Intracellular domains of amyloid precursor-like protein 2 interact with CP2 transcription factor in the nucleus and induce glycogen synthase kinase-3 β expression

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Amyloid precursor protein (APP) is a member of a gene family that includes two APP-like proteins, APLP1 and 2. Recently, it has been reported that APLP1 and 2 undergo presentiin-dependent y-secretase cleavage, as does APP, resulting in the release of an \sim 6 kDa intracellular C-terminal domain (ICD), which can translocate into the nucleus. In this study, we demonstrate that the APLP2-ICDs interact with CP2/LSF/LBP1 (CP2) transcription factor in the nucleus and induce the expression of glycogen synthase kinase 3β (GSK- 3β), which has broad-ranged substrates such as τ - and β -catenin. The significance of this finding is substantiated by the in vivo evidence of the increase in the immunoreactivities for the nuclear C-terminal fragments of APLP2, and for GSK-3 β in the AD patients' brain. Taken together, these results suggest that APLP2-ICDs contribute to the AD pathogenesis, by inducing GSK-3 β expression through the interaction with CP2 transcription factor in the nucleus.

Cell Death and Differentiation (2007) 14, 79–91. doi:10.1038/sj.cdd.4401928; published online 28 April 2006

The pathology of Alzheimer's disease (AD) is characterized by the deposition of neuritic plaques, of which the principal components are 40 and 42 amino-acid amyloid beta peptides $(A\beta)$ derived from amyloid precursor protein (APP).¹ APP belongs to a family of conserved type I transmembrane proteins including Apl-1 in Caenorhabditis elegans,² Appl in Drosophila,³ and APP, APP-like protein 1 (APLP1)⁵ and APLP2^{6,7} in mammals.

APLP1 and 2 display substantial amino-acid and domain homologies with APP. Although they do not have an $A\beta$ domain, their intracellular C-terminal domains (ICDs) are highly similar to that of APP.⁸ APLP1 is known to be implicated in synaptogenesis,9,10 and recombinant APLP2 possesses an ability to promote neurite outgrowth.¹¹ However, their physiological roles in neurons are still not fully understood. In the brains of AD patients, APLP2 immunoreactivity has been detected in a subset of neuritic plaques,¹² suggesting that APLP2 might be involved in the pathogenesis of AD.

APLP2 matures through the same secretory/cleavage pathway as APP⁷ and previously, APLP1 and 2 were reported to be processed by γ -secretase in a presenilin 1-dependent manner.^{8,10} The APLP family can be also cleaved by ε secretase to generate the ICD composed of the last 50 amino acids of APLPs C-terminus (C50).^{13,14} Moreover, the ICDs of APLP1 and APLP2 produced by y-secretase enhanced Fe65dependent gene transcriptional activation, as have been reported for the APP intracellular domain (AICD).¹⁰

Fe65, which is known to be ones of the adaptor proteins for APP, contains three protein-protein interaction domains, a WW and two phosphotyrosine binding (PTB) domains. The PTB2 domain, which is located in the C-terminal half of the molecule, is responsible for the interaction between Fe65 and the cytosolic tail of APP through the YENPTY domain of APP.¹⁵

Here, we demonstrate that the ICDs of APLP2 (APLP2-ICDs: C57, C50) interact with CP2 transcription factor in the nucleus and induce glycogen synthase kinase 3β (GSK- 3β) expression, whereas their point mutants with Y738G in the ⁷³⁸YENPTY⁷⁴³, an Fe65 binding domain, do not. CP2 is a ubiquitously expressed 66-kDa protein, which regulates several genes including GSK-38.16 Polymorphic variation in the CP2 gene has been known to be one of the AD risk factors.¹⁷

We also provide in vivo evidence that the immunoreacitivities for C-terminal fragments (CTFs) of APLP2 detected by C12 antibody, which specifically recognizes the last 12 amino acids of APLP2 and for GSK-3 β are significantly upregulated in the AD brains.

Results

APLP2-ICDs are translocated into the nucleus in a Fe65dependent manner. At first, we confirmed whether APLP2-ICDs were detectable in the nucleus after transfecting C57 or

Received 04.8.05; revised 26.1.06; accepted 01.3.06; Edited by P Nicotera; published online 28.4.06

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Keywords: amyloid precursor-like protein 2; Alzheimer's disease: CP2 transcription factor; Fe65; GSK-3β; τ

Abbreviations: APLP, amyloid precursor like protein; APP, amyloid precursor protein; AICD, APP intracellular domain; A β , amyloid beta protein; AD, Alzheimer's disease; PTB, phosphotyrosine binding domain; GSK-3 β , glycogen synthase kinase-3 β ; CTFs, C-terminal fragments; C57, C50, 57, 50 amino-acid residues carboxy terminal fragment of APLP2; C57^{Y738G}, C50^{Y738G}, Y738G point mutated C57, C50; HEK, human embryonic kidney; DAPT, N-[N-(3,5-difluorophenacetyl)-L-alanyl]-Sphenylglycine *τ*-butyl ester; NGF, nerve growth factor; NFT, neurofibrillary tangle

C50 (Figure 1B (a)) at 48-h post-transfection. APLP2-ICDs were detected in the nuclear fractions of C57- or C50transfected HEK293 cells using C12 antibody, which specifically recognizes the last 12 amino acids of the C-terminus of APLP2, not crossreacting with the C-terminus of APP¹⁸ or anti-GFP antibody (Figure 1B (a)). In addition, we also examined the distribution of APLP2-ICDs in the nuclear and cytosolic fractions of HEK293 cells co-transfected with Fe65 and either APLP2-ICDs or their point mutants (Y738G). The point mutation of Y738 to G in the ⁷³⁸YENPTY⁷⁴³ domain was reported to significantly impair Fe65 binding to APLPs.¹⁰ APLP2-ICDs were more abundantly detected in the nuclear fractions of the mock or the Y738G point mutants transcfected cells. By contrast, in the cytosolic fractions, relatively small amount of APLP2-ICDs was detected in the cells overexpressing APLP2-ICDs compared to the cells overexpressing their point mutants (Figure 1B (b and c)). This result illustrated in Figure 1B (b) suggests that binding with Fe65 plays a role in nuclear translocation of APLP2-ICDs, as proposed before.⁸ Next, the presence of APLP2-ICDs was also found in the nuclear fractions of HEK293 cells overexpressing full-length APLP2-751 (Figure 1B (d)). The purity of the nuclear or cytosolic fraction was checked using anti-calnexin, anti- β -tubulin and anti-histone 3 antibodies (Figure 1B (a–d)).

We then investigated the subcellular localizations of APLP2-ICDs and their colocalization with Fe65 by confocal

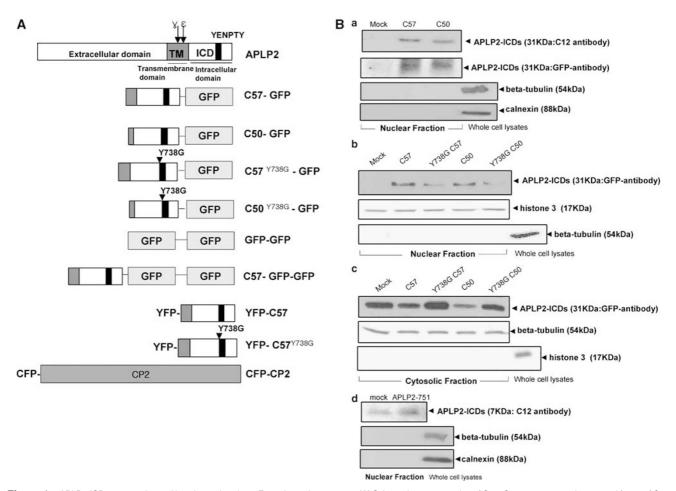
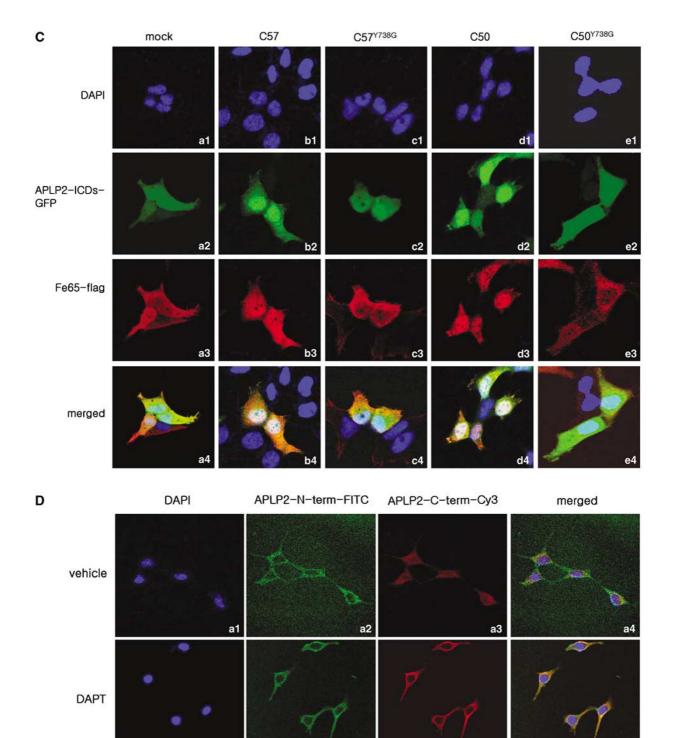


Figure 1 APLP2-ICDs are translocated into the nucleus in an Fe65-dependent manner. (**A**) Schematic representation of C57, C50 constructs, point mutated forms of C57 and C50 (C57^{Y736G}, C50^{Y736G}, C10^{Y736G}, C10^{Y736G}

laser scanning microscopy in HEK293 cells co-transfected with APLP2-ICDs (C57, C50, C57^{Y738G} or C50^{Y738}) and Fe65. GFP and Cy3 fluorescence signals indicating APLP2 and Fe65 immunoreactivities, respectively, showed that APLP2-ICDs (C57, C50) and Fe65 were colocalized mainly in the nucleus of the HEK293 cells and also in the cytoplasm

(Figure 1C (b1–b4 and d1–d4)), whereas the GFP signals corresponding to the empty vector or point mutated APLP2-ICDs ($C57^{Y738G}$, $C50^{Y738}$) were observed evenly throughout the cytoplasm and nucleus (Figure 1C (a1–a4, c1–c4 and e1–e4)). Higher level of Fe65 immunoreactivity in the cytoplasm of cells expressing mock or Y738G mutated



b

APLP2-ICDs led us to hypothesize that the nuclear localization of Fe65 is also dependent on APLP2-ICDs (Figure 1C (a1–a4, c1–c4 and e1–e4)), consistent with other previous report.¹⁰

The C57 or C50-GFP fusion proteins have an approximately 31 kDa molecular weight, which is below the exclusion limit of size to pass the nuclear membrane by passive

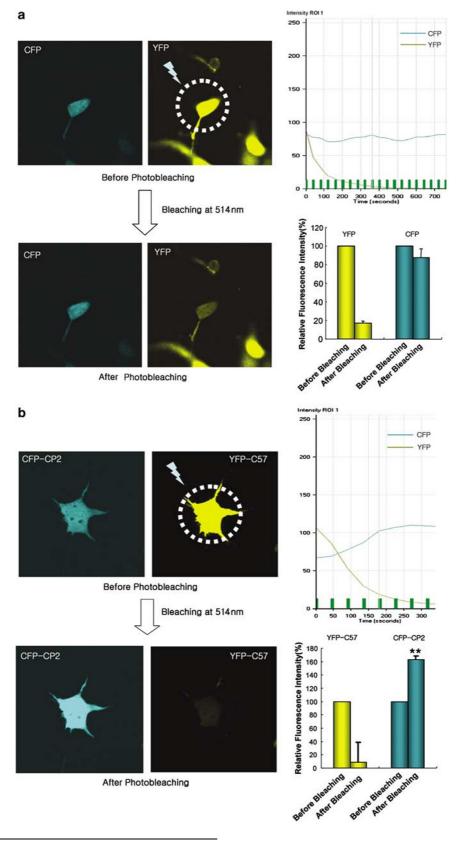
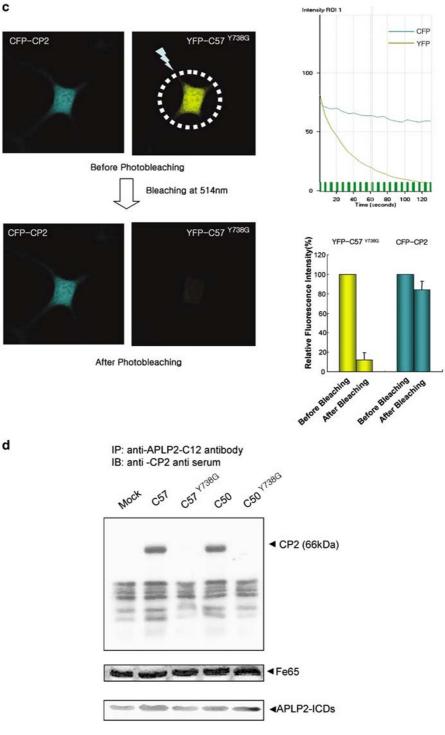


Figure 2 (Continued)

npg 82 diffusion. To investigate whether APLP2-ICDs could translocate into the nucleus by an active transport system, we generated construct encoding fusion protein composed by two consecutive GFP units fused to C57 (Figure 1A). As shown in Supplementary Figure S1a, GFP-GFP fusion protein is mostly found in the cytoplasm, as expected from its molecular size, whereas the C57-GFP-GFP is observed mainly in the nucleus despite of its large molecular size (Supplementary Figure S1b).

In Figure 1D, HEK293 cells were co-transfected with fulllength APLP2-751 in the pCB6 vector and Fe65 in the pcDNA3 flag vector. The transfected cells were incubated in the presence or in the absence of DAPT {N-[N-(3,5difluorophenacetyl)-L-alanyl]-S-phenylglycine t-butyl ester}, an inhibitor of γ -secretase, at 2 μ M for 48 h, and then labeled with C12 antibody and APLP2 (C17) antibody, which specifically recognizes the APLP2 N-terminal extracellular domain, conjugated with Cy3 or FITC, respectively.

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Vehicle-treated cells showed APLP2-CTFs immunoreactivity in the nucleus (Figure 1D (a1, a3 and a4)). By contrast, the APLP2-CTFs immunoreactivity represented by Cy3 fluorescence was not observed in the nucleus of the cells treated with DAPT (Figure 1D (b1, b3 and b4)). The FITC fluorescence representing the presence of APLP2 N-terminal extracellular domain was not observed in the nucleus in the presence or absence of DAPT treatment (Figure 1D (a1, a2, a4, b1, b2 and b4)).

Molecular interaction between C57 CP2 and transcription factor by fluorescence resonance energy transfer (FRET) analysis. Fe65 is reported to interact with CP2 transcription factor through its PTB1 domain.¹⁹ Here, we examined whether C57 interacts with CP2 transcription factor by employing an FRET method (Figure 2(a-c)). HEK293 cells on coverslips in six-well plates were transiently co-transfected with one of the following combinations: (1) pEYFP vector and pECFP vector, (2) YFP-C57 and CFP-CP2, (3) YFP-C57^{Y738G} and CFP-CP2. At 24 h after transfection, the coverslips were mounted on Axiovert 200 inverted microscope (Zeiss, Jena Germany). The energy transfer was detected as an increase in donor fluorescence (ECFP) after complete photobleaching of the acceptor molecules (EYFP). Cells showing both of ECFP and EYFP signals were photobleached at 514 nm (laser power 100%) to destroy the acceptor molecules. The cells were rescanned using 458-nm light and an increase of the ECFP signal within the photobleached area was used as a measure of FRET. As shown in Figure 2a, in the cells co-expressing pECFP and pEYFP vectors only, CFP donor signal was rather slightly decreased to $87.8\pm3.1\%$, after photobleaching of YFP signal at 514 nm. In Figure 2b, CFP donor signal was significantly increased to 163.3±13.2% (P < 0.01, n = 7), after complete photobleaching of YFP signal at 514 nm, indicating that FRET is present between CFP-CP2 and YFP-C57. By contrast, FRET was not observed between YFP-APLP2-C57^{Y738G} and CFP-CP2, suggesting that YENPTY, an Fe65 interacting domain is critically required for the interaction between C57 and CP2 transcription factor (Figure 2c). HEK293 cells are expressing endogenous Fe65, so it remains to be determined whether C57 and CP2 interact directly or indirectly through Fe65.

We confirmed these results by using an immunoprecipitation method (Figure 2d). APLP2-ICDs tagged with a GFP sequence and Flag-tagged Fe65 were transiently co-expressed in HEK293 cells, and it was confirmed that all transfected proteins were equally expressed in all the groups (Figure 2d). The immunoprecipitation of APLP2-ICDs by C12 antibody and immunoblotting for CP2 demonstrated that the transfected APLP2-ICDs (C57, C50) and endogenously existing CP2 interacted in the nuclear fraction (Figure 2d), whereas the Y738G point mutated APLP2-ICDs (C57^{Y738G}, C50^{Y738G}) did not (Figure 2d). Our findings demonstrate that the APLP2-ICDs and CP2 interact in the nucleus and that interactions between these proteins are governed by YENPTY domain in APLP2-ICDs.

APLP2-ICDs induce GSK-3 β expression, and τ phosphorylation in NGF-differentiated PC12 cells and rat primary cortical neurons whereas the Y738G point mutants do not. Based on the finding that APLP2-ICDs and CP2 transcription factor interact in the nucleus, we examined changes in GSK-3 β , one of the genes regulated by CP2 transcription factor. GSK-3 β protein was upregulated in C57- or C50 expressing HEK293, NGF-differentiated PC12 cells and in rat primary cortical neurons, but not in cells expressing C57^{Y738G}, C50^{Y738G} (Figure 3a). We also checked for changes in Tyr²¹⁶ phosphorylated GSK-3 β , an active form of this enzyme. Phosphorylated GSK-3 β at Tyr²¹⁶ was increased in C57 or C50 expressing NGF-differentiated PC12 cells, but was not increased in the Y738G point mutated APLP2-ICDs (C57^{Y738G}, C50^{Y738G}) transfected cells (Figure 3a). It has been reported that the activity of GSK-3 β is mediated by the phosphorylation of two distinct sites, and by transcriptional regulation.²⁰ Tyr²¹⁶ phosphorylation, which is necessary for its activity, is known to increase in response to apoptotic stimuli in neuronal cells.²¹ The GFP tagged APLP2-ICDs were confirmed to be equally expressed in mock, APLP2-ICDs and Y738G mutated APLP2-ICDs transfected cells, by using anti-GFP antibody (data not shown).

The mRNA level of GSK-3 β was also increased in APLP2-ICDs transfected PC12 cells (Figure 3b). However, forms containing the Y738G point mutation in the ⁷³⁸YENPTY⁷⁴³ domain of APLP2-ICDs (C57^{Y738G}, C50^{Y738G}) did not upregulate the GSK-3 β at either the protein or mRNA level in PC12 cells, indicating that YENPTY, an Fe65 interacting domain is critical for the induction of GSK-3 β expression (Figure 3 (a and b)).

Figure 2 Molecular interaction between APLP2-C57 and CP2 transcription factor examined by FRET. HEK293 cells on coverlips in six well plates were transiently cotransfected with one of the following combinations: (a) pEYFP vector and pECFP vector, (b) YFP-C57 and CFP-CP2, (c) YFP-C57^{Y738G} and CFP-CP2. At 24 h after cotransfection, the coverslips were mounted on Axiovert 200 inverted microscope (Zeiss, Jena Germany). An initial scan was obtained at low energy using the 458 line of the argon laser to record the ECFP signal. A second scan was performed with the 568 line, and cells of colocalization was noted. Cells were then photobleached with intense 514 nm light (laser power 100%) to destroy the acceptor molecules, EYFP. The energy transfer was detected as an increase in donor fluorescence (ECFP) after complete photobleaching of the acceptor molecules (EYFP) at 514 nm. The amount of energy transfer was calculated as the percent increase in donor fluorescence after acceptor photobleaching. An increase of the ECFP signal within the photobleached area was used as a measure of the amount of FRET present. Figures in the left panel show representative results from the seven independent experiments. The right upper panel shows a representative graph indicating the time-lapse fluorescence changes in the region of interest (ROI). The green line represents the ECFP fluorescence; gray, EYFP throughout the progression of successive bleaching marked as vertical green bars. In the right lower panel, relative fluorescence intensity is denoted after standardization of intensities before and after photobleaching of EYFP at 514 nm. **P*<0.05 *versus* fluorescence intensity before photobleaching of YFP at 514 nm (by one-way ANOVA). To confirm the interactions between APLP2-ICDs and CP2, we performed immunoprecipitated with anti-APLP2 (C12) antibody, immunoblotting with anti-CP2 antiserum. The expression levels of APLP2-ICDs in pEGFP-N1 in HEK293 cells, were checked using anti-GFP antibody

To investigate whether endogenously generated APLP2-ICDs affect GSK-3 β expression, we examined the effect of DAPT on GSK-3 β levels after transfecting full-length APLP2 into HEK293 cells and found that the GSK-3 β was not upregulated in the DAPT treated cells, indicating that the ICDs generation by γ -secretase activity is critical for GSK-3 β expression (Figure 3c). We confirmed a significant decrease in ICD production by DAPT treatment both in the nuclear fraction and in whole-cell lysates of HEK293 cells over-expressing full-length APLP2 (data not shown).

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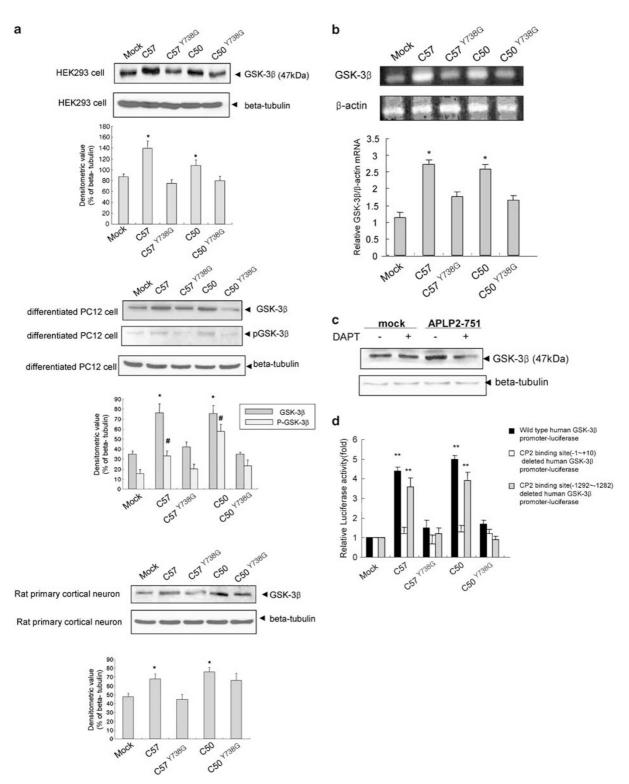


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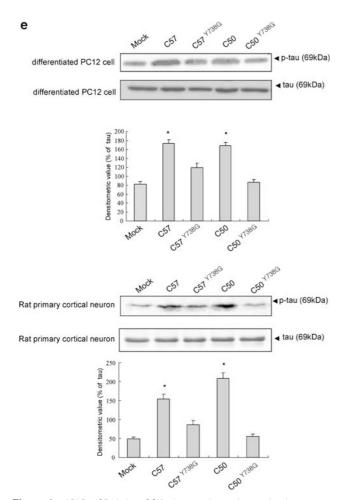


Figure 3 APLP2-ICDs induce GSK-3 β expression, and upregulate its promoter activity and τ phosphorylation in NGF-differentiated PC12 cells, rat primary cortical neurons and HEK293 cells. (a) At 48-h post-transfection, the changes in GSK-3 β protein and p-GSK-3 β at Tyr²¹⁶ were checked by Western blotting with anti-GSK-3 β or anti-p-GSK-3 β at Tyr²¹⁶ antibodies in HEK293 cells, PC12 cells and rat primary cortical neurons transfected with APLP2-ICDs. The graph shows the densitometric value of the GSK-3 β or p-GSK-3 β versus β -tubulin (* (GSK-3 β), #P<0.05 (p-GSK- 3β) by one-way ANOVA). (b) At 48-h post-transfection, GSK- 3β mRNA levels were determined by RT-PCR in PC12 cells transfected with APLP2-ICDs. The relative ratios of the levels of GSK-3 β and β -actin mRNA were normalized using the M4 image analysis program (*P < 0.05, by one-way ANOVA). (c) At 48-h posttransfection, GSK-3 β levels were examined in HEK293 cells expressing full-length APLP2, which had been incubated in the presence or absence of DAPT, a ysecretase inhibitor. (d) GSK-3 β promoter activity was measured by luciferase activity assay in PC12 cells cotransfected with APLP2-ICDs and wild-type human GSK-3 β promoter or deletion mutants of human GSK-3 β promoter without nt -1292 to ~ 1282 or nt -1 to $\sim +10$ in the pGL2 vector. Data represent the means \pm S.E. of four separate experiments. (**P < 0.01 by one-way ANOVA). (e) Changes in phosphorylated tau at the Ser²⁰² and Thr²⁰⁵ residues were checked using AT8 antibody in PC12 cells and rat primary neurons expressing APLP2-ICDs. The graph shows the densitometric value of P-tau versus tau (*P<0.05 by one-way ANOVA)

In order to confirm the effects of APLP2-ICDs on GSK-3 β induction, GSK-3 β promoter activity was measured by luciferase activity assay after co-transfecting APLP2-ICDs and human GSK-3 β promoter in the pGL2 vector. GSK-3 β promoter activity was significantly increased by about four- to five-fold in PC12 cells expressing C57 or C50. No significant

increase was observed in the cells expressing APLP2-ICDs containing a Y738G point mutation (Figure 3d).

GSK-3 β promoter has potential c-Myb, CRE, AP1, AP2, Tst-1, Sp1 and MZF-1 binding sites in addition to two CP2 binding sites. Putative CP2 binding sites were identified at nt -1292 to ~ -1282 and nt -1 to $\sim +10$ by Lau *et al.*,²⁰ and it has been reported that very low reporter gene expression is found when the region between nt -171 and +29 was deleted, indicating that essential elements had been removed. In our study, we prepared deletion mutants without nt -1292 to -1282 (GCGCACACCAA) or nt -1 to $\sim +10$ (GCCCGGGCCAA). Promoter activity was determined by luciferase assay after co-transfecting APLP2-ICDs (C57, C50, C57^{Y738G} and C50^{Y738G}) with one of these deletion mutants of human GSK-3 β promoter in the pGL2 vector. As shown in Figure 3d, deletion of the CP2 binding site nt -1 to \sim + 10 inhibited the promoter activity upregulation by APLP2-ICDs (C57, C50), whereas deletion of CP2 binding site nt -1292 to ~ -1282 did not affect the upregulation of the human GSK-3 β promoter activity by APLP2-ICDs (C57, C50). Our results show that CP2 binding site located at the nt -1 to \sim + 10 sequence in the human GSK-3 β promoter region are essential for the gene transcription by APLP2-ICDs.

τ, whose hyperphosphorylated form is a main component of neurofibrillary tangles (NFTs), is a substrate of GSK-3β.^{22,23} At 48 h post-transfection, we checked the phosphorylation level of τ, at its Ser²⁰² and Thr²⁰⁵ residues by Western blotting in APLP2 ICDs-transfected PC12 cells and in rat primary neurons using AT8 antibody. In APLP2 ICDs-transfected cells, the level of τ phosphorylation at Ser²⁰² and Thr²⁰⁵ residues was found to be significantly increased without changes in nonphosphorylated form of τ. However, no significant increase in τ phosphorylation was found for Y738G mutated APLP2 ICDs-transfected groups (Figure 3e).

The overexpression of APLP2-ICDs decreases cell viabilities in NGF-differentiated PC12 cells, and in HEK293 cells. We evaluated the viabilities of C57, C50, C57^{Y738G}, C50^{Y738G}-transfected differentiated PC12 cells (Figure 4a) and HEK293 cells (Figure 4b) by an LDH release assay. At 48-h post-transfection, cells expressing C57 or C50 showed increased LDH release by about 30% (*versus* the nontransfected controls), while cells expressing Y738G point mutated C57 or C50 (C57^{Y738G}, C50^{Y738G}) showed significantly lower LDH release than those expressing C57 or C50 (Figure 4 (a and b)), indicating that YENPTY, an Fe65 binding domain contributes to cellular toxicity.

To examine whether GSK-3 β is responsible for the reduced cell viability by C57 or C50, we checked the effects of two GSK-3 β inhibitors. LiCl and indirubin-3'-monoxime, 5-iodo were treated for 48 h post-transfection at 10 mM and at 9 nM, respectively, and the cell viabilities were evaluated NGF-differentiated PC12 and HEK293 cells overexpressing APLP2-ICDs. These treatments significantly prevented cellular death induced by APLP2-ICDs (Figure 4 (a and c)), indicating that the GSK-3 β induction by APLP2-ICDs is related to the reduced cell viability. Next, we investigated whether APLP2-ICDs induced apoptosis in HEK293 cells using Hoechst 33258 dye staining. The cells transfected with C57 or C50 showed significant nuclear condensation as

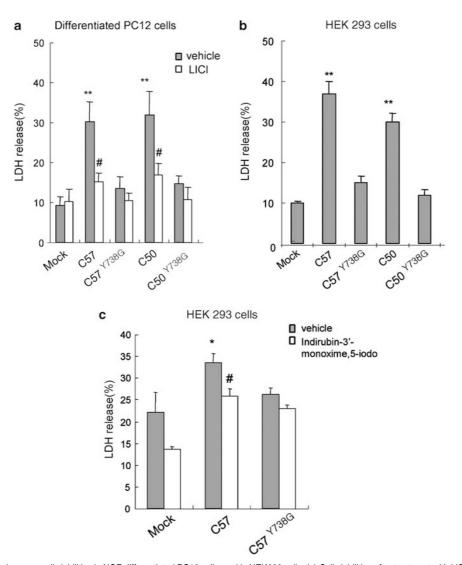


Figure 4 APLP2-ICDs decrease cell viabilities in NGF-differentiated PC12 cells, and in HEK293 cells. (a) Cell viabilities after treatment with LiCl (10 mM) were measured by a LDH assay in PC12 cells expressing APLP2-ICDs. Data represent the means \pm S.E.M. of four separate experiments. (**P<0.01 *versus* mock transfected cells, #P<0.05 *versus* vehicle treated APLP2-ICDs transfected cells, by one-way ANOVA). (b) The cell viabilities of mock, C57-, C50 or Y738G point mutated APLP2-ICDs (C57^{Y738G})-transfected HEK293 cells were assessed by an LDH assay (**P<0.01, *P<0.05, by one-way ANOVA). (c) Cell viabilities after treatment with indirubin-3'-monoxime, 5'iodo (9 nM), were measured by an LDH assay in PC12 cells expressing APLP2-ICDs. Data represent the means \pm S.E.M. of four separate experiments (*P<0.01 *versus* mock transfected cells, #P<0.05 *versus* vehicle treated APLP2-ICDs. Data represent the means \pm S.E.M. of four separate experiments (*P<0.01 *versus* mock transfected cells, #P<0.05 *versus* vehicle treated APLP2-ICDs. Data represent the means \pm S.E.M. of four separate experiments (*P<0.01 *versus* mock transfected cells, #P<0.05 *versus* vehicle treated APLP2-ICDs transfected cells, by one-way ANOVA). (d) HEK293 expressing mock or APLP2-ICDs were stained with Hoechst 33258 dye

shown in Supplementary Figure S2; this finding is consistent with the report that conditional GSK-3 β transgenic mice show TUNEL positive and cleaved caspase-3 immunostained neurons.²⁴

Immunoreacitivities for APLP2-CTFs and GSK-3 β are elevated in the AD brain. The localization of APLP2-CTFs and GSK-3 β in a total of five AD and four nondemented age-matched control cases was examined. Immunostaining of APLP2-CTFs using C12 antibody showed that the cytoplasmic and the nuclear immunoreactivities of APLP2-CTFs are present in the neurons of hippocampus, dentate gyrus and parahippocampal cortex of the human AD brain

(Braak stage V,²⁵) (Figure 5A (a, c, e, g and i)). Especially, nuclear immunoreactivities for APLP2-CTFs in CA1, 2, 3 and parahippocampal cortex were significantly stronger than the same regions of age-matched control brains (Figure 5A (a, c, e and i)). However, the APLP2-CTFs immunoreactivities in dentate gyrus were not significantly different from the age-matched control brains (Figure 5A (g and h)).

The GSK-3 β immunoreactivity was greatly increased in the neurons of the hippocampus (Figure 5B (a and c)), in the neurons of the dentate gyrus (Figure 5B (e)), and in the parahippocampal cortex (Figure 5B (g)) of the AD brains *versus* the normal age-matched control brains (Figure 5B (b and d) (hippocampal neurons), f (dentate gyrus) and h (parahippocampal cortex)).

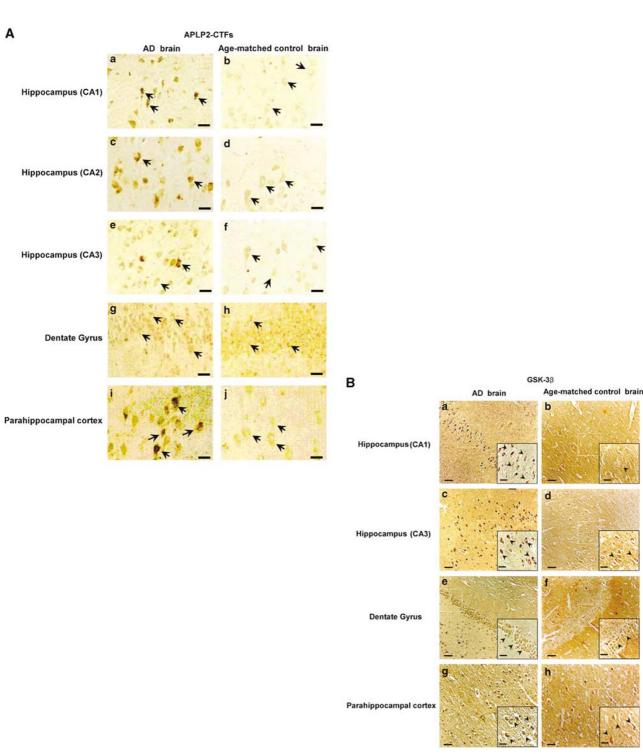


Figure 5 APLP2-CTFs and GSK-3 β immunoreactivities are elevated in the AD brain. The immunoreactivities of APLP2-CTFs (**A**) and GSK-3 β (**B**) were examined in the hippocampus (**A** (a, c, e); **B** (a, c)), the dentate gyrus (**A**(g), **B**(e)), and in the parahippocampal cortex (**A**(i),**B**(g)) of AD brain and normal age-matched control brain (hippocampus, **A**(b, d, f), **B**(b, d); dentate gyrus, **A**(h), **B**(f); parahippocampal cortex, **A**(j), **B**(h)) using C12 antibody which specifically recognizes the last 12 amino acids of APLP2, and anti-GSK-3 β antibody, respectively. Scale bar in (**A**) indicates 50 μ m, in (**B**); lower magnification photographs (\times 100) indicates 100 μ m; in inset (\times 400), 50 μ m. Arrowheads indicate APLP2-CTFs and GSK-3 β , in the cytoplasm; the arrow indicates APLP2-CTFs in the nucleus

Discussion

In the present study, we examined the pathophysiological roles for APLP2-ICDs (C57, C50), generated by γ - or ϵ -

secretases in AD pathogenesis. Among the three members of the APP gene family, the pathophysiological significance of APLP1 and APLP2 in relation to AD has not been adequately investigated until recently.

npg 88 APP has been shown to undergo extensive proteolytic processing by two major pathways.^{26,27} APP, which is a type I transmembrane protein as well as APLPs, is cleaved consecutively, first at the extracellular juxtamembrane region by α - or β -secretase and secondly at the intramembrane region by γ -secretase.²⁶

Similarly with APP, the N-terminal ectodomain of APLP2 is shedded by β -secretase^{28,29} as well as by ADAM 10 and 17,³⁰ whereas the C-termini remain in the membrane^{10,28} and can be further processed by γ - or ε -secretases^{13,14} to release ICDs with signaling properties.^{10,28}

The APLP2-ICDs are known to interact with Fe65 via a YENPTY domain, which APP, APLP1 and APLP2 have in common in their C-terminus, translocating into the nucleus by Fe65-dependent manner⁸ and show Fe65-dependent gene transcriptional activity by heterologous reporter genes.¹⁰ Neprilysin gene transcription is reported to be upregulated by APLP-ICDs as well as the ICDs of APP and APLP1.³¹

In the present study, based on the previous report that the Fe65 PTB1 domain interacts with the CP2 transcription factor,¹⁹ we reasoned that APLP2-ICDs might interact with the CP2 transcription factor in the nucleus. Therefore, we investigated the interaction of APLP2-ICDs with CP2. APLP2-ICD was found to interact with the CP2 transcription factor in the nucleus by employing FRET and immunoprecipitation methods.

CP2 is the prototypical member of a novel mammalian protein family sharing a high degree similarity to Elf-1, a *Drosophila melanogaster* tissue-specific factor.³² By interacting with a hyphenated sequence composed of two directly repeated four base pair motifs repeated by a 6-bp linker (CNRG-N6-CNR [G/C]), a ubiquitously expressed CP2 transcription factor regulates some genes including GSK-3 β , α -macroglobulin and certain factors involved in the inflammatory response that exert T-cell responses when subjected to mitogenic stimulation.^{20,32} A previous report showed that the A allele of the 3'-UTR CP2 gene polymorphism has been reported to increase the risk of sporadic AD.³³

We examined the effects of APLP2-ICDs on the expression of GSK-3 β , a gene known to be regulated by the CP2 transcription factor. The APLP2-ICDs increased both GSK-3 β protein and mRNA levels in HEK293, NGF-differentiated PC12 cells and in rat primary cortical neurons. By contrast, the Y738G point mutated APLP2-ICDs did not affect GSK-3 β expression, indicating that the interaction of ICDs with Fe65 via the YENPTY domain in the C-terminus of APLP2 is essential for the gene-inducing effects of APLP2-ICDs.

GSK-3 β is an important regulator of several cellular processes, which include transcription, cell cycle progression, cell survival and cytoskeletal organization.^{34–37} Increased levels of GSK-3 β have been found in the AD brain, and active GSK-3 β was found to accumulate in pretangle neurons.³⁸ Transgenic mice overexpressing GSK-3 β were shown to display τ hyperphosphorylation, disrupted microtubules, and apoptotic neurons.²⁴ These results imply that alterations in the control of GSK-3 β expression may occur in the AD brain. Our results also show that immunoreactivities for the nuclear APLP2-CTFs, and for GSK-3 β were upregulated in in the AD patients' brain (Figure 5).

NFTs, one of the neuropathological hallmarks of AD, are composed of bundles of paired helical filaments, the major protein subunit of which is abnormally phosphorylated τ , the microtubule-associated protein.³⁹ Hyperphosphorylated τ forms NFTs, one of the hallmarks of AD, and leads to apoptosis by disrupting cytoskeletal and axonal transport.⁴⁰ GSK-3 β has been identified as a primary kinase in this process among several protein kinases including cdc2, cdk5 and MAP kinases.²³ The changes in the levels of phosphorylated τ were examined by using AT8 antibody, which recognizes phosphorylated τ at its Ser²⁰² and Thr²⁰⁵ sites. APLP2-ICDs were found to increase τ phosphorylation in PC12 cells and rat primary cultured neurons, whereas the Y738G point mutated APLP2-ICDs (C57^{Y738G}, C50^{Y738G}) did not.

Here, we demonstrate that APLP2-ICDs induce GSK-3 β expression, through interaction with the CP2 transcription factor in the nucleus, followed by subsequent τ phosphorylation. These findings suggest that APLP2-ICDs may contribute to AD pathogenesis.

Materials and Methods

Materials. Lithium chloride and Hoechst 33258 were purchased from Sigma-Aldrich (MO, USA); Indirubin-3'-monoxime, 5-iodo and DAPT {*N*-[*N*-(3, 5-diffuorophenacetyl)-L-alanyl]-*S*-phenylglycine *t*-butyl ester}, from Calbiochem (Darmstadt, Germany). Anti-GFP monoclonal antibody was obtained from BD Biosciences (CA, USA); anti-APLP2, C-terminal polyclonal antibody (C12; against last 12 amino acids of the C-terminus of APLP2) was a generous gift from Dr Gopal Thinakaran at the University of Chicago. Anti-CP2 serum was a generous gift from Dr. Roeder of Rockefeller University. Anti-Flag M5 monoclonal antibody was purchased from Sigma-Aldrich (MO, USA); anti-human p- τ monoclonal antibody, clone AT8, from Innogenetics (Belgium), anti-APLP2 (C-17) antibody, which recognizes APLP2 N-terminal extracellular domain, anti-GSK-3 β , anti- τ antibodies were obtained from Santa Cruz Biotechnology, Inc. (CA, USA).

DNA constructs and mutagenesis. Mouse APLP2-751 cDNA in the pCB6 vector was a kind gift from Dr Gopal Thinakaran at the University of Chicago. Rat Fe65 cDNA in the pcDNA3 vector was kindly provided by Dr Joseph, D Buxbaum at Mount Sinai School of Medicine. The cDNA constructs encoding different lengths of the APLP2-ICDs were produced as follows (Schematic diagrams of the APLP2-ICDs are shown in Figure 1A). The constructs were generated by PCR from mouse APLP2-751 cDNA to encompass the last 57 or 50 amino-acid residues. The amplified cDNAs were subcloned into Xhol and BamHI sites of the pEGFP-N1 eukaryotic expression vector (BD Biosciences, CA, USA). To construct Y738G of C57 or -C50, we used a QuickChange Site-Directed Mutagenesis Kit (Stratagene, CA, USA). The GFP-GFP expression vector was generated by cloning in the BamHI-Agel sites of the pEGFP-N1 vector (BD Biosciences, CA, USA). The EGFP cDNA was obtained by direct amplification from the same vector. The C57-GFP-GFP was obtained by cutting C57 cDNA from XhoI and BamHI sites and inserted into the same sites of the GFP-GFP vector prepared as described above. YFP-C57 or YFP-C57 with point mutation Y738G was prepared by transferring the Xhol, BamHI cut fragment of C57 or its point mutation cloned in the pEGFP-N1 vector to the pEYFP-C1 vector (BD Biosciences, CA, USA). CFP-CP2 construct was generated by PCR from mouse CP2 in the pRc vector. All constructs were sequenced using an ABI310 Sequencer.

Cell culture and transfection. SH-SY5Y and HEK293 cells were plated in a six-well plate maintained in Dulbecco's modified Eagle's medium (DMEM, Life Technologies, NY, USA) supplemented with 10% fetal bovine serum (FBS) and 0.3% antibiotics at 37°C and 5% CO₂. PC12 cells originating from rat pheochromocytoma were plated on a polyethyleneimine (PEI, 0.2 mg/ml, Sigma-Aldrich, MO, USA) coated plates and maintained in DMEM supplemented with 10% FBS (GIBCO BRL, NY, USA) and 0.3% antibiotics at 37°C in 5% CO₂. PC12 cells were treated with nerve growth factor (NGF, 50 ng/ml) from Calbiochem (Darmstadt,

Germany) to differentiate for 4 days. The cerebral cortex was dissected out of Sprague–Dawley rat embryo at embryonic day 17 and dissociated by gentle trituration. Cells were plated on PEI coated plates and after overnight incubation in DMEM supplemented with 10% FBS, the medium was changed to a serum-free medium as defined for neurons (neurobasal medium supplemented with B27 and penicillin–streptomycin–amphotericin B mixture (Gibco BRL)). Experiments were performed in 14- to 15-day cultures. HEK293 cells, NGF differentiated PC12 cells and rat cortical primary neurons were transiently transfected with 2 μ g of plasmid DNA and 3 μ l of Fugene 6 (Roche Molecular Biochemicals, Germany) in 1 ml of growing medium, according to the manufacturer's instruction.

Nuclear and cytosolic fractionation. At 48-h post-transfection, cells plated on six-well plates were washed with ice-cold PBS and then nuclear and cytoplasmic fractionation was performed, according to the manufacturer's protocol (Nuclear and Cytoplsmic Extraction Reagents; PIERCE, IL, USA).

Immunoprecipitation. A measure of 100 μ g of samples in 1 ml lysis buffer were rotated with Protein G Plus/Protein A agarose (Oncogene, Darmstadt, Germany) for 1 h at 4°C, and then centrifuged at 14 000 r.p.m. for 20 min. Supernatants were incubated with 1 μ g of primary antibody with rotation for 1 h at 4°C, treated with Protein G Plus/Protein A agarose for 1 h at 4°C, and centrifuged at 14 000 r.p.m. for 20 min. The pellet was boiled for 7 min in sample buffer (0.2 M Tris-HCl pH 6.8, 10% SDS, 25% glycerol and 0.01% bromophenol blue) and then centrifuged at 13 000 r.p.m. for 15 min. The supernatant was subjected to SDS polyacrylamide gel electrophoresis.

Western blotting. Protein was resolved in SDS polyacrylamide gel and electrophoresis was performed at $30-50 \ \mu g$ of protein/lane, and transferred onto a nitrocellulose membrane (Amersham Pharmacia, Buckinghamshire, UK). The protein blot was proved with appropriate antibodies and then detected with horseradish peroxidase-conjugated secondary antibody (Amersham Pharmacia). The immunoreactive bands were visualized using an ECL enhanced chemiluminescence system (ECL; Amersham Pharmacia, Buckinghamshire, UK).

Fluorescence resonance energy transfer. FRET measurements were observed using Zeiss LSM 510 confocal microscope mounted on Zeiss Axiovert 200 inverted microscope. FRET was measured employing a method developed for laser scanning confocal microscopy using an argon laser to excite EYFP or ECFP. 41-43 HEK293 cells on six-well plates were co-transfected with YFP vector, YFP-C57, or YFP-C57^{Y738G} and CFP-CP2 using Fugene 6 (Roche Molecular Biochemicals, Germany). At 24 h after transfection, the coverslips were mounted on Axiovert 200 inverted microscope (Zeiss, Jena Germany). Initial scan was obtained at low energy using the 458 nm line of the argon laser to record the ECFP signal. A second scan was performed with the 568 nm line, and cells of colocalization was noted. Cells was then photobleached with intense 514 nm light (laser power 100%) to destroy the acceptor molecules. The energy transfer was detected as an increase in donor fluorescence (ECFP) after complete photobleaching of the acceptor molecules (EYFP) at 514 nm. The amount of energy transfer was calculated as the percent increase in donor fluorescence after acceptor photobleaching. An increase of the ECFP signal within the photobleached area was used as a measure of the amount of FRET present.

Cell viability assay. LDH activities in the conditioned media were measured using a CytoTox 96 Non-Radioactive Cytotoxicity Assay kit (Promega, WI, USA), according to the manufacturer's instructions and absorbance was measured at 490 nm using a microplate reader (Molecular Devices, Sunnyvale, CA, USA).

Evaluation of apoptosis with Hoechst dye staining. Hoechst 33258 (Sigma-Aldrich, MO, USA) was used to assess chromatin condensation as published, previously.⁴⁴

Immunocytochemistry. Cells were fixed in 4% paraformaldehyde for 1 h at 4°C and then for 30 min at room temperature. After quenching the cells in 50 mM NH₄CI–PBS for 10 min, they were washed in PBS and permeabilized for 30 min at room temperature in PBS containing 0.1% Triton X-100 and 1 mg/ml bovine serum albumin (permeabilization buffer). The following steps were performed at room temperature in permeabilization buffer. Cells were incubated with appropriate primary antibodies for 1 h. After three washes, primary antibodies were revealed by incubating the cells for 1 h with FITC or Cy3 conjugated secondary antibodies

 random primer (Gibco BRL, NY, USA) and Superscript II reverse transcriptase (Gibco BRL, NY, USA). PCR was then carried out using a Gene Cycler (BioRad, Hercules, CA, USA), with a denaturation step of 94°C/5 min followed by 25 cycles of 95°C/1 min, 57°C/1 min and 72°C/2 min and a final extension step of 72°C/10 min. The primer pairs used to amplify rat GSK-3β were forward: 5′ GGA TCT GCC ATC

(LSM 510, Zeiss, Germany).

Luciferase activity assay. Human GSK-3 β promoter (nt -2090 to \sim nt + 1035) in the pCAT vector was a kind gift from Dr PC Shaw at the Chinese University of Hong Kong. The promoter region in the vector was excised at *KPn*I and *Bg1*II sites and inserted into the *KPn*I and *Bg1*II sites of PGL2 promoter plasmid (Promega, WI, USA). For co-transfection experiments, 4 μ g of human GSK-3 β promoter in the PGL2 vector and APLP2-ICDs (C57, C50, C57^{Y738G} or C50^{Y738G}) in the pEGFPN1 vector were added to the PC12 cells. Luciferase activity was measured using a Biocounter M1500 luminometer (Lumac, GE Groningen, Netherlands). Luciferase activities were normalized *versus* the obtained protein concentrations.

(Jackson ImmunoResearch, PA, USA). After three washes in permeabilization

buffer and one wash in PBS, the cells were mounted on microscope slide in

mounting medium (DAKO, Carpinteria, CA, USA), including DAPI (1 µM) for

mounting and nucleus staining. Cells were observed under a confocal microscopy

Reverse transcription polymerase chain reaction (RT-PCR)

analysis. Total RNA was isolated using the easy-BLUE™ RNA Extraction Kit

(iNtRON Biotechnology, Seoul, Korea) from cells. RNA was reverse transcribed with

GAG ACA TT3'; reverse: 5'CCA ACT GAT CCA CAC CAC TG3'.

Immunohistochemistry. AD and age-matched control brains were obtained from the autopsy service at University of Oklahoma Heath Sciences Center, USA and Korea Scientific Investigation Center. A neuropathological diagnosis of AD was confirmed according to the criteria of Braak.²⁵ Blocks of AD or control tissue were fixed in 10% neutral buffered formalin for 48 h. After fixation, the brain tissue was dehydrated and embedded in paraffin. Immunohistochemistry was performed using a Vectastain avidin–biotin complex (ABC) elite kit (Vector, Burlingame, CA, USA) as described previously.⁴⁵ Images were captured using an optical microscope (BX51TF, Olympus Optical Co., Japan) equipped with an exposure control unit (PM-CB20) and a camera (CPM-C35DX).

Statistical analysis. Results are presented as means \pm s.e.m. All statistical analysis was performed using SPSS for Windows 10.0.7 software (SPSS Inc., Chicago, USA). One-way ANOVA test was used to study the relationship between the different variables. p < 0.05 was considered to be significant.

Acknowledgements

We are grateful to Drs. Gopal Thinakaran and Vincenzo Casolaro at the University of Chicago and Johns Hopkins University for providing the APLP2 cDNA, antiserum and CP2 cDNA, respectively. We also thank Drs Joseph D Buxbaum at Mount Sinai School of Medicine and Pang-Chui Shaw for providing rat Fe65 cDNA and human glycogen synthase kinase 3β promoter, respectively. This work was supported by a National Creative Research Initiative Grant (2003–2005) from MOST, in part by the BK21 Human Life Sciences program and by a grant from Seoul National University Bundang Hospital Research Fund.

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Supplementary Information accompanies the paper on Cell Death and Differentiation website (http://www.nature.com/cdd)