Amyloid- β Peptide A β 3pE-42 Induces Lipid Peroxidation, Membrane Permeabilization, and Calcium Influx in Neurons^{*}

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Pyroglutamate-modified amyloid- β (pE-A β) is a highly neurotoxic amyloid- β (A β) isoform and is enriched in the brains of individuals with Alzheimer disease compared with healthy aged controls. Pyroglutamate formation increases the rate of AB oligomerization and alters the interactions of A β with Cu²⁺ and lipids; however, a link between these properties and the toxicity of pE-A β peptides has not been established. We report here that A β 3pE-42 has an enhanced capacity to cause lipid peroxidation in primary cortical mouse neurons compared with the fulllength isoform (A β (1-42)). In contrast, A β (1-42) caused a significant elevation in cytosolic reactive oxygen species, whereas A β 3pE-42 did not. We also report that A β 3pE-42 preferentially associates with neuronal membranes and triggers Ca²⁺ influx that can be partially blocked by the N-methyl-D-aspartate receptor antagonist MK-801. Aß3pE-42 further caused a loss of plasma membrane integrity and remained bound to neurons at significantly higher levels than A $\beta(1-42)$ over extended incubations. Pyroglutamate formation was additionally found to increase the relative efficiency of AB-dityrosine oligomer formation mediated by copper-redox cycling.

 $A\beta^3$ peptides are found in every human brain; however, the concentration and composition of $A\beta$ peptide isoforms are distinctly different in healthy individuals and people with AD (1–3). Amino-truncated $A\beta$ peptides are abundant in the AD brain (4, 5) and increase in prevalence with disease progression (6). The process of $A\beta$ amino-truncation can occur via the actions of aminopeptidases on full-length $A\beta$ pep-

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tides (7, 8), via altered cleavage of amyloid precursor protein in the generation of A β (9–11), and potentially by A β -copper-redox cycling reactions (12). As a consequence, aminotruncation can expose glutamate residues (positions 3 and 11 of A β) to cyclization by the action of glutaminyl cyclase (QC), forming the highly amyloidogenic pyroglutamate-A β (pE-A β) peptides A β 3pE-40, A β 3pE-42, A β 11pE-40, and A β 11pE-42 (7, 13).

Pyroglutamate formation significantly increases the hydrophobicity of A β , causing the peptide to aggregate more rapidly and form oligomers at lower concentration thresholds (5, 14, 15). pE-A β peptides also demonstrate increased β -sheet (aggregate structure) stability (16, 17), differences in fibril ultrastructure (18, 19), and altered interactions with copper ions (20, 21) and synthetic lipid membranes (22, 23). Notably, trace quantities of A β 3pE-42 have been observed to dramatically enhance the aggregation and neurotoxicity of $A\beta(1-42)$ (24), prompting descriptions of pE-A β as "prionlike." Still, it remains unclear as to the cytotoxic potency of pE-AB peptides compared with their full-length AB counterparts. Some studies have demonstrated pE-A β peptides to have enhanced toxicity (24-26), although others have reported no difference in toxicity between the isoforms (27-30). Methodological differences may account somewhat for variability in the relative toxicities reported (Table 1), yet molecular mechanisms to explain changes in cytotoxicity have not been defined.

One mechanism through which A β peptides cause cytotoxicity is by production of reactive oxygen species (ROS) via facile copper-redox cycling (31–33), which can in turn effect oxidative damage to neuronal proteins and lipids (34). Imbalances in ROS production and detoxification are strongly implicated in AD neurodegeneration, reflected by cerebral elevations in oxidized DNA, lipids, and proteins (35-37). Pyroglutamate formation alters $A\beta$ -Cu²⁺ coordination modes (20, 21), although it is not known whether this affects the capacity of pE-A β peptides to undertake redox cycling and produce cytotoxic ROS. We therefore aimed to determine whether full-length $A\beta$ and pE-A β possess differences in their capacity to alter ROS flux and cause oxidative damage to neurons in vitro. Additionally, A β isoforms were compared for their capacity to form oligomers and covalent tyrosine-tyrosine bonds (dityrosine) as a result of A β -copper-redox cycling. The capacity for A β to form dityrosine has previously been correlated with neurotoxicity (38), although recent reports have found that $A\beta$ fibrils within



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³ The abbreviations used are: Aβ, amyloid-β; NMDAR, N-methyl-D-aspartate receptor; AD, Alzheimer disease; ROS, reactive oxygen species; QC, glutaminyl cyclase; pE-Aβ, pyroglutamate-Aβ; HBSS, Hanks' balanced salt solution; BisTris, 2-[bis(2-hydroxyethyl)amino]-2-(hydroxymethyl)propane-1,3-diol; DIV, days *in vitro*; DPPC, diphenyl-1-pyrenylphosphine; DCF, dichlorofluorescein; 3-CCA, coumarin-3-carboxylic acid; SEC, size-exclusion chromatography; AFM, atomic force microscopy.

TABLE 1

Overview of publications comparing the cytotoxicity of pE-A β and full-length A β peptides in vitro

The following abbreviations are used in table: CSF, cerebrospinal fluid; DMSO, dimethyl sulphoxide; HFIP, 1,1,1,3,3,3-hexafluoro-2-propanol; LDH, lactate dehydrogenase; PBS, phosphate buffered saline.

Reference	Aβ Stock Preparation	Aβ Treatment	Toxicity Measurements	Results
Tekirian et al, 1999 (27)	Lyophilized in 1 mM HCl then redissolved in cell culture media or PBS (pH 7.4)	Fresh peptides dissolved in cell culture media to $5 - 50 \mu$ M. Aged peptides (250μ M) incubated in PBS for 1 week at 37 °C, diluted into media.	Phase-contrast micrographs of primary cortical rat neurons taken after 0, 3, 6, 12 and 24 h treatment. Cells that were smooth, round and vacuole free scored as 'alive'.	No difference in toxicity between A β 1-40 and A β 3pE-40, nor A β 1-42 and A β 3pE-42 at any time-point, fresh or aged.
Russo et al, 2002 (49)	DMSO, 1 mM Aβ stock	A β oligomerized for 3 days (37 °C) in cell media and applied to cells at 0.1, 1.0 or 10 μ M for 24 h.	Cortical astrocytes and hippocampal neurons viability measured by mitochondrial function (MTT assay) and cell permeability (LDH leakage)	A β 3pE-40 more toxic than all other peptides on astrocytes & neurons (MTT assay), no difference between other peptides. A β 3pE-40 & A β 3pE-42 induce more LDH leakage from astrocytes than A β 1-40 & A β 1-42 - no difference for neurons.
Shirotani et al, 2002 (28)	Expression of APP constructs to produce different Aβ isoforms in primary cortical mouse neurons	Neurons predominantly expressed either $A\beta I$ -X, $A\beta 3$ -X or $A\beta 3$ pE-X, where X= residues 34 – 42 of $A\beta$ sequence.	Cell viability measured by cellular dehydrogenase activity (WST-8 assay). Tolerance to H ₂ O ₂ . Calpain activation.	Same decrease in cell viability between all constructs compared to control after 3 days of expression. No difference in tolerance to H_2O_2 or levels of Calpain activation between APP constructs.
Youssef et al, 2008 (29)	Peptides dissolved to 5 mg/mL in HFIP, dried, then re- dissolved in cell culture media.	A β peptides incubated in cell culture media for 1 h (room temperature). Incubated on cells at 1 μ M (24 – 48 h) or injected in brains (500 pmol total, for 2 – 14 days).	Primary cortical mouse neurons viability measured by caspase induction, MTT and calcein assays. ROS measured by dichlorofluoroscein assay 2 days post- injections, behavior day 14.	Aβ1-42 and Aβ3pE-42 oligomers showed same levels of neurotoxicity, ROS production, caspase induction and impairment of spatial memory.
Nussbaum et al, 2012 (24)	A β dissolved in HFIP to 1 mM, diluted in cell culture media and HFIP removed by air stream.	A β (5 μ M) in culture media incubated 24 h at 4 °C, that 24 h on cells at ≤ 1 μ M. 1:19 mixtures of A β 3pE-42:A β 1-42 made from peptides either incubated together, or separately then mixed, before applying to cells.	Primary cortical mouse neuron viability tested after 24 h treatment by mitochondrial function (XTT assay) and calcein-AM vital stain.	Aβ3pE-42 and 1:19 mixtures (3pE-42:1-42) significantly more toxic than Aβ1-42 (only when peptides incubated together as a mixture before applying to cells). Oligomer mixtures maintained neurotoxicity after serial dilution in fresh Aβ1-42 to contain only 1 part Aβ3pE-42 per 8000 Aβ1-42.
Galante et al, 2012 (25)	Aβ dissolved in DMSO, lyophilized by vacuum centrifugation then resuspended in 1% NH₄OH and diluted in PBS	Peptides separated into small (4.5 kDa), medium (14 kDa) and large (99 kDa) aggregates, then incubated on cells at 100 nM for up to 4 days	SH-SY5Y human neuroblastoma viability measured by MTT assay and xCELLigence (Roche) real-time viability monitoring.	Medium and large aggregates of both $A\beta1-42$ and $A\beta3pE-42$ were significantly toxic to neurons, however $A\beta3pE-42$ small aggregates were additionally toxic while $A\beta1-42$ small aggregates were not.
Schlenzig et al, 2012 (26)	Aβ dissolved in HFIP, dried by evaporation then redissolved in DMSO.	HFIP treated peptides re-dissolved to 50 μ M in DMSO then diluted in artificial-CSF to 250 or 500 nM and incubated on hippocampal slices.	Mouse hippocampal slice long-term potentiation (LTP) measured following 2 h exposure to synthetic peptides in carbogenated artificial-CSF.	A β 3pE-40 significantly more inhibitory to LTP than A β 1-40 at 500 nM. Likewise, A β 3pE-42 significantly more inhibitory than A β 1-42 at 250 nM.
Bouter et al, 2013 (30)	A β dissolved in HFIP, frozen, then HFIP evaporated. Stocks dissolved in NaOH (100 mM) to 2 mg/mL stored at -80 °C until use	A β stocks freshly diluted to 1, 5 or 10 μ M in DMEM/F12 media and applied to cell cultures for 24 h.	Cortical rat neurons cultured in 48-well plates, viability measured by calcein-AM vital stain using fluorescence plate reader.	Identical toxicity observed for $A\beta I$ -42, $A\beta 3pE$ -42 and $A\beta 4$ -42 at all concentrations tested.

amyloid plaques contain intense dityrosine immunoreactivity (39), indicating that dityrosine formation may be associated with AD amyloidogenesis. Further comparisons were made between the peptides for their capacity to perturb neuronal membranes and induce changes in neuronal ion homeostasis (Ca^{2+} flux).

Experimental Procedures

Materials— $A\beta$ peptides were purchased from the ERI Amyloid Laboratory, LLC, and purified by reversed-phase HPLC to >95% purity. All chemicals used were analytical grade and purchased from ChemSupply (Australia) unless otherwise stated. Thioflavin-T, coumarin-3-carboxylic acid (3-CCA), butylated hydroxytoluene, bovine serum albumin, and paraformaldehyde were purchased from Sigma. Diamsar (1,8-diaminosarcophagine) was synthesized as reported previously (40).

Preparation of Aβ Solutions and Cu²⁺-oxidized Aβ Oligomers—Aβ stock solutions were prepared by dissolving lyophilized peptides to 5 mg/ml in NaOH (60 mM) and incubating at ambient temperature for 5 min to dissociate aggregated material. Solutions were then diluted to 1 mg/ml in MilliQ H₂O and 10× PBS (PBS is defined as 50 mM sodium phosphate, 2.7 mM KCl, 137 mM NaCl) at a v/v/v ratio of 2:7:1 (NaOH, H₂O, 10× PBS). The preparation was sonicated for 10 min in an iced water bath and then centrifuged at 16,500 × g for 10 min at 4 °C. Supernatants (upper 80% of solution) were removed to prechilled tubes on ice; pH was adjusted by addition of NaH₂PO₄ (0.5 M) to 1.0% v/v and then kept on ice for immediate use. Aβ stock concentrations were determined by UV spectrometry by measuring absorbance at 214 nm (A_{214}) and applying the following extinction coefficients (M^{-1} cm⁻¹) determined from UV scans and amino acid analysis: A β (1–40) = 91,462; A β 3pE-40 = 89,705; A β (1–42) = 94,526; and A β 3pE-42 = 90,925.

A β oligomers were generated by reacting A β (10 μ M) with Cu²⁺ (10 μ M, as CuCl₂·glycine₆) and ascorbate (100 μ M) at 37 °C on a vertically rotating wheel at 20 rpm, in 2-ml round-bottom tubes (catalog no. 0030123.344, Eppendorf). Reactions were halted with the addition of EDTA (to 250 μ M) and placed on ice. Control incubations of A β only were incubated and sampled under identical conditions for comparison. A β fibrillization assays (thioflavin-T assays) were performed as described previously (41).

Detection of Cell-free Hydroxyl Radical Production-Hydroxyl radical production in AB-copper-redox cycling reactions was assessed with the fluorescent probe coumarin-3carboxylic acid (3-CCA) using methodology adapted from Manevich *et al.* (42). Reactions containing A β peptides (20 μ M) and 3-CCA (100 μ M) in the presence or absence of Cu²⁺ (20 μ M, as CuCl₂·glycine₆) and ascorbate (300 μ M) in PBS, pH 7.4, were incubated in black-walled clear bottom microplates (PerkinElmer Life Sciences) at 37 °C in a heated Flexstation III spectrophotometer (Molecular Devices). Fluorescence signals were read from the bottom of the microplate every 30 s using excitation λ 395 nm and measuring emission at λ 450 nm. Rate constants for the 3-CCA reactions were determined from one-phase association regressions of logY transformed data (data manipulations performed in GraphPad Prism version 6.0).



Aβ-Dityrosine Determination and Hydrophobic Index Calculation—The dityrosine content of Cu²⁺-reacted Aβ samples was determined by fluorescence spectrophotometry (λ 320 nm excitation and λ 420 nm emission), as described previously (43). Reaction half-times and rate constants ($K = \min^{-1}$) were calculated from one-phase association regressions of fluorescence data (GraphPad Prism version 6.0). Calculation of Aβ hydrophobic scores were determined from the sum of the amino acid residue hydropathic indexes (44). Pyroglutamyl residues were assigned a hydropathy value of -1.0.

*A*β *Detection by Western Blot*—Tissue extracts and synthetic peptides were separated by SDS-PAGE using 4-12% XT Bis-Tris gels (Criterion, Bio-Rad) according to the manufacturer's instructions. Samples were transferred to pre-assembled PVDF membrane stacks using a Trans-Blot® semi-dry transfer apparatus (Bio-Rad). Blots were blocked in TBS-T (10 mM Tris-HCl, 50 mM NaCl, 0.1% v/v Tween 20, pH 8.0) containing 5% w/v skim milk. Primary antibodies were incubated on blots for at least 1 h at room temperature or overnight at 4 °C. HRP-conjugated rabbit anti-mouse or goat anti-rabbit immunoglobulins (Dako) were diluted 1:10,000 in TBST and incubated for 1 h at room temperature. All antibodies were diluted in TBS-T containing 5% skim milk and 0.05% w/v sodium azide. Blots were washed four times for 10 min in TBS-T after each primary and secondary antibody binding step. Chemiluminescence signals were captured after application of ECL (Immobilon, Millipore) with an LAS3000 detector and analyzed using MultiGauge software (Fujifilm). A β peptides were detected using the monoclonal mouse antibodies 4G8 (Covance) or 6E10 (Signet Laboratories) diluted to 1 μ g/ml. The dityrosine modification was detected using the 1C3 antibody raised against synthetic dityrosine (catalog no. NWA-DIY020, Northwest Life Science Specialties), at a 1:1000 dilution.

Size-exclusion Chromatography (SEC) and Atomic Force Microscopy (AFM)—A β oligomers in HBSS buffer were prepared as above but at twice the concentration (20 μ M A β , 20 μ M Cu²⁺, 200 μ M ascorbate) to provide adequate signals for measurement.

SEC analysis was performed using a BioLogic DuoFlow QuadTec 40 system (Bio-Rad) fitted with a Superdex 75 10/300 column (catalog no. 17-5174-01, GE Healthcare). Both equilibration and operation of the column were in Tris-buffered saline (20 mM Tris, 200 mM NaCl, pH 8.0, 0.2 μ m filtered and de-gassed) at a flow rate of 0.5 ml/min and ambient temperature. The absorbance at 214, 260, and 280 nm was monitored, collecting 5 data points/s. Samples were injected onto the column (0.5 ml per run, ~45 μ g of A β) immediately after collection at indicated time points.

AFM analyses were performed on A β reactions prepared in Neurobasal medium (catalog no. 21103-049, Gibco) at 10 μ M. Samples (10 μ l) were collected and spotted on freshly cleaved mica, dried at room temperature for 5 min in a laminar flow hood, and rinsed with 2 ml of de-ionized water (Milli-Q, Millipore). The sample was blown dry with nitrogen (Coregas Nitrogen 4.0) before being transferred to the AFM sample stage (Asylum Research Cypher AFM). Images were acquired in alternating current (tapping) mode in air using Tap300-G silicon AFM probes (Budget Sensors) with scan rates of 1.5–2.5 Hz; drive amplitude was kept to a minimum with amplitude set-points of 60-80%. The mask threshold was set to 250 pm for image analysis.

Primary Neuronal Cell Culture and $A\beta$ Clearance Assays— All experiments involving animals were conducted in accordance with the Australian Code of Practice for the Use of Laboratory Animals and were approved by the Institutional Animal Experimentation Ethics Committee.

Mouse cortical neuronal cultures isolated from C57Bl/6 E14 embryos were prepared as described previously (45). Cells were plated in poly-D-lysine-coated 48- or 96-well plates at a density of 150,000 cells/cm². All cell culture materials were purchased from Gibco/Thermo Fisher unless otherwise stated. Cells were grown in Neurobasal medium (NB) containing B27 supplements, gentamicin, and 0.5 mM GlutaMAXTM. Fresh NB medium containing B27 minus antioxidants (B27-AO) and cytosine- β -D-arabinofuranoside (2 μ M; catalog no. C1768, Sigma) were applied after 6 days *in vitro* (DIV). Neurons were further incubated until DIV 8 or 9 prior to applying treatments.

Levels of cell-bound A β were measured following exposure to A β peptides applied in NB medium containing B27-AO and cytosine- β -D-arabinofuranoside. Cells (DIV 8 or 9) were treated for 48 h with A β mixtures (10 μ M total). The media were removed, and cells were washed twice with Dulbecco's PBS (catalog no. 14190-144, Gibco), and then the cells were extracted with M-PER reagent (catalog no. 78501, Thermo-Scientific). For fractionation studies, cells were scraped into TBS, pH 7.5, containing protease inhibitors (cOmplete, catalog no. 11873580001, Roche Applied Science) and probe-sonicated by two rounds of 10 bursts (40% power, 0.5 s each) with a Sonifier S-250D (Branson). Lysates were centrifuged at 100,000 \times g, and supernatants (TBS phase) were collected to fresh tubes. The pellets were extracted with an equal volume of Na₂CO₃ (100 mM, pH 12) for 1 h of incubation on ice and then briefly vortexed and centrifuged again to collect supernatants (carbonate phase). Pellets were resuspended in an equal volume of urea buffer (7 M urea, 2 M thiourea, 4% CHAPS, 1% DTT, 50 mM Tris, pH 8.0). Western blots were performed on 5 μ g of total protein based on BCA assays (catalog no. 23225, Thermo-Scientific) of TBS phases, loading equal volumes for carbonate and urea phases.

Cell Viability Measurement-M17 neuroblastomas were cultured in DMEM containing fetal bovine serum (FBS; 10% v/v) and penicillin/streptomycin (catalog no. 15140, Gibco). Cells were plated in 6-well culture plates (catalog no. 140675, Nunc) at 40,000 cells/cm² and incubated overnight (37 °C, 5% CO₂). Fresh medium supplemented with retinoic acid (10 μ M) was applied, and the cells were allowed to differentiate for 48 h. Freshly prepared A β solutions (10 μ M) or synthetic lipid peroxide (1 μ M) were applied to cells for 4 h and then the cells were collected using trypsin and resuspended in 1 ml of PBS containing 2% FBS. Propidium iodide was added to a final concentration of 1 μ g/ml. Flow cytometric analysis was performed on a Beckman Coulter CyAn ADP analyzer with Summit 4.0.3 acquisition software. Debris and cell aggregates were excluded by gating. Live and dead cells were then identified based on exclusion and inclusion, respectively, of propidium iodide. Analysis of data were performed using FCS Express 4 analysis software.

Detection of Neuronal ROS Flux and Lipid Peroxidation— Neuronal ROS flux was measured using the redox-sensitive probe 2,7-dichlorofluorescin diacetate (catalog no. D399, Molecular Probes), essentially as described previously (46). Briefly, 2,7-dichlorofluorescin diacetate (100 μ M) was applied to cells (DIV 9) in NB medium supplemented with B27-AO (37 °C, 5% CO₂) for 60 min; the media were then removed, and the cells were washed with Hanks' balanced salt solution (HBSS; catalog no. 14175-079, Gibco). Freshly prepared A β peptides were diluted to 10 μ M in HBSS and applied to the neurons for a further 60 min of incubation (37 °C, 5% CO₂, light protected) until reading DCF fluorescence (λ 485 nm excitation, λ 530 nm emission, and λ 515 nm emission cutoff filter) with a Flexstation III spectrophotometer (Molecular Devices) equilibrated to 37 °C.

Lipid peroxidation was measured using the lipid hydroperoxide probe diphenyl-1-pyrenylphosphine (DPPP; catalog no. 62237, Cayman Chemicals), with methodology adapted from Ref. 47. DPPP (stock dissolved in N₂-purged DMSO to 25.88 mm) was diluted to 50 μ m in NB media and applied to cells (DIV 8-9) in 96-well black-walled microplates (Greiner) and then incubated for 60 min at 37 °C, 5% CO₂ in the dark. The DPPPcontaining media were removed; the cells were rinsed twice with HBSS and then equilibrated in HBSS at 37 °C and 5% CO₂ for 30 min before treatment. Freshly prepared A β peptides were applied to cells for 4 h until reading fluorescence (λ 340 nm excitation and λ 380 nm emission) with an EnSpire multimode spectrophotometer (PerkinElmer Life Sciences). Positive control wells were treated with a synthetic lipid hydroperoxide standard (catalog no. 705014; Cayman Chemicals) for comparison. Care was taken at all steps to minimize artifactual oxidation of DPPP by reducing exposure to light.

Neuronal Calcium Flux Measurement-Cortical neurons plated at 230,000 cells/cm² in 96-well black-walled microplates (Greiner) were loaded with the fluorescent Ca²⁺ sensor Fluo4-AM (catalog no. F14201, Molecular Probes) at DIV9 for 30 min at 37 °C, 5% CO₂, and then equilibrated to room temperature for 30 min. Fluorescence (excitation λ 490 nm and emission λ 520 nm) was measured at base line for 10 reads at 27-s intervals followed by a further 10 reads after injection of glutamate or A β (final concentrations of 1 and 10 μ M, respectively) using a Fluostar plate reader (BMG Labtech). Ca²⁺ flux values ($\Delta F/F_0$) were expressed as the difference between mean baseline and immediately following glutamate or A β application. Experiments determining the effect of metal depletion utilized neuronal cultures pre-treated for 1 h with Diamsar (10 µM). To assess the contribution of NMDAR signaling in A β -induced Ca²⁺ flux, cultures were pre-treated for 15 min with the NMDAR antagonist MK-801 (1 mM). In separate experiments the supplied assay buffer (HBSS) was replaced with Ca^{2+}/Mg^{2+} -free HBSS to determine the source of Ca^{2+} entering the cytosol.

Results

Pyroglutamate Formation Alters the Production of ROS and Dityrosine by $A\beta$ —pE-A β peptides demonstrated greatly increased fibrillization rates compared with the respective fulllength isoforms (Fig. 1*A*), consistent with previous reports (14, 18). $A\beta(1-40)$, $A\beta(1-42)$, $A\beta$ 3pE-40, and $A\beta$ 3pE-42 were also compared for their capacity to produce ROS upon reaction with Cu^{2+} and ascorbate. The rate of 'OH production was significantly higher for A β (1–42) compared with A β 3pE-42 (Fig. 1*B*). In contrast, A β 3pE-40 produced more 'OH than A β (1–40) and indeed all other A β isoforms. The kinetics of 'OH production showed a similar sigmoidal pattern for all peptides with signals reaching plateau after ~20 min.

Differences in redox cycling and 'OH production between AB peptides were further studied to compare formation of dityrosine bonds. Dityrosine formation rates partially reflected 'OH production rates whereby Aβ3pE-40 formed dityrosine more rapidly than $A\beta(1-40)$ (half-times of 5.44 and 33.64 min, respectively), whereas A $\beta(1-42)$ formed dityrosine more rapidly than A β 3pE-42 (half-times of 5.38 and 7.31 min, respectively) (Fig. 1, *C* and *D*). Total A β -dityrosine content, however, did not mirror the levels of 'OH produced by each peptide; $A\beta(1-42)$ formed more than double the amount of dityrosine than all other peptides. A direct relationship was not observed between the dityrosine formation rate and relative A β hydrophobicity (Fig. 1E); however, the efficiency of dityrosine formation (calculated as a ratio of dityrosine formed per unit of 'OH generated) was increased for both pE-A β peptides relative to their full-length counterparts (Fig. 1F).

Comparison of the Oligomerization of Pyroglutamate-A β and Full-length A β in the Presence and Absence of Cu²⁺—Previous studies have shown differences between A β (1–40) and A β (1– 42) in oligomerization and dityrosine formation when reacted with Cu²⁺ and biological reductants (48). We compared the profiles of full-length A β and pE-A β oligomers generated in the presence or absence of Cu²⁺ and ascorbate. A β (1–42) and A β 3pE-42 rapidly formed SDS-stable oligomers within 5 min of reaction with Cu²⁺, whereas increases in A β (1–40) and A β 3pE-40 oligomers were observed at 15 min (Fig. 2A). Significant formation of SDS-stable oligomers was not observed for A β incubated in the absence of Cu²⁺ and ascorbate over the same time period.

Dityrosine cross-links were also detected in $A\beta(1-40)$ and $A\beta$ 3pE-40 reactions by immunostaining with a dityrosine-specific antibody (1C3), revealing predominantly dimer and trimer-sized (~8/12 kDa) oligomers and confirming the enhanced capacity for $A\beta$ 3pE-40 dityrosine formation compared with $A\beta(1-40)$ (Fig. 2*B*). The 1C3 antibody, however, did not show significant immunoreactivity to Western blots of Cu²⁺-reacted $A\beta(1-42)$ and $A\beta$ 3pE-42 peptides (data not shown).

We additionally examined the oligomer states of $A\beta(1-42)$ and $A\beta$ 3pE-42 in non-denaturing conditions by SEC and AFM. After a 4-h incubation in PBS, the $A\beta(1-42)$ oligomer profile did not change considerably from the freshly prepared solution, resolving predominantly as a single low-mass peak (<14 kDa) by size exclusion, with minor signals corresponding to oligomers of ~40-75 kDa (Fig. 3*A*). Addition of Cu²⁺ and ascorbate to $A\beta(1-42)$ caused a sizable increase in the abundance of oligomers but did not appear to alter their mass. By comparison, $A\beta$ 3pE-42 ternary states changed considerably during the 4-h incubation, observed as a dramatic loss of low-mass peak (<14 kDa) in either the presence or absence of Cu²⁺ and ascorbate (Fig. 3*B*). After a 4-h incubation, residual signals remaining in





FIGURE 1. **Comparison of the aggregation, redox cycling, and dityrosine formation capacities of A** β **peptides.** *A*, A β (5 μ M) fibrillization kinetics measured by thioflavin-T assay (data cropped at start of plateau due to signal decay as A β sediments in microplate wells). *B*, kinetics of 'OH formation in reactions of A β (20 μ M) with CuCl₂ (20 μ M) and ascorbate (300 μ M) as measured by 3-CCA fluorescence assay. Kinetics of dityrosine formation in reactions of A β (10 μ M) with CuCl₂ (10 μ M) and ascorbate (100 μ M) A β (1–40) and A β 3pE-40 peptides (*C*), A β (1–42) and A β 3pE-42 peptides (*D*), measured by fluorescence spectrometry of dityrosine bond (320 nm excitation and 420 nm emission). *E*, relationship between A β hydrophobicity and dityrosine formation was further calculated as a ratio of the amount of dityrosine formed per unit of 'OH produced, plotted against A β hydrophobicity. All measurements were repeated a minimum of three times over separate days using freshly prepared A β solutions. Data shown are mean \pm S.E.

the A β 3pE-42 reactions without Cu²⁺ resolved predominantly as an oligomeric peak at ~30-45 kDa. In the A β 3pE-42 reaction containing Cu²⁺ there was some preservation of the monomeric signal after the 4-h incubation, with an additional polydisperse peak across the 30-75-kDa range.

To simulate A β oligomerization in cell culture conditions, reactions were conducted in neurobasal media over an extended incubation period, and the peptide structures were assessed by AFM (Fig. 3*C*). Little difference in peptide structure was observed during a 48-h incubation of A β (1–42) in unsupplemented neurobasal media (maximum *z* <1 nm). In the A β (1–42) reactions supplemented with Cu²⁺/ascorbate, there was similarly no noticeable change after 4-h incubation, although a large increase in the size and abundance of oligomers was observed after 48 h (maximum *z* = 4.9 nm). Likewise, A β 3pE-42 structures did not demonstrate a noticeable size difference over a 48-h incubation in neurobasal media in the

absence of supplemental Cu²⁺/ascorbate (maximum z < 1 nm), although the reactions supplemented with Cu²⁺/ascorbate showed a stepwise size increase over the incubation period (maximum z = 0.4, 1.0, and 1.8 nm at 0, 4, and 48 h, respectively).

 $A\beta 3pE-42$ Predominantly Associates with Neuronal Membranes and Is Resistant to Clearance—pE-A β has been found to resist proteolytic clearance in astrocyte cultures (49, 50). We therefore compared the residual levels of $A\beta(1-42)$ and $A\beta 3pE-42$ following extended exposures to cortical neuron cultures, additionally assessing the capacity for minor quantities of $A\beta 3pE-42$ to affect the clearance of $A\beta(1-42)$. Neuronal $A\beta$ levels were strikingly different after 48 h of treatment with the two peptides; relative levels of $A\beta 3pE-42$ were 4.5-fold higher than cells treated with $A\beta(1-42)$ (Fig. 4A). A trend toward higher mean levels of residual $A\beta$ was observed when $A\beta(1-42)$ solutions were supplemented with either 50 or 25%



FIGURE 2. *A*, Western blots stained with 4G8 anti-A β antibody demonstrating rapid formation of SDS-stable low molecular weight A β oligomers catalyzed by reaction with Cu²⁺ and ascorbate (*Asc.*). *B*, comparison of A β (1–40) and A β 3pE-40 dityrosine oligomer profiles. Western blots were stained with 1C3 anti-dityrosine antibody.

(mol/mol) A β 3pE-42; however, this effect was not observed when A β 3pE-42 represented only 5% of the total A β concentration (Fig. 4*A*).

From experiments utilizing sequential extraction of cells following A β exposure, both the A β (1–42) and A β 3pE-42 peptides were predominantly found in membrane fractions, with relatively little A β in the soluble (TBS) phase observed qualitatively by Western blot (Fig. 4*B*). A β in cell extracts resolved primarily as monomers on SDS-PAGE; however, bands corresponding to low-mass oligomers (~8–20 kDa) were observed in longer blot exposures.

Toxicity of Pyroglutamate-A β Is Associated with Membrane Damage but Not Cytosolic ROS Production—The capacities of each peptide to produce elevations in cytoplasmic and membrane ROS were studied using primary cortical mouse neurons. The fluorescent ROS probe DCF was used to measure cytoplasmic ROS flux over a 1-h treatment with freshly prepared A β peptides. A β (1–42) induced an approximate 10-fold increase in DCF fluorescence compared with control cells, whereas A β (1–40), A β 3pE-40, or A β 3pE-42 treatments were not found to be statistically different from controls (Fig. 5, A and B).



FIGURE 3. Comparison of oligomer structures formed in the presence or absence of Cu²⁺ and ascorbate. A β (1–42) (20 μ M) (A) and A β 3pE-42 (20 μ M) (B) solutions incubated at 37 °C for 4 h in HBSS, pH 7.4, \pm Cu²⁺ (20 μ M) and ascorbate (200 μ M) were assessed for oligomer formation by SEC using a Superdex 75 column. *Main figure* shows the region of >14 kDa; *insets* show the full chromatogram with molecular mass standards (214 nm UV detection). *C*, AFM images of A β peptides (10 μ M) incubated for indicated times in Neurobasal medium. Each image displays a 2 × 2 μ m field and 10 nm total *z*-range; *insets* display a 200 × 200 nm field and 2 nm total *z*-range. SECs represent the same profile from a minimum of n = 3 independent replicates. AFM analyses were repeated on independent reactions over separate days; representative data are shown.

Minor quantities of A β 3pE-42 were found to significantly disrupt the neuronal ROS-inducing capacity of A β (1–42); the ROS flux induced by solutions of A β (1–42) was diminished by 47% when A β 3pE-42 was present at 5% (mol/mol) of total A β (Fig. 5*B*).

To assess $A\beta$ induction of ROS at neuronal membranes, we directly measured lipid peroxidation in living cortical mouse neurons using the fluorescent probe DPPP over a 4-h treatment. In direct contrast to the DCF assays, $A\beta(1-42)$ (10 μ M) did not increase neuronal DPPP fluorescence, whereas cells treated with $A\beta$ 3pE-42 (10 μ M) demonstrated a statistically sig-





FIGURE 4. Freshly prepared Aß peptides were applied to cortical mouse neurons and levels of residual A β remaining cell-bound and intact after 48 h of incubation were measured by Western blot (4G8 antibody) to model A β proteolytic clearance/resistance. A, A β (1–42), A β 3pE-42, or mixtures thereof were quantified by densitometric analysis of bands at 4-6 kDa; percentages indicated in *column titles* represent the proportion of A β 3pE-42 mixed with A β (1-42) to give a final concentration of 10 μ M. A β levels are reported as relative densitometry signals compared with cells treated with $A\beta(1-42)$ alone, as $A\beta$ signals were not observed in vehicle control wells in these conditions. B, cell extracts 48 h after application of freshly prepared A β peptides to cortical neurons were fractionated to separate soluble A β (TBS extracted), peripheral membrane-bound A β (carbonate phase), and integral membrane-bound A β (urea phase). A β monomers are apparent at the approximate molecular mass of A β (4 –5 kDa), and A β oligometric signals are identified by arrowheads (contrast enhanced panel). 4G8-reactive signals above 35 kDa were present in all groups (including vehicle control) suggesting that these signals are not A β but may correspond to the amyloid precursor protein and its cleavage products in the cortical neurons. All densitometry signals were corrected for protein loading by re-probing blots for actin. Samples shown are representative of three separate fractionations per group; panels have been cropped from the same blot image to show only relevant lanes. Data are mean \pm S.E., statistical significance is relative to A β (1-42)treated group.** represents $p \le 0.01$.

nificant 35% increase in lipid peroxidation, comparable with levels induced by 1 μ M synthetic lipid hydroperoxide (Fig. 5*C*). Further contrasting the DCF assays, minor quantities of A β 3pE-42 had no effect on the lipoperoxide-inducing capacity of A β (1–42).

To determine whether lipid peroxidation was associated with a loss of plasma membrane integrity, differentiated M17 neuroblastomas were measured for propidium iodide uptake by flow cytometry following a 4-h A β treatment. Consistent with the lipid peroxidation experiments, A β 3pE-42 induced significantly higher levels of membrane damage than A β (1–42) (Fig. 5*D*). A β (1–42) solutions seeded with 5% (mol/mol) A β 3pE-42 induced practically identical levels of plasma membrane damage as $A\beta(1-42)$ alone.

 $A\beta 3pE-42$ Induces Rapid Ca^{2+} Influx in Primary Cortical *Neurons*—Recent reports of pE-A β interactions with synthetic lipid membranes have suggested that pE-A β neurotoxicity is exerted via formation of ion-permeable membrane pores (22, 23). We compared the capacity of $A\beta(1-42)$ and $A\beta$ 3pE-42 to alter cellular ion homeostasis by measuring Ca²⁺ flux in cortical neurons immediately following application of freshly prepared A β . A β 3pE-42 (10 μ M) effected a rapid and significant elevation in neuronal Ca²⁺ levels as measured with the Fluo-4 fluorescent Ca^{2+} sensor, comparable with levels induced by 1 μ M glutamate (Fig. 6, A and B). By comparison, A β (1–42) did not cause significant elevation in Ca²⁺ levels above those of the vehicle control. When neurons were pre-treated with the NMDAR antagonist MK-801, there was an approximate 50% decrease in Ca²⁺ flux induced by both glutamate and A β 3pE-42, indicating that A β 3pE-42-induced Ca²⁺ flux is at least partially attributed to NMDAR activation (Fig. 6B). To determine whether A β 3pE-42-induced Ca²⁺ flux was dependent on A β metal interactions, the neuronal culture media were depleted of bioavailable first row transition metals with the chelator Diamsar. Rapid induction of Ca^{2+} flux by A β 3pE-42 was not affected by Diamsar pre-treatment, suggesting that the membrane perturbation was independent of $A\beta$ -copper-redox cycling (Fig. 6*C*). Aβ3pE-42 did not induce significant elevations in cytosolic Ca^{2+} when provided in Ca^{2+}/Mg^{2+} -free media, indicating that A β 3pE-42 causes Ca²⁺ to enter from the extracellular space and not via release from intracellular stores (Fig. 6D).

Discussion

 $A\beta$ in the human brain is represented by a heterogeneous and dynamic mixture of isoforms, with significant compositional variation between individuals (1–3, 5, 51). The predominance of pE-A β in the central core of plaques suggests an early involvement in amyloid deposition in the AD brain (52), whereas correlation between pE-A β and a decline in Mini Mental State Examination scores implicate pE-A β cytotoxicity in AD neurodegeneration (53, 54). In parallel, markers of oxidative stress are among the earliest detectable pathological changes in transgenic AD mouse models (55) and the human AD brain (36, 56, 57), with numerous lines of evidence implicating A β as a central contributor to oxidative stress in AD (58, 59).

Our data indicate that full-length $A\beta$ and pE- $A\beta$ exert different oxidative insults upon neurons, representing different mechanisms of neurotoxicity. $A\beta(1-42)$ was found to significantly increase cytosolic ROS levels, whereas $A\beta$ 3pE-42 induced lipid peroxidation in the absence of cytosolic ROS flux. Additionally, we found that that the neuronal membrane damage caused by $A\beta$ 3pE-42 results in a functional loss of plasma membrane integrity. These findings are consistent with recent publications demonstrating the capacity for $A\beta$ 3pE-42 to form membrane-disrupting pores in synthetic lipid bilayers (22, 23). Similarly, previous studies have shown that pE- $A\beta$ oligomers disrupt lysosome membrane integrity in cultured neurons (50) and cause lactate dehydrogenase leakage from cultured astrocytes (49).



FIGURE 5. **Comparison of the neurotoxicity profiles of freshly prepared** $A\beta$ **peptides.** Levels of ROS flux in cortical neurons loaded with the fluorescent probe dichlorodihydrofluorescein diacetate measured 60 min after treatment with $A\beta(1-40)$ and A3pE-40 peptides (A) or $A\beta(1-42)$ and $A\beta3pE-42$ peptides (B) (total $A\beta = 10 \ \mu$ M; percentages indicate amount of $A\beta3pE-40$ and $A\beta3pE-42$ peptides seeded into solutions of $A\beta(1-40)$ and $A\beta(1-42)$, respectively). *C*, levels of cortical neuron lipid peroxidation (*LPO*) following 4 h of treatment with $A\beta$ (10 μ M total) or synthetic lipid peroxide (1 μ M) measured with the DPPP fluorescent probe, represented as difference from baseline readings. *D*, cell viability (membrane integrity) of differentiated M17 neuroblastomas treated with $A\beta$ peptides (10 μ M) or lipid peroxide (1 μ M) for 4 h, measured by flow cytometry analysis of propidium iodide uptake. Each data set is representative of a minimum of three experiments performed over separate days (shown as mean \pm S.E.). *, $p \le 0.05$; **, $p \le 0.01$; *ns*, not significant, p > 0.05.

The earliest detectable effect of A β 3pE-42 on neuronal homeostasis that we observed was the capacity to cause rapid Ca²⁺ influx, an effect that was partially ameliorated by pretreatment of neurons with the NMDAR antagonist MK-801. The Ca²⁺ flux induced by A β 3pE-42 appears to be separate from its capacity to undergo copper-redox cycling as the effect persisted when media were depleted of row 1 transition metals. Changes in cellular Ca^{2+} homeostasis induced by A β have previously been implicated in A β toxicity and are thought to occur via multiple mechanisms such as pore formation and NMDAR activation (60, 61). Importantly, however, the aggregation state of the A β preparation significantly affects the capacity to induce Ca²⁺ flux; monomeric (freshly prepared) and fibrillar $A\beta(1-42)$ is not found to induce Ca²⁺ flux in SH-SY5Y neuroblastomas, whereas oligomeric preparations agonize NMDAR and trigger rapid Ca^{2+} flux (61, 62). Similarly, we found freshly prepared $A\beta(1-42)$ to cause only modest elevation in cortical neuron Ca^{2+} levels above controls, which is contrasted by the much larger Ca^{2+} flux induced by A β 3pE-42. Collectively, these data indicate that the unique neurotoxicity of pE-A β peptides is exerted through multiple interactions at the cell surface, including activation of NMDAR pathways, subsequently followed by peroxidation of membrane lipids and a loss of membrane integrity.

The potential for amino-truncated A β and pE-A β to alter the oligomerization and toxicity of full-length A β has been an area

of recent debate and speculation. Aß3pE-42 has a demonstrated capacity to dramatically enhance the aggregation of fulllength A β (14), as does A β 3-42 (41). A β 3pE-42 has further been reported to enhance the toxicity of $A\beta(1-42)$ in a prionlike seeding mechanism in cortical neuron cultures (24). Due to the strong membrane association of A β 3pE-42, we initially predicted A β 3pE-42 seeds to shift the ROS generation of A β (1– 42) to a lipid compartment, thereby increasing $A\beta(1-42)$ -induced lipoperoxidation; however, AB3pE-42 was not found to increase A β (1–42) ROS production in either the cytosolic or membrane fractions. In contrast, the capacity for A β 3pE-42 seeds to significantly reduce $A\beta(1-42)$ cytosolic ROS production indicates that A β 3pE-42 does not enhance A β (1-42) toxicity via ROS production, yet it suggests significant interaction between the peptides in neuronal cultures. This effect may be a result of accelerated A $\beta(1-42)$ aggregation in the presence of A β 3pE-42, as previous studies have found that A β -ROS production decreases with aggregation (32, 63). In the study by Nussbaum *et al.* (24), trace quantities of A β 3pE-42 were found to enhance the toxicity of $A\beta(1-42)$ when co-aggregated prior to cell treatment; however, when A β 3pE-42 was seeded into $A\beta(1-42)$ solutions immediately before applying to cells (as in our experiments), the toxicity of the seeded mixture was identical to the individual A $\beta(1-42)$ treatment. It is therefore likely that A β aggregates possess different cytotoxic properties depending on both the composition of the A β mixture and the





FIGURE 6. **Measurement of Ca²⁺ flux in cortical mouse neurons using the Fluo-4 Ca²⁺ sensor.** *A*, representative kinetics during the acute exposure of neurons to freshly prepared A β (10 μ M). *B*, levels of Ca²⁺ flux induced by exposure to A β (1–42) and A β 3pE-42 (10 μ M each peptide) relative to glutamate-treated cells (*Glu*; 1 μ M). Neurons were additionally pre-treated for 15 min with or without the NMDAR antagonist MK-801 (1 mM) to assess the contribution of NMDAR activation in A β 3pE-42-induced Ca²⁺ flux. *C*, effect of transition metal depletion on A β -induced Ca²⁺ flux in neurons pre-treated for 1 h with the chelator Diamsar. *D*, separate Ca²⁺ flux assays were performed in HBSS buffer devoid of Ca²⁺ and Mg²⁺ to determine the source of Ca²⁺ moving into the cytosol. Each treatment was tested in a minimum of four replicate wells per assay and the experiments repeated at a minimum of three times over separate days. Data are represented by *box* and *whiskers* (Tukey method); statistical significance is shown *above bars* as relative to the vehicle treatment (PBS) or between treatments as indicated. *, $p \le 0.05$; **, $p \le 0.001$; ***, $p \le 0.001$; ns, not significant, p > 0.05.

timing of the A β isoform interactions. The possibility also exists that A β 3pE-42 seeded mixtures cause oxidative damage that escape detection by the DCF and DPPP fluorescence ROS probes or possess different redox cycling capacities in other cerebral cell types (*e.g.* astrocytes and microglia). Other prevalent amino-truncated and pE-A β peptides found in AD brain tissues, such as A β (3–42), A β (4–42), and A β 11pE-42 (2, 52), require further investigation as they also demonstrate enhanced amyloid-seeding capacity and could potentially alter A β -ROS dynamics.

AFM and SEC analysis revealed that $A\beta(1-42)$ and A β 3pE-42 oligomers differed not only in the rate of formation but also in size and structure. Previous studies have similarly reported differences between $A\beta(1-42)$ and $A\beta$ 3pE-42 in the profile of oligomers and fibril ultrastructures formed in aqueous buffers (18, 30). Lee *et al.* (22) report that A β 3pE-42 forms larger oligomers in synthetic lipid membranes than $A\beta(1-42)$ and with faster kinetics of assembly, although it remains to be determined how this relates to the relative level of toxicity. A β 3pE-42 neurotoxicity has also been demonstrated with a broad range of size fractions (monomers to >100 kDa), whereas the toxicity of A $\beta(1-42)$ fractions was isolated to oligomers with an observed mass larger than 14 kDa (25). This is consistent with our SEC findings demonstrating the relative stability of low-mass A β (1–42) species (<14 kDa) in aqueous solution and the observation that freshly prepared A $\beta(1-42)$ is

less neurotoxic than A β 3pE-42. The metastable nature of A β peptides, however, presents many technical challenges in delineating "toxic" fractions from "benign" fractions as purification processes undoubtedly alter oligomerization kinetics. Likewise, A β peptides undergo significant structural changes in extended cell culture incubations; hence, it is pertinent to correlate toxicity markers with time-matched biophysical characterizations.

The A β 3pE-40 and A β 3pE-42 peptides have been found to resist proteolysis in astrocyte cultures (49) and accumulate in the lysosomes of astrocytes in cell culture and the AD temporal cortex (50). Consistent with these reports, we observed A β 3pE-42 to resist clearance in neuronal cultures, remaining at significantly higher levels than $A\beta(1-42)$ over extended incubations. A β 3pE-42 was, however, not found to prevent the clearance of A $\beta(1-42)$ from neural cultures when present at minor quantities (5% mol/mol), suggesting that A β 3pE-42 does not transfer protease resistance to $A\beta(1-42)$ when applied to cells as fresh preparations. This does not exclude the possibility that pE-A β may affect the clearance of full-length A β when the peptides are aggregated, which will require further investigation. The capacity for AB3pE-42 to resist proteolytic degradation in both neurons and astrocytes is highly relevant given that pE-A β peptides are found in the cores of amyloid plaques in the AD brain (52), suggesting that pE-A β peptides, once formed, are long-lived neurotoxins.

Dityrosine is another post-translational protein modification that confers resistance to cellular catabolism. Total dityrosine levels are elevated in the AD hippocampus, neocortex, and ventricular cerebrospinal fluid compared with cognitively healthy individuals (64), yet the contribution of $A\beta$ to cerebral dityrosine formation is not well understood. Dityrosine crosslinked A β fibrils are resistant to formic acid digestion, and sections of AD brains display intense dityrosine immunoreactivity within plaques (39). The pE-A β isoforms demonstrated increased efficiency for dityrosine oligomer formation compared with full-length $A\beta$. This observation can likely be attributed to increased hydrophobicity and the propensity to oligomerize; dityrosine formation is dependent on both the production of A β -tyrosyl radicals and the close proximity of A β molecules to allow tyrosine-tyrosine coupling (33). It is reasonable to speculate that the capacity for amyloid plaques to resist solubilization and clearance is due to the contribution of both the pE and dityrosine modifications to $A\beta$, which may account for the abundance of pE-A β in plaque cores (52).

Our data demonstrate clear differences in the neurotoxic mechanisms of pE-A β and full-length A β . The toxicity of pE-A β has recently been highlighted in mouse models overexpressing the soluble isoform of QC, demonstrating exacerbated neurodegeneration and behavioral deficits when crossed with mouse lines overexpressing $A\beta$ or amyloid precursor protein transgenes (65, 66). The specific targeting of pE-A β via inhibition of QC-catalyzed pyroglutamate synthesis has demonstrated promising results; transgenic AD mice treated with a QC inhibitor show reduced plaque load, reduced gliosis, and an improvement in context memory and spatial learning (67). QC knock-out does not completely inhibit pE-A β formation in transgenic AD mouse models, however (68), suggesting that multiple pathways of pE-A β formation may exist. Antibodies currently under development as passive vaccine therapies for AD show significant differences in their capacity to bind amino-truncated A β species (69) and thus may fail to remove pE-A β and its precursors (A β 3-40/42 and A β 11-40/42). Individuals with AD may therefore require different therapeutic interventions to target distinct $A\beta$ isoforms and their specific mechanisms of toxicity. A β ROS production, oligomerization, and dityrosine formation are potential therapeutic targets for AD; metal-protein attenuating compounds that inhibit these reactions are found to effect a marked decrease in amyloid deposition and improvement in cognitive deficits in transgenic AD mice (70, 71). Our observations indicate that the separate pathways of oxidative damage and neurotoxicity exerted by Aß3pE-42 potentially have a unique contribution to AD pathology. These findings elicit further consideration of pE-A β peptides as targets in the pursuit of biomarkers and disease-modifying therapeutics for AD.

Author Contributions—A. P. G. and J. A. D. wrote the manuscript. A. P. G. designed and conducted the experiments. B. X. W. assisted in the conduct of the lipid peroxidation assays, SEC, and AFM analyses. T. J. assisted in the conduct of the calcium flux experiments. J. G. performed the AFM imaging and analysis. C. L. M., K. J. B., A. I. B., and R. A. C. provided critical revision of the manuscript. All authors reviewed the results and approved the final version of the manuscript. Acknowledgments—We thank Dr. Paul Donnelly (School of Chemistry, University of Melbourne, Australia) for kindly providing Diamsar and Dr. Vanta Jameson (Melbourne Brain Centre Flow Cytometry Facility, Australia) for assistance with flow cytometry analysis. AFM was performed in collaboration with the Materials Characterization and Fabrication Platform at the University of Melbourne and the Victorian node of the Australian National Fabrication Facility. We thank Dr. Lauren Hyde and Prof. Ray Dagastine for their input into the design and conduct of the AFM analyses.

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