

# Amyloid- $\beta_{42}$ signals tau hyperphosphorylation and compromises neuronal viability by disrupting alkylacylglycerophosphocholine metabolism

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**Perturbation of lipid second messenger networks is associated with the impairment of synaptic function in Alzheimer disease. Underlying molecular mechanisms are unclear. Here, we used an unbiased lipidomic approach to profile alkylacylglycerophosphocholine second messengers in diseased tissue. We found that specific isoforms defined by a palmitic acid (16:0) at the *sn*-1 position, namely 1-O-hexadecyl-2-acetyl-*sn*-glycero-3-phosphocholine (C16:0 PAF) and 1-O-hexadecyl-*sn*-glycero-3-phosphocholine (C16:0 *lyso*-PAF), were elevated in the temporal cortex of Alzheimer disease patients, transgenic mice expressing human familial disease-mutant amyloid precursor protein, and human neurons directly exposed to amyloid- $\beta_{42}$  oligomers. Acute intraneuronal accumulation of C16:0 PAF but not C16:0 *lyso*-PAF initiated cyclin-dependent kinase 5-mediated hyperphosphorylation of tau on Alzheimer disease-specific epitopes. Chronic elevation caused a caspase 2 and 3/7-dependent cascade resulting in neuronal death. Pharmacological inhibition of C16:0 PAF signaling, or molecular strategies increasing hydrolysis of C16:0 PAF to C16:0 *lyso*-PAF, protected human neurons from amyloid- $\beta_{42}$  toxicity. Together, these data provide mechanistic insight into how disruptions in lipid metabolism can determine neuronal response to accumulating oligomeric amyloid- $\beta_{42}$ .**

Alzheimer disease | glycerophosphocholine | lipidomics |

The aberrant processing of the amyloid precursor protein to different assemblies of amyloid  $\beta$  ( $A\beta$ ) peptides ranging from 37 to 42 amino acids is an early and necessary prerequisite for the development of Alzheimer disease (AD) (1). The “amyloid cascade hypothesis” defines generation of these smaller, toxic  $A\beta$  fragments, specifically soluble  $A\beta_{42}$  oligomers, as the root cause of AD (1). The severity of AD progression, however, is highly correlated with the rate of abnormal tau processing (2). Underlying molecular mechanisms linking  $A\beta_{42}$  biogenesis to the aggregation of normally soluble tau proteins into hyperphosphorylated oligomers remain elusive.

$A\beta_{42}$  can activate cytosolic phospholipase  $A_2$  (cPLA<sub>2</sub>) (3, 4), a Group IVa PLA<sub>2</sub> that preferentially hydrolyzes arachidonic acid from the *sn*-2 position of 1-*O*-alkyl-2-arachidonoyl- and 1-*O*-acyl-2-arachidonoyl-glycerophospholipids (5). Inhibiting cPLA<sub>2</sub> activation completely attenuates  $A\beta_{42}$  neurotoxicity; blocking the different metabolic arms of the arachidonic acid cascade confers only partial protection (3, 4, 6). Little is known about the fate of the glycerophospholipid backbone following the release of arachidonic acid by cPLA<sub>2</sub>, although accumulation of choline-containing lipids is associated with accelerated cognitive decline in AD (7, 8). The alkyl-*lyso*-glycerophosphocholines and *lysophosphatidylcholines* (LPCs) are of particular interest (Fig. S1). These metabolites are biologically active in their own right and can be further modified by *lysophosphatidylcholine* acyltransferases (LPCATs). LPCAT activity also increases in AD (9), notably in the posterior-temporal

entorhinal cortex, a region characterized by the earliest tau pathology (2). Transfer of a long-chain acyl group to the *sn*-2 position by LPCAT1 and LPCAT2 regenerates structural membrane lipids, whereas addition of a small acetyl group produces a family of powerful lipid second messengers known as platelet activating factors (PAFs) (10, 11).

Here, we used an unbiased lipidomics approach to identify metabolic disruptions in alkylacylglycerophosphocholine second messengers in the posterior-entorhinal cortex of individuals with AD, TgCRND8 transgenic mice, as well as human neurons directly exposed to soluble  $A\beta_{42}$  oligomers. We found that  $A\beta_{42}$  triggers a selective destabilization in Land's cycle metabolites defined by a palmitic acid (16:0) at the *sn*-1 position. The acute accumulation of 1-O-hexadecyl-2-acetyl-*sn*-glycero-3-phosphocholine (C16:0 PAF), but not its immediate precursor and metabolite 1-O-hexadecyl-*sn*-glycero-3-phosphocholine (C16:0 *lyso*-PAF), signals the phosphorylation of tau at AD-specific epitopes. Chronic elevation activates an endoplasmic reticulum (ER)-associated calpain and caspase cascade compromising neuronal viability. Strategies that either promote the hydrolysis of C16:0 PAF to C16:0 *lyso*-PAF or inhibit downstream signal transduction pathways protect neurons from  $A\beta_{42}$  toxicity and prevent aberrant tau processing.

## Results

**Alkylacylglycerophosphocholine Metabolism Is Disrupted in AD.** Glycerophospholipids were extracted postmortem from the posterior/entorhinal cortex of AD patients and control subjects (Fig. 1A). PAF isoforms were profiled by high-performance liquid chromatography electrospray ionization mass spectrometry (LC-ESI-MS) (12). Lipids with mass-to-charge ratios (*m/z*) of 450–600 were analyzed in positive ion mode by MS scan for a protonated molecule at expected *m/z* followed by precursor ion scan for a diagnostic phosphocholine product ion at *m/z* 184. Peak intensities were standardized against C13:0 LPC, a synthetic internal standard added at the time of lipid extraction. Nineteen alkylacylglycerophosphocholine species were identified (Figs. S2 and S3). Three of these species were significantly elevated in AD cortex: 1-O-hexadecyl-2-acetyl-*sn*-glycero-3-phosphocholine (C16:0 PAF), its immediate metabolite/precursor 1-O-hexadecyl-*sn*-glycero-3-phosphocholine (C16:0 *lyso*-PAF), and 1-O-oleyl-2-*lyso*-*sn*-glycero-

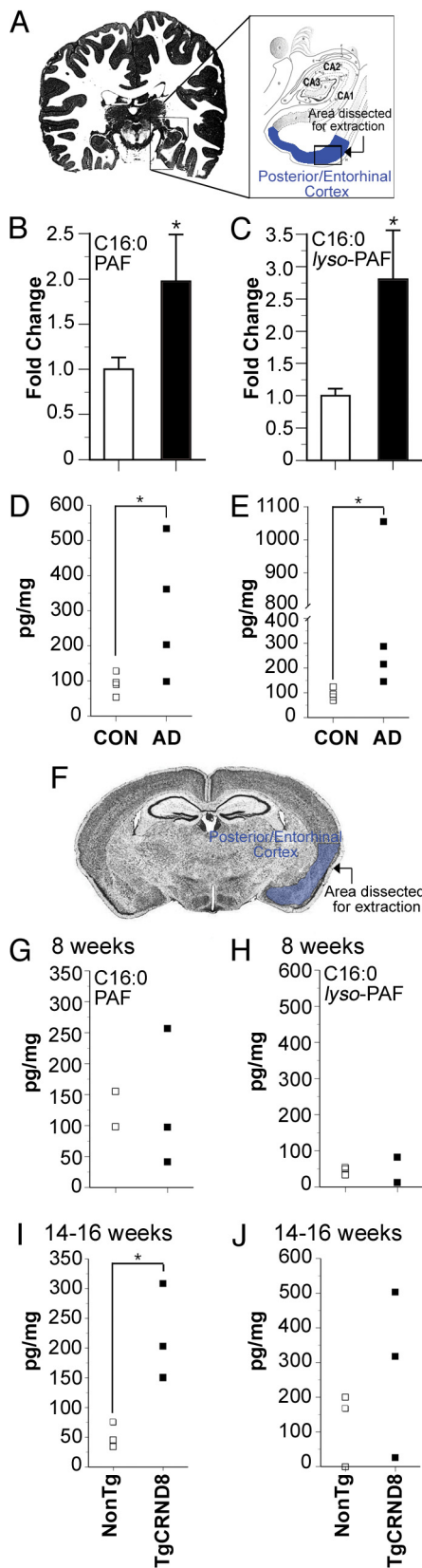
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**Fig. 1.** C16:0-PAF and its immediate precursor and metabolite C16:0-lyso-PAF accumulate in posterior temporal/entorhinal AD cortex and TgCRND8 mice. (A) Alkylacylglycerophosphocholines in postmortem human posterior-entorhinal cortex were profiled by LC-ESI-MS (Figs. S2 and S3). (B) C16:0 PAF and (C) C16:0 lyso-PAF levels were elevated in AD tissue ( $n = 4$  individuals/condition). Data are expressed as fold change relative to controls  $\pm$  SEM.

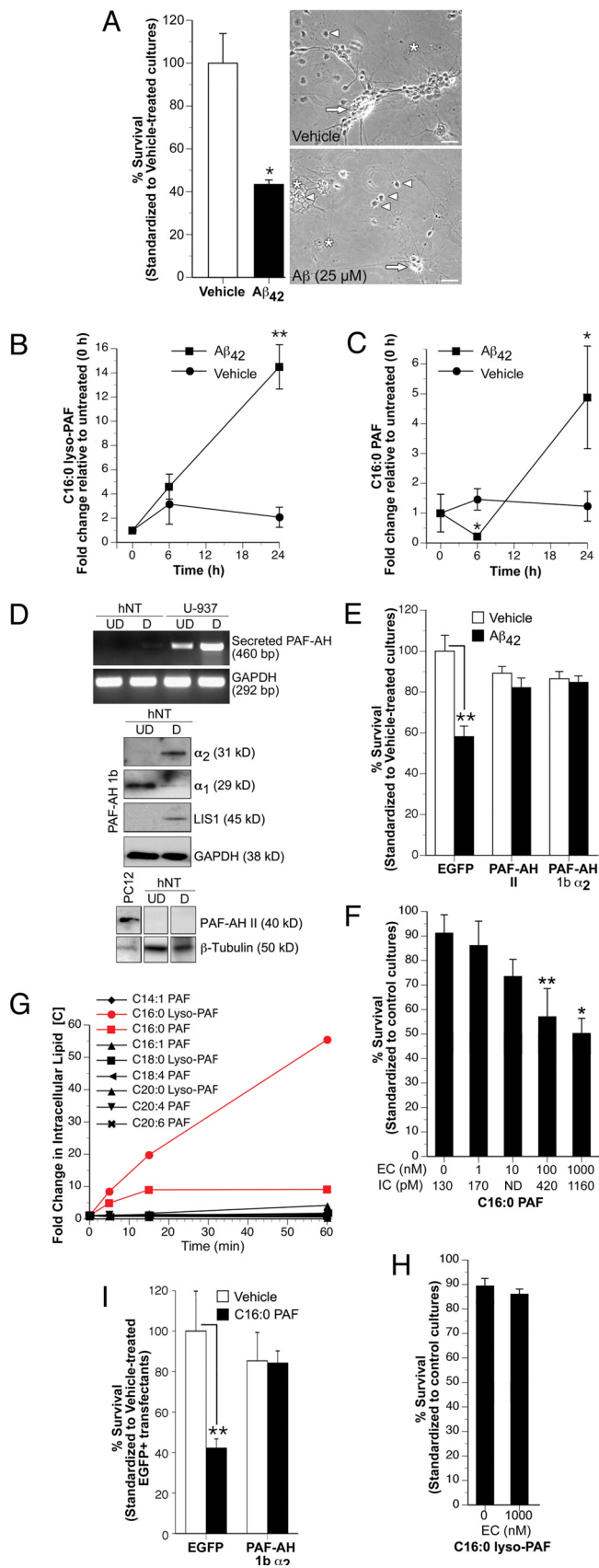
3-phosphocholine (C18:1 lyso-PAF) (Fig. 1 B and C and Figs. S2 and S3). As elevations in both C16:0 PAF and C16:0 lyso-PAF suggested a specific disruption in the remodeling of palmitic acid containing-PAFs, we used deuterated standards to quantify tissue concentrations. Clear separation was obtained between C16:0 PAF and isobaric C18:0 LPC with the isoform altered in AD co-eluting with  $d_4$ -C16:0 PAF (Fig. S4). Tissue concentrations, expressed as pg/mg in Fig. 1 D and E, represent approximate molar concentrations of  $258 \pm 30$  pM increased to  $639 \pm 205$  pM for C16:0 PAF and  $316 \pm 25$  pM increased to  $1035 \pm 455$  pM for C16:0 lyso-PAF in control and AD cortex, respectively.

**Disruption of C16:0 PAF Metabolism Is Detected in Symptomatic TgCRND8 Mice.** To establish the kinetics of palmitic-PAF metabolic disruption, we analyzed the posterior temporal/entorhinal cortex of TgCRND8 transgenic mice (Fig. 1F). TgCRND8 mice express a double mutant form (KM670/671NL+V717F) of the human amyloid precursor protein gene under the control of the prion protein promoter. Increases in cortical  $A\beta_{42}/A\beta_{40}$  ratios are initially observed at 10–12 weeks of age coincident with learning and memory impairment (13). We found that cortical levels of C16:0 PAF and C16:0 lyso-PAF of TgCRND8 mice were comparable to nontransgenic (NonTg) littermates at 8 weeks of age (Fig. 1 G and H), yet were 4-fold and 2-fold higher, respectively, by 14–16 weeks of age (Fig. 1 I and J). Tissue concentrations, expressed as pg/mg in Fig. 1 I and J, represent approximate molar concentrations of  $104 \pm 25$  pM C16:0 PAF in NonTg increased to  $407 \pm 105$  pM in TgCRND8 and  $268 \pm 135$  pM C16:0 lyso-PAF in NonTg increased to  $597 \pm 320$  pM in 14- to 16-week-old TgCRND8 mice.

**Intracellular Accumulation of C16:0 PAF Signals  $A\beta_{42}$  Neurotoxicity.** To determine whether  $A\beta_{42}$  directly disrupts C16:0 PAF metabolism in human neurons, hNT neurons were treated with  $A\beta_{42}$  (25  $\mu$ M) prepared as soluble oligomers (14). A 24-h exposure signaled neuronal loss (Fig. 2A). Over the course of treatment, neuronal C16:0 lyso-PAF concentrations increased steadily rising to 15-fold that of control cultures ( $12 \pm 0.1$  pM to  $180 \pm 8$  pM) (Fig. 2B). Endogenous neuronal C16:0 PAF levels initially dropped from approximately  $127 \pm 27$  pM to  $28 \pm 4$  pM over the first 6 h of treatment and then increased to 5-fold that of control levels within 24 h ( $619 \pm 115$  pM) (Fig. 2C). These data are consistent with a specific increase in Land's cycle remodeling of palmitic acid-PAFs in response to oligomeric  $A\beta_{42}$ . Also intriguing was the transient decrease in endogenous C16:0 PAF levels suggesting a compensatory increase in intracellular catabolism following acute  $A\beta_{42}$  exposure.

To provide mechanistic insight, we sought to modulate the  $A\beta_{42}$ -triggered increase in C16:0 PAF metabolism with minimal perturbation of other lipid networks. Because genetic, molecular, and pharmacological strategies targeting cPLA<sub>2</sub>, LPCAT1, or LPCAT2 will also impinge upon the arachidonic acid cascade, phosphatidylcholine and lysophosphatidylglycerol remodeling, as well as structural membrane lipid composition, we used a gain-of-function approach to promote hydrolysis of C16:0 PAF to C16:0 lyso-PAF. PAF-AHs cleave the *sn*-2 acetyl chain of PAF alkylacylglycerophosphocholines generating alkyl-lyso-glycerophosphocholines (lyso-PAFs) (Fig. S1). There are three PAF-AH isoforms. The 45-kDa secreted PAF-AH isoform is not expressed by hNT neurons

Tissue concentrations, expressed as pg/mg tissue wet weight, were calculated for (D) C16:0 PAF and (E) C16:0 lyso-PAF in comparison to deuterated samples spiked at the time of analysis (Fig. S4). Each square represents an individual patient (\* indicates  $P < 0.05$ , Student's *t* test). (F) Dissection coordinates (blue) of regions analyzed by LC-ESI-MS in TgCRND8 and NonTg control mice. Quantification of C16:0 PAF and C16:0 lyso-PAF species was performed at 8 weeks of age in panels G and H and at 14–16 weeks of age in panels I and J by standard addition ( $n = 15$  measurements/sample, \*,  $P < 0.05$ , Student's *t* test,  $n = 3$  mice/condition).



**Fig. 2.** C16:0 PAF accumulates in human neurons treated with soluble A $\beta_{42}$  oligomers compromising neuronal viability (A) A 24-h exposure to soluble A $\beta_{42}$  oligomers (25  $\mu$ M) elicited neuronal death as assessed by TUNEL (Left, \*,  $P < 0.05$ , Student's  $t$  test,  $n = 6$ –10 independent cultures/condition). The

(Fig. 2D). Cytosolic PAF-AH II is also not expressed by hNT neurons (Fig. 2D). Cytosolic PAF-AH Ib is an acetylhydrolase-specific complex composed of two 29-kDa  $\alpha_1$  and/or 30-kDa  $\alpha_2$  catalytic subunits and two 45-kDa regulatory LIS1 subunits (15). We have previously shown that overexpression of the  $\alpha_2$  catalytic subunit enhances PAF hydrolysis in cells that endogenously express all three PAF-AH Ib subunits (16). As observed over the course of CNS development (17), the  $\alpha_1$  catalytic subunit was detected at higher levels before differentiation, while the  $\alpha_2$  catalytic and LIS1 regulatory subunits were maximally expressed following terminal differentiation of hNT neurons (Fig. 2D). Based on this profile, neurons were transfected with EGFP and empty vector, EGFP and PAF-AH II, or EGFP and PAF-AH I  $\alpha_2$ . Neurons ectopically expressing PAF-AH II or overexpressing the PAF-AH I  $\alpha_2$  catalytic subunit were protected from A $\beta_{42}$  (Fig. 2E).

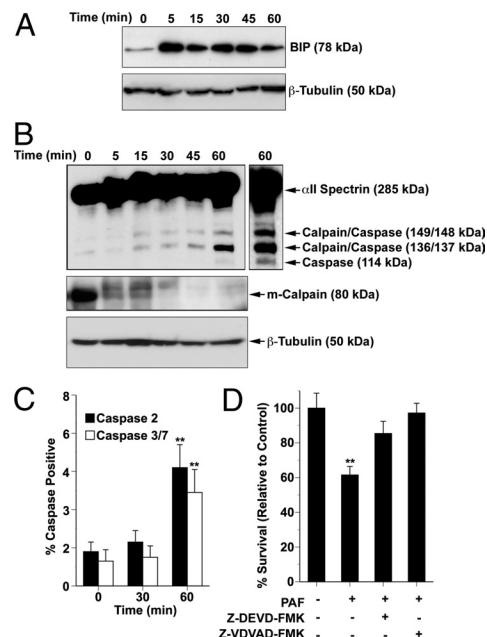
These findings implicated C16:0 PAF and not C16:0 lyso-PAF in A $\beta_{42}$  toxicity. To test this directly, we used a biochemical loading approach to elevate intraneuronal C16:0 PAF concentrations in the absence of any other toxic stimuli. PAFs can be internalized following binding to PAFR and clatherin-mediated endocytosis, transglutaminase-dependent transmembrane bilayer movement, engagement with low affinity binding sites at the plasma membrane, or direct physicochemical effects at the plasma membrane (16, 18, 19). As hNT neurons do not express PAFR (Fig. S5a), we established the kinetics of PAFR-independent PAF lipid internalization using a Bodipy fluorophore-labeled isoform [1-(O-[11-(4,4-difluoro-5,7-dimethyl-4-bora-3a,4a-diaza-*s*-indacene-3-propionyl)amino]undecyl)-2-acetyl-*sn*-glycero-3-phosphocholine; C11:0 B-PAF]. A progressive increase in cell-associated C11:0 B-PAF was observed over the first 60 min of lipid addition to media (Fig. S6a). Thin layer chromatography (TLC) of lipid loosely associated with the neuronal plasma membrane (acid-labile fractions), intracellular lipid (acid-resistant fractions), and lipid present in extracellular

morphology of single hNT neurons (arrowhead) or aggregated in bundles (arrows) on non-neuronal cells (asterisk) is shown after 24-h treatment with vehicle (Top Right) or A $\beta_{42}$  (Bottom Right). (Scale bar, 50  $\mu$ m.) (B) Treatment with 25  $\mu$ M A $\beta_{42}$  oligomers increased intraneuronal C16:0 lyso-PAF levels. (C) A transient decrease in C16:0 PAF was detected within 6 h of treatment followed by a progressive increase (\*,  $P < 0.05$ , \*\*,  $P < 0.01$ , ANOVA, post-hoc Dunnett's  $t$  test). Data are expressed as fold change relative to untreated hNT control cultures grown in complete media  $\pm$  SEM ( $n = 4$  cultures/condition). (D) Undifferentiated (UD) and differentiated (D) hNT cultures do not express secreted/plasma PAF-AH mRNA (RT-PCR). Positive controls were human U-937 cells (UD) and 12-*O*-tetradecanoylphorbol-13-acetate-treated U-937 cells differentiated to a macrophage lineage (D). GAPDH was amplified from the same random-primed cDNA to confirm template integrity. All three PAF-AH I subunits were detected in both undifferentiated (UD) and differentiated (D) hNT cultures by Western analysis. Only optimal exposure times are presented to show differences in expression between UD and D cultures. GAPDH was used as a loading control (Lower). PAF-AH II protein was not detected in hNT cultures. The positive control (+) was PC12 cells.  $\beta$ -tubulin was used as a loading control (Lower). (E) Transfection of hNT neurons with either PAF-AH II or PAF-AH I  $\alpha_2$  and EGFP, but not empty vector, and EGFP-protected hNT neurons from a 24-h exposure to 25  $\mu$ M A $\beta_{42}$  (\*,  $P < 0.05$ , ANOVA, post-hoc Dunnett's  $t$  test). (F) Elevating intracellular C16:0 PAF concentrations to >400 pM/cell in the absence of any other toxic stimuli compromised neuronal viability as assessed by TUNEL 24 h after treatment. EC, extracellular, IC, intracellular concentration. ICs were determined by LC-ESI-MS. Viability was standardized to untreated-control cultures (\*,  $P < 0.05$ , \*\*,  $P < 0.01$ , ANOVA, post-hoc Dunnett's  $t$  test). (G) Intraneuronal alkylacylglycerophosphocholine metabolites were profiled by LC-ESI-MS following direct exposure to 1  $\mu$ M C16:0 PAF. No significant change in any other PAF species was observed in C16:0 PAF-treated hNT neurons beyond the expected elevation in intracellular C16:0 PAF and a progressive increase in C16:0 lyso-PAF. (H) C16:0 lyso-PAF did not compromise neuronal viability as assessed by TUNEL. (I) Increasing cytosolic PAF hydrolysis protected hNT neurons from 24 h C16:0 PAF challenge. Cultures were transfected with EGFP and empty vector or EGFP and PAF-AH I  $\alpha_2$  (\*\*,  $P < 0.01$ , ANOVA, post-hoc Dunnett's  $t$  test). All treatments were carried out in serum-free media containing 0.025% BSA. No degradation of exogenous PAF in media was detected under these conditions (Fig. S8a).

milieu revealed that cell-associated C11:0 B-PAF was rapidly internalized within 5 min (Fig. S6b). An increase in hydrolysis was observed within 15 min with the C11:0 B-*lyso*-PAF metabolite released into the extracellular milieu (Fig. S6b). This balance between lipid internalization and lipid hydrolysis maintained a constant elevated intracellular concentration of C11:0 B-PAF for at least 60 min (Fig. S6b). Conversely, when hNT neurons were loaded with *lyso*-PAF, C11:0 B-*lyso*-PAF was rapidly internalized but converted to C11:0 alkylacylglycerophosphocholine and not C11:0 B-PAF (Figs. S6c and S7). Preferential conversion of C11:0 *lyso*-PAF to C11:0 B-PAF was only observed when neurons were treated in the presence of A $\beta$ <sub>42</sub> (Fig. S7).

We exploited this loading approach to elevate acutely C16:0 PAF and C16:0 *lyso*-PAF levels in hNT neurons without directly impacting on other glycerophosphocholine second messengers. Intracellular lipid concentrations were determined by LC-ESI-MS and neuronal viability was assessed by TUNEL after a 24-h exposure. Significant neuronal death was only observed when intracellular C16:0 PAF levels reached molar concentrations comparable to those detected in postmortem AD tissue, 14- to 16-week-old TgCRND8 mice, and A $\beta$ <sub>42</sub>-treated hNT neurons (i.e., >400 pM/cell) (Fig. 2F). To establish specificity, fold changes in other changes in other endogenous alkylacylglycerophosphocholine concentrations were profiled and quantified by LC-ESI-MS. Nine PAF species were identified in human neurons. No significant change in any other isoform was observed beyond the expected sustained elevation in the target lipid and a progressive increase in C16:0 *lyso*-PAF likely the result of compensatory hydrolysis (Fig. 2G). Only direct 24-h application of C16:0 PAF (Fig. 2F) not C16:0 *lyso*-PAF (Fig. 2H) signaled neuronal death. To demonstrate that neurotoxicity was not the result of nonspecific lytic effects of extracellular PAF, plasma membrane integrity was assessed using ethidium bromide homodimer (EthD-1), a membrane impermeant dye. Direct application of C16:0 PAF to hNTs did not increase EthD-1 uptake over a 24-h treatment period (Fig. S8b). Loss in membrane integrity was only observed using methyl-carbamyl PAF (mc-PAF), a PAF analog resistant to hydrolysis, applied to hNT neurons at concentrations that exceeded critical micelle concentration (2.5–5  $\mu$ M) (Fig. S8c). These data provided converging evidence that intraneuronal accumulation of C16:0 PAF signals a loss of viability in human neurons. To confirm these data, hNT neurons were transfected with EGFP and empty vector or EGFP and PAF-AH I  $\alpha$ <sub>2</sub> and treated with 1  $\mu$ M C16:0 PAF. As observed in A $\beta$ <sub>42</sub>-treated cultures (Fig. 2E), enhancing cytosolic hydrolysis of C16:0 PAF to C16:0 *lyso*-PAF prevented neuronal loss (Fig. 2J).

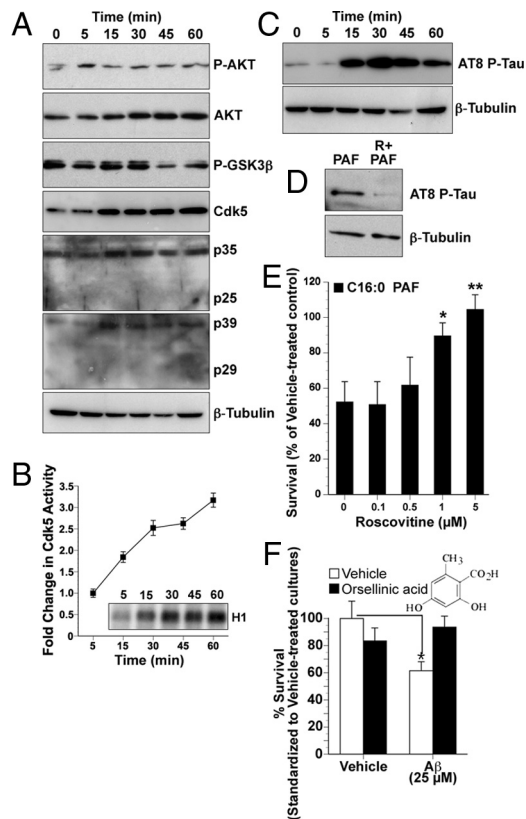
**Chronic C16:0 PAF Accumulation Induces an ER Stress-Dependent Calpain/Caspase Cascade.** To establish how intraneuronal C16:0 PAF accumulation signals A $\beta$ <sub>42</sub> toxicity, we first assessed calcium influx. A 180-s exposure to C16:0 PAF or C16:0 *lyso*-PAF did not acutely elevate intraneuronal calcium levels in the absence of PAFR as would be expected by a receptor-mediated event (Fig. S5b). We next assessed ER stress responses. In murine neurons, A $\beta$ <sub>42</sub> activates both calpains and ER-associated caspase-12 (20). The human homolog of caspase-12 has a frameshift mutation resulting in a premature stop codon and an amino acid substitution within a critical site for caspase enzymatic activity, rendering it catalytically inactive (21). Caspase 2 is a candidate human ER caspase, while caspase 7 translocates to the ER following apoptotic challenge (22). To test whether rising C16:0 PAF levels impair ER function, we assessed changes in BIP (GRP78) expression. BIP increased rapidly in hNTs in response to C16:0 PAF loading (Fig. 3A). ER stress-associated calpain/caspase activation was next assessed by examining spectrin cleavage. Spectrin is a scaffold protein that anchors the plasma membrane to the cytoskeleton. The kinetics of cleavage can be used to track sequential calpain/caspase activation. Proteolytic processing by calpains ( $\mu$  and  $m$ ) yields 149- and 136-kDa fragments, whereas caspase 2, 3, and 7 cleavage yields 148- and 137-kDa



**Fig. 3.** Intracellular accumulation of C16:0 PAF elicits ER stress and activates *m*-calpain and caspases 2 and 3/7. (A) Representative immunoblots of three independent experiments showing induction of BIP within 5 min of hNT exposure to 1  $\mu$ M C16:0 PAF. (B) The same hNT protein lysates were immunoblotted for  $\alpha$ II-spectrin and *m*-calpain using  $\beta$ -tubulin as a loading control. Cleavage fragments generated by calpains (149, 136 kDa) and caspases (148, 137, 114 kDa) are indicated. (C) hNT neurons were treated with 1  $\mu$ M PAF for 0, 30, and 60 min, and caspase activity was assessed using FAM-VDVAD-FMK (caspase 2) or FAM-DEVD-FMK (caspase 3/7). Data represent the percentage of cells positive for activated caspases (\*\*,  $P < 0.01$ , ANOVA, post-hoc Dunnett's *t* test). (D) Functional dependence of C16:0 PAF-induced neuronal apoptosis on caspase activation was measured by LIVE/DEAD assay. Inhibiting caspase activation protected hNT neurons from elevated intraneuronal accumulation of C16:0 PAF (\*,  $P < 0.05$ , ANOVA, post-hoc Dunnett's *t* test).

fragments. Both the calpain/caspase-generated 148/149-kDa fragments can be further processed to 114-kDa and 34-kDa products by caspases 3/7 but not caspase 2 (23). We observed an increase in spectrin cleavage, generating both 148/149-kDa and 136/137-kDa fragments, 15 min after C16:0 PAF application (Fig. 3B). A further increase in the levels of the 136/137-kDa fragment was observed within 60-min coincident with the appearance of the caspase 3/7-specific product (Fig. 3B). To confirm these data, calpain and caspase activation were assessed directly. Activation of *m*-calpain is associated with cleavage of the 80-kDa calpain pro-protein (24). Using an antibody that detects the inactive pro-protein but not the cleaved product, a marked decrease in pro-*m*-calpain levels was detected 5–45 min after C16:0 PAF challenge consistent with activation (Fig. 3B). By direct enzyme assay, significant caspase 2 and caspase 3/7 activation was only observed at 60 min, following calpain activation (Fig. 3C). To establish whether this calpain-caspase signaling pathway was sufficient to compromise neuronal viability, hNT neurons were treated with C16:0 PAF in the presence of the caspase 2 inhibitor Z-VDVAD-FMK or the caspase 3/7 inhibitor Z-DEVD-FMK. Inhibiting either caspase 2 or caspase 3/7 protected hNTs from C16:0 PAF neurotoxicity (Fig. 3D).

**Acute Intracellular Accumulation of C16:0 PAF Activates Tau Kinases and Signals Tau Phosphorylation at AD-Specific Phosphoepitopes.** Calpains can activate “tau kinases,” including glycogen synthase kinase 3 $\beta$  (GSK3 $\beta$ ) and cyclin-dependent kinase 5 (Cdk5) (25, 26). Alkylacylglycerophosphocholine second messengers have already been shown to stimulate GSK3 $\beta$  activity (27). Their impact on Cdk5 or tau has not been examined. Here, we assessed indicators of both



**Fig. 4.**  $A\beta_{42}$ -induced accumulation of C16:0 PAF induces Cdk5 and GSK3 $\beta$  activity leading to increased phosphorylation of tau on AD-associated epitopes with associated neurotoxicity inhibited by orsellinic acid. (A) Protein (15  $\mu$ g) from hNT cells treated with 1  $\mu$ M C16:0 PAF for 0 (control), 5, 30, 45, or 60 min were immunoblotted to assess kinase expression and posttranslational modifications of effector proteins. Increases in protein expression of Cdk5 and its activators p35 and p39 were detected within 15 min. A decrease in phospho-GSK3 $\beta$ , associated with activation, was observed by 45 min. A transient increase in phosphorylation of AKT at Thr-308 was observed within 5 min. (B) A progressive increase in Cdk5 activity in hNTs following treatment with 1  $\mu$ M C16:0 PAF was directly determined by *in vitro* assay of  $P^{32}$ -labeled histone 1 by immunoprecipitated Cdk5 protein (*inset*). Quantification was performed by densitometry. (C) Phosphorylation of tau on AD-associated epitopes was observed by Western analysis following C16:0 PAF treatment using the AT8-phospho-specific antibody. (D) Inhibition of Cdk5 with 5  $\mu$ M roscovitine prevented C16:0 PAF-mediated AT8-reactive tau phosphorylation.  $\beta$ -tubulin immunoblots were used as loading controls. Data are representative of replicate experiments. (E) Roscovitine protected hNT neurons from a 24-h treatment with 1  $\mu$ M C16:0 PAF as assessed by LIVE/DEAD assay ( $n = 15$  wells per culture conducted in replicate experiments, \*\*,  $P < 0.01$ , \*,  $P < 0.05$ , ANOVA, post-hoc Dunnett's *t* test). (F) The C16:0 PAF inhibitor orsellinic acid (5  $\mu$ M) protected hNT cultures from a 24-h treatment with 25  $\mu$ M  $A\beta_{42}$  ( $n = 20$  coverslips per culture conducted in replicate experiments, ANOVA, post-hoc Tukey test, \*\*,  $P < 0.01$  vs. vehicle control).

tau kinase activation and inhibition in hNTs loaded with C16:0 PAF. A transient increase in phosphorylation of the GSK3 $\beta$  regulator AKT at Thr-308, commonly associated with the inhibition of GSK3 $\beta$  activity, was observed after 5 min (Fig. 4A). These changes were followed by an activity-associated decrease in GSK3 $\beta$  phosphorylation at Ser-9 after 45 min (Fig. 4A). These data indicate that GSK3 $\beta$  is not activated within the first 45 min of intracellular C16:0 PAF accumulation. More rapid increases in the levels of Cdk5 and its activators p35 and p39 were observed within 15 min of C16:0 PAF application (Fig. 4A). Despite apparent *m*-calpain activation (Fig. 3B), cleavage of p35 and p39 to the more potent p25 and p29 Cdk5 activators was not detected (Fig. 4A). This p25/p29-independent induction of Cdk5 activity, as a result of acute C16:0 PAF accumulation, was validated by assessing transfer of radiola-

beled  $P^{32}$  to histone-1 following immunoprecipitation of Cdk5 (Fig. 4B).

To evaluate AD-associated tau phosphorylation, we examined two of the main sites aberrantly phosphorylated in AD brain (Ser-202 and Thr-205) (28). AT8-reactive sites were phosphorylated within 15 min of C16:0 PAF exposure (Fig. 4C). To establish whether phosphorylation was mediated by Cdk5, cultures were treated with a general Cdk5/p35 inhibitor roscovitine. Roscovitine prevented the C16:0 PAF-dependent phosphorylation of tau on AD-specific epitopes (Fig. 4D) and protected hNT neurons from C16:0 PAF toxicity (Fig. 4E).

**Pharmacological Inhibition of Intracellular C16:0 PAF Signaling Protects Human Neurons from  $A\beta_{42}$  Oligomers.** Taken together, these data provide strong evidence that a pathogenic accumulation of C16:0 PAF in response to oligomeric  $A\beta_{42}$  participates in signaling the hyperphosphorylation of tau and, ultimately, causes caspase-dependent neuronal death. However, pharmacological strategies targeting C16:0 PAF must also consider the amnesiac impact of PAFR inhibition on learning and memory (29, 30). PAFR is expressed by subpopulations of neurons, notably in the hippocampus (30). Thus, strategies to inhibit C16:0 PAF signaling must be able to pharmacologically distinguish between PAFR-dependent (neuroprotective/functional) and PAFR-independent (neurotoxic) PAF pathways. Orsellinic acid is a fungal-derived benzoic acid that inhibits C16:0 PAF-induced ER caspase activation without affecting PAFR-neuroprotective signal transduction (31). As with gain-of-function approaches to promote C16:0 PAF hydrolysis and restore physiological intracellular concentrations (Fig. 2 E and I), inhibition of intracellular PAF signal transduction with orsellinic acid protected hNT neurons from  $A\beta_{42}$  toxicity (Fig. 4F). As a control, we treated cells with the PAFR antagonist (and structural PAF analog) CV3988. CV3988 did not impact on C16:0-mediated tau phosphorylation in PAFR-negative human neurons or  $A\beta_{42}$  toxicity (Fig. S5d).

## Discussion

In this study, we elucidate a pathway of neuronal dysfunction triggered by aberrant alkylacylglycerophosphocholine metabolism in response to oligomeric  $A\beta_{42}$ . Our unbiased lipidomics approach revealed a selective disruption of C16:0 PAF and C16:0 *lyso*-PAF metabolism in the posterior/entorhinal cortex of AD patients. We found that this disruption occurred following pathogenic increases in cortical  $A\beta_{42}/A\beta_{40}$  ratios (13) in TgCRND8 transgenic mice and could be elicited directly in human neurons by exposure to soluble  $A\beta_{42}$  oligomers. Acute elevations of C16:0 PAF signaled tau phosphorylation on AD-specific phospho-epitopes. Chronic elevations initiated calpain and caspase-dependent neuronal death. Through gain-of-function and pharmacological inhibitory approaches, we found that human neurons can be protected from  $A\beta_{42}$  either by enhancing hydrolysis of C16:0 PAF to C16:0 *lyso*-PAF or inhibiting C16:0 PAF-dependent intracellular signal transduction.

Although consistent with the upregulation in cPLA $_2$  and LPCAT activities observed in AD and in response to  $A\beta_{42}$  (3, 4, 9), the specificity of this metabolic disruption was nonetheless surprising. Disruption in the Land's cycle predicts a general increase in PAF metabolism not the hydrolysis of structural lipids with specific carbon chain lengths (Fig. S1). It may be that the accumulation of hexadecyl-PAFs indicates a shift in substrate availability in diseased tissue consistent with a larger metabolic disruption in lipid metabolism. The loss of apolipoprotein E, for example, a risk factor associated with AD, alters the traffic of polyunsaturated lipids from astrocytes to neurons, biasing the composition of long-chain fatty acids in synaptosomes phosphocholines toward hexadecyl (16:0) species in null-mutant mice (32). Furthermore, the calcium-independent PLA $_2\gamma$  (iPLA $_2\gamma$ ) isoform, capable of hydrolyzing both *sn*-1 and *sn*-2 unsaturated fatty acid chains, shows specificity for

palmitic acid-containing carbon chains. iPLA<sub>2</sub> activity is downregulated in some AD patients, and this could lead to an enrichment in hexadecyl-containing precursor lipids (33, 34). Moreover, LPCAT1 and LPCAT2 preferentially use 18:2- and 18:3-CoA (LPCAT1) or 20:4-CoA (LPCAT2) over acetyl substrates (35) and thus are more likely to regenerate structural membrane lipids than produce PAF second messengers under physiological conditions. We show here that, in the absence of A $\beta$ <sub>42</sub>, hNT neurons preferentially convert internalized *lyso*-PAF to alkylacylglycerophosphocholines as expected, while in the presence of A $\beta$ <sub>42</sub>, C16:0 PAF accumulation is favored. Thus, it is likely that exposure to pathogenic concentrations of A $\beta$ <sub>42</sub> alters LPCAT substrate preference toward acetyl-CoA generating PAF.

With respect to underlying signaling mechanisms, we show that rising intraneuronal concentrations of C16:0 PAF in response to A $\beta$ <sub>42</sub> target the ER to sequentially activate *m*-calpain, Cdk5, and GSK3 $\beta$ , leading to AT8-reactive phosphorylation of human tau. While PAFs have previously been shown to activate GSK3 $\beta$  (27), our data demonstrate that alkylacylglycerophosphocholine metabolites can also signal through Cdk5. Pharmacological inhibition of Cdk5 prevented both C16:0 PAF-induced tau phosphorylation and C16:0 PAF-induced caspase-dependent neuronal apoptosis, indicating that this kinase is likely the primary effector. Both pharmacological intervention with orsellinic acid, a PAF inhibitor that targets the PAF-induced ER caspase pathway (31) and molecular gain-of-function approaches to enhance hydrolysis of C16:0 PAF to C16:0 *lyso*-PAF were sufficient to inhibit A $\beta$ <sub>42</sub> neurotoxicity, providing further evidence that these pathways can be potentially modulated in AD. Collectively, these data provide insight into how disruption of glycerophosphocholine metabolism can define neu-

ronal response to accumulating oligomeric A $\beta$ <sub>42</sub> identifying potential targets for therapeutic intervention.

## Methods

**Tissue.** Flash-frozen parahippocampal gyri without fixation from deceased AD patients (74  $\pm$  8 years) and controls (78  $\pm$  16 years) who suffered sudden death due to non-neurological complications were obtained from the Douglas Hospital Research Centre Brain Bank (Montreal, Canada). Postmortem delay was between 5 and 25 h. TgCRND8 mice (13) and NonTg littermate controls were killed at 8 or 14–16 weeks of age. All animal manipulations were performed according to the strict ethical guidelines for experimentation established by the Canadian Council for Animal Care.

**Lipid Analysis.** Glycerophospholipids were extracted according to a modified Bligh and Dyer procedure (16). LC-ESI-MS and TLC analyses were performed as described in *SI Materials and Methods*.

**Neuronal Culture and Assays.** NT2/D1 cells were terminally differentiated to hNT neurons as described in ref. 36. RT-PCR conditions and primer sequences were as described in ref. 16. All treatment details are detailed in *SI Materials and Methods*. Transfections, quantification, calcium imaging methodologies, activity assays, and antibodies used for Western analyses are described in *SI Materials and Methods*.

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# Supporting Information

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## SI Materials and Methods

**Lipid Extraction and TLC Analyses.** Tissue and cell pellets were weighed and placed in 1 mL ice-cold methanol acidified with 2% acetic acid and extracted as described in ref. 1. For LC-ESI-MS, a 46- $\mu$ L sample, consisting of 10  $\mu$ L lipid extract, 5  $\mu$ L deuterated standards (2.5 ng/species), and 31  $\mu$ L 0.1% formic acid in H<sub>2</sub>O, was analyzed per MS run exactly as described in ref. 2. To control for extraction efficiency, human samples were spiked with 187.5 ng C13:0 LPC at time of extraction. Where relative estimation is indicated, concentrations were determined in comparison to 1.25 ng d<sub>4</sub>-C16:0 PAF or d<sub>4</sub>-C<sub>16</sub>-lyso-PAF spiked at the time of MS analysis. Tissue wet weight in grams was divided by the specific gravity of brain tissue (1.050) to obtain the volume in milliliters for molar calculations (3). For TLC, lipid extraction from media and of acid-labile and acid-resistant fractions was performed as described in ref. 1. Briefly, hNTs, terminally differentiated for 6 weeks as described in ref. 4 and seeded at a final density of  $1 \times 10^7$  cells/10-cm dish, were incubated at 37 °C with 1  $\mu$ M B-PAF in DMEM/F12 media containing 0.025% BSA for 0, 2, 5, 15, 30, or 60 min. At each time point, four plates/condition were removed from the incubator and placed on ice. Media was removed, and 1 mL ice-cold methanol acidified with 2% acetic acid was added to this extracellular fraction. Cells were washed with 200 mM Na acetate, pH 4.5, 25 mM NaCl, 1% BSA to remove B-PAF present at the plasma membrane or bound to cell surface proteins. This acid-labile fraction was collected and extracted. The remaining monolayer of cells was collected in 1 mL acidified methanol by scraping the plate with a cell lifter (Fisher) representing the intracellular fraction resistant to the 200 mM Na acetate, pH 4.5, 25 mM NaCl, 1% BSA. Lipids were extracted from all three fractions at each time point by the Bligh and Dyer method, developed on TLC plates (20  $\times$  20 cm Silica Gel 60; Fisher) in a solvent system of chloroform/methanol/acetic acid/water (50:30:8:5, vol/vol) using B-PAF and B-lyso-PAF (Molecular Probes) as authentic markers. Fluorescent lipids were visualized under UV light using AlphaImager-1220 software (Alpha Innotech). Fluorescence intensity corresponding to lipid yield was determined by densitometry using the Advanced Measurement Module of OpenLab version 3.5 expressed as the percentage of total lipids. Live-cell imaging following exposure to 1  $\mu$ M C11:0 B-PAF was performed as described in ref. 1.

**Neuronal Culture, Treatment, Death, mRNA Analyses, Transfections, and Protein Assays.** Cells were treated for up to 24 h with C16:0 PAF (0.01–1  $\mu$ M), C16:0 lyso-PAF (1  $\mu$ M), mc-PAF (0.01–5  $\mu$ M) all from Biomol Research Laboratories, or A $\beta$ <sub>42</sub> (25  $\mu$ M; Sigma) prepared as soluble oligomers (5). All treatments were performed in serum-free media containing 0.025% BSA devoid of PAF-AHs (Fig. S8a). Some cultures were pretreated for 15 min with roscovitine (0.1–5  $\mu$ M; Sigma) or vehicle (0.1% DMSO). Treatments were carried out for up to 24 h as indicated. Caspase activity in live cells was measured using CaspaTag (caspase 3/7; Chemicon) or FLICA (caspase 2; Intergen) assays according to the manufacturers' protocol. Cdk5 activity was determined as described in ref. 6. To assess caspase dependence, hNTs were treated with C16:0 PAF (1  $\mu$ M) or vehicle in the presence and absence of the caspase 2 inhibitor Z-VDVAD-FMK (50  $\mu$ M) or 3/7 inhibitor Z-DEVD-FMK (50  $\mu$ M). Cell survival was assessed by LIVE/DEAD Viability/Cytotoxicity assay (Molecular Probes) or TUNEL (Roche). To test functional impact, hNTs were treated with C16:0 PAF (1  $\mu$ M) in the

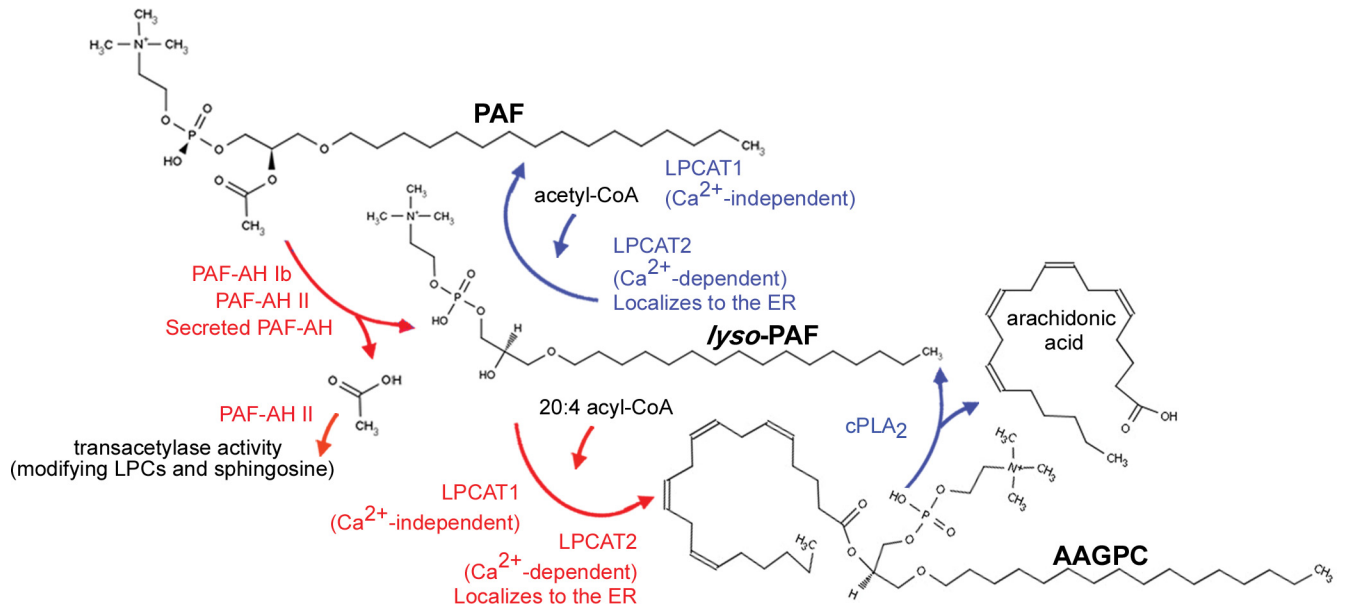
presence or absence of the caspase 2 inhibitor Z-VDVAD-FMK (50  $\mu$ M) or the caspase 3/7 inhibitor Z-DEVD-FMK (50  $\mu$ M) for 24 h. Data were expressed as percent survival relative to vehicle-treated cells. PAF-AH I  $\alpha_2$  in pcDNA 3.1 (kindly provided by Dr. H. Arai, University of Tokyo), or PAF-AH II cloned in our laboratory into pcDNA 3.1 (1) were cotransfected into hNT cells with EGFP in pcDNA3.1 (Invitrogen) by calcium phosphate transfection. Cell survival in serum-free media following C16:0 PAF (1  $\mu$ M) or A $\beta$ <sub>42</sub> (25  $\mu$ M) treatment was calculated as described in ref. 7. Where transfection of PAF-AH II or PAF-AH  $\alpha_2$  with EGFP or empty vector and EGFP is indicated, all treatments were carried out 72 h after transfection when maximal EGFP was observed. Transfection efficiency ( $\approx$ 22%) was comparable in all conditions as established by counting EGFP-positive cells immediately before treatment in each experiment as well as, in separate experiments, PAF-AH II immunoreactive cells. Cell survival was calculated as the EGFP-positive cell number following treatment/mean EGFP-positive cell number in the vehicle control times 100. Data were analyzed by Student's *t* test or ANOVA followed by post-hoc Dunnett's *t* tests or Tukey tests as applicable using Instat v3.0.

**Antibodies Used.** Western blot analysis was performed using antibodies raised against  $\alpha_1$ ,  $\alpha_2$ , or PAF-AH II (1:1,000; Dr. H. Arai, University of Tokyo), GAPDH (1:2,000; IMGENEX), LIS1 clone 210 (1:5,000; Dr. O. Reiner, Weizmann Institute of Science), BIP/GRP78 (1:1,000; Cell Signaling),  $\alpha$ II-spectrum (1:1,000; Sigma), *m*-calpain (1:500; Sigma), tau (AT8, 1:1,000; Sigma),  $\beta$ -tubulin (1:10,000; Sigma), Cdk5 (C-8, 1:1,000; Santa Cruz), p35 (C-19, 1:500; Santa Cruz), p39 (C-20, 1:500; Santa Cruz), phospho-GSK3 $\beta$  (Ser-9, 1:1,000; Cell Signaling), AKT (1:1,000; Cell Signaling), and phospho-AKT (Thr-308, 1:1,000). Secondary antibodies were horseradish peroxidase-conjugated anti-mouse IgG (1:2,000), anti-rabbit IgG (1:5,000), and anti-goat IgG (1:800; all from Jackson Immunolabs) visualized using SuperSignal West Pico (MJS BioLynx). Quantification of protein expression was performed by densitometric analysis of individual bands using ImageJ analysis software [v1.42; National Institutes of Health (NIH)].

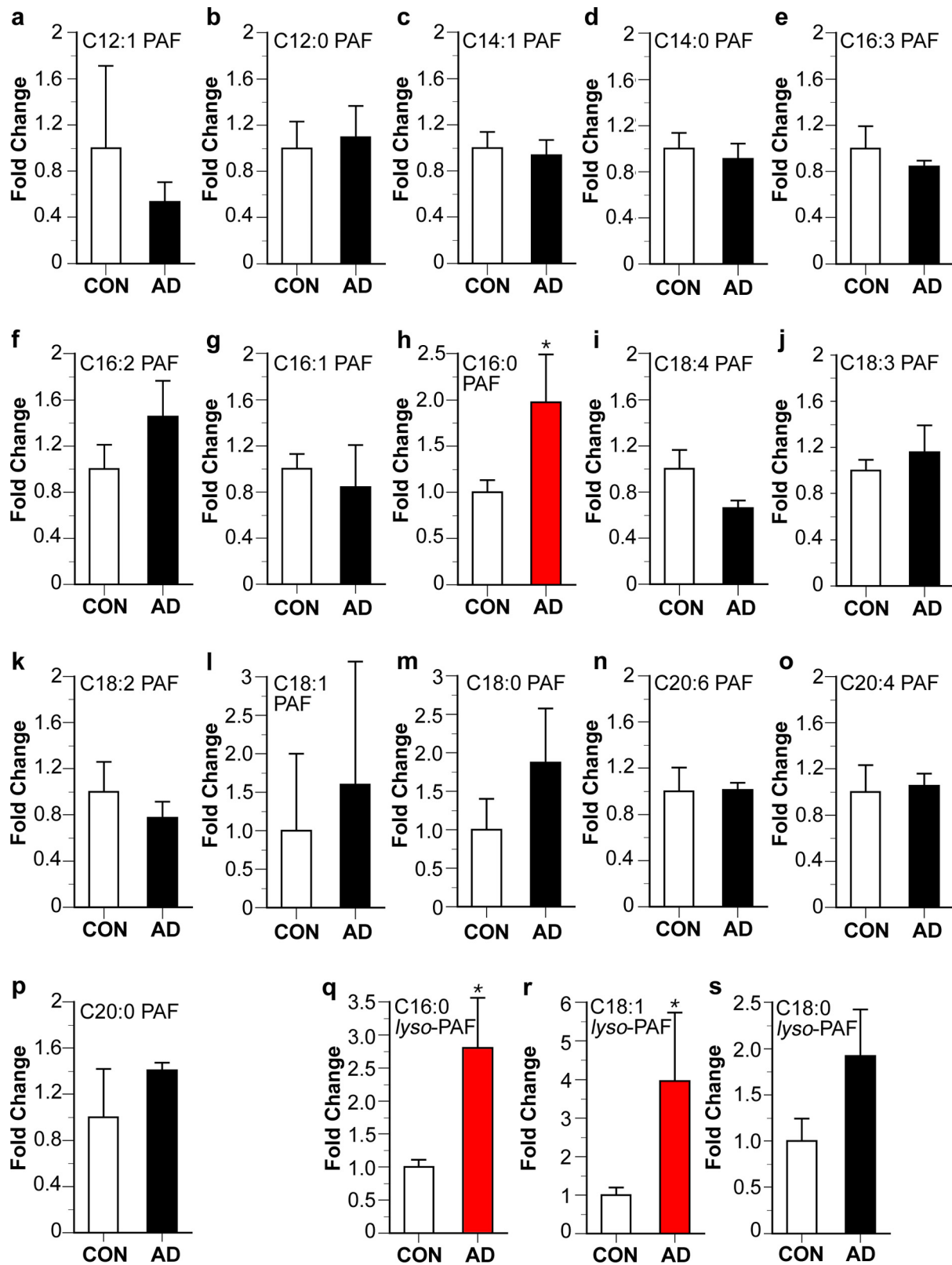
**Ratiometric Determination of Intracellular Calcium Mobilization.** Briefly, hNT neurons on glass coverslips were loaded with 10  $\mu$ M fura-2-AM plus 0.02% pluronic F-127 (Molecular Probes) for 30 min at 37 °C. After rinsing with PSS Mg<sup>2+</sup>-free buffer containing 2 mM HEPES, pH 7.2, 140 mM NaCl, 5 mM KCl, 2.3 mM CaCl<sub>2</sub>, and 10 mM glucose cells were allowed 5 min in the same buffer to stabilize. Coverslips were placed in a microperfusion system housed on the stage of an LSM-410 Zeiss (Carl Zeiss) inverted laser scanning microscope equipped with an argon-krypton ion laser and with a Fluor 40 $\times$ /0.75W acroplan objective. After a region of interest was chosen, fura-2 fluorescence was measured at 510 nm emission with 340/380 nm dual excitation. The basal level of calcium was recorded for 120 s, followed by the application of the control PSS buffer, supplemented with 0.025% BSA (vehicle) for another 240 s. C16:0 PAF (1  $\mu$ M) was then applied for 180 s in control buffer followed by a subsequent 60-s wash out with control buffer alone. C16:0 lyso-PAF (1  $\mu$ M) in control buffer was then applied for 180 s followed again by a 60-s wash out with control buffer alone. Finally, K<sup>+</sup> (10 mM) was applied as a positive control and the recording was continued for another 160 s. The 340–380 nm fluorescence ratio (R<sub>340/380</sub>) for five cells was measured on five independent coverslips.

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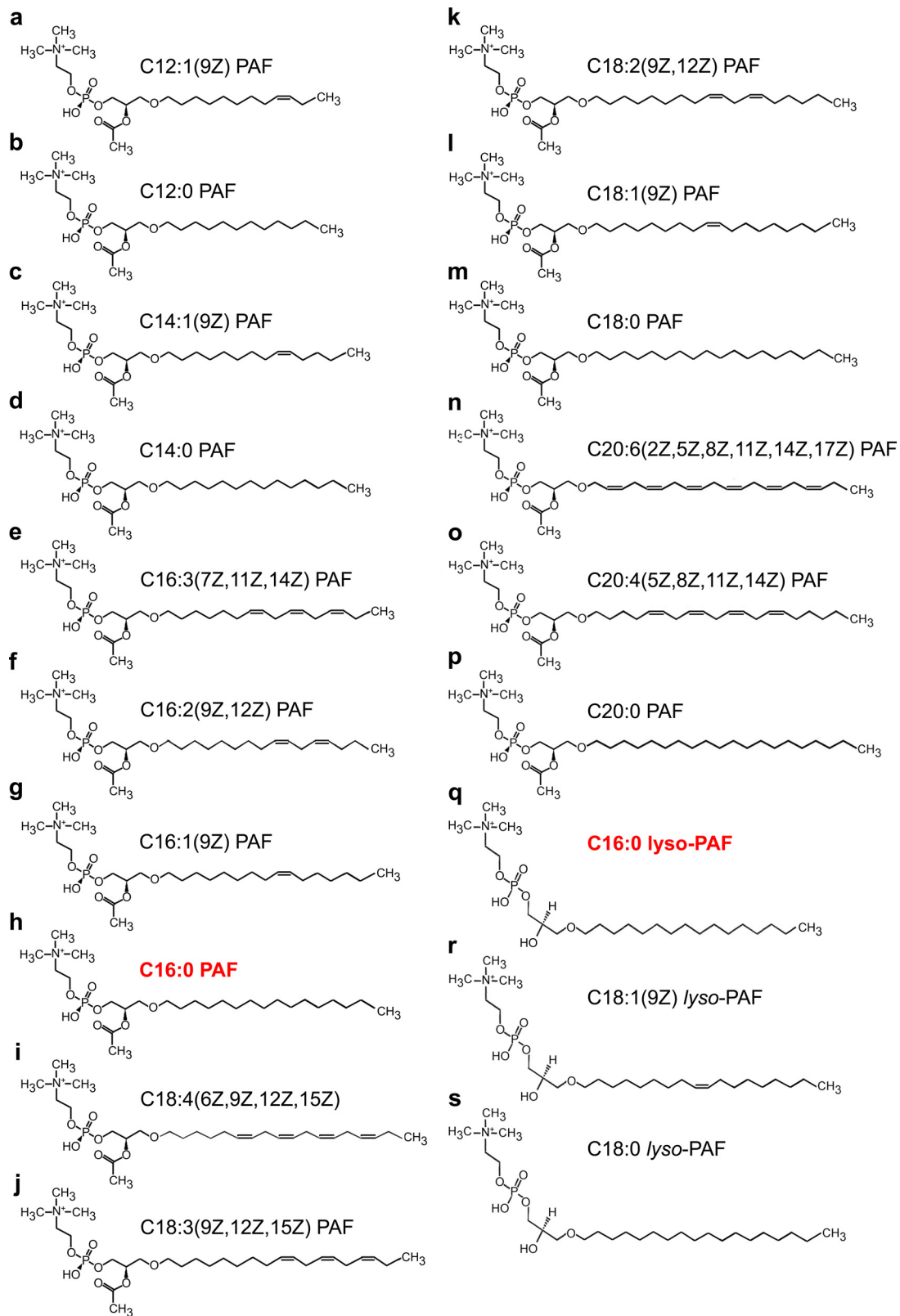




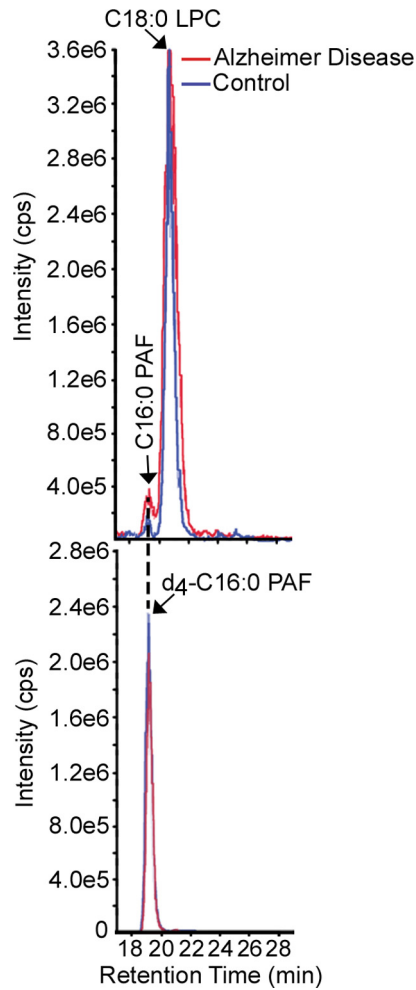
**Fig. S1.** The alkylacylglycerophosphocholine remodeling pathway (Lands' cycle). Alkylacylglycerophosphocholines can be remodeled to PAFs through the actions of cPLA<sub>2</sub> releasing arachidonic acid and LPCAT1 and LPCAT2 with acetyl-CoA as the acyl donor (blue). Individual PAF isoforms are defined by the length and degree of unsaturation of the long chain fatty acid at the *sn*-1 position. PAFs are themselves remodeled back to structural membrane lipids by the actions of any one of three PAF-AH isoforms followed by LPCAT1 or LPCAT2 using acyl-CoA donors (red). In brain, PAF-AH I is neuron-specific. PAF-AH II has additional transacetylase activity that can transfer the free acetyl group to the *sn*-2 position of LPCs and sphingosines. Abbreviations: AAGPC, 1-O-alkyl-2-arachidonoyl-glycerophosphocholine; cPLA<sub>2</sub>, cytosolic phospholipase A<sub>2</sub>; LPC, *lysophosphatidylcholine*; LPCAT, LPC-acyltransferase; PAF, platelet activating factor; PAF-AH, PAF acetylhydrolase.



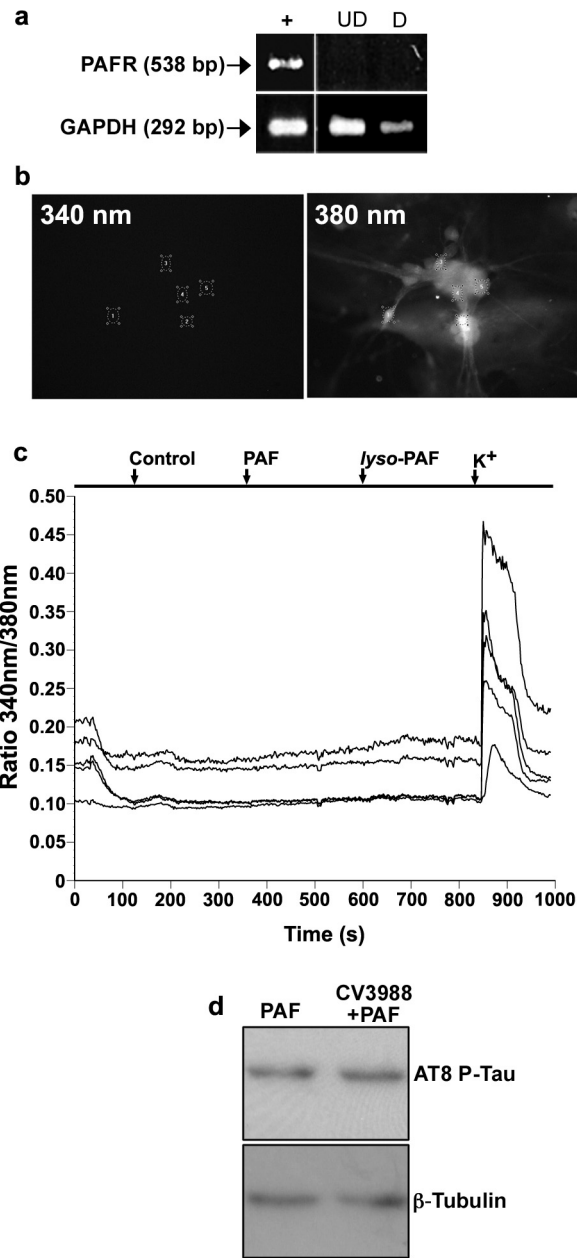
**Fig. S2.** Specific PAF metabolites accumulate in AD tissue. (a–p) Sixteen PAF species and (q–s) three *lyso*-PAF species were detected in human posterior/entorhinal cortex by LC-ESI-MS. Predicted structures are indicated in Fig. S3. Species were identified on the basis of mass and retention time. Significant changes in (h) C16:0 PAF, (q) C16:0 *lyso*-PAF, and (r) C18:1 *lyso*-PAF levels were detected in AD tissue ( $n = 4$  individuals/condition). Data are expressed as fold change relative to controls + SEM (\*,  $P < 0.05$ , Student's  $t$  test).



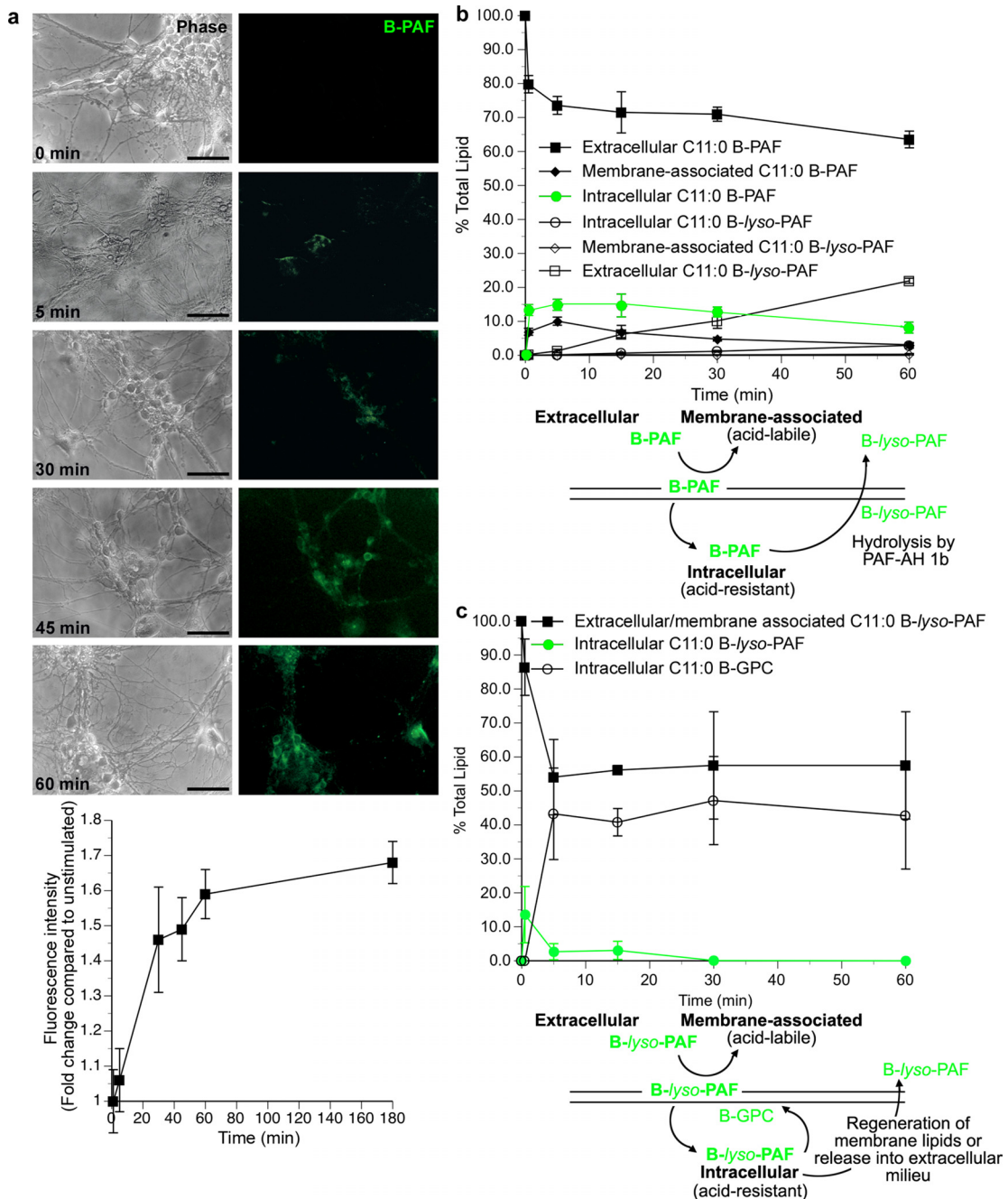
**Fig. S3.** Predicted PAF structures identified in control and AD cortex. (a–r) The predicted structures of the 16 (a–r) PAF species and (q–s) three *lyso*-PAF species detected in human posterior/entorhinal cortex by LC-ESI-MS with the likely position of double bonds are indicated.



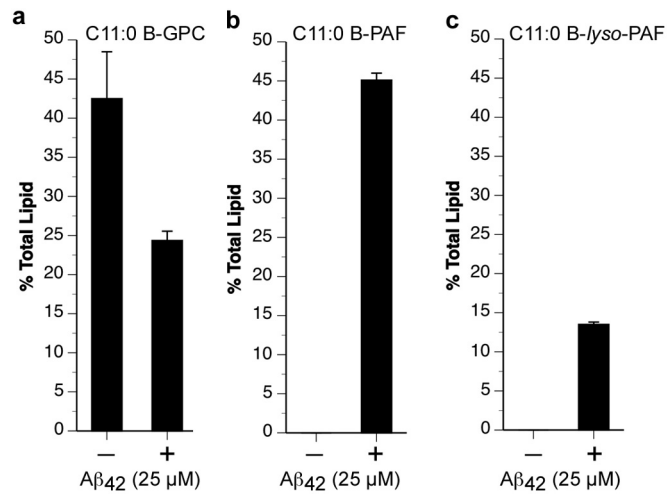
**Fig. 54.** The identities of C16:0 PAF and C16:0 *lyso*-PAF were verified by co-elution with deuterated standards. As PAF species can be isobaric with other phosphatidylcholines, analytes were reanalyzed following spike with deuterated d4-C16:0 PAF or d4-C16:0 *lyso*-PAF. Representative extracted ion chromatograms at  $m/z$  524 (*Upper*) or 528 (*Lower*) with a phosphocholine product ion at  $m/z$  184.0 detected by precursor ion scan in positive ion mode are presented. C16:0 PAF clearly separates from isobaric C18:0 LPC (*Upper Inset*) and coelutes with d4-C16:0 PAF (*Lower Inset*).



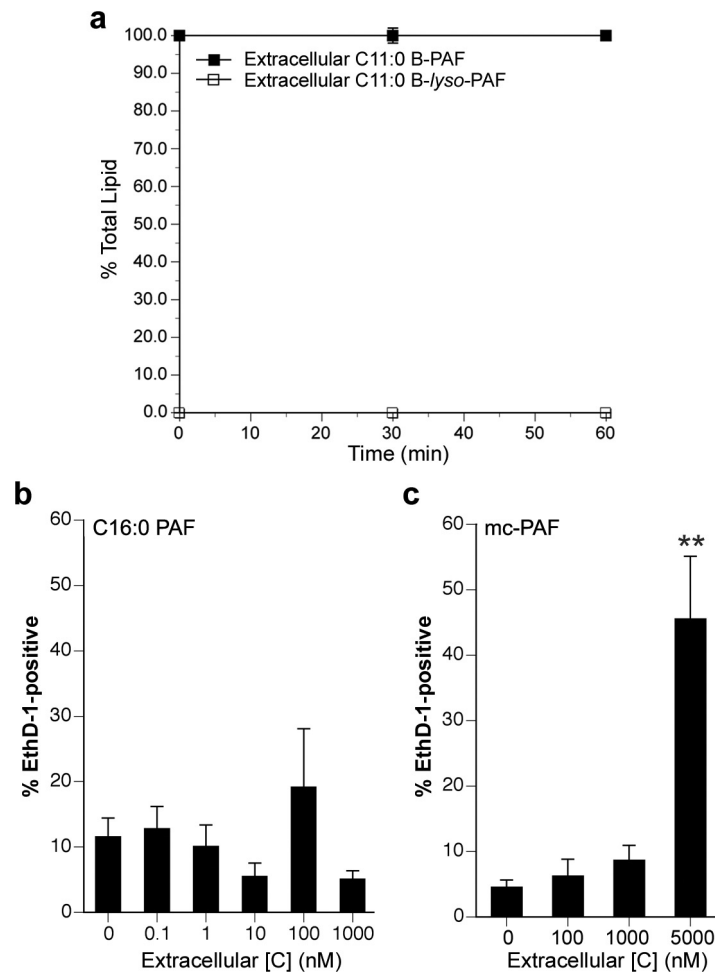
**Fig. S5.** C16:0 PAF and C16:0 *lyso*-PAF do not elicit signals associated with PAFR activation in PAFR-negative human neurons. (a) Undifferentiated NT2/D1 precursor cells (UD) and terminally differentiated hNT neurons (D) do not express PAFR mRNA as determined by RT-PCR. PC12 cells stably transfected with human PAFR(+) were used as a positive control. GAPDH was amplified from the same random-primed cDNA to confirm template integrity. (b) C16:0 PAF and C16:0-*lyso*-PAF do not acutely mobilize calcium. Fluorescent micrographs depict hNT neurons loaded with the calcium-sensitive indicator Fura-2 AM. Neurons on which quantification was performed are identified numerically as 1–5. *Left* depicts cells during the period of C16:0 PAF exposure. *Right* depicts cells following application of K<sup>+</sup> (10 mM) (c) Fura-2 fluorescence was measured at 510 nm emission with 340/380 nm dual excitation. The basal level of calcium was recorded for 120 s, followed by the application of the control PSS buffer, supplemented with 0.025% BSA (vehicle) for another 240 s. C16:0 PAF, C16:0 *lyso*-PAF, and finally K<sup>+</sup> was then sequentially applied for 180 s in control buffer followed by a subsequent 60-s wash out with control buffer alone. The 340/380 nm fluorescence ratio (R340/380) for five cells was measured on five independent coverslips. Of 25 cells stimulated with C16:0 PAF over five different experiments, none exhibited a calcium response. All of these neurons stimulated with K<sup>+</sup> exhibited a robust response. A representative experiment of five replicates is depicted. (c) Treatment with 10 μM CV3988, a PAFR-specific antagonist, did not inhibit C16:0 PAF-mediated AT8-reactive tau phosphorylation.



**Fig. S6.** Internalization and metabolic fate of C11:0 B-PAF and C11:0 B-lyso-PAF administered to hNT neurons. (a) The kinetics of B-PAF internalization were assessed by live-cell imaging. hNT neurons were incubated with 1  $\mu$ M B-PAF at 37  $^{\circ}$ C for 0, 5, 30, 45, 60, and 180 min in serum-free treatment media containing 0.025% BSA. At each time point, B-PAF-containing media were removed and set aside. Cells were washed with PBS plus 1% BSA to remove phospholipids loosely bound to the extracellular face of the plasma membrane, imaged, and the B-PAF-containing media replaced. Data are reported as the mean  $\pm$  SEM in fluorescence intensity standardized to background autofluorescence before treatment. Cell-associated fluorescence increased over the first 60 min of exposure. (Scale bar, 50  $\mu$ m.) (b) The fate of internalized B-PAF was determined by TLC. hNT neurons were treated with 1  $\mu$ M B-PAF. After 0, 1, 5, 15, 30, and 60 min of exposure, lipids were extracted from three fractions: Extracellular (media), membrane-associated, acid-labile (lipids loosely bound to the plasma membrane), and internalized, acid-resistant (lipids extracted from hNT neurons). Fractions were separated by TLC and species identified in comparison to labeled standards run concurrently. Data represent the relative fluorescence present in each fraction expressed as a percentage of the total labeled lipid recovered and are the mean  $\pm$  SEM of two independent experiments conducted in replicate. C11:0 B-PAF was rapidly internalized by hNT neurons and hydrolyzed to C11:0 B-lyso-PAF. The labeled metabolite was released into the extracellular media. Elevated intracellular levels of C11:0 B-PAF remained constant for the first 30 min, dropping gradually once the rate of hydrolysis exceeded the rate of internalization. (c) C11:0 B-lyso-PAF was also rapidly internalized. Extracellular C11:0 B-PAF associated with the plasma membrane and was internalized within 1 min of exposure. Intracellular lipid was, however, converted to C11:0 B-alkylacylglycerophosphocholines (C11:0 B-GPC) and not C11:0 B-PAF in the absence of any other toxic stimuli.



**Fig. S7.** Treatment of hNT neurons with A $\beta_{42}$  induces conversion of internalized C11:0 B-lyso-PAF to C11:0 B-PAF. The acid-resistant fractions representing internalized lipid in hNT neurons were analyzed by TLC. In control cultures (-), all of the internalized C11:0 B-lyso-PAF was converted to C11:0 B-alkylacylglycerophosphocholine (GPC) (a) or released as C11:0 B-lyso-PAF (acid-labile fraction; Fig. S6c). In cultures treated with A $\beta_{42}$  (+), more internalized lipid was retained (c) and was converted to C11:0 B-PAF (b) as compared to C11:0 GPC (a).



**Fig. S8.** Direct exposure of hNT neurons in treatment media to C16:0 PAF does not compromise plasma membrane integrity at submicellar concentrations. (a) The absence of PAF-AH activity in serum-free treatment media containing 0.025% BSA was confirmed by the addition of a BODIPY fluorophore-labeled C11:0 PAF isoform [1-(O-[11-(4,4-difluoro-5,7-dimethyl-4-bora-3a,4a-diaza-s-indacene-3-propionyl)amino]undecyl)-2-acetyl-sn-glycero-3-phosphocholine] directly to media with extraction and analysis by TLC. No degradation of C11:0 B-PAF to C11:0 B-lyso-PAF was detected when lipids were incubated for up to 60 min at 37 °C in treatment media. (b) C16:0 PAF did not cause EthD-1 uptake compared to vehicle-treated control cultures indicating membrane integrity was not impaired. (c) Loss of membrane integrity was only observed when PAF hydrolysis was suppressed and extracellular PAF lipid concentrations exceeded critical micelle concentration demonstrated using the PAF-AH-resistant PAF analog mc-PAF.