0.011 (-0.103,0.080)	0.806
PL-PX	0.836 (-10.256,11.927)	0.882	-9.159 (-28.192,9.873)	0.342	0.033 (-0.041,0.107)	0.383
PL/PE	-12.040 (-22.932,-1.148)	0.031	6.949 (-12.185,26.083)	0.473	-0.068 (-0.141,0.005)	0.068

P-values were estimated from linear regression models and adjusted for age and sex. Results from 3A were further adjusted for APOE ϵ 4 carrier status. Due to small number of subjects with genotyping in UPenn cohort, results presented in Table 3B are not adjusted for APOE ϵ 4. Above results were not adjusted for concomitant medications. Three indices were calculated to assess the following: a) **PL-PX**: index of biosynthesis and/or remodeling of ethanolamine plasmalogens using mean of PL226/PL224, PL205/PL224, and PL205/PL226; b) **PL/PE**: index of ratio of omega-3 fatty acid-containing plasmalogens to the DHA-containing phosphatide PtdEtn 16:0/22:6 using mean of PL226/PE226 and PL205/PE226; and c) **PBV**: overall composite index of plasmalogen biosynthesis and/or remodeling using a mean of all five ratios contributing to index (a) and index (b).
Abbreviations: CSF: cerebrospinal fluid; Ttau: total-Tau; A β 1-42: amyloid beta peptide, residues 1-42.

Figure 1

