

## RESEARCH ARTICLE

# Oleic Acid Ameliorates Amyloidosis in Cellular and Mouse Models of Alzheimer's Disease

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## Keywords

Alzheimer's disease, amyloid, Cos-7 cells, oleic acid, plaques, transgenic mice.

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Received 19 August 2010; accepted 29 September 2010.

doi:10.1111/j.1750-3639.2010.00449.x

## Abstract

Several lines of evidence support protective as well as deleterious effects of oleic acid (OA) on Alzheimer's disease (AD) and other neurological disorders; however, the bases of these effects are unclear. Our investigation demonstrates that amyloid precursor protein (APP) 695 transfected Cos-7 cells supplemented with OA have reduced secreted amyloid-beta ( $A\beta$ ) levels. An early-onset AD transgenic mouse model expressing the double-mutant form of human APP, Swedish (K670N/M671L) and Indiana (V717F), corroborated our *in vitro* findings when they were fed a high-protein, low-fat (18% reduction), cholesterol-free diet enriched with OA. These mice exhibited an increase in  $A\beta_{40}/A\beta_{42}$  ratio, reduced levels of beta-site APP cleaving enzyme (BACE) and reduced presenilin levels along with reduced amyloid plaques in the brain. The decrease in BACE levels was accompanied by increased levels of a non-amyloidogenic soluble form of APP (sAPP $\alpha$ ). Furthermore, the low-fat/+OA diet resulted in an augmentation of insulin-degrading enzyme and insulin-like growth factor-II. These results suggest that OA supplementation and cholesterol intake restriction in a mouse model of AD reduce AD-type neuropathology.

## INTRODUCTION

The brains of Alzheimer's disease (AD) patients are characterized by the deposition of amyloid plaques (26). The amyloid plaques consist of different forms of the amyloid-beta ( $A\beta$ ) derivatives of the amyloid precursor protein (APP) (28), which are produced by the cleavage of APP by two proteolytic enzymes,  $\beta$ - and  $\gamma$ -secretases. Out of various  $A\beta$  peptides produced, products of 40 ( $A\beta_{40}$ ) and in particular 42 ( $A\beta_{42}$ ) residues are the most common constituents of amyloid plaques (37). Moreover, soluble form of oligomeric  $A\beta$  are widely accepted as the prime trigger for AD pathogenesis, resulting in neuronal loss with consequential dementia and death [reviewed in (27)].

There is a substantial amount of data that conferred a protective (11, 32, 33) as well as deleterious effects (21, 25, 43) of oleic acid (OA; 18:1 n-9) on AD and other neurological disorders. For instance, a significant decrease in monounsaturated fatty acids (FAs), including OA, has been reported in the frontal cortex and hippocampus of AD brains (11, 33). OA has also been shown to significantly inhibit the activity of prolyl endopeptidase (32); prolyl endopeptidase levels are significantly increased in AD brains, suggesting their functional role in brain amyloidogenesis (3). However, the biochemical and pathological bases whereby OA affects AD remain unclear. In addition, none of the previous studies had examined cholesterol-free diets in combination with low-fat intake and enriched OA.

In this study for the first time, we show that the brains of AD mice on low-fat/+OA diet show multifactorial protective effects in

pathways important to the development of amyloid plaques. These results suggest that regulation of dietary OA intake may offer a new tool to reduce the risk of AD.

## MATERIALS AND METHODS

### Cell culture

Cos-7 cells were stably transfected with pCEP-APP695 (encoding human APP695) plasmid maintained in DMEM (Sigma, Germany) with 10% fetal calf serum (FCS, PAN, Germany) and 200  $\mu$ g/mL hygromycin (14). Cells were cultured in 10-cm dishes, washed three times with FCS-free media (to get rid of the lipids present in the FCS) and incubated for 12 h with delipidated-FCS DMEM containing 0.01–25  $\mu$ M of OA–bovine serum albumin (BSA) complexes. OA containing DMEM were renewed every 4 h to maintain steady concentration of lipid. The 8- to 12-h conditioned media were analyzed. Cells grew equally well as determined by cell protein concentration and maintained normal morphology at all concentrations of OA used.

### OA preparation

OA (Nu-Chek-Prep, Inc., USA), organic solvents and BSA solution were purged with argon extensively, prior to use. The protocols to form a fresh OA–BSA (2:1) complex are based on established procedures (2).

## Mice, treatment and tissue preparation

All experimental procedures were performed according to the animal care guidelines of the University of Western Ontario (approval ID: 2004-065-06). All studies were performed on 5-month-old congenic C57BL/6J male mice heterozygous for TgCRND8, which express a double-mutant form of human APP695: Swedish (K670N/M671L) and Indiana (V717F) (6). TgCRND8 mice demonstrate severe AD-like amyloid pathogenesis, including A brain deposits as early as 3 months of age (6). Mice used in these studies were housed on 12 h light/dark cycles. At 3 weeks of age, TgCRND8 mice were provided *ad libitum* access to water and one of three diets ( $n = 6$  for each diet) until 20–21 weeks of age, at which time they were sacrificed by cervical decapitation. Brains were rapidly removed and sectioned sagittally into hemibrains. One hemibrain was used for either biochemical [protein, RNA and enzyme-linked immunosorbent assays (ELISAs)] or lipid (OA) analyses. The other hemibrain was fixed in 10% neutral buffered formalin and 0.1 M Phosphate buffered saline (PBS) for 48 h and then stored at 4°C in PBS and 1% sodium azide until it is used for immunohistochemistry.

## Dietary manipulations

The experimental diets involved the following three groups: (i) conventional mouse chow (PicoLab Mouse Chow Diet 20, Purina Mills, St. Louis, MO, USA); (ii) soybean oil (1%), high-protein, cholesterol-free control diet (low fat/–OA; L10047, Research Diets, New Brunswick, NJ, USA) and (iii) soybean oil (1%), high-protein, cholesterol-free diet supplemented with 2% OA (low fat/+OA). The compositions of ii and iii diets were as follows: alcohol extracted casein (304.7 g/kg), DL-methionine (4.6 g/kg), sucrose (279.3 g/kg), maltodextrin (Fro-Dex) (284.4 g/kg), vitamin mix (V10001, AIN-76A) (10.2 g/kg), mineral mix (S10001, AIN-76A; Research Diets, New Brunswick, NJ, USA) (35.0 g/kg), cellulose (40.6 g/kg), xanthan gum (8.6 g/kg) and soybean oil (10.2 g/kg) with and without OA (20.0 g/kg). Chow diet was used to compare the effects of low-fat/–OA diet alone on AD parameters.

## Quantitative Western blotting

Western blot to analyze proteins has been described previously (5). Blots were probed with anti-APP (A-8717, 1:15 000) and anti-nicastrin (NCT, N-1660) from Sigma, anti- $\beta$  site APP cleaving enzyme (BACE)–C-terminal fragment (CTF) (1:1000, a gift from Dr Michael Willem), anti- $\beta$ -actin (1:5000) from Santa Cruz (sc-1616), anti-insulin-degrading enzyme (IDE)-1 (1:4000) and anti-soluble APP $\alpha$  (sAPP $\alpha$ ) (1:5000, gifts from Dr Dennis Selkoe), anti-presenilin 1-N terminal fragment (PS1-NTF) (1:2500, a gift from Dr Sam Gandy), anti-transferrin (TTR) (1:500, a gift from Dr Sancia Gaetani), anti-insulin-like growth factor-II (IGF-II) (1:300, a gift from Dr Victor Han) and anti-prion protein (PrP) (1:5000, a gift from Dr Alex Strom). Secondary antibodies were from Amersham, USA (1:15 000 for goat anti-rabbit, 1:5000 for goat anti-mouse). Data analysis was performed using the Molecular Analyst II densitometric software (Bio-Rad, Hercules, CA, USA).

## Real-time RT-PCR

RNA isolation, reverse transcription and PCR conditions have been described previously (5). The primers used are: TgAPP, forward: 5'-GAT GAC GTC TTG GCC AAC ATG-3' and reverse: 5'-CGG AAT TCT GCA TCC AGA TTC AC-3' (308-bp product); murine APP, forward: 5'-CCG ACG ATG TCT TGG CCA AC-3' and reverse: 5'-CCG AAT TCT GCA TCC ATC TTC AC-3' (310-bp product); PrP, forward: 5'-GGG GAC AAC CTC ATG GTG GTA GT-3' and reverse: 5'-TCC ACT GGC CTG TAG TAC ACT TGG-3' (283-bp product) and  $\beta$ -actin, forward: 5'-TCG TGG GCC GCT CTA GGC ACC A-3' and reverse: 5'-GTT GGC CTT AGG GTT CAG GGG GG-3' (256-bp product). Data analysis was performed using the BioRad GeneX (Bio-Rad, Hercules, CA, USA).

## A $\beta$ sandwich ELISAs

Quantification of A $\beta$ 40 and A $\beta$ 42 production in conditioned media (18) and in brain (Signet Laboratories, Dedham, MA, USA) was carried out by standard ELISA methods.

## Fatty acid analysis

FAs in the brain were isolated and methylated according to Moser and Moser (29). The fatty acid methyl ester (FAME) mixture was then re-suspended in hexane and analyzed by gas chromatography–mass spectroscopy (GC-MS). GC-MS analysis was performed on a Hewlett-Packard Series II 5890 gas chromatograph (Hewlett-Packard, Wilmington, DE, USA), coupled to an HP-5971 mass spectrometer equipped with a Supelcowax SP-10 capillary column (Supelco, Bellefonte, PA, USA). FAME mass was determined by comparing areas of unknown FAMES to those of a fixed concentration of 17:0 internal standard.

## Immunohistochemistry and image analysis

Plaque density was measured using immunohistochemistry with a biotinylated-4G8 monoclonal antibody (Signet Laboratories), directed against amino acids 17–24 of the  $\beta$ -amyloid peptide. The mean area of the field (0.307 mm<sup>2</sup>) was examined in three regions of neocortex, hippocampus and amygdala: 0.5–0.9 mm, 0.9–1.35 mm, and 1.35–1.8 mm mediolateral from the bregma (Mouse Brain Atlas, Franklin and Paxinos 2007) on 40  $\mu$ m thick rostral (cross) sections of each hemisphere. Plaque density was calculated by dividing the total area of A $\beta$ -positive structures by the total area of the regions analyzed (in square micrometres).

## Statistics

All values were presented as mean  $\pm$  standard error of the mean (S.E.M.). Protein densitometric, A $\beta$ 40 and 42 levels, and plaque density were compared by Student's unpaired two-tailed *t*-tests. In vitro A $\beta$  levels and ratios of CTFs to full-length APP were analyzed using one-way analysis of variance. The significance level was chosen to be 0.05 ( $P \leq 0.05$ ). Correlations between parameters were tested by linear regression analysis.

## RESULTS

### Effects on CNS lipid profile

Central nervous system (CNS) lipid profile showed a very significant increase in the levels of OA up to twofold ( $12.01 \pm 2.0\%$  distribution or  $2842.35 \pm 514.59$  nmol/g,  $P < 0.01$ ) compared to low-fat/–OA control ( $6.4 \pm 2.3\%$  distribution or  $1403.28 \pm 474.59$  nmol/g) and chow mice ( $8.2 \pm 1.1\%$  distribution or  $2037.4 \pm 6012$  nmol/g).

### Effects on *in vitro* A $\beta$ levels

OA supplementation resulted in lower levels of both A $\beta$  peptides in the conditioned media of APP695 transfected Cos-7 cells starting at as low as 0.01  $\mu$ M compared to the cells supplemented with BSA (Figure 1A).

### Effects on *in vivo* A $\beta$ levels

The hippocampus and the adjacent cortices of low-fat/+OA animals demonstrated a non-significant increase in the A $\beta$ 40 levels ( $2.241 \pm 0.83$ ,  $1.00 \pm 0.41$ ,  $1.4 \pm 0.19$ , ng/mL) and a very significant >30% decrease in the A $\beta$ 42 levels ( $2.08 \pm 0.14$ ,  $4.0 \pm 0.7$ ,  $5.5 \pm 0.40$ , ng/mL,  $P < 0.001$ ) compared to low-fat/–OA mice and chow mice, respectively (Figure 1B).

### Effects on plaque pathology

The decrease in A $\beta$  levels following dietary OA enrichment also led to histopathological changes, specifically that of (>30%) decreased amyloid plaque number in the neocortical ( $4.21 \pm 0.23$ ,  $6.22 \pm 0.63$ ,  $8.7 \pm 0.11$ ,  $P < 0.05$ ), hippocampal ( $3.88 \pm 0.20$ ,  $5.65 \pm 0.22$ ,  $6.6 \pm 0.43$ ,  $P < 0.05$ ) and amygdala ( $1.89 \pm 0.09$ ,  $2.75 \pm 0.15$ ,  $2.96 \pm 0.18$ ,  $P < 0.05$ ) regions of low-fat/+OA mice brains when compared to low-fat/–OA and chow mice, respectively (Figure 1C).

### Effects on *in vivo* $\alpha$ -secretase pathway

Western blot showed over 20% decrease in the levels of mature membrane-bound amyloid precursor protein (APPm;  $6.25 \pm 0.59$  vs.  $8.12 \pm 0.62$ ,  $P < 0.05$ ) and ~50% increase in the levels of neuroprotective sAPP $\alpha$  ( $9.4 \pm 0.31$  vs.  $6.0 \pm 0.65$ ,  $P < 0.05$ ) in the cortices of low-fat/+OA mice compared to low-fat/–OA mice, respectively (Figure 2A).

### Effects on *in vivo* $\beta$ -secretase pathway

The levels of BACE were >20% lowered in low-fat/+OA mice ( $6.83 \pm 0.39$  vs.  $9.0 \pm 0.4$ ,  $P < 0.01$ ) compared to low-fat/–OA mice (Figure 2A). APP–CTFs levels ( $2.56 \pm 0.49$  vs.  $3.85 \pm 0.29$ ,  $P < 0.05$ ) were >30% lowered in low-fat/+OA mice.

### Effects on *in vivo* PS/ $\gamma$ -secretase enzyme complex

The PS/ $\gamma$ -secretase enzyme complex generates A $\beta$  from APP–CTF $\beta$ . PS1-NTF levels ( $5.59 \pm 0.28$  vs.  $6.29 \pm 0.18$ ,  $P < 0.05$ )

were >10% lowered in low-fat/+OA brains compared to low-fat/–OA controls (Figure 2B). There was no significant change in NCT levels in either low-fat/–OA or low-fat/+OA groups of mice.

### Effects on *in vivo* A $\beta$ metabolism

We next checked the levels of IDE and TTR (proteins responsible for the clearance of A $\beta$ ) as well as IGF-II as a function of diet. Low-fat/+OA mice showed >50% increases in the IDE levels ( $6.8 \pm 0.40$  vs.  $4.37 \pm 0.81$ ,  $P < 0.01$ ), >20% increases in IGF-II levels ( $4.01 \pm 0.40$  vs.  $3.1 \pm 0.22$ ,  $P < 0.05$ ) and >40% increases in the TTR levels ( $4.81 \pm 0.31$  vs.  $3.25 \pm 0.02$ ,  $P < 0.05$ ) compared to low-fat/–OA controls, respectively (Figure 2B).

### Effects on PrP levels

The CRND8 transgenic APP (Tg) expression is driven by the PrP promoter; to support the absence of the effect of the different diets on PrP activity, the protein levels of PrP among different groups were investigated. As shown in Figure 2B, the different dietary regimens did not cause any significant change in PrP levels, thereby supporting that both the low-fat/–OA and low-fat/+OA diets did not influence the levels of transgene expression.

### Effects on secretase products

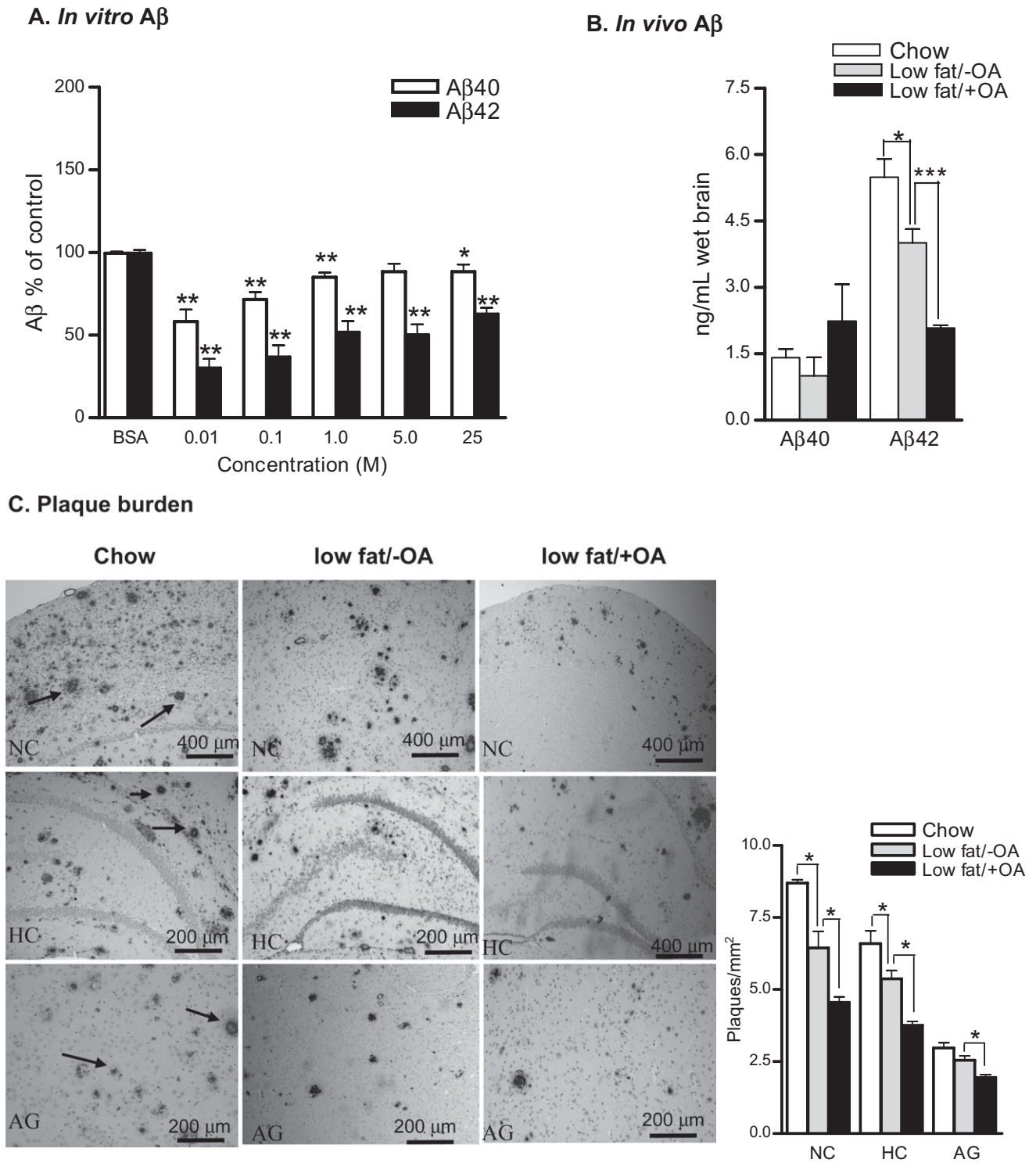
Significant direct correlation was found between increasing levels of OA (percentage distribution) and CTFs/APP ratios ( $R^2 = 0.991$ ,  $P < 0.001$ ) (Figure 2C). Together, these results suggest that there is an association between OA levels and secretase activity.

### Effects on WT and transgene expressions

Both the low-fat/–OA and low-fat/+OA diets did not have any effect on the expression levels of either CRND8 transgene-derived APP or endogenous murine APP (Figure 3A). Similar to PrP protein levels, the levels of PrP gene were comparable across the groups (Figure 3B).

## DISCUSSION

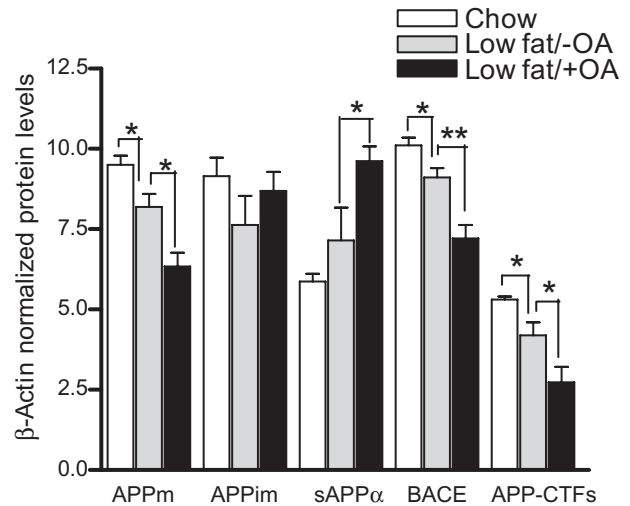
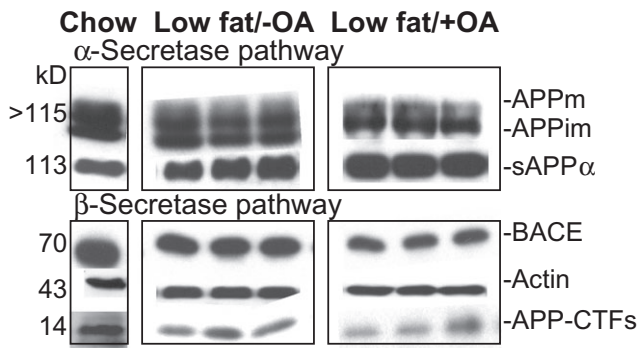
As cholesterol-enriched diets have been shown to increase BACE level and activity, APP processing, and A $\beta$ 42 levels (reviewed in reference 44), we therefore designed a diet depleted in cholesterol and enriched in OA to find out if the protective effects of cholesterol depletion could be enhanced by OA supplementation. Instead of doing a blinded, full-blown animal study, in pilot experiments we screened OA in the cell culture AD model to generate the pilot data as well as to set up the direction for future *in vivo* experiments and to do the power analysis. In the cell culture model that produces high levels of A $\beta$  in the medium to mimic transgene-derived A $\beta$  overproduction in TgCRND8 mice, OA tends to have inhibitory effects on the levels of both A $\beta$  species in the conditioned media. However, as the supplemental concentration of OA is increased, the inhibition of A $\beta$  levels is decreased, most probably attributed to the increased rigidity of the membrane bilayer or altered cellular pH as a result of OA enrichment. Membrane-rigidifying properties of OA at higher concentrations (when it exceeded 20% of the total fat) have been



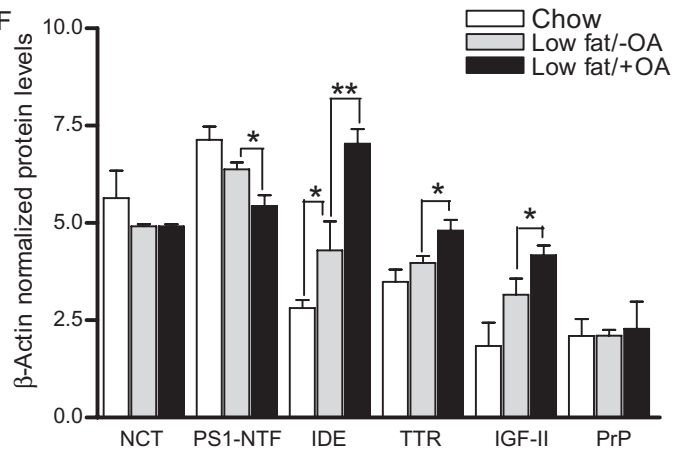
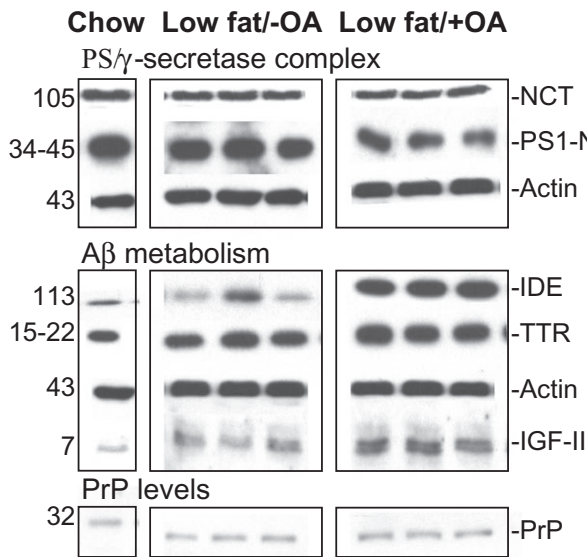
**Figure 1.** In vitro and in vivo effects of oleic acid (OA) supplement on amyloid-beta (Aβ) levels and amyloid plaque burden. **A.** BSA control normalized Aβ levels in the conditioned media after 12-h incubation with OA (*n* = 3). **B.** Enzyme-linked immunosorbent assay results for Aβ40 and Aβ42 levels in the hippocampus and adjacent cortices of chow, low-fat/-OA and low-fat/+OA mice. **C.** Photomicrographs of Aβ stained amyloid

plaque burden (number and size) in the neocortex (NC), hippocampus (HC) and amygdala (AG) of chow, low-fat/-OA and low-fat/+OA mice. Arrows indicate Aβ stained plaques. The plot shows the quantitative analysis of plaque density. The mean ± standard error of the mean are shown for all plots (*n* = 6 for each experiment). \*\*\**P* < 0.001, \*\**P* < 0.01 and \**P* < 0.05.

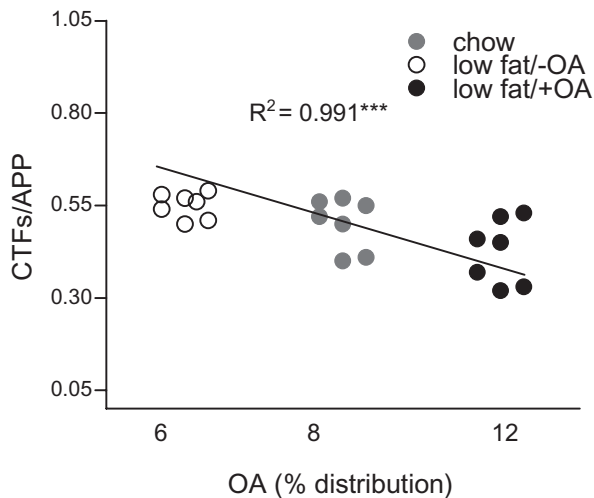
**A. Protein levels**



**B. Protein levels**

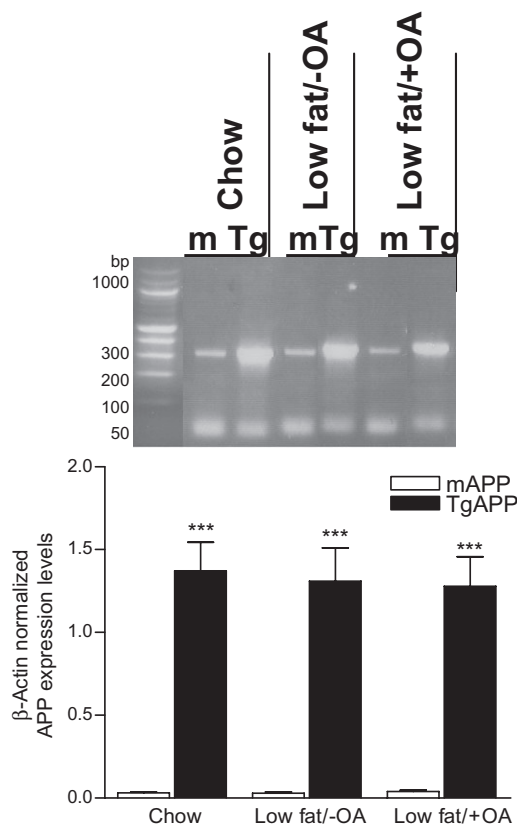


**C. OA vs APP-CTFs**

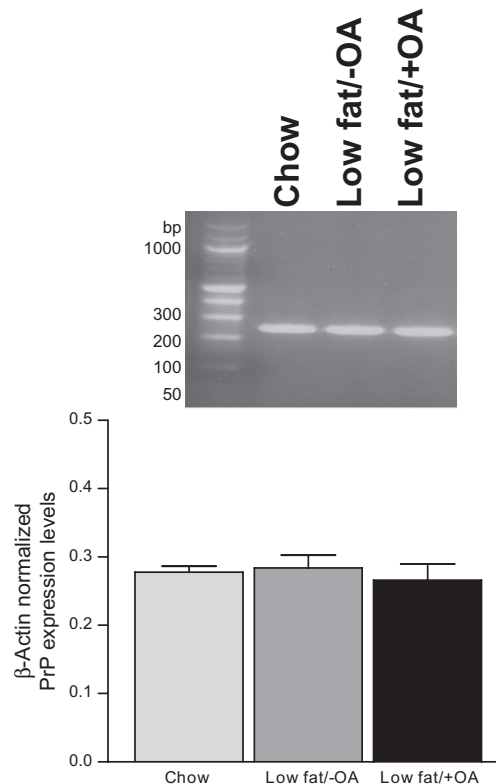


**Figure 2.** In vivo effects of oleic acid (OA) supplement on brain protein levels. Western blots of mature amyloid precursor protein (APPm), immature amyloid precursor protein (APPim), soluble APPα (sAPPα), β-site APP cleaving enzyme (BACE), β-actin and APP-C-terminal fragments (CTFs) (A) nicastrin (NCT), presenilin 1-N terminal fragment (PS1-NTF), insulin-degrading enzyme (IDE), transthyretin (TTR), insulin-like growth factor-II (IGF-II), prion protein (PrP) and β-actin protein levels (B) in the frontal cortices of chow, low-fat/-OA and low-fat/+OA mice, (n=6 for each experiment; out of six, only one animal is shown for chow, and three animals are shown for low-fat/-OA and low-fat/+OA mice). Respective molecular weights (kD) are shown on the left. Plots on the right show the quantitative analysis of Western blots. C. Plot shows secretase-generated APP-C-terminal fragment (α/β) ratios vs. OA percentage distribution. The mean ± standard error for the mean are shown for all plots. \*\*P<0.01 and \*P<0.05. Abbreviation: Aβ = amyloid-beta.

## A. APP gene expression



## B. PrP gene expression



**Figure 3.** Effects of oleic acid (OA) supplement on gene expression levels. Real-time RT-PCR of endogenous murine (m), CRND8 transgene-specific (Tg) amyloid precursor protein (A) and prion protein (B) in chow, low-fat/-OA and low-fat/+OA mice, (n=6 for each experiment, out of 6 only 1 animal is shown for each group). Plots show the quantitative analysis (mean  $\pm$  S.E.M.) of  $\beta$ -actin-normalized gene expression levels, \*\*\* $p < 0.001$ .

shown to impair the transport of A $\beta$  from the membranes and to be positively correlated with reference memory errors in AD rats (43). Furthermore, higher concentrations of OA are expected to increase the acidic pH of the cells, and the highest  $\beta$ -secretase activity is detected at acidic pH (12, 13, 22, 46). In addition, A $\beta$  degradation is mediated by several proteases; given the intracellular localizations and pH optima, these proteases are unlikely to operate at highly acidic pHs (30). We therefore speculate that at a higher OA concentration, the cellular pH is altered, which then exerts a significant biological effect on the intracellular processing of APP, resulting in increased A $\beta$  levels. Correspondingly, OA has been shown to promote the aggregate formation of superoxide dismutase 1 mutants *in vitro* (21) and caused an elevation in PS1,  $\gamma$ -secretase and consequently, A $\beta$  levels in Chinese hamster ovary (CHO) PSwt-1 cells (25) when administered at 2500 and 400  $\mu$ M, respectively. It is important to mention that transfected Cos-7 cells are well established as models for examining the effects of AD proteins/genes on a number of parameters (24, 36, 40). Our results also demonstrate that APP protein could be effectively expressed in Cos-7 cells as already shown (14, 17, 24, 36, 40) and that the transfected Cos-7 cells could be used as a model of APP protein overexpression to examine the function of APP in

mammalian cells (24, 36, 40). In line with our *in vitro* data, we decided to maintain a lower concentration of OA in the experimental mice diet to enhance its membrane-fluidizing effects. Here, we show that dietary OA enrichment indeed provides protection against AD-type neuropathology in the brains of TgCRND8 mice, provided fat intake is lowered (approximately 18% reduction) and cholesterol is depleted from the diet. This outcome appears to be the result of a combination of several underlying factors.

Low-fat/+OA diet resulted in an overall increase of the alpha (sAPP $\alpha$ ) and a decrease in beta (combined APP-CTFs, although the predominant form lowered is CTF $\beta$  and not CTF $\alpha$  as a result of >50% elevated sAPP $\alpha$ ) proteolytic processing of the membrane-bound mature APP and corresponding A $\beta$ 42 peptides, implying that OA may downregulate A $\beta$  generation either by altering APP trafficking to secretase-containing compartments of the membrane or the secretase enzymatic activity itself. Although the cause of this shift is unclear, a decrease in primary brain BACE levels is likely to contribute. Though not investigated here, it would be of interest to assess whether the OA-mediated regulation of BACE levels is stearoyl CoA desaturase (SCD-1) dependent. SCD-1 is the enzyme that catalyzes the conversion of long-chain saturated FA to oleate

(39, 41). There have been reports stating that the induction of SCD-1 activity leads to an increase in cholesterol, plasma lipids and lipoproteins (20, 31). As a feedback mechanism, high OA may down-regulate SCD-1 and may consequently lower cholesterol and corresponding diminished BACE levels and activity. In this regard, SCD-1 has been shown to be upregulated by A $\beta$  treatment (41). Alternatively, the shift could be explained by the altered processing of APP, where the alpha pathway is augmented, resulting in a reciprocal diminishment of the beta pathway.

The decrease in the levels of BACE (and consequently A $\beta$ 42 peptides and amyloid plaque burden) also reflects a compensatory or regulatory feedback mechanism stemming from the lower levels of its substrates, APP (47). Cholesterol reduction also appears to augment OA's direct effect on APP processing, as cholesterol reduction has been shown to promote the  $\alpha$ -secretase pathway, elevate sAPP $\alpha$  (23) and reduce 12-kDa CTF $\beta$  (10). Conversely, cholesterol retention enhances  $\beta$ -secretase (19, 42) and  $\gamma$ -secretase activities and corresponding A $\beta$  production (9, 45). While the overall reduction of A $\beta$ 42 peptides following OA enrichment should be enough to diminish the brain amyloid plaque burden, the concurrent decrease in the levels of PS1-NTF observed following OA enrichment also supplemented this outcome. In addition to the effects of OA on decreasing PS1-NTF levels, a previous study showed that lysosomal cholesterol levels are the strong inducers of PS2 transcription (7); therefore, lower dietary cholesterol in a low-fat/OA diet may also have suppressed the levels of PS1-NTF. Nevertheless, as NCT is likely the limiting factor in functional PS/ $\gamma$ -secretase complex assembly rather than PS1 (which could be present in excess without any effect on  $\gamma$ -secretase activity) (5), this suggests that the activity of the PS/ $\gamma$ -secretase complex should not be necessarily reduced with a decrease in PS1 levels. Although no significant change was observed on the levels of NCT, nevertheless, OA's effects on other members of the  $\gamma$ -secretase complex, such as Pen2 or Aph-1, cannot be ruled out.

In order to find out the involvement of additional factors in decreasing A $\beta$  levels and amyloid plaques, we next checked the proteins responsible for A $\beta$  degradation, such as IDE, insulin-like growth factors (IGFs) and TTR. A reduction in A $\beta$  levels is likely to be partially enhanced by an increased degradation of these peptides as a consequence of an over 50% increment in brain IDE levels following OA enrichment. This agrees well with the diminishment in the levels of IDE that is inversely correlated with increased levels of A $\beta$ 42 in transgenic mice (16) and human mutant PS2 and PS1 (V97L) expressing cells (34), and perhaps has resulted from the reduced levels of its substrate (35), insulin caused by a low-fat diet (1) in low-fat/OA mice.

Elevated levels of circulating insulin (such as in diabetes) provoke amyloid accumulation by directly competing with A $\beta$  for the IDE, thereby limit A $\beta$  degradation by IDE (15). The significant elevation in IGF-II levels in low-fat/OA mice supports and extends earlier findings, where IGFs (IGF-1 & IGF-II) have been demonstrated to protect against A $\beta$  toxicity *in vitro* (8). In addition, OA, along with LA, has been shown to enhance the growth-promoting effects of IGF-I (4). Alternatively, or in conjunction with OA's direct effect on IGF-II, increased levels of sAPP $\alpha$  have also been shown to drive the expression of IGF-II and to protect against A $\beta$ -induced neuronal death (38).

Finally, an interesting and important observation from these studies was that cholesterol depletion (low fat/OA) alone

imparted significant protective effects on the levels of factors directly or indirectly related to APP processing or metabolism. The AD brains on a low-fat/OA diet had dramatic decreases of BACE, A $\beta$ 42 peptides and amyloid plaque burden, which may reflect a compensatory or regulatory feedback mechanism stemming from the lower levels of its substrates, APP. Along similar lines, the AD brains on a low-fat/OA diet showed significantly higher levels of sAPP $\alpha$  and IDE. While the exact underlying basis for this particular effect is unknown, cholesterol deficiency and a low-fat diet are most likely the factors responsible for the shift in the non-amyloidogenic processing of APP (reviewed in reference 44) and elevated IDE levels (1), respectively.

We argue that the OA metabolism either directly or indirectly caused by SCD-1 led to a decrease in the levels of BACE and consequently, CTF $\beta$  and A $\beta$ 42. At the same time, OA, along with a low-fat diet, decreased circulating insulin levels and increased the availability of IDE to metabolize A $\beta$ . Therefore, the major factor leading to decreased brain levels of A $\beta$  in low-fat/OA transgenic mice is not only its decreased production (APP, BACE, PS1 levels) but also the decreased tissue accumulation (increased IDE and IGF-II). Dietary strategies aimed at reducing A $\beta$  levels should take into account interactions of dietary components and the metabolic outcomes, in particular, levels of protein, fat, total calories and enrichment of beneficial FAs.

## ACKNOWLEDGMENTS

Grants from the London and Middlesex Alzheimer's Association (Marion and Chester Fish Alzheimer's Disease Research Grant), the Canadian Institutes of Health Research (MOP49546) to R. F. R., Alberta Heritage Foundation for Medical research to D.W. and an Ontario Mental Health Foundation award to Z. A. funded this research. The authors declare that they have no conflict of interest.

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