

ported by two findings. First, Ig- $\alpha$ -CD8 chimeric heterodimers but not Ig- $\beta$ -containing chimeras are able to transduce signals leading to interleukin-2 production in the B cell lymphoma, A20 (24). Second, stimulation through isolated CD3  $\epsilon$  or TCR  $\zeta$  chains results in distinct patterns of phosphorylation (9) and Ca<sup>2+</sup> mobilization responses (25).

The observation that Ig- $\alpha$  and Ig- $\beta$  can bind to different effector molecules may have implications for the function of other oligomeric receptors that can utilize alternate ARH1-containing chains to assemble different signal transduction complexes. For example, TCRs that contain  $\delta$ - $\epsilon$  heterodimers may use different signal transduction pathways than those that contain  $\gamma$ - $\epsilon$  heterodimers (26). Differences in lymphocyte responses to antigen, which occur as a function of maturation, could be determined by such expression of alternative signal transduction complexes (27).

The Src family tyrosine kinases and pp40 and pp42 have different patterns of association with the ARH1 motifs of Ig- $\alpha$  and Ig- $\beta$ , which suggests that they may interact with distinct sites within the cytoplasmic domains. These interactions, which are stable in NP-40, are reminiscent of other kinase-linked receptors such as the PDGFR, in which discrete regions mediate the receptor's association with distinct effector molecules (21, 28). Presumably, chain-specific residues, in the context of the antigen receptor's ARH1 motifs, determine the specificity of these interactions.

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7 April 1992; accepted 1 July 1992

## Production of the Alzheimer Amyloid $\beta$ Protein by Normal Proteolytic Processing

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The 4-kilodalton (39 to 43 amino acids) amyloid  $\beta$  protein ( $\beta$ AP), which is deposited as amyloid in the brains of patients with Alzheimer's disease, is derived from a large protein, the amyloid  $\beta$  protein precursor ( $\beta$ APP). Human mononuclear leukemic (K562) cells expressing a  $\beta$ AP-bearing, carboxyl-terminal  $\beta$ APP derivative released significant amounts of a soluble 4-kilodalton  $\beta$ APP derivative essentially identical to the  $\beta$ AP deposited in Alzheimer's disease. Human neuroblastoma (M17) cells transfected with constructs expressing full-length  $\beta$ APP and M17 cells expressing only endogenous  $\beta$ APP also released soluble 4-kilodalton  $\beta$ AP, and a similar, if not identical, fragment was readily detected in cerebrospinal fluid from individuals with Alzheimer's disease and normal individuals. Thus cells normally produce and release soluble 4-kilodalton  $\beta$ AP that is essentially identical to the 4-kilodalton  $\beta$ AP deposited as insoluble amyloid fibrils in Alzheimer's disease.

In the brains of patients with Alzheimer's disease (AD), amyloid is deposited in senile plaques (1, 2) and, in many cases, in the walls of cerebral and meningeal blood vessels (3). The amyloid fibrils in the brains of

patients with AD are composed of a 39- to 43-amino acid amyloid  $\beta$  protein ( $\beta$ AP) (4-8) that is derived from a set of 695- to 770-amino acid precursor proteins (7, 9-11) collectively referred to as the amyloid  $\beta$  protein precursor ( $\beta$ APP). In each of the 695- to 770-residue  $\beta$ APPs, the  $\beta$ AP is an internal peptide that begins 99 residues from the COOH-terminal end of the  $\beta$ APP and extends from the extracellular-intraluminal region into the middle of the hydrophobic, membrane-spanning domain. The  $\beta$ APP is normally processed in constitutive secretory (12-15) and endosomal-lysosomal (16-20) pathways. In the secretory pathway, the  $\beta$ APP is cleaved within the  $\beta$ AP (21, 22) to produce a large secreted derivative and a small membrane-associated fragment, neither of which can produce amyloid because they do not contain the entire

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$\beta$ AP. The endosomal-lysosomal system processes a complex set of COOH-terminal derivatives that includes potentially amyloidogenic forms with the entire  $\beta$ AP at or near their  $\text{NH}_2$ -terminus (18, 20, 23).

To determine whether  $\beta$ AP can be produced from  $\beta$ AP-bearing COOH-terminal  $\beta$ APP derivatives (18), human mononuclear leukemic (K562) cells were stably transfected with  $\text{LC}_{99}$ , a construct (Fig. 1A) that begins with the 17-amino acid  $\beta$ APP signal peptide, continues with the leucine that normally follows the  $\beta$ APP signal peptide, and ends with the 99 amino acids at the COOH-terminal end of the  $\beta$ APP ( $\beta\text{AP}_{1-43}$  followed by  $\text{C}_{56}$ ). As expected (Fig. 1B), lysates of these cells have

large amounts of an 11.4-kD protein that is immunoprecipitated both by an antiserum to the COOH-terminal end of the  $\beta$ APP and by SGY2134, an antiserum raised to  $\beta\text{AP}_{1-40}$  that contains antibodies directed primarily at  $\beta\text{AP}_{1-16}$  (18). The size of this 11.4-kD protein is precisely that expected after the signal peptide has been removed to produce  $\text{LC}_{99}$ , and this 11.4-kD protein is rapidly metabolized after metabolic labeling for 1 hour (Fig. 1B).

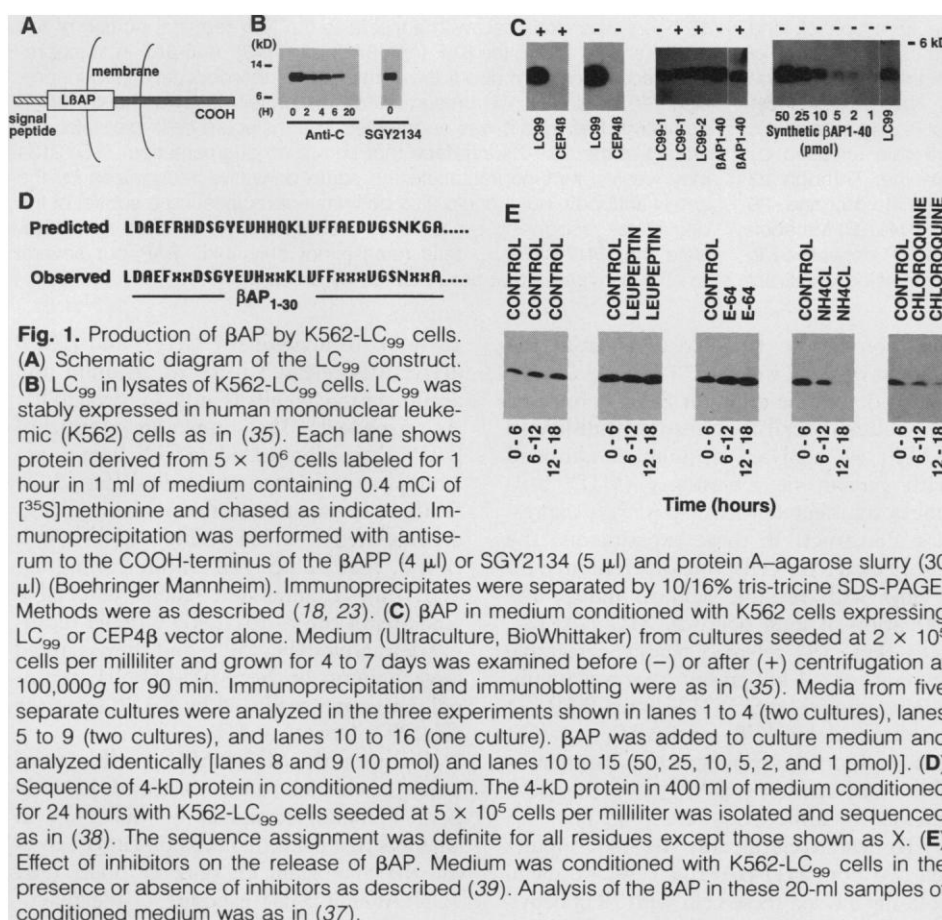
In an effort to identify  $\beta$ AP released from K562- $\text{LC}_{99}$  cells, we used SGY2134 to immunoprecipitate the protein in medium conditioned either with K562- $\text{LC}_{99}$  cells or with cells (K562-CEP4 $\beta$ ) transfected with vector alone. The immunoprecipitated pro-

tein was separated on 10 to 16% tris-tricine gels (18), transferred to immobilon P, and labeled with 4G8, a mouse monoclonal antibody to  $\beta\text{AP}_{17-24}$  (24). As a control, synthetic  $\beta\text{AP}_{1-40}$  (1, 2, 5, 10, 25, or 50 pmol) was added to the culture medium and analyzed identically. A 4-kD protein that comigrated with synthetic  $\beta\text{AP}_{1-40}$  was identified in medium conditioned with K562- $\text{LC}_{99}$  cells but not in medium conditioned with K562 cells transfected with vector alone (Fig. 1C).

Virtually all K562- $\text{LC}_{99}$  cells (>95%) were viable at the end of conditioning as judged by trypan blue exclusion. Nonetheless, to exclude the possibility that the externalized 4-kD protein was associated with sloughed membranes (possibly from dying cells), we compared the immunoprecipitates obtained from medium before and after centrifugation at 100,000g for 90 min. Neither synthetic  $\beta\text{AP}_{1-40}$  nor the 4-kD protein released from K562- $\text{LC}_{99}$  cells was removed appreciably by high speed centrifugation (Fig. 1C).

For definitive identification, the 4-kD protein in 400 ml of medium conditioned for 24 hours with  $2 \times 10^8$  K562- $\text{LC}_{99}$  cells was concentrated, immunoprecipitated with SGY2134, transferred to immobilon, excised after visualization with Coomassie blue, and sequenced. The sequence (Fig. 1D), which could be read as far as cycle 31, showed the 4-kD protein to be  $\beta$ AP with a leucine at its  $\text{NH}_2$ -terminus. Thus our data show that cells transfected with a model  $\beta$ AP-bearing, COOH-terminal derivative process that derivative to release  $\beta$ AP into the medium.

Because our previous data indicated that  $\beta$ AP-bearing COOH-terminal derivatives are produced in the endosomal-lysosomal system (23), we investigated the effect of inhibitors of this system (leupeptin, E-64, ammonium chloride, and chloroquine) on the production of  $\beta$ AP from  $\text{LC}_{99}$  (Fig. 1E). Control cells showed a slight progressive increase in the  $\beta$ AP released in the 0- to 6-, 6- to 12-, and 12- to 18-hour intervals, most likely due to continuing cell division. Cells treated with leupeptin and E-64 behaved similarly. Thus proteases inhibited by these agents are not involved in the production of  $\beta$ AP from  $\text{LC}_{99}$ . However, ammonium chloride, added when the medium was changed after a control conditioning period of 6 hours, caused a marked reduction in the production of  $\beta$ AP in the 6- to 12-hour interval, and it virtually abolished production in the 12- to 18-hour interval. Chloroquine had a similar but less dramatic effect. Thus the two lysosomotropic agents, which block proteolysis by alkalizing acidic compartments, reduce the production of  $\beta$ AP. These data suggest that the  $\beta$ AP may be produced, at least in part, by lysosomal processing of  $\beta$ AP-bearing, COOH-terminal derivatives and that the



**Fig. 1.** Production of  $\beta$ AP by K562- $\text{LC}_{99}$  cells.

(A) Schematic diagram of the  $\text{LC}_{99}$  construct. (B)  $\text{LC}_{99}$  in lysates of K562- $\text{LC}_{99}$  cells.  $\text{LC}_{99}$  was stably expressed in human mononuclear leukemic (K562) cells as in (35). Each lane shows protein derived from  $5 \times 10^6$  cells labeled for 1 hour in 1 ml of medium containing 0.4 mCi of [ $^{35}\text{S}$ ]methionine and chased as indicated. Immunoprecipitation was performed with antiserum to the COOH-terminus of the  $\beta$ APP (4  $\mu\text{l}$ ) or SGY2134 (5  $\mu\text{l}$ ) and protein A-agarose slurry (30  $\mu\text{l}$ ) (Boehringer Mannheim). Immunoprecipitates were separated by 10/16% tris-tricine SDS-PAGE. Methods were as described (18, 23). (C)  $\beta$ AP in medium conditioned with K562 cells expressing  $\text{LC}_{99}$  or CEP4 $\beta$  vector alone. Medium (Ultraculture, BioWhittaker) from cultures seeded at  $2 \times 10^5$  cells per milliliter and grown for 4 to 7 days was examined before (–) or after (+) centrifugation at 100,000g for 90 min. Immunoprecipitation and immunoblotting were as in (35). Media from five separate cultures were analyzed in the three experiments shown in lanes 1 to 4 (two cultures), lanes 5 to 9 (two cultures), and lanes 10 to 16 (one culture).  $\beta$ AP was added to culture medium and analyzed identically [lanes 8 and 9 (10 pmol) and lanes 10 to 15 (50, 25, 10, 5, 2, and 1 pmol)]. (D) Sequence of 4-kD protein in conditioned medium. The 4-kD protein in 400 ml of medium conditioned for 24 hours with K562- $\text{LC}_{99}$  cells seeded at  $5 \times 10^5$  cells per milliliter was isolated and sequenced as in (38). The sequence assignment was definite for all residues except those shown as X. (E) Effect of inhibitors on the release of  $\beta$ AP. Medium was conditioned with K562- $\text{LC}_{99}$  cells in the presence or absence of inhibitors as described (39). Analysis of the  $\beta$ AP in these 20-ml samples of conditioned medium was as in (37).

**Fig. 2.**  $\beta$ AP in CSF. Analysis of 3-ml samples was as in (37). (A) Autopsy CSF. The age and post-mortem interval (mean  $\pm$  SEM) of the histopathologically confirmed AD patients were  $74.6 \pm 1.8$  years and  $7.5 \pm 2.5$  hours, respectively. The age and postmortem interval of patients shown histopathologically to be without central nervous system disease were  $68.7 \pm 4.3$  years and  $8.9 \pm 1.6$  hours, respectively. (B) CSF from living patients. The diagnosis of AD in the living patients was established clinically by reviewing established criteria (40–42). All patients included were classified as having probable AD. Patients classified in the non-AD category were evaluated similarly. The ages (mean  $\pm$  SEM) of the AD and non-AD patients were  $73.2 \pm 1.3$  and  $73.6 \pm 0.7$  years, respectively.

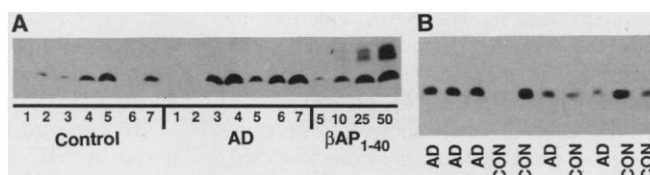
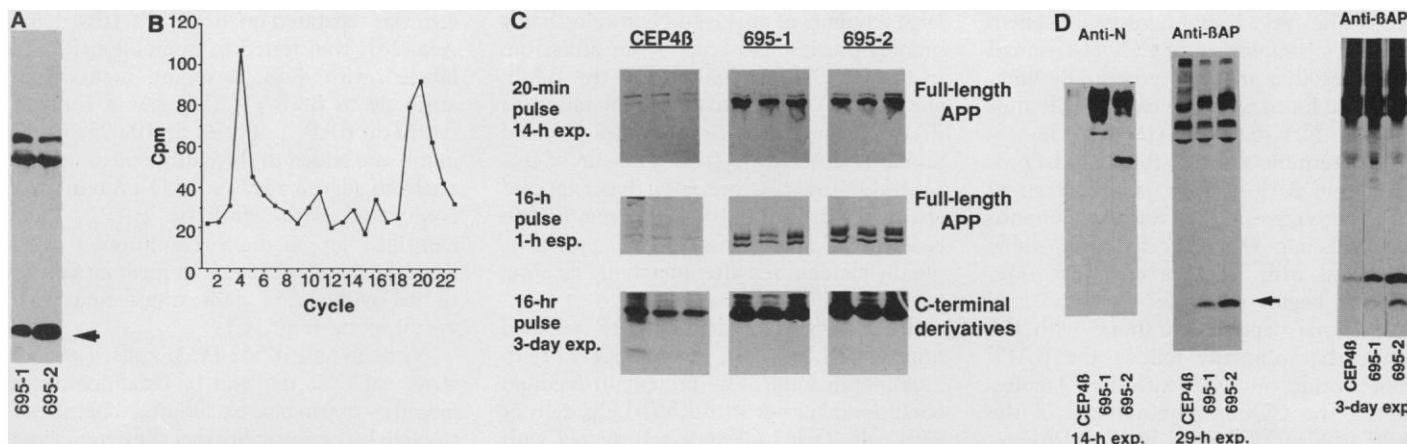


Figure 2A shows an autoradiograph of an immunoblot for  $\beta$ AP in autopsy CSF. The lanes are labeled Control, AD, and  $\beta\text{AP}_{1-40}$ . The AD lane shows a strong band, while the Control and  $\beta\text{AP}_{1-40}$  lanes show much weaker bands. Figure 2B shows an autoradiograph of an immunoblot for  $\beta$ AP in CSF from living patients. The lanes are labeled AD, CON, and CON. The AD lane shows a strong band, while the CON lanes show much weaker bands.



**Fig. 3.** Production of  $\beta$ AP by M17 cells. **(A)** Immunoprecipitation and immunoblotting of the 4-kD protein released by M17- $\beta$ APP<sub>695</sub> cells. Two M17 lines stably transfected to express  $\beta$ APP<sub>695</sub> as in (18) were treated with retinoic acid (1  $\mu$ M) for 7 days to induce the formation of long neurites. Confluent (10-cm) dishes were then used to condition 10 ml of medium containing 1  $\mu$ M retinoic acid for 3 days. Immunoprecipitation with SGY2134 and labeling with 4G8 were as in (37). Note the antibody-related artifact at the 10/16% interface of the gels that is also seen in (D) (lanes 3 to 5). **(B)** Radiosequencing of the 4-kD protein released by M17- $\beta$ APP<sub>695</sub> cells after labeling with [<sup>3</sup>H]phenylalanine. Through 23 cycles, [<sup>3</sup>H]phenylalanine was detected at positions 4, 19, and 20 indicating that the 4-kD protein is  $\beta$ AP. For details see (44). **(C)** Metabolically labeled full-length  $\beta$ APP and COOH-terminal  $\beta$ APP derivatives in M17 cell lines stably transfected with a  $\beta$ APP<sub>695</sub> expression construct

(695-1, 695-2) or vector alone (CEP4 $\beta$ ). For details see (43). **(D)** Immunoprecipitation of metabolically labeled derivatives released into the medium. Medium from the cells in (C) that were metabolically labeled for 16 hours was examined with antisera to the NH<sub>2</sub>-terminal portion of the  $\beta$ APP (anti-N) and to the  $\beta$ AP (anti- $\beta$ AP) as in (43). Anti- $\beta$ AP immunoprecipitated a protein of about the same size as the derivative immunoprecipitated by anti-N. We presume that this protein is an authentic  $\beta$ APP derivative, because it was augmented in transfected cells (see also the artifact at the 10/16% interface that shows no augmentation). SGY2134 may weakly immunoprecipitate the same derivative recognized by the anti-N antibody, but it could also be immunoprecipitating a subset of the derivatives recognized by anti-N or even an entirely different derivative. Note that M17- $\beta$ APP<sub>695</sub> cells release not only 4-kD  $\beta$ AP but several smaller derivatives that are detected by anti- $\beta$ AP.

$\beta$ AP may be a relatively indigestible  $\beta$ APP fragment that is then externalized (25, 26). Additional data are clearly needed, however, to establish the role of lysosomal processing in  $\beta$ AP production and to rule out the possibility that chloroquine and ammonium chloride are interfering with a different processing pathway.

To determine whether  $\beta$ AP is also released into human cerebrospinal fluid (CSF) and by cultured cells expressing full-length  $\beta$ APP, we used the same criteria that were successful in assaying synthetic  $\beta$ AP<sub>1-40</sub> and the  $\beta$ AP released from K562-LC<sub>99</sub> cells: immunoprecipitation by SGY2134 of a 4-kD protein that can be detected on 4G8 immunoblots. In our initial experiment on CSF, we analyzed 3-ml samples obtained at autopsy from seven patients with AD and seven control subjects (Fig. 2A). Strong signals were obtained in CSF from five of the seven autopsy-confirmed AD patients, but considerable  $\beta$ AP was also present in samples from three of the seven control subjects. We also examined 3-ml samples of CSF from living patients, five from patients with probable AD and five from age-matched patients without AD (Fig. 2B). Again, there was considerable overlap in the amount of  $\beta$ AP observed in the AD and non-AD group. Thus our initial survey indicates that there is (i) readily detectable  $\beta$ AP in the CSF of AD and control patients, (ii) considerable apparent interindividual variation in the amount of  $\beta$ AP in CSF, and (iii) no obvi-

ous correlation between AD and the amount of  $\beta$ AP in CSF.

To determine whether  $\beta$ AP is released by cultured cells expressing full-length  $\beta$ APP, we analyzed medium conditioned with human neuroblastoma (M17) cells stably transfected with a  $\beta$ APP<sub>695</sub> expression construct. In these experiments, the transfected M17 cells were differentiated for 7 days with retinoic acid to induce the formation of long neurites. M17- $\beta$ APP<sub>695</sub> cells (Fig. 3A) released readily detectable amounts of a 4-kD protein that was immunoprecipitated by SGY2134 (anti- $\beta$ AP<sub>1-16</sub>) and labeled by 4G8 (anti- $\beta$ AP<sub>17-24</sub>). On immunoblots this protein was indistinguishable from synthetic  $\beta$ AP<sub>1-40</sub> or the  $\beta$ AP released by K562-LC<sub>99</sub> cells (Fig. 1).

To definitively identify the 4-kD protein released from M17- $\beta$ APP<sub>695</sub> cells, we metabolically labeled these cells with [<sup>3</sup>H]phenylalanine for 16 hours, isolated the 4-kD protein released into the culture medium, and radiosequenced it. The  $\beta$ AP, which begins with the sequence Asp-Ala-Glu-Phe, has phenylalanine at positions 4, 19, and 20. Radiosequencing of the 4-kD protein released by M17- $\beta$ APP<sub>695</sub> cells showed [<sup>3</sup>H]phenylalanine at precisely those positions (Fig. 3B), indicating that this protein is  $\beta$ AP.

In order to better quantitate the  $\beta$ AP released from these cells, M17- $\beta$ APP<sub>695</sub> cells and M17 cells transfected with vector alone (M17-CEP4 $\beta$ ) were metabolically labeled with [<sup>35</sup>S]methionine for 20 min to analyze the rate of  $\beta$ APP synthesis and for

16 hours to analyze the  $\beta$ APP and  $\beta$ APP derivatives released into the medium and accumulating in cells (Fig. 3, C through E). As expected, the immature form of  $\beta$ APP<sub>695</sub> predominated in cell lysates examined after 20 min of labeling (Fig. 3C). With 16 hours of continuous labeling, cells accumulated considerable full-length  $\beta$ APP, particularly the mature form (Fig. 3C); they accumulated readily detectable amounts of the 8- to 12-kD COOH-terminal derivatives (Fig. 3C); and they secreted large amounts of the ~105-kD NH<sub>2</sub>-terminal derivative produced by  $\beta$ APP secretase (Fig. 3D, lanes 2 and 3).

Significantly, M17- $\beta$ APP<sub>695</sub> cells continuously labeled for 16 hours released the 4-kD  $\beta$ AP immunoprecipitated by SGY2134 in amounts that permitted detection after exposing the fluorogram for only 29 hours (Fig. 3D). After a 3-day exposure of the fluorograms (Fig. 3D), it was evident that M17 cells transfected with vector alone (which endogenously express much more  $\beta$ APP than K562-CEP4 $\beta$  cells) also release the 4-kD  $\beta$ AP. Thus even M17 cells expressing only endogenous  $\beta$ APP release the 4-kD  $\beta$ AP.

To compare the various derivatives produced by M17- $\beta$ APP<sub>695</sub> cells, we analyzed them with phosphorimaging technology. Since good estimates of the number of radio-labeled amino acids in each derivative can be made, the relative number of moles of derivative present in each dish can be accurately assessed by dividing the per dish signal obtained for each derivative by the number of

radiolabeled amino acids that it contains. This analysis showed that, in M17- $\beta$ APP<sub>695</sub> cells labeled continuously for 16 hours, the 4-kD  $\beta$ AP accumulates in the medium at about 40% of the amount of the major ~105-kD derivative. Thus, M17 cells externalize a significant amount of the ~4-kD  $\beta$ AP.

Our analysis of human CSF and transfected cells indicates that the  $\beta$ APP is normally cleaved both in its membrane-spanning domain on the COOH side of the  $\beta$ AP, and in its extracellular-intraluminal region on the NH<sub>2</sub> side of the  $\beta$ AP to release a soluble 4-kD derivative essentially identical to the  $\beta$ AP that forms amyloid in AD (27). Thus it is likely that amyloid deposition in AD involves pathways that normally produce extracellular  $\beta$ AP and that the amount of amyloid deposited depends on the rate of  $\beta$ AP production, its rate of removal, and the rate at which soluble  $\beta$ AP forms insoluble amyloid fibrils. It will, therefore, be important to exploit cultured cells to identify (i) the specific pathway and proteases that produce and release  $\beta$ AP, (ii) factors that foster  $\beta$ AP production, and (iii) potential therapeutic agents that can prevent  $\beta$ AP production. In addition, it will be important to identify the mechanism or mechanisms that remove soluble extracellular  $\beta$ AP and to clarify the factors (28–32) that cause soluble, extracellular  $\beta$ AP to form amyloid fibrils.

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- On the basis of its size and solubility, the 4-kD  $\beta$ AP identified here appears to be most closely related to the  $\beta$ AP<sub>1–39</sub> that forms cerebrovascular amyloid fibrils (8) and the 4-kD  $\beta$ AP recently identified (33, 34) as the major constituent of diffuse plaques (preamyloid deposits) found in the AD brain and to a lesser extent in the normal aged brain. The amyloid in classic senile plaques, which is less soluble than preamyloid or vascular amyloid, consists primarily of  $\beta$ AP<sub>1–42</sub> or  $\beta$ AP<sub>1–43</sub>. The  $\beta$ APP processing that produces the soluble 4-kD  $\beta$ AP identified in this report may produce  $\beta$ AP<sub>1–42</sub> or  $\beta$ AP<sub>1–43</sub>, but this peptide would be less soluble and thus more difficult to detect than the slightly shorter  $\beta$ AP that forms cerebrovascular amyloid fibrils or preamyloid.
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- LC<sub>99</sub> was expressed by using an episomal replicon (CEP4 $\beta$ ) similar to that described (36) except that expression was driven by a CMV promoter. To assemble CEP4 $\beta$ -LC<sub>99</sub>, we used a  $\beta$ APP<sub>695</sub> template and appropriate oligonucleotides to amplify by the polymerase chain reaction (PCR) two  $\beta$ APP cDNAs. Using oligos 1 and 2, we amplified a cDNA with a Hind III restriction site followed by bases –30 to 54 and bases 1789 to 1809; with oligos 3 and 4, we amplified a cDNA with bases 39 to 54, followed by bases 1789 to 2119 and a Bam HI restriction site [numbering is according to (7)]. The two amplified cDNAs were gel-purified and mixed in equimolar amounts, and the hybrid was amplified in a second PCR. The amplified cDNA was cut with Hind III and Bam HI and cloned into CEP4 $\beta$  to produce a vector containing untranslated 5' sequence (–30 to –1) and the adjacent sequence (1 to 54) encoding  $\beta$ APP<sub>1–18</sub> (signal peptide plus leucine) followed by sequence encoding C<sub>99</sub> (1789 to 2085) and 3' untranslated sequence (2086 to 2119). The oligos employed were (1) CCCAAGCTTCACAGCT-AGCGCACTCGGTGCCCGCGCAGGGTCTG, (2) GTCATGTCTGGAATTCTGCATCCAGCGCCCGA-GCCGT, (3) TGGACGGCTCGGGCGCTGGATG-CAGAATTCCGACAT, and (4) GACTCGAGTCTGACGGATCTGTCCAACTTCAGAGGCTGC. Stable lines expressing CEP4 $\beta$  or CEP4 $\beta$ -LC<sub>99</sub> were established by transfection with lipofectin (BRL).
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- For immunoprecipitation of  $\beta$ AP, 5 to 20 ml of medium was made 1× in RIPA (150 mM NaCl, 10 mM tris, pH 8.0, 1% Nonidet P-40, 0.5% cholic acid, 0.1% SDS, and 5 mM EDTA)–LPT (1  $\mu$ g/ml leupeptin, 0.1  $\mu$ g/ml pepstatin, 2 mM phenylmethyl sulfonylfluoride, and 1  $\mu$ g/ml 7-amino-1-chloro-3-tosylamido-2-heptanone) buffer; SGY2134 (20  $\mu$ l) and RIPA-LPT-washed protein A-agarose slurry (40  $\mu$ l) were added; and the medium was rocked at 4°C for 24 hours. After washing three times with RIPA-LPT and once with TBS (10 mM tris, pH 8, 150 mM NaCl), the immunoprecipitated protein was separated by 10/16% tris-tricine SDS–polyacrylamide gel electrophoresis (PAGE), transblotted to Immobilon P (Millipore) at 100 V for 2 hours, incubated with 5% dried milk for 1 hour, and labeled with 4G8 (1:500) for 3 hours. Fluorograms were then prepared with an ECL Western Blotting Kit (Amersham) according to the manufacturer's specifications.
- Medium (400 ml) conditioned with K562-LC<sub>99</sub> cells was concentrated to 11 ml (Amicon concentrator, 3-kD cutoff filter). The 4-kD  $\beta$ AP was then immunoprecipitated with anti- $\beta$ AP<sub>1–40</sub> antibody (SGY2134), separated by tris-tricine SDS-PAGE, and transblotted to Immobilon P as in (37). After visualization with Coomassie blue R-250, the 4-kD protein was excised and thoroughly washed with distilled H<sub>2</sub>O, and the sequence shown in Fig. 1D was obtained on a 477A Sequencer with 120A-PTH analyzer (Applied Biosystems).
- Each of five 20-ml aliquots of medium were conditioned with K562-LC<sub>99</sub> cells at 5 × 10<sup>5</sup> cells per milliliter for 6 hours, and the control medium (0 to 6 hours) from each set of cells was removed. Fresh control medium was added to the first set of cells; medium containing leupeptin (100  $\mu$ g/ml), E-64 (100  $\mu$ M), and ammonium chloride (20 mM) or chloroquine (100  $\mu$ M) was added to the other sets of cells. After 6 hours of conditioning, this medium (6 to 12 hours) was removed. A fresh 20-ml aliquot of the identical medium was then added for six additional hours of conditioning (12 to 18 hours).
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- Two M17 lines stably transfected with  $\beta$ APP<sub>695</sub> (different from those in Fig. 3A) and a line stably transfected with CEP4 $\beta$  alone were generated, selected, and maintained with Hygromycin (500  $\mu$ g/ml) as in (18). Triplicate 10-cm dishes containing each of these lines were treated with retinoic acid (1  $\mu$ M) for 7 days. Confluent dishes were then labeled for 20 min (Fig. 3C) or 16 hours (Fig. 3, C and D) in 5 ml of methionine-deficient medium (Gibco) containing 0.8 mCi of [<sup>35</sup>S]methionine. For Fig. 3C, the cells in each 10-cm dish were lysed and immunoprecipitated for 3 hours at 4°C with antiserum to the COOH-terminus of the  $\beta$ APP (5  $\mu$ l) and protein A-agarose slurry (25  $\mu$ l) as in (18). For Fig. 3D, the medium covering the triplicate dishes in Fig. 3C (16-hour pulse) was pooled to give 15 ml of conditioned medium. Immunoprecipitation was performed as in (37). From each pool (CEP4 $\beta$ , 695-1, and 695-2), 1 ml was immunoprecipitated for 3 hours at 4°C with 10  $\mu$ l of anti-N and 25  $\mu$ l of protein G-agarose slurry (lanes 1 to 3), and 6 ml were immunoprecipitated for 24 hours at 4°C with 20  $\mu$ l of anti- $\beta$ AP and 40  $\mu$ l of protein A-agarose (lanes 4 to 6). Anti-N is a goat polyclonal antiserum (207) raised to a truncated form of  $\beta$ APP<sub>770</sub> ( $\beta$ APP<sub>18–688</sub>). Anti- $\beta$ AP is a rabbit polyclonal antiserum (SGY2134) raised to synthetic  $\beta$ AP<sub>1–40</sub> that contains antibodies directed primarily to  $\beta$ AP<sub>1–16</sub>. Immunoprecipitates in Fig. 3, C and D, were separated by 10/16% tris-tricine SDS-PAGE.
- For radiosequencing, each of four confluent 10-cm dishes of M17- $\beta$ APP<sub>695</sub> cells was labeled for 16 hours in 5 ml of phenylalanine-deficient medium (Gibco) containing 1.0 mCi of [<sup>3</sup>H]phenylalanine. The labeled 4-kD protein in each dish was immunoprecipitated, separated, and transferred to Immobilon P as in (37). After autoradiographic visualization, the 4-kD protein excised from each of four lanes was pooled and sequenced as in (38), and the <sup>3</sup>H signal released in each cycle was analyzed.
- We thank S. G. Speciale for providing the CSF samples from living patients, A. LeBlanc for anti-C<sub>21</sub>, K. S. Kim for 4G8, B. Greenberg for 207, and B. Spengler and J. Biedler for the M17 subclone of their BE(2) neuroblastoma line. Supported by NIH grants AG05891 and AR02594 (B.F.) and AG06656 (S.G.Y.) and an ADRDA Zenith award (S.G.Y.).

7 August 1992; accepted 11 September 1992