

Amyloid β Protein (1–40) Forms Calcium-Permeable, Zn^{2+} -Sensitive Channel in Reconstituted Lipid Vesicles^{†,‡}

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ABSTRACT: Amyloid β protein (A β P) forms senile plaques in the cerebrocortical blood vessels and brain parenchyma of patients with Alzheimer's disease (AD). The nonfamilial or sporadic AD (SAD), the most prevalent form of AD, has been correlated with an increased level of 40-residue A β P (A β P_{1–40}). However, very little is known about the role of A β P_{1–40} in AD pathophysiology. We have examined the activity of A β P_{1–40} reconstituted in phospholipid vesicles. A combined light fluorescence and atomic force microscope (AFM) was used to image the structure of reconstituted vesicles and ⁴⁵Ca²⁺ uptake was used as an assay for calcium permeability across the vesicular membrane. Vesicles reconstituted with fresh and globular A β P_{1–40} contain a significant amount of A β P and exhibit strong immunofluorescence labeling with an antibody raised against the N-terminal domain of A β P, suggesting the incorporation of A β P_{1–40} peptide in the vesicular membrane. Vesicles reconstituted with A β P_{1–40} exhibited a significant level of ⁴⁵Ca²⁺ uptake. The vesicular calcium level saturated over time, showing an important ion channel characteristic. The ⁴⁵Ca²⁺ uptake was inhibited by (i) a monoclonal antibody raised against the N-terminal region of A β P and (ii) Zn^{2+} . However, a reducing agent (DTT) did not inhibit the ⁴⁵Ca²⁺ uptake, indicating that the oxidation of A β P or its surrounding lipid molecules is not directly involved in A β P-mediated Ca²⁺ uptake. These findings provide biochemical and structural evidence that fresh and globular A β P_{1–40} forms calcium-permeable channels and thus may induce cellular toxicity by regulating the calcium homeostasis in nonfamilial or sporadic Alzheimer's disease.

Molecular mechanisms and the cause of Alzheimer's disease (AD)¹ are poorly understood. Alzheimer's disease (AD) is associated with depositions of amyloid β proteins (A β P), 39–43-amino acid peptides, as morphologically heterogeneous neuritic plaques and cerebrovascular deposits (1, 2). These peptides are derived from a proteolytic cleavage of the β -amyloid precursor protein (A β PP), a widely expressed but conserved integral membrane protein with a single membrane-spanning peptide. The subtype and amount of A β P polypeptides vary among various forms of ADs. There is a differential accumulation of A β P_{1–40} and A β P_{1–42} in sporadic Alzheimer's disease (SAD) and nondemented brain samples (3), and a mutation in presenilins is linked with an increased ratio of A β P_{1–42} to A β P_{1–40} in familial Alzheimer's disease (FAD) (4–7). The nonfamilial or sporadic AD (SAD), the most prevalent form of AD, is correlated with an increased level of 40-residue A β P (A β P_{1–40}). However, very little is known about the role of

A β P_{1–40} in such pathology and its mechanism(s) of action.

Amyloid β protein plays an early and *causative* role in the AD pathogenic cascade (8–11). Proposed mechanisms of A β P toxicity include its interaction with cell-surface receptors such as the tachykinin receptors (12), changing the cellular ion concentration via plasma membrane channels (13–17), and activating oxidative pathways and increasing the responsiveness to oxidative stress (for reviews, see refs 18 and 19). However, the roles of the reactive oxygen mechanism and the tachykinin receptor pathway are indirect and less definitive. Moreover, A β Ps of differing sizes associated with the reactive oxygen hypothesis have produced conflicting effects on cytoskeletal organization and cell lysis (20–25), and the role of the tachykinin neuropeptide pathway in A β P toxicity has been inconsistent.

An altered cellular calcium level is a common denominator underlying A β P toxicity (17, 26–29). Such a calcium change could be caused either indirectly via modulating the existing Ca²⁺ channel or directly by A β P-formed channels. A β P_{1–40} elicits cation-selective currents when reconstituted in artificial lipid bilayers (13–15) and when incubated with membrane patches excised from hypothalamic GnRH neurons (30). Low-angle X-ray diffraction studies have shown that A β P_{25–35} is highly lipophilic and spontaneously inserts into the membrane hydrocarbon core (21, 31). Durell et al. (32) have proposed theoretical models for the structure of the ion channel formed by the membrane-bound A β P_{1–40}. However, no direct structural data from EM, X-ray diffraction, NMR,

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¹ Abbreviations: A β P, amyloid β protein; AD, Alzheimer's disease; APP, amyloid precursor protein; AFM, atomic force microscope.

or other microscopic techniques are available to support the presence of the A β P channel. Moreover, other groups have often found conflicting results from electrical recordings (33, 34).

We have used an integrated atomic force and light fluorescence microscope (35) and $^{45}\text{Ca}^{2+}$ uptake assay (16) to examine calcium permeability in lipid vesicles reconstituted with A β P_{1–40}. Commonly used electrical current measurement across lipid membranes often includes contaminating currents when reconstituted with small hydrophobic peptides. $^{45}\text{Ca}^{2+}$ uptake measures Ca^{2+} transport across the lipid vesicle bilayer. $^{45}\text{Ca}^{2+}$ uptake study is not subjected to such contaminations. Moreover, it allows examination of the dose-response relationship of $^{45}\text{Ca}^{2+}$ uptake, and thus, various pharmacological studies are feasible. Consistent with the possibility of fresh A β Ps forming ion-permeable channels, we show that (i) A β P incorporates in reconstituted lipid vesicles, (ii) vesicles reconstituted with A β P exhibit a significant level of $^{45}\text{Ca}^{2+}$ uptake which is blocked by anti-A β P antibody and Zn^{2+} , but not blocked by an antioxidant, and (iii) the kinetics of calcium uptake is consistent with that of ion channel activity.

EXPERIMENTAL PROCEDURES

Materials. 1,2-Dioleoyl-*sn*-glycero-3-phosphatidylethanolamine (DOPE) and 1,2-dioleoyl-*sn*-glycero-3-phosphoserine (DOPS) were obtained from Avanti Polar Lipids (Alabaster, AL). Synthetic β -amyloid_{1–40} peptide was purchased from Bachem (Torrance, CA) and also obtained from Y. Hirakura (University of California, Los Angeles, CA) which was synthesized at Yale University (New Haven, CT). A monoclonal antibody against the N-terminus of A β P [3D6, anti-A β P 1–5 (DAEFR)] was kindly provided by R. Rydel at Athena Neurosciences, Inc. (South San Francisco, CA). Goat anti-mouse IgG conjugated with cy3 was purchased from Jackson Immuno Research Laboratories (West Grove, PA). A β P_{1–40} was dissolved in H₂O at a concentration of 5 mg/mL and stored at -20°C . Before use, A β P_{1–40} was thawed and bath sonicated for 5 min to break up any cluster formation.

Liposome Preparation. Equal portions of DOPE and DOPS (10–25 mg/mL) in chloroform were mixed together and dried under a stream of argon gas, and the organic solvent residue was removed by vacuuming with a speed vac for 1 h. Dried lipids were resuspended in a 20 mM Hepes (pH 7.4) aqueous solution at a concentration of 1.11 mg/mL. The lipid mixture was bath sonicated for 20 min. To reconstitute A β P into liposomes, 1 part of A β P (5 mg/mL) was mixed with 9 parts (volume) of the lipid mixture and bath sonicated for 20 min. To obtain the control liposomes without A β P, $1/9$ of a volume of H₂O was added to the lipid mixture and bath sonicated for 20 min. The final liposome buffer contains 20 mM NaHepes (pH 7.40) and 1 mg/mL lipids (1:1 DOPS/DOPE); the A β P concentration was 0.5 mg/mL.

The percentage of A β P incorporation in liposomes was estimated with a method described by Mason et al. (31). Briefly, liposomes reconstituted with A β Ps were centrifuged at 35000g for 90 min at 5°C . The relative A β P contents in the pellets (liposome) and supernatant were measured with the amido black dye binding protein assay. To ensure that

the measured A β P content was associated with liposomes and not unassociated large A β P aggregates, 20 mM Hepes buffer containing the same amount of A β P without lipid was centrifuged in parallel to determine the quantity of unincorporated A β P in the pellet. Very little of the A β P dissolved in aqueous buffer was sedimented (less than 0.5%) after centrifugation for 90 min at 35000g, indicating that little soluble A β P formed very large aggregates.

Immunolabeling with Anti-A β P Antibody. Liposomes reconstituted with or without A β Ps as well as buffer containing only soluble A β Ps were centrifuged at 35000g for 90 min at 5°C (very little A β Ps was pelleted from buffer containing only soluble A β Ps). Pellets were rinsed and resuspended in 20 mM Hepes, and portions of each sample were deposited on glass coverslips for 20 min followed by rinses with phosphate-buffered saline (PBS, Life Technologies, Rockville, MD) and fixation for 10 min with 4% paraformaldehyde. Each sample was washed with PBS and with PBS containing 3% bovine serum albumin (BSA) and 1% goat serum. The samples were incubated with 3D6 anti-A β P antibody in PBS (2 $\mu\text{g}/\text{mL}$) containing 3% BSA and 1% goat serum, for 1 h at room temperature. After washing with PBS, the sample was incubated with cy3-conjugated goat anti-mouse IgG (1:500 dilution) under the same condition that was used for primary antibody incubation. Fluorescence images were captured using 60 \times objective lenses with a Bio-Rad MRC-1024 laser confocal microscope.

Imaging Liposomes with Atomic Force Microscopy. AFM images were obtained as previously described (16) using a prototype of Bioscope AFM (36) and a Multimode AFM (Digital Instruments, Santa Barbara, CA). A contact mode AFM was used for most of the images. Oxide-sharpened silicon nitride tips with a nominal spring constant of about 0.06 N/m (Digital Instruments) were used for most of the experiments. All AFM imaging was conducted on liposomes in aqueous solutions. For liposomes reconstituted with or without A β P_{1–40}, 20–50 μL of liposomes was absorbed on a clean glass coverslip or glass Petri dish or freshly cleaved mica for 30 min. The substrate was then rinsed with a buffer [20 mM Hepes (pH 7.4)] and imaged in the buffer. The imaging force was regularly monitored and kept to a minimum, unless specified. The imaging force generally varied from a sub-nanonewton (nN) to tens of nanonewtons. All imaging was performed at room temperature (22 – 24°C). The liposomes were also imaged in deionized water as described previously (16).

Measurement of the Rate of $^{45}\text{Ca}^{2+}$ Uptake into Liposomes. The $^{45}\text{Ca}^{2+}$ uptake assay was based on a method previously described (16). Briefly, 25 μL of liposomes reconstituted with or without A β P_{1–40} was incubated with 75 μL of a buffer containing 20 mM Hepes (pH 7.4), 0.5 mM Ca^{2+} , and 1 μCi of $^{45}\text{Ca}^{2+}$ for a specific period of time, at room temperature. For each sample, $^{45}\text{Ca}^{2+}$ uptake was stopped with 300 μL of blocking buffer containing 10 mM Tris-HCl (pH 7.4), 20 mM ZnCl_2 , 0.5 mM CaCl_2 , and 15 $\mu\text{g}/\text{mL}$ 3D6 antibody. The 400 μL mixture of liposomes and the blocking buffer was promptly loaded onto a Chelex 100 column (Bio-Rad, Hercules, CA; bead volume of 3 mL) which had been pre-equilibrated with a buffer containing 200 mM sucrose, 20 mM Tris-HCl (pH 7.4), and 0.3% BSA. The liposomes were immediately eluted from the column with 5 mL of a buffer containing 200 mM sucrose and 20 mM Tris-HCl (pH 7.4).

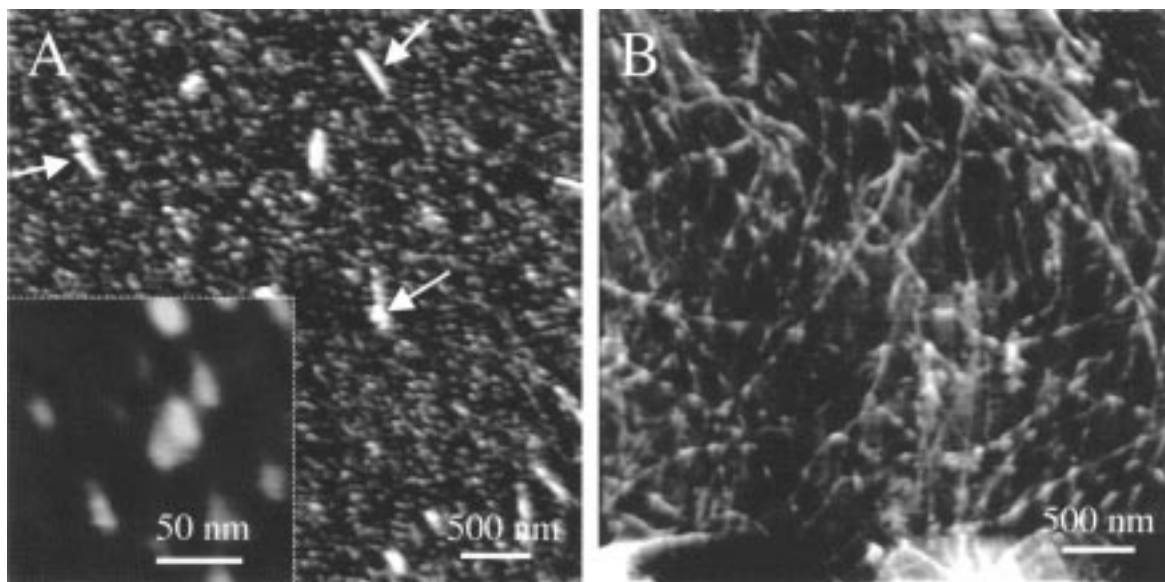


FIGURE 1: Images of fresh and aged A β P₁₋₄₀ produced with an AFM. AFM height mode images (surface plot) of A β Ps adsorbed on mica and imaged in PBS. (A) Image of fresh A β P₁₋₄₀ adsorbed on mica. Primarily, globular A β Ps with a few short fibrils (denoted by the arrows) are present. The average size of the individual globules is 8–10 nm, though larger conglomerates are also visible. The inset gives an enlarged view of some globular aggregates. No attempt was made to discern the high-resolution structure of individual subunits. (B) Image of aged A β P₁₋₄₀ (after incubation for 24 h), where long fibrils are present.

Fractions (0.1–0.2 mL) of the eluted liposome containing buffers were counted with a Beckman liquid scintillation counter to determine the relative $^{45}\text{Ca}^{2+}$ content of the liposomes.

To confirm that the saturation level of intravesicular $^{45}\text{Ca}^{2+}$ had equilibrated with the external $^{45}\text{Ca}^{2+}$ level, the intravesicular $^{45}\text{Ca}^{2+}$ level after a prolonged period (>10 min) of uptake was compared with the internal $^{45}\text{Ca}^{2+}$ level in liposomes reconstituted in the presence of A β Ps and 1 μCi of $^{45}\text{Ca}^{2+}$. This also provides a good estimate of the total internal volume of the liposomes, by comparing the $^{45}\text{Ca}^{2+}$ counts in the liposomes versus the total $^{45}\text{Ca}^{2+}$ counts in the buffer.

RESULTS

AFM Imaging of A β Ps. The structural conformation of A β P such as globular versus fibrillar is reported to correlate with A β P toxicity. We imaged fresh as well as aged A β P₁₋₄₀ adsorbed on a freshly cleaved mica surface with an AFM in aqueous solutions. AFM images of freshly prepared A β Ps reveal the presence of both small particles with an average diameter of 1–2 nm and larger globular aggregates with an average diameter of 8–10 nm. Figure 1A shows representative larger globular aggregates with a nominal size of 8–10 nm. No attempt was made to image the fine structure in these globular aggregates, but they could potentially form A β P channels and exhibit ion channel-like activity when incorporated into a lipid bilayer. Freshly prepared A β Ps also contain a very small percentage of 200–300 nm short fibers (less than 1%) as indicated by the arrows in Figure 1A. In contrast to the AFM images of fresh A β Ps, AFM images of aged A β Ps (A β Ps stored for ≥ 24 h at room temperature) revealed a mixed population of mostly long fibrous strands (>90%) and some small globular particles (Figure 1B).

AFM Imaging of Liposomes and Immunolocalization of A β P in Reconstituted Liposomes. Liposomes reconstituted with or without A β P₁₋₄₀ were imaged with an AFM.

Moreover, to examine whether A β P₁₋₄₀ was incorporated in reconstituted liposomes, we examined immunofluorescence labeling of liposomes reconstituted with or without A β P₁₋₄₀ and quantified the content in the reconstituted vesicles with a quantitative protein assay.

Liposomes reconstituted with A β Ps have a well-defined vesicular structure as revealed by AFM imaging (Figure 2A–C). Vesicles reconstituted without A β Ps are smaller but more spherical compared to the vesicles reconstituted with A β Ps. Vesicles reconstituted with A β P ranged mostly in size from approximately 0.2 to 0.5 μm in diameter, with a few larger vesicles up to 1.0 μm in diameter. The larger vesicles were often flattened in a disk shape. The vesicular flattening and larger size are probably due to the protein–lipid interactions as is the case with other membrane proteins such as gap junctions and acetylcholine receptors (37, 38). The height of the flattened vesicles was between 20 and 30 nm, which is typical for unilamellar vesicles composed of upper and bottom lipid bilayers (39). When imaged with a higher imaging force, which can be used to dissect away the upper bilayer, the AFM images of vesicles clearly showed that the vesicles were bound by a single lipid bilayer wall (Figure 2C).

The quantitative protein assay showed that after reconstitution, $15 \pm 1\%$ ($n = 3$, mean \pm standard error of the mean) of all A β P was associated with the liposomes, suggesting the incorporation of A β P in the liposomes.

Liposomes reconstituted with or without fresh A β P₁₋₄₀ were also immunolabeled with an anti-A β P antibody (3D6). Figure 3A shows immunolabeling of liposomes reconstituted with A β P₁₋₄₀. On the other hand, little immunolabeling was observed in samples prepared from resuspended pellets of A β P₁₋₄₀ (Figure 3B), which is not surprising since very little A β P was sedimented by centrifugation (<0.5%). There was also no immunofluorescence observed in liposomes reconstituted without A β P₁₋₄₀ (Figure 3C). These results indicate that A β P₁₋₄₀ is incorporated in the liposome membrane. We

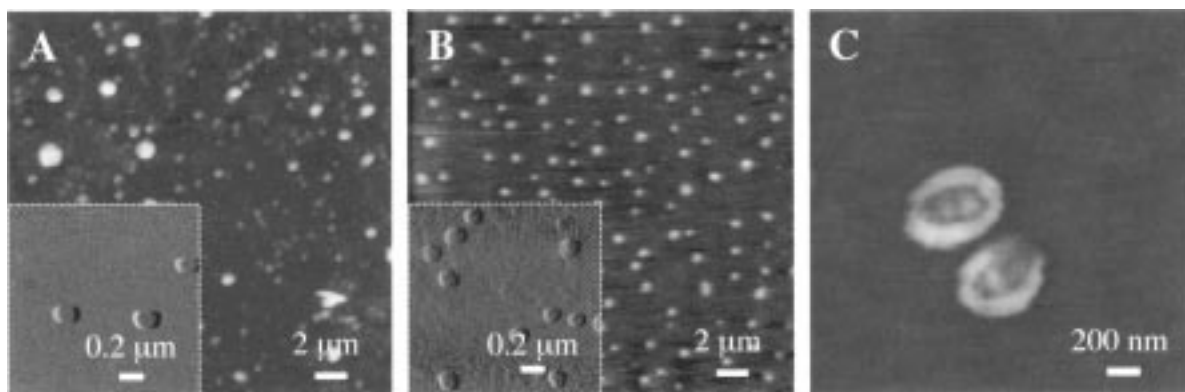


FIGURE 2: AFM images of reconstituted vesicles. Height mode images of DOPS/DOPE liposomes prepared by bath sonication, reconstituted (A) with or (B) without A β P. Insets in panels A and B show closeup error mode AFM images of representative liposomes with or without A β P, respectively. Panel C is a height mode image of liposomes reconstituted with A β P₁₋₄₀ imaged with a higher imaging force, a force which can be used to dissect the upper vesicular bilayer. The vesicles are clearly unilamellar and surrounded by a single bilayer. The thickness of a single bilayer is approximately 6 nm.

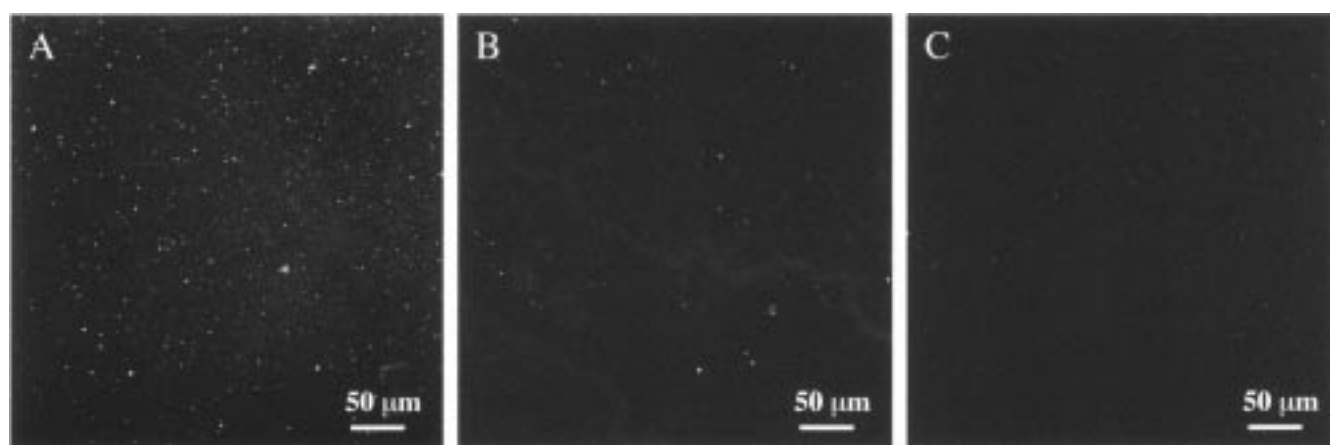


FIGURE 3: Confocal images of reconstituted liposomes fluorescently immunolabeled with anti-A β P antibody 3D6. Liposomes reconstituted with and without A β P₁₋₄₀ as well as soluble A β P were centrifuged, and pellets were resuspended and absorbed onto glass coverslips for immunolabeling. (A) Liposomes reconstituted with A β P₁₋₄₀ exhibit punctate immunofluorescence labeling for A β P. Little immunolabeling was observed in samples of either resuspended sediment of centrifuged soluble A β P₁₋₄₀ (B) or liposomes reconstituted without A β P₁₋₄₀ (C). All three images are projections from multiple vertical optical sections. A strong fluorescence signal suggests that there could be a large amount of A β P incorporated in the vesicles, although the brightness (and the spatial dimension) of the fluorescence signal depends on several factors, including the number of fluorophores and antibodies and the amount and conformations of A β P molecules.

also examined whether fibrillar A β Ps are reconstituted in the lipid vesicles. Strikingly, no significant level of immunolabeling was observed in the vesicles reconstituted with aged A β Ps, suggesting a lack of any incorporation into lipid vesicles (data not shown).

A β P Channel-Specific Calcium Uptake. Liposomes reconstituted with A β P₁₋₄₀ (Figure 4, \blacktriangle) have a level of $^{45}\text{Ca}^{2+}$ uptake that is 6–10 times higher than that of liposomes reconstituted without A β P₁₋₄₀ (Figure 4, \bullet). The extent of $^{45}\text{Ca}^{2+}$ uptake by liposomes reconstituted with fresh A β P₁₋₄₀ exhibits a time-dependent saturating pattern (Figure 4, \blacktriangle), consistent with the channel-like activity of reconstituted A β Ps. The higher level of $^{45}\text{Ca}^{2+}$ uptake is not caused by the direct binding of $^{45}\text{Ca}^{2+}$ to incorporated A β P peptides or A β P–lipid molecule complexes since A β P₁₋₄₀ alone (without any lipids) did not exhibit any significant $^{45}\text{Ca}^{2+}$ radioactivity (Figure 4, \blacksquare). $^{45}\text{Ca}^{2+}$ uptake is partially inhibited by a monoclonal antibody (3D6, 20 $\mu\text{g}/\text{mL}$) raised against the N-terminal region of A β P (Figure 4, \circ), showing that the level of $^{45}\text{Ca}^{2+}$ uptake is A β P-specific. Furthermore, liposomes reconstituted with bovine serum albumin alone (without A β Ps) did not exhibit any significant $^{45}\text{Ca}^{2+}$ uptake

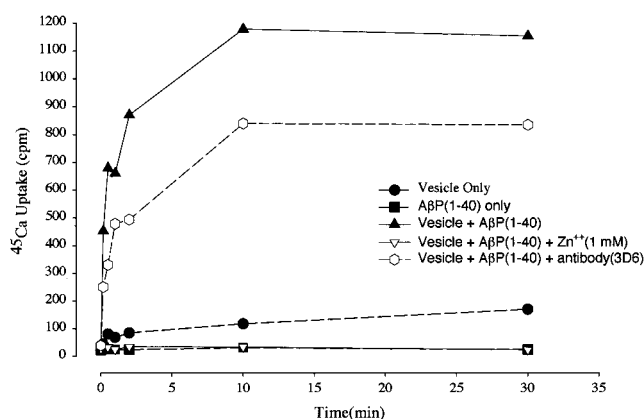


FIGURE 4: Time-dependent uptake of $^{45}\text{Ca}^{2+}$ under different conditions: $^{45}\text{Ca}^{2+}$ uptake in liposomes reconstituted with lipids only (\bullet) and liposomes reconstituted with A β P₁₋₄₀ (\blacktriangle). $^{45}\text{Ca}^{2+}$ uptake in liposomes reconstituted with A β P₁₋₄₀ was partially inhibited by an anti-A β P monoclonal antibody (3D6, 20 $\mu\text{g}/\text{mL}$) (\circ) and completely blocked by 1 mM Zn²⁺ (∇). In the absence of lipids, A β P₁₋₄₀ (0.5 mg/mL) alone did not bind to $^{45}\text{Ca}^{2+}$ (\blacksquare).

above the levels observed for the liposomes reconstituted with lipids alone (data not shown).

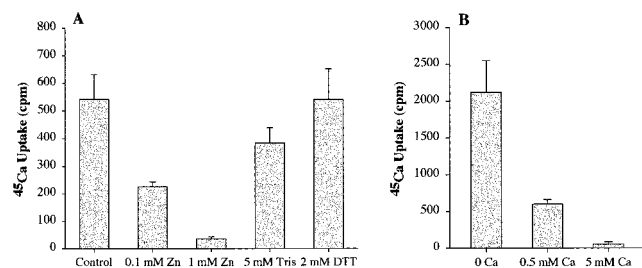


FIGURE 5: ⁴⁵Ca²⁺ uptake into liposomes reconstituted with A β P₁₋₄₀. (A) Level of ⁴⁵Ca²⁺ uptake after incubation for 2 min under control conditions and in the presence of 0.1 mM Zn²⁺, 1 mM Zn²⁺, 5 mM Tris, or 2 mM dithiothreitol. (B) Level of ⁴⁵Ca²⁺ uptake after incubation for 1 min in the presence of 0, 0.5 (the control condition), and 5 mM nonradioactive Ca²⁺. The data in these graphs are the average of triplicate samples, and the error bars represent the standard error of the mean.

To ensure that all liposomes reconstituted with A β P took up ⁴⁵Ca²⁺ and the saturation ⁴⁵Ca²⁺ level in reconstituted liposomes reached equilibrium with the external ⁴⁵Ca²⁺ level, we measured the ⁴⁵Ca²⁺ level in vesicles reconstituted in the presence of 1 μ Ci of external ⁴⁵Ca²⁺. Under such a condition, the concentration of ⁴⁵Ca²⁺ in the vesicles should be the same as the ⁴⁵Ca²⁺ concentration in the external buffer. For a normalized liposome concentration (1 mg/mL phospholipid), the amount of ⁴⁵Ca²⁺ in vesicles reconstituted in external ⁴⁵Ca²⁺ was $2.1 \pm 0.3\%$ ($n = 4$) of the total ⁴⁵Ca²⁺ in the buffer, which is not statistically significantly different from the saturation levels of ⁴⁵Ca²⁺ uptake in liposomes reconstituted with A β P. In a paired experiment, after incubation for 30 min in the ⁴⁵Ca²⁺ buffer, the amount of ⁴⁵Ca²⁺ in vesicles reconstituted in the absence of external ⁴⁵Ca²⁺ was $1.7 \pm 0.3\%$ ($n = 4$) of the total ⁴⁵Ca²⁺ in the buffer. These results indicate that the saturation ⁴⁵Ca²⁺ level in liposomes was near equilibrium with the external ⁴⁵Ca²⁺ level in the buffer and essentially all liposomes contained at least one A β P-formed Ca²⁺ channel. These results also give an estimate of the total volume of the vesicles as approximately 2% of the buffer volume, which is typical for unilamellar liposomes (40).

We examined the mechanisms of A β P₁₋₄₀-specific calcium uptake. In this study of A β P₁₋₄₀, ⁴⁵Ca²⁺ uptake was blocked by 1 mM Zn²⁺ (Figure 4, ∇), which may also compete against binding of ⁴⁵Ca²⁺ to negatively charged phospholipid headgroups. The ⁴⁵Ca²⁺ uptake in liposomes reconstituted with A β P₁₋₄₀ was not inhibited by DTT (2 mM), an antioxidant (Figure 5A), but was partially inhibited by Tris (5 mM, pH 7.4). Figure 5A summarizes the level of ⁴⁵Ca²⁺ uptake by liposomes reconstituted with A β P₁₋₄₀ after incubation for 2 min with ⁴⁵Ca²⁺ in the presence of (i) 0.1 mM Zn²⁺, (ii) 1 mM Zn²⁺, (iii) 5 mM Tris (pH 7.4), and (iv) 2 mM DTT. Moreover, ⁴⁵Ca²⁺ uptake is competitively inhibited by increasing the concentration of nonradioactive Ca²⁺. Figure 5B summarizes the level of ⁴⁵Ca²⁺ uptake in liposomes after incubation for 1 min in buffers containing (i) 0, (ii) 0.5 mM (control), and (iii) 5 mM nonradioactive Ca²⁺. The data points in these graphs are the averages of triplicate samples for each point, and the error bars represent the standard error of the mean.

DISCUSSION

We have examined functional properties of A β P₁₋₄₀ reconstituted in lipid vesicles using complementary tech-

niques of atomic force microscopy, light immunofluorescence microscopy, and ⁴⁵Ca²⁺ uptake. Vesicles reconstituted with fresh and globular A β P₁₋₄₀ contain a significant amount of A β Ps and exhibit strong immunofluorescence labeling with an antibody raised against the N-terminal domain of A β P, suggesting the incorporation of A β P₁₋₄₀ in the vesicular membrane. Vesicles reconstituted with A β P₁₋₄₀ exhibited a significant level of ⁴⁵Ca²⁺ uptake that saturated over time, an important ion channel characteristic. The ⁴⁵Ca²⁺ uptake was inhibited by (i) a monoclonal antibody raised against the N-terminal region of A β P and (ii) Zn²⁺. However, a reducing agent (DTT) did not inhibit the ⁴⁵Ca²⁺ uptake, indicating that the oxidation of A β P or its surrounding lipid molecules is not directly involved in the A β P-mediated Ca²⁺ uptake.

AFM Imaging of Isolated A β Ps. AFM images of freshly prepared A β P₁₋₄₀ show the presence of both small particles with an average diameter of 1–2 nm and larger globular aggregates with an average diameter of 8–10 nm. The small diameter particles may represent monomeric and dimeric A β P polypeptides. The larger globular aggregates with an average size of 8–10 nm could represent the channel-forming portion of A β Ps and exhibit ion channel-like properties after proper incorporation into a lipid bilayer. Such a globular size is consistent with the size of A β P channels as proposed in the model of its three-dimensional structure (32). The model predicts channels to be made of four to eight monomeric subunits and is supported by electrical measurements in reconstituted bilayer studies (13–15, 34). In the study presented here, we did not intend to decipher the channel-like structural arrangement of the globular A β Ps; such study will require extensive and sub-nanometer resolution structural examination of reconstituted vesicles with A β Ps.

Roher et al. (41) have recently imaged A β P₁₋₄₂ with an AFM and showed granular ellipsoidal particle aggregates with average size of 3–4 or 7–8 nm and with occasional aggregation into larger conglomerates. The average size difference of A β P₁₋₄₂, the 3–4 nm diameter as observed by Roher et al. (41) versus the 1–2 nm diameter observed in our study, may reflect the differences in the imaging conditions [hydrated A β Ps imaged in PBS in our study vs dry A β Ps imaged in 2-propanol and air in the study by Roher et al. (41)] and/or the difference in the packing of residues 1–40 and 1–42 of A β Ps. Fresh A β Ps, negatively stained and imaged with an EM, were globular particles with an average diameter of 30 nm (24). In our AFM studies, though the smaller particles (2–10 nm) predominate, a few particles with a larger diameter were also apparent. Such a distribution of particle size as observed by AFM and EM studies may reflect the varying degrees of constraints imposed on sample preparations and imaging by these techniques; AFM imaging requires very little sample preparation, and the imaging is less invasive. Aged A β Ps exhibit fiber-like features with some globular structures. Previous EM images of aged A β Ps exhibit similar fiber-like structures (24). Significantly though, in that study, all samples derived from the populations with high toxicity contained small globular A β P particles.

A β P in Reconstituted Liposomes. Our protein assay experiment provides evidence that A β P₁₋₄₀ was incorporated in the liposomes, though the rate of incorporation was lower than that of A β P₂₅₋₃₅ (31). In addition, anti-A β P antibody immunolabeling was observed on vesicles reconstituted with

A β P_{1–40}, but not in vesicles reconstituted without A β P_{1–40} or in isolated A β P_{1–40} adsorbed to the substrate (Figure 3). The site-directed antibody used in our study is specific to the extramembranous amino terminus portion of the A β P polypeptide. Thus, the immunolabeling results suggest that the amino-terminal portion of the A β P polypeptide is exposed to the exterior of the lipid vesicles and the polypeptide is not entirely encapsulated in the aqueous interior of liposomes. This is consistent with the results of the low-angle X-ray diffraction study of A β P_{25–35} and lipid membrane interaction (31), which showed that A β P_{25–35} is highly lipophilic and spontaneously intercalates into the membrane bilayer and suggested that peptide residues 29–35 are imbedded in the hydrocarbon core of the lipid bilayer.

Recently, Choo-Smith et al. (42) have reported a requirement for the presence of gangliosides for the interaction of A β P with vesicles containing zwitterionic or acidic phospholipids. We have not systematically examined A β Ps reconstituted in zwitterionic phospholipids. Our results from the ⁴⁵Ca²⁺ uptake assay and immunolabeling of A β Ps in liposomes, however, suggest that A β Ps can directly interact with negatively charged phospholipids and do not require any gangliosides (31). The presence of A β P in phospholipid vesicles is consistent with the possibility of A β P_{1–40} forming cation-selective channels in membranes (13–15, 30). The relatively high level of A β P incorporation in the vesicular membrane is a desirable feature for imaging of A β P channels in the membrane. Further study is underway in an effort to examine the three-dimensional structure of the putative A β P channel.

Since A β PP is a membrane protein, it was believed that the membrane-bound A β P portion would not be found as a free peptide except when there is membrane injury leading to a proteolytic cleavage of A β PP. In addition, it was believed that the self-aggregating and hydrophobic nature of A β P would prohibit it from existing as a soluble, circulating peptide in normal biological fluids. Subsequent studies, however, revealed soluble A β Ps being secreted into the conditioned media of a variety of primary or transfected A β PP-expressing cells under normal metabolic conditions (43–45). Our result and other recent studies (31) suggest that the soluble A β Ps could directly interact with lipid membrane. Whether they form a channel-like three-dimensional structure is still under investigation, but the increasing evidence suggests the presence of such channels (see the following section).

A β P_{1–40} Channel-Specific Calcium Uptake. Liposomes reconstituted with A β P_{1–40} exhibit a significant level of ⁴⁵Ca²⁺ uptake. The calcium uptake is dose-dependent and exhibits a time-dependent saturation pattern. The calcium uptake was inhibited by zinc, but not by physalamin and DTT. Recent studies suggest that A β Ps of different sizes (i.e., A β P_{1–42}, A β P_{25–35}, and A β P_{1–40}) form cation-selective channels whose activity can be inhibited by zinc, Tris, and other related compounds (16). A β P_{1–40}-specific cationic channels are reported in AD-free fibroblasts, endothelial cells, and neuronal patches (26–30), and A β P_{1–40}-specific calcium uptake has been reported in AD-free fibroblasts (preliminary results; 17).

It has recently been shown that A β P binds specifically and saturably with zinc in a biphasic mode: high-affinity binding ($K_D = 107$ nM) and low-affinity binding ($K_D = 5.2$

μ M) (46). The zinc-binding site has been mapped to a stretch of contiguous residues between positions 6–28 of the A β P sequence. Zinc, at a concentration of more than 1 mM, facilitates the precipitation of soluble A β P from aqueous buffer (47, 48). This property could be pathologically important because zinc is abundant in the same neocortical regions where A β P deposits are most commonly found. Moreover, the concentration of zinc reaches a high micromolar range during glutamatergic neurotransmission (49, 50), providing one possible explanation for the propensity of A β P to deposit close to the neocortical synaptic vicinity.

An antioxidant (DTT) did not inhibit calcium uptake, which indicates that A β P_{1–40}-dependent ⁴⁵Ca²⁺ uptake is not due to the oxidation of the A β P_{1–40} peptide (12). Antioxidants presumably interact with the membrane lipid components and change the membrane permeability (for reviews, see refs 16–18). Our results strongly suggest that the antioxidants have little or no effect on A β P_{1–40} channel-mediated ionic exchange in vitro. Such a finding is consistent with our recent observation that antioxidants have no effect on calcium permeability through channels formed by A β P_{1–42} (16). At the cellular level, A β Ps of differing sizes, reported to be associated with the reactive oxygen hypothesis, have produced conflicting effects on the cytoskeletal organization and cell lysis (20–25). In AD-free aged fibroblasts, Trolox and DTT did not prevent A β P_{1–40}-induced cell degeneration (preliminary results; 17); a similar observation was made in endothelial cells as well (preliminary results; 29).

For the channel hypothesis to be relevant in cellular toxicity, the efficiency of channel formation should be high. The concentration of A β P_{1–40} for channel formation in our study is very comparable to the concentrations at which others have described channel formation both by A β P_{1–40} and A β P_{1–42} and by other amyloid proteins such as amylin, PRP(106–126) (13, 30, 33, 51), and serum amyloid A (52), as well as C type natriuretic peptide (53), β 2 microglobulin, and transthyretin.

Because of the nature of the aggregation process, it is likely that only a small fraction of the total A β P is available for channel formation. Monomeric or very small aggregates [dimers or trimers (32)] of A β P probably cannot form channels. Very large aggregates of A β Ps probably also cannot form channels. So, it is consistent with what we know about aggregation that only a small percentage of the total protein is available for channel formation.

The concentration of A β P required to form channels is much higher than those required for other channel formers such as colicin or diphtheria toxin. However, it is clear that the amyloid peptides are not proteins designed to be toxins, but rather aberrant proteins with a weak toxic property which nonetheless turns out to be significant under conditions where disease processes lead to abnormal accumulation or perhaps abnormally high concentrations of these proteins.

Although the concentrations of A β P used in reconstitution studies are higher than the physiologic concentration of A β P in cerebrospinal fluid, it is certainly possible that A β P attains higher concentrations in localized areas of affected tissues. Indeed, there is some evidence that A β P accumulation occurs in intracellular compartments such as endosomes and lysosomes (54) and could be released under appropriate pathophysiologic conditions. Our preliminary results show that

A β P₁₋₄₀ causes a significant increase in the cellular level of free calcium and the eventual cellular degeneration.

In summary, our study strongly suggests that A β P₁₋₄₀ forms calcium-permeable, Zn²⁺-sensitive channels in vitro and allows calcium transport across lipid membranes. Such calcium exchange across the cell plasma membrane could destabilize cellular calcium homeostasis and induce cellular toxicity.

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