

Stimulation of β -Amyloid Precursor Protein Trafficking by Insulin Reduces Intraneuronal β -Amyloid and Requires Mitogen-Activated Protein Kinase Signaling

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Alzheimer's Disease (AD) is characterized by cerebral accumulation of β -amyloid peptides ($A\beta$), which are proteolytically derived from β -amyloid precursor protein (β APP). β APP metabolism is highly regulated via various signal transduction systems, e.g., several serine/threonine kinases and phosphatases. Several growth factors known to act via receptor tyrosine kinases also have been demonstrated to regulate $s\beta$ APP secretion. Among these receptors, insulin and insulin-like growth factor-1 receptors are highly expressed in brain, especially in hippocampus and cortex. Emerging evidence indicates that insulin has important functions in brain regions involved in learning and memory. Here we present evidence that insulin significantly reduces intracellular accumulation of $A\beta$ and that it does so by accelerating β APP/ $A\beta$ trafficking from the trans-

Golgi network, a major cellular site for $A\beta$ generation, to the plasma membrane. Furthermore, insulin increases the extracellular level of $A\beta$ both by promoting its secretion and by inhibiting its degradation via insulin-degrading enzyme. The action of insulin on β APP metabolism is mediated via a receptor tyrosine kinase/mitogen-activated protein (MAP) kinase kinase pathway. The results suggest cell biological and signal transduction mechanisms by which insulin modulates β APP and $A\beta$ trafficking in neuronal cultures.

Key words: β -amyloid; β -amyloid precursor protein; insulin; MAPK; Alzheimer's disease; diabetes mellitus; intracellular trafficking; endoplasmic reticulum; trans-Golgi network; plasma membrane

Neuropathological hallmarks of Alzheimer's Disease (AD) include deposition of β -amyloid ($A\beta$) plaques, neurofibrillary tangles, and neuronal cell loss in vulnerable brain regions. Plaques contain an aggregated population of heterogeneous $A\beta$ peptides derived from β -amyloid precursor protein (β APP). Full-length β APP undergoes proteolytic β -secretase and γ -secretase activities to generate $A\beta$ 40 and $A\beta$ 42 peptides, the predominant $A\beta$ variants. In addition to these amyloid-generating activities, full-length β APP can undergo alternative processing by an enzymatic activity termed " α -secretase" that cleaves within the $A\beta$ region. This activity releases a soluble fragment ($s\beta$ APP α) extracellularly and precludes $A\beta$ formation. Several studies indicate that $A\beta$ is toxic to neurons. Accumulation of $A\beta$ peptides within the brain is believed to initiate the pathological cascade culminating

in clinical AD, a hypothesis supported by the development of early-onset familial AD (FAD) within pedigrees harboring autosomal dominant gene mutations in β APP that lead to the excessive generation of $A\beta$ (for review, see Selkoe, 1998). Cell biological studies have demonstrated that both $A\beta$ 40 and $A\beta$ 42 are produced intracellularly (Cook et al., 1997; Xu et al., 1997; Lee et al., 1998; Skovronsky et al., 1998; Greenfield et al., 1999). Moreover, recent evidence raises the possibility that intracellular $A\beta$ 42 may play a direct pathogenic role in AD neuropathology (for review, see Wilson et al., 1999; Gouras et al., 2000).

β APP metabolism is highly regulated via various signal transduction systems, e.g., various serine/threonine kinases and phosphatases (for review, see Mills and Reiner, 1999). Several growth factors known to act via receptors with intrinsic tyrosine kinase activity also have been demonstrated to regulate $s\beta$ APP secretion (Refolo et al., 1989; Schubert et al., 1989). Among these receptors, insulin and insulin-like growth factor-1 (IGF-1) receptors are highly expressed in brain, particularly in hippocampus and cortex (Werther et al., 1987). Recently, it has been demonstrated that insulin receptors are concentrated at the synaptic level and that they are a component of postsynaptic densities in cultured hippocampal neurons (Abbott et al., 1999). Moreover, insulin can recruit GABA_A receptors to the postsynaptic domain (Wan et al., 1997), suggesting a role for this hormone in synaptic plasticity. Emerging evidence indicates that insulin has important functions in brain regions involved in learning and memory (Wickelgren, 1998; Zhao et al., 1999). Recent findings demonstrated insulin receptor upregulation and reduced insulin receptor-mediated ty-

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rosine kinase activity in AD brains (Frolich et al., 1998, 1999). Insulin and IGF-1 have been shown to regulate tau phosphorylation via GSK-3 β , suggesting that neurofibrillary tangle formation in AD may be downstream of insulin signaling (Hong and Lee, 1997; Lesort et al., 1999; Lesort and Johnson, 2000). Recently, investigators have begun to address the potential role of insulin in β APP metabolism. It was shown that insulin elevates s β APP secretion in SH-SY5Y cells (Solano et al., 2000). Moreover, it was demonstrated that insulin-degrading enzyme (IDE) degrades extracellular A β in microglial and neuronal cultures and that insulin can prevent this degradation, thereby impairing the clearance of extracellular A β (Qiu et al., 1998; Vekrellis et al., 2000).

Here we report that insulin decreases intracellular levels and increases extracellular levels of both A β 40 and A β 42. These effects of insulin are associated with accelerated β APP/A β trafficking from the Golgi/trans-Golgi network (TGN) to the plasma membrane and are prevented by inhibitors of tyrosine kinase and MAP kinase kinase.

MATERIALS AND METHODS

Cell cultures. N2a neuroblastoma cells transfected with human β APP695 were maintained in medium containing 50% DMEM and 50% Opti-MEM, supplemented with 5% FBS, 200 μ g/ml G418, and antibiotics (Life Technologies, Gaithersburg, MD). Primary neuronal cultures were derived from the cerebral cortices of day 17 (E17) embryos obtained from timed pregnant Sprague Dawley rats as described previously (Gouras et al., 1998). Neurons were used for experiments after 4–5 d in culture.

Pulse-chase experiments and insulin treatment. N2a cells (80% confluent in 10 cm dish) or primary neuronal cultures (10^7 cells/10 cm dish) were labeled for 20 min with 750 μ Ci/ml [35 S]methionine (NEN-DuPont, Boston, MA) in methionine-free DMEM supplemented with L-glutamine (Life Technologies). Cells were chased at 37°C in serum-free DMEM (Life Technologies) or in serum- and glucose-free DMEM supplemented with 50 mM 2-deoxy-D-glucose in the absence or presence of insulin (Sigma, St. Louis, MO). In some experiments 0.3–30 μ M glucagon, 0.1–1 μ M IGF-1, 2.5–25 ng/ml EGF, 25–250 ng/ml NGF, 1–10 ng/ml PDGF, 10–100 ng/ml aFGF, or 10–100 ng/ml bFGF (BD Transduction Laboratories, Franklin Lakes, NJ; gifts of Dr. J. Schlessinger, New York University Medical Center) was added during the chase. For continuous metabolic labeling the cells were incubated for 4 hr with 750 μ Ci/ml [35 S]methionine in methionine-free DMEM supplemented with L-glutamine in the absence or presence of 1 μ M insulin. For steady-state experiments the cells were treated for 4–16 hr with 0.3–1 μ M insulin in the presence or absence of 10–25 μ M tyrphostin-25, 500 nM wortmannin, 25 μ M U73122, or 10 μ M PD98059; the samples were analyzed by Western blot.

A β degradation assay. N2a cells expressing wild-type human β APP695 were labeled continuously for 4 hr with 750 μ Ci/ml [35 S]methionine, and medium was collected to serve as the source for labeled A β and s β APP α proteins. Nonlabeled serum-free conditioned medium (CM) was collected after the incubation of primary neuronal cultures for 16 hr. This CM was mixed with the labeled A β /s β APP α -containing medium and incubated further at 37°C for 16 hr in the absence or presence of 1 μ M insulin and/or 1 mM 1,10-phenanthroline. In some experiments IDE was eliminated from the cold CM by immunodepletion, using an anti-IDE monoclonal antibody (9B12) (Shii and Roth, 1986). The amounts of labeled A β or s β APP α remaining in each sample after incubation were determined by immunoprecipitation with antibody 4G8 (for A β) or 6E10 (for s β APP α) (Senetek PLC, St. Louis, MO), followed by 10–20% Tris/tricine (A β) or 4–12% Tris/glycine (s β APP α) SDS-PAGE.

Immunoprecipitation and Western analysis. Media were collected, centrifuged briefly to remove cell debris, and sequentially immunoprecipitated first with 4G8 for A β and then with 6E10 for s β APP α secreted from N2a cells or 22C11 (Roche Pharmaceuticals, Nutley, NJ) for s β APP secreted from rat neurons. s β APP β secreted from N2a cells was determined by immunoprecipitation with 22C11 antibody after immunodepletion of s β APP α with 6E10 antibody. Immunoprecipitation of IDE was performed with 28H1 monoclonal antibody (Shii and Roth, 1986). Cells

($1\text{--}2 \times 10^7$) were scraped from plates in ice-cold PBS with a rubber policeman. After centrifugation the pellets were treated with 250 μ l of 3% SDS in PBS containing 10 μ l/ml of β -mercaptoethanol and subjected to vortexing and heating at 95°C for 10 min, followed by sonication and centrifugation at 100,000 \times g for 10 min. The resultant supernatants were diluted 1:4; adjusted to a final concentration of 2% Triton X-100 and (in mM) 190 NaCl, 20 Tris-HCl, pH 8.8, and 2 EDTA; and subjected to immunoprecipitation and SDS-PAGE analysis that used 10–20% Tris/tricine gels (for A β) or 4–12% Tris/glycine gels (for s β APP α detection). For Western blot the samples were transferred to polyvinylidene difluoride (PVDF) membranes (Millipore, Bedford, MA or Bio-Rad, Hercules, CA), and the membranes were boiled in PBS for 5 min. A β and s β APP α were detected by using 6E10 monoclonal antibodies. For Western blot analysis of full-length β APP and presenilin-1 (PS-1) N-terminal fragment, the samples were analyzed on 12% SDS-PAGE gels, transferred to PVDF membrane (Millipore), and immunoblotted with antibody 369 (Buxbaum et al., 1990) and antibody Ab14 (Seeger et al., 1997), respectively. Anti-insulin receptor antibody was from Neomarkers (Fremont, CA). Intracellular IDE was detected by immunofluorescence as described previously (Greenfield et al., 1999) with 9B12 and 28H1 monoclonal antibodies (Shii and Roth, 1986).

Immunoprecipitation–mass spectrometry analysis. Media or cell lysates were immunoprecipitated with 4G8 antibody and protein A/G-agarose beads. The molecular masses and concentrations of immunoprecipitated A β species were measured by matrix-assisted laser desorption/ionization time-of-flight mass spectrometry analysis, as described previously (Xu et al., 1998). For analysis, A β 12–28 and insulin internal standards were added to the samples.

Sucrose gradient fractionation. Fractionation by sucrose gradient was performed as described previously (Greenfield et al., 1999). After 16 hr of incubation in the absence or presence of 1 μ M insulin, N2a cells were homogenized by using a stainless steel ball bearing homogenizer in 5 vol of 0.25 M sucrose, 10 mM Tris-HCl, pH 7.4, 1 mM MgAc $_2$, and a protease inhibitor mixture. The homogenate was loaded on top of a step gradient comprised of 1.5 ml of 2 M sucrose, 4 ml of 1.3 M sucrose, 3.0 ml of 1.16 M sucrose, and 2.0 ml of 0.8 M sucrose. All sucrose solutions contained 10 mM Tris-HCl, pH 7.4, and 1 mM MgAc $_2$. The gradients were centrifuged for 2.5 hr at 100,000 \times g in a Beckman SW41Ti rotor. Fractions were collected and assayed for total protein with BCA assay. A β and β APP were assayed as described above. Proteins from each fraction were analyzed by Western blot with antibodies against calnexin, γ -adaptin (BD Transduction Laboratories), or ARF3 (Affinity BioReagents, Neshanic Station, NJ) to identify the fractions containing, respectively, endoplasmic reticulum (ER), TGN, and cytosol/post-TGN vesicles. To determine the fractions enriched in plasma membrane proteins, we fractionated N2a cells after biotinylation of surface proteins and detected biotinylated β APP as described below.

Cell surface biotinylation. Biotinylation was performed on confluent monolayer N2a cells overexpressing β APP695 by using sulfo-NHS-LC-biotin [sulfo-succinimidyl-6-(biotinamido)-hexanoate; Pierce, Rockford, IL]. The reagent was dissolved in PBS with calcium and magnesium, pH 7.2, at 0.5 mg/ml and added twice to the cultures for 20 min at 4°C. After thorough washing, the cells were lysed with 3% SDS as described above. β APP was immunoprecipitated by using 369 antiserum and was analyzed by Western blot. Biotinylated β APP was detected by using HRP-conjugated streptavidin and reaction with a chemiluminescent substrate (NEN-DuPont).

Endoglycosidase-H Digestion. To evaluate the effect of insulin on ER-to-Golgi trafficking of β APP, we pulse-labeled N2a cells for 5 min with 750 μ Ci/ml [35 S]methionine (NEN-DuPont) in methionine-free DMEM and chased the cells for 5–45 min in the absence or presence of 1 μ M insulin as described above. [35 S]-labeled β APP was immunoprecipitated from the SDS-soluble lysates by 369 antiserum. The immune complexes were boiled for 5 min in a buffer containing 50 mM Tris-HCl, pH 7.6, 0.5% SDS, and 1% β -mercaptoethanol to dissociate β APP from the antibody; sodium citrate was added to a final concentration of 0.05 M. Then the samples were incubated in the presence or absence of 50 U of endoglycosidase-H (New England Biolabs, Beverly, MA) at 37°C for 16 hr. β APP isoforms were separated on a 6% Tris/glycine gel, transferred to PVDF membrane, and detected by autoradiography.

Quantification and densitometry. Gels were exposed to x-ray film, and the films were scanned with an Agfa Arcus II scanner. Band intensities were analyzed and quantified by using NIH Image Quant software, version 1.52. Statistical analysis was performed with ANOVA, followed by a *post hoc* test.

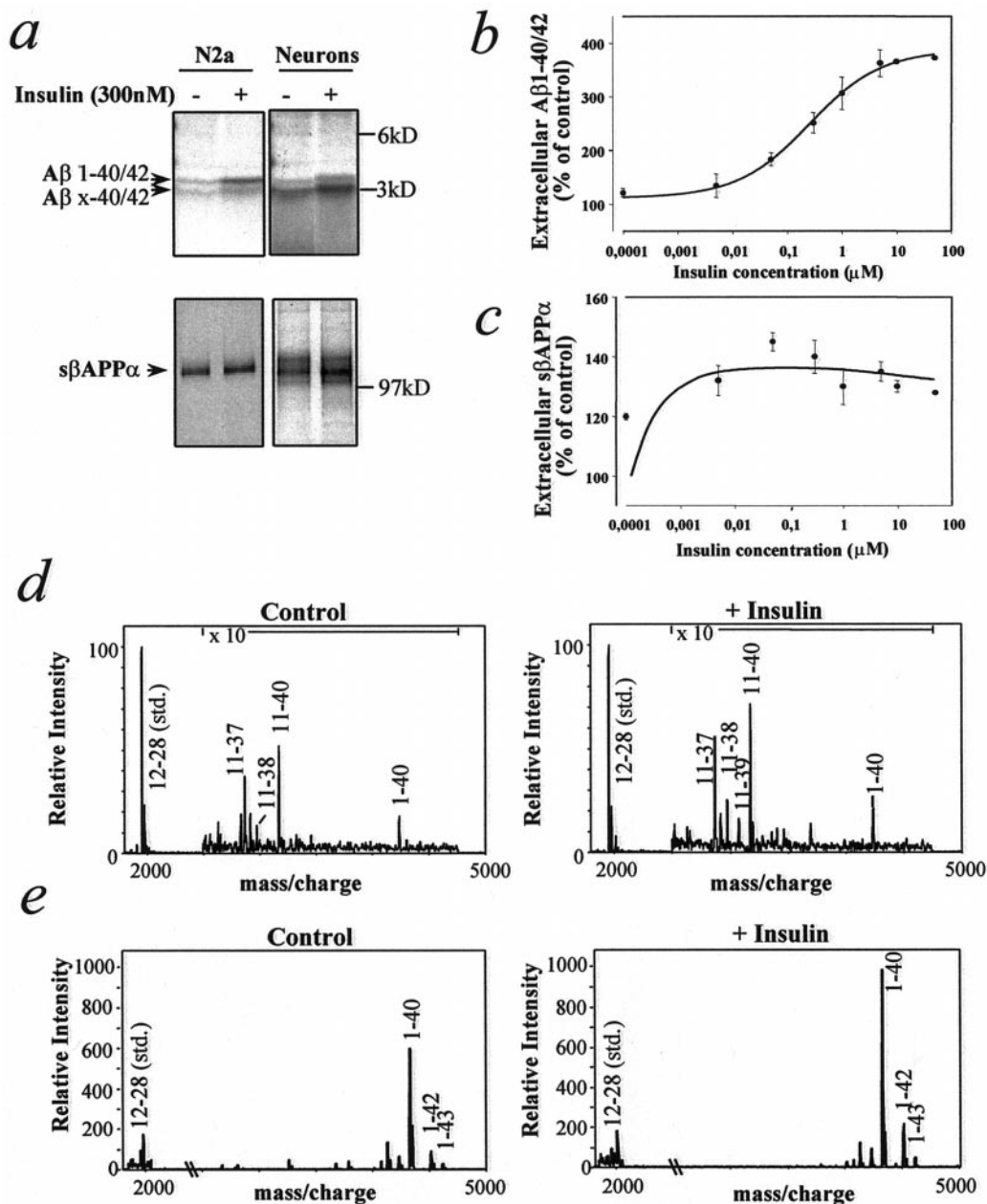


Figure 1. Effect of insulin on the extracellular levels of A β and sAPP α in murine neuroblastoma cells (N2a) and primary rat cortical neuronal cultures. Cells were pulse-labeled for 20 min with [35 S]methionine and incubated in serum-free medium in the absence or presence of various concentrations of insulin for 4 hr. *a*, Representative autoradiographic analysis of extracellular A β (top) and sAPP α (bottom) after incubation with or without 300 nM insulin. *b*, Extracellular levels of A β from N2a cells as a function of insulin concentration. *c*, Extracellular levels of sAPP α from N2a cells as a function of insulin concentration. For *b* and *c* the data represent means \pm SD; $n = 3$. *d*, *e*, IP-MS analysis of extracellular A β from primary neurons (*d*) and N2a cells (*e*) after incubation in the absence or presence of 1 μ M insulin for 4 hr.

RESULTS

Insulin increases extracellular levels of A β 40 and A β 42

To examine the effect of insulin on β APP metabolism, we pulse-labeled neuroblastoma (N2a) cells overexpressing human β APP695 or primary cultures of rat cortical neurons with [35 S]methionine for 20 min and chased the cells in the absence or presence of insulin for 4 hr. Cells that were treated with insulin showed a three- to fourfold increase in extracellular levels of both 4 kDa A β 1–40/42 and 3 kDa N-terminally truncated A β x-40/42 peptides, mainly composed of A β 11–40/42 (Gouras et al., 1998)

(Fig. 1*a*, top). The effect of insulin on extracellular levels of A β 1–40/42 was concentration-dependent, with a minimal effective concentration of \sim 50 nM and a half-maximal effect at \sim 300 nM (Fig. 1*b*). Insulin caused a \sim 1.4-fold increase in sAPP α extracellular levels (Fig. 1*a*, bottom, *c*), with a small increase occurring at the lowest concentration (0.2 nM) that was tested. Secretion of sAPP β was not altered by insulin treatment (data not shown). Immunoprecipitation–mass spectrometry (IP-MS) analysis revealed increases in extracellular A β 1–40 in both primary neurons and N2a cells (Fig. 1*d,e*) and increases of A β 1–42

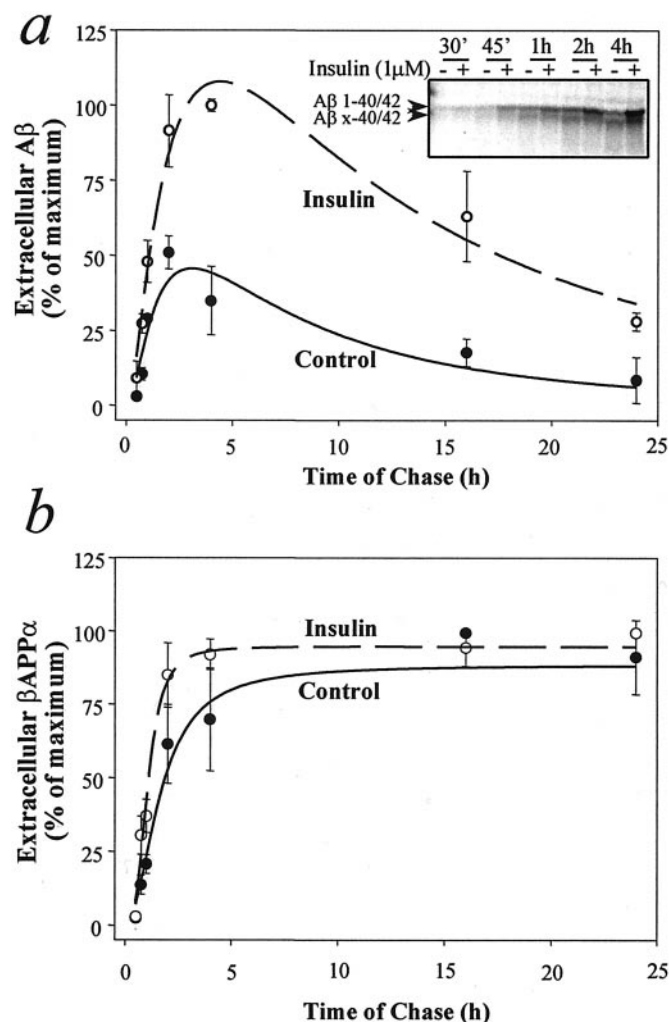


Figure 2. Time course of A β (*a*) and sAPP α (*b*) extracellular levels from N2a cells in the absence or presence of insulin. Cells were pulse-labeled with [35 S]methionine for 20 min and incubated in serum-free medium in the absence or presence of 1 μ M insulin for different intervals. Data represent means \pm SD; $n = 3$. Inset in *a* shows a representative autoradiographic analysis of extracellular A β after periods of chase from 30 min to 4 hr.

in N2a cells (Fig. 1*e*). The effect of insulin on the extracellular levels of A β 1–40/42 and A β x-40/42 in cultures of N2a cells was evident at all incubation times from 30 min to 24 hr (Fig. 2*a*). A β secretion in the absence or presence of insulin peaked at 4 hr and decreased thereafter, indicating that A β is susceptible to degradation by proteases released from the cells. Extracellular sAPP α was increased slightly by insulin at early, but not at late, incubation times (Fig. 2*b*) and reached a plateau level because of its resistance to the degrading activity of proteases. β APP levels were not altered by insulin during pulse–chase experiments. Glucose-containing medium was required for the action of insulin on A β . Glucose concentrations higher than the standard growing concentration did not alter basal or insulin-stimulated A β levels (data not shown).

Insulin increases extracellular A β both by reducing IDE-mediated A β degradation and by stimulating A β secretion

To investigate whether the increased level of extracellular A β was attributable at least in part to the inhibition of extracellular

degradation, we examined the effect of IDE on the levels of extracellular A β . IDE was found intracellularly by immunocytochemistry (data not shown), and pulse–chase experiments demonstrated a substantial amount of IDE in the medium of primary neuronal cultures after a 16 hr chase (Fig. 3*a*). It was reported previously that IDE can cause the rapid degradation of A β 1–40/42, but not A β x-40/42, and that the enzymatic activity of IDE against A β 1–40/42 could be prevented by insulin, phenantroline, or immunodepletion with an IDE-specific antibody (Qiu et al., 1998; Vekrellis et al., 2000). We have confirmed those earlier studies in the present investigation (Fig. 3*b,c*). Insulin and phenantroline each acted as a potent inhibitor of the degradation of A β by IDE in our *in vitro* setting (Fig. 3*c*). However, insulin, phenantroline, or immunodepletion of IDE from our system did not rescue A β completely from degradation, suggesting that other proteases may be involved in A β breakdown. To investigate whether insulin can stimulate the secretion of A β in addition to preventing its degradation, we treated N2a cells with or without insulin in the absence or presence of the IDE inhibitor 1,10-phenantroline for various time intervals (Fig. 3*d*). Phenantroline alone increased the extracellular levels of A β . However, the effect of phenantroline was less than that of insulin alone and was significant only at later intervals (4–16 hr). When phenantroline was present during the incubation period to inactivate IDE, insulin further enhanced the extracellular levels of A β . These results indicate that insulin can stimulate A β secretion in addition to inhibiting IDE-mediated A β degradation. In addition, the secretion of the 3 kDa N-terminally truncated A β x-40/42 peptides, which are resistant to degradation by IDE (Qiu et al., 1998; Vekrellis et al., 2000) (Fig. 3*b*), was stimulated by insulin (see Figs. 1*a*, 2*a*).

Insulin reduces intracellular A β 40 and A β 42

To examine further the role of insulin in β APP metabolism, we examined the effect of insulin on intracellular A β , which recently has been hypothesized to be important in AD (Wild-Bode et al., 1997; Skovronsky et al., 1998; Chui et al., 1999; Wilson et al., 1999; Gouras et al., 2000; Mochizuki et al., 2000). N2a cells were treated with 1 μ M insulin for 4 or 16 hr; A β was extracted with SDS and analyzed by immunoprecipitation and SDS-PAGE. Intracellular A β was reduced by 20% after 4 hr and by 45% after 16 hr of insulin treatment (Fig. 4*a,b*). Immunoprecipitation–mass spectrometry analysis revealed that intracellular A β 40 and A β 42 both were reduced after insulin treatment (Fig. 4*c*). No change was detected in the levels of full-length β APP or PS-1 N-terminal fragment after insulin treatment for 4 hr (data not shown) or 16 hr (Fig. 4*d,e*).

Insulin reduces A β in the Golgi/TGN by accelerating β APP/A β transport to the plasma membrane

β APP resides predominantly in the Golgi/TGN, which is also the main site of A β generation (Cook et al., 1997; Hartmann et al., 1997; Xu et al., 1997; Lee et al., 1998; Skovronsky et al., 1998; Greenfield et al., 1999). Insulin is known to stimulate protein transport from TGN or post-TGN vesicles to the plasma membrane (Pessin et al., 1999). To define the subcellular compartments in which insulin exerts its action on β APP and A β trafficking, we performed subcellular fractionation by using a well characterized sucrose gradient procedure (Greenfield et al., 1999) after 16 hr of incubation in the absence or presence of insulin. Insulin reduced A β in membrane fractions collected from

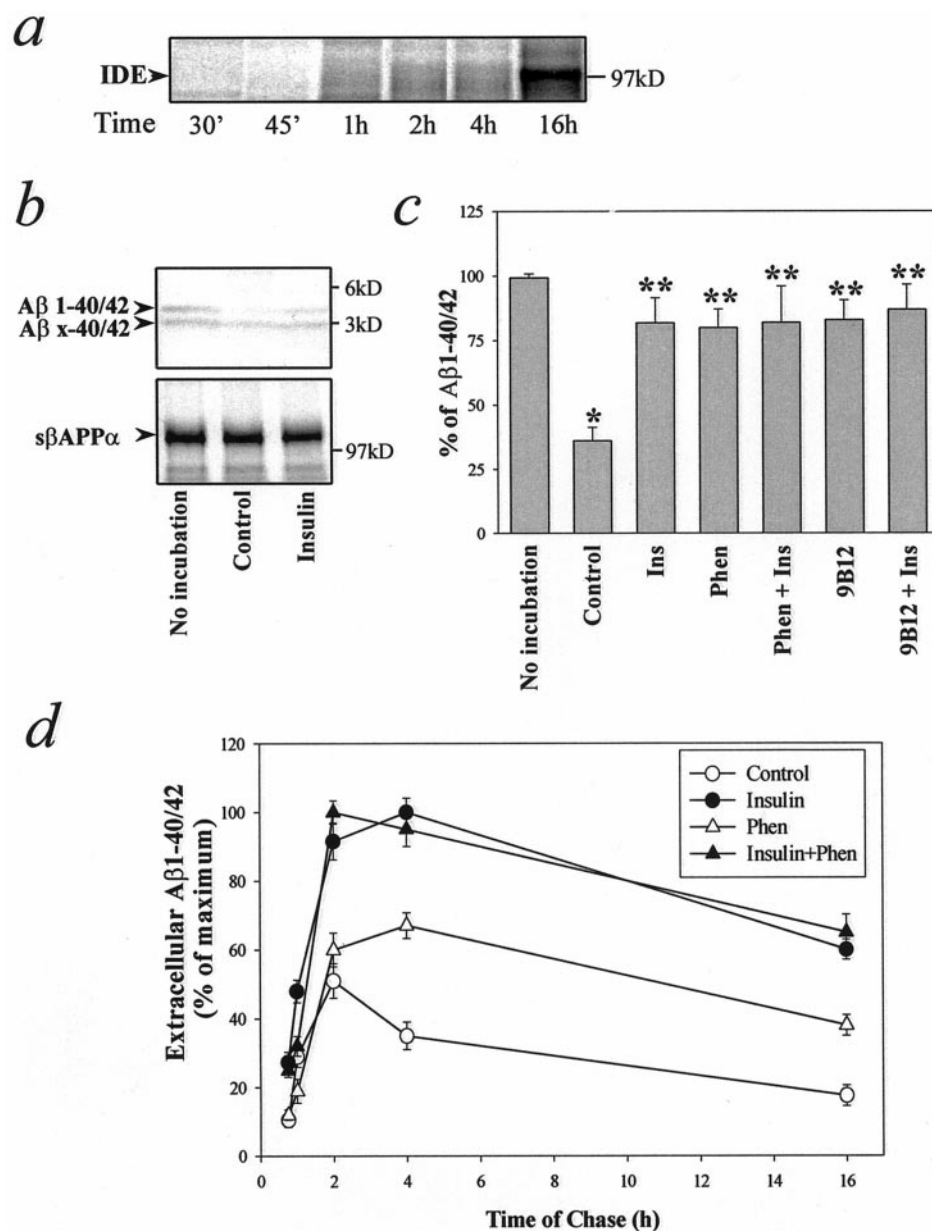


Figure 3. Insulin inhibits A β degradation via IDE and stimulates the secretion of A β . *a*, IDE released in the medium from cultured neurons as a function of time. Primary cortical neurons were pulse-labeled for 20 min and chased for different intervals up to 16 hr. IDE was immunoprecipitated with 28H1 monoclonal anti-IDE antibody and analyzed on SDS-PAGE. *b*, *c*, N2a cells were incubated for 4 hr with [35 S]methionine to produce labeled A β and sAPP α . Serum-free conditioned medium was collected from cultured primary neurons, which had been incubated for 16 hr at 37°C. Then this medium was mixed with the medium containing labeled A β or sAPP α and incubated for a further 16 hr in the absence or presence of the indicated substances (see Materials and Methods). *b*, Insulin inhibits A β degradation. Shown is an autoradiographic analysis of labeled A β (top) and sAPP α (bottom). Media were collected before or after *in vitro* incubation in the absence or presence of 1 μ M insulin, immunoprecipitated with anti-A β (4G8) or anti-sAPP (22C11) antibodies, and analyzed on SDS-PAGE. Insulin caused a marked inhibition of A β 1-40/42 degradation. *c*, IDE-mediated A β degradation was inhibited by 1 μ M insulin (*Ins*), 1 mM 1,10-phenanthroline (*Phen*), or immunodepletion of IDE with a monoclonal anti-IDE antibody (9B12; see Materials and Methods). Data represent means \pm SD; $n = 3$. * $p < 0.01$ with respect to the sample with no incubation; ** $p < 0.01$ with respect to control sample. *d*, Insulin stimulates A β secretion. N2a cells were pulse-labeled for 20 min and chased for various times in the absence or presence of 1 μ M insulin, 1 mM 1,10-phenanthroline (*Phen*), or a combination of the two compounds. Data represent means \pm SD; $n = 3$.

the interfaces corresponding to secretory vesicles and Golgi/TGN, but not from those corresponding to heavy membranes such as the ER, plasma membranes, and lysosomes (Fig. 5*a,b*). Moreover, after a 2 hr preincubation of cells at 20°C to accumulate newly synthesized [35 S]-labeled β APP in the TGN, insulin-dependent secretion of A β and sAPP α was observed within 7.5–15 min of the addition of the hormone (data not shown), supporting the idea that insulin stimulates A β and sAPP α trafficking from the TGN. After insulin treatment, full-length β APP also is reduced in vesicle and Golgi-enriched fractions, whereas it is increased in the heavy membrane fractions (Fig. 5*c,d*). The total amount of β APP was not changed overall by insulin treatment (see Fig. 4*d*). Calnexin (ER), γ -adaptin (TGN), ARF3 (post-TGN vesicles, cytosol), and surface biotinylated β APP (plasma membranes) were assayed to identify fractions that are enriched in these organelles (Fig. 5*e*). Our data indicate that insulin reduces intracellular A β and stimulates its secretion by increasing the β APP/A β egress from the Golgi/TGN and from

post-TGN vesicles. To determine whether the effect of insulin on trafficking was reflected in the number of β APP molecules on the plasma membrane, we treated N2a cells for 4 hr with insulin and then labeled them with biotin. Insulin treatment resulted in an almost twofold increase in β APP molecules at the cell surface (Fig. 5*f*).

Insulin does not affect ER-to-Golgi transport

It has been shown that insulin stimulates the export of leptin from the ER in rat adipocytes (Barr et al., 1997). However, an effect of insulin on ER-to-Golgi transport has not been reported for any other protein. To investigate whether insulin could affect β APP trafficking between ER and Golgi, we pulse-labeled N2a cells for 5 min and chased them for up to 45 min in the absence or presence of insulin. Immunoprecipitated β APP molecules were subjected to endoglycosidase-H (Endo-H) digestion, and the rate of appearance of Endo-H-resistant, N-linked oligosaccharide-modified isoforms of β APP was evaluated. Newly synthesized

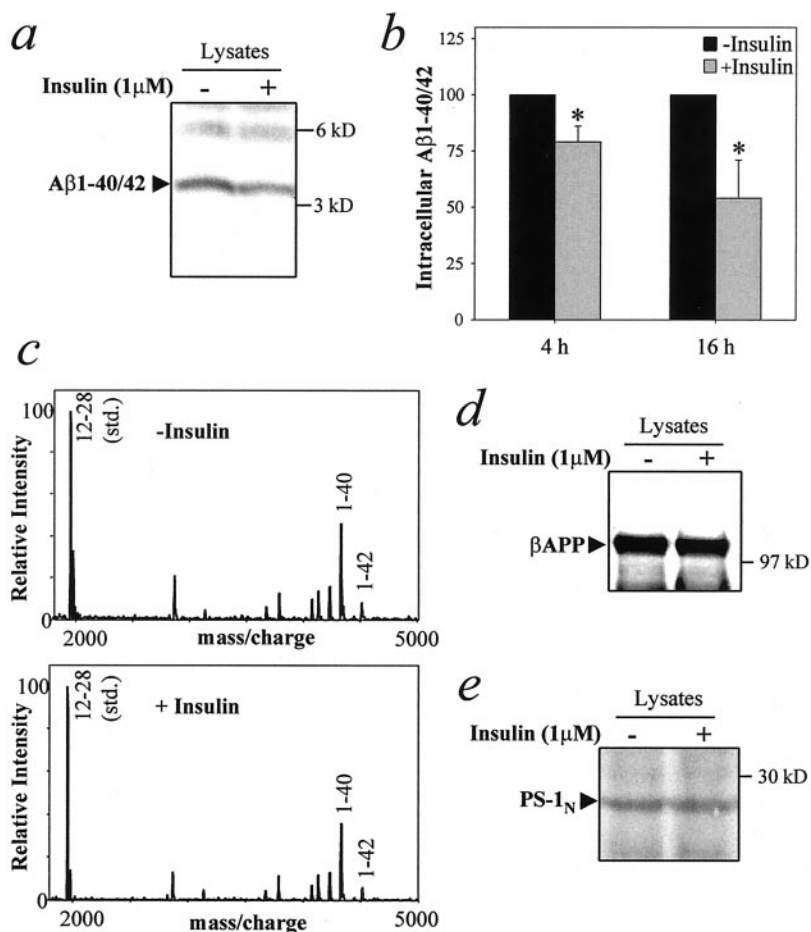


Figure 4. Insulin reduces intracellular levels of A β in N2a cells. Cells were treated for 4 or 16 hr with or without 1 μ M insulin and lysed in SDS. *a*, *b*, Intracellular A β was detected by immunoprecipitation with 4G8, followed by SDS-PAGE and Western blotting, using 6E10 monoclonal antibody, which recognizes only A β 1–40/42. *a*, Representative autoradiographic analysis of intracellular A β after 16 hr of treatment in the absence or presence of 1 μ M insulin. *b*, Quantitative analysis of intracellular A β after treatment with insulin for 4 or 16 hr. Data represent means \pm SD; $n = 5$. * $p < 0.05$ versus control. *c*, IP-MS analysis of intracellular A β 40/42 levels after 4 hr of treatment with 1 μ M insulin. *d*, *e*, Western blot analysis for full-length β APP (*d*) and the PS-1 N-terminal fragment (*e*) after 16 hr of treatment with 1 μ M insulin.

β APP (~105 kDa) is sensitive to digestion by Endo-H (Fig. 6; $t = 0$ min). After 5–10 min of chase, a ~115 kDa β APP form appeared that was resistant to Endo-H digestion. Clearly, insulin treatment did not alter β APP maturation significantly or the pattern of Endo-H resistance, an indicator of ER-to-Golgi trafficking, at any time of chase up to 45 min.

Insulin regulates β APP processing via a receptor tyrosine kinase

Insulin receptors are present in both N2a and primary neuronal cultures (Fig. 7*a*). To determine whether the action of insulin involves a tyrosine kinase, we inhibited intrinsic tyrosine kinase activity with tyrphostin-25, a nonselective inhibitor of receptor and nonreceptor tyrosine kinases, and measured its effect on insulin-stimulated A β secretion in N2a cells. Tyrphostin-25 (25 μ M) abolished the effect of insulin on A β secretion, which was accompanied by a small reduction in basal secretion (Fig. 7*b*). At a concentration of 10 μ M, tyrphostin-25 partially inhibited the effect of insulin on A β levels without any effect on basal secretion (data not shown). These results indicate that tyrosine kinase activity is essential for the effect of insulin on A β trafficking. In addition, the effect of insulin on A β was mimicked by IGF-1, but not by glucagon, EGF, NGF, PDGF, aFGF, or bFGF (data not shown), indicating the specificity of the insulin effect. These various results indicate that the effect of insulin is mediated via interaction with the insulin/IGF-1 receptor.

Insulin regulates β APP processing via MEK/MAP kinase cascade

To investigate the insulin pathway downstream of the insulin receptor, we studied the effect of selective inhibitors of three known insulin-activated signal transduction cascades. PD98059, a highly selective inhibitor of MAP kinase kinase activation (Alessi et al., 1995; Dudley et al., 1995), abolished the effect of insulin on A β secretion (Fig. 7*c*) and intracellular A β (data not shown) without altering basal levels of these parameters. In contrast, wortmannin, an inhibitor of PI-3 kinase, and U73122, an inhibitor of phospholipase C, caused a nonselective inhibition of both basal- and insulin-stimulated secretion of A β after incubation for 4 hr (Fig. 7*c*).

DISCUSSION

The present observations, that intracellular A β decreases whereas extracellular A β increases in response to insulin, could be explained by a mechanism involving insulin-stimulated intracellular trafficking of β APP and A β . This proposal is supported by substantial evidence showing that insulin selectively stimulates protein transport through the secretory pathway (Bogan and Lodish, 1999). In fact, we report here that insulin (1) accelerates β APP/A β trafficking from the TGN, the main site of A β generation, to the plasma membrane; (2) increases the number of β APP molecules on the plasma membrane; (3) increases the extracellular levels of A β even when IDE is inhibited by phenantroline; and (4) increases the

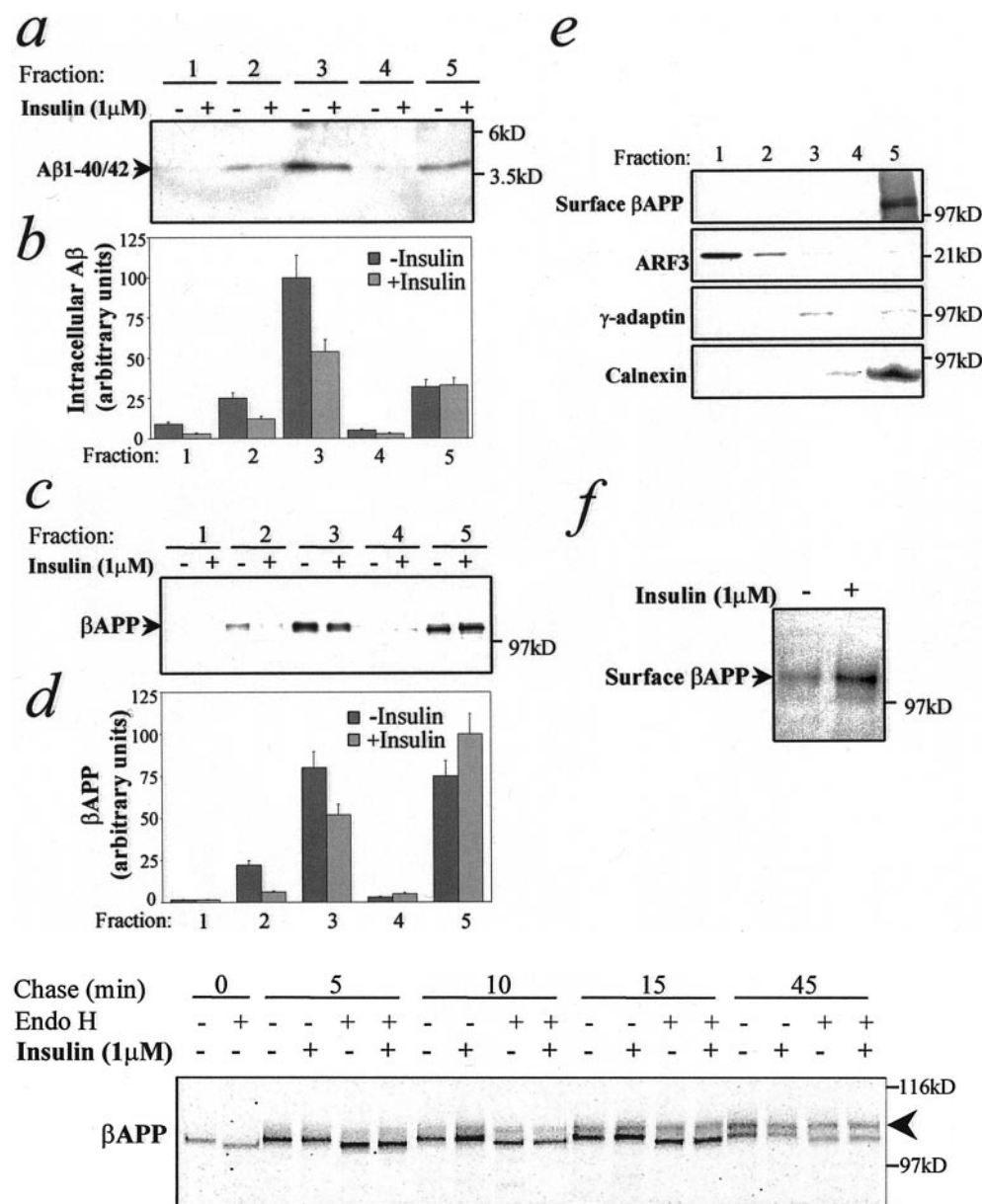


Figure 5. Insulin influences A β and β APP trafficking in N2a cells. Cells were treated for 16 hr in the absence or presence of 1 μ M insulin, homogenized, and fractionated on an equilibrium flotation sucrose gradient (see Materials and Methods). *a*, Representative autoradiographic analysis and quantitative analysis (*b*) of A β subcellular distribution after insulin treatment. *c*, Representative autoradiographic analysis and quantitative analysis (*d*) of intracellular β APP subcellular distribution after insulin treatment. *e*, Markers for subcellular compartments. Proteins from each fraction were precipitated by trichloroacetic acid and analyzed by Western blot, using the antibodies anti-calnexin (ER), anti- γ -adaptin (TGN), or anti-ARF3 (post-TGN vesicles, cytosol). Surface β APP (plasma membrane) was determined as described (see Materials and Methods). *Fraction 1* = 0.25 M sucrose solution (loading, cytosol). *Fractions 2–5* correspond, respectively, to interfaces between 0.25/0.8 M (post-TGN vesicles), 0.8/1.16 M (Golgi/TGN), 1.16/1.3 M, and 1.3 M/2 M (heavy membranes such as ER, plasma membranes) sucrose solutions. *f*, N2a cells were treated for 4 hr in the absence or presence of 1 μ M insulin. Surface proteins were labeled with biotin. Biotinylated β APP was analyzed by immunoprecipitation with 369 antibody and Western blot with HRP-conjugated streptavidin.

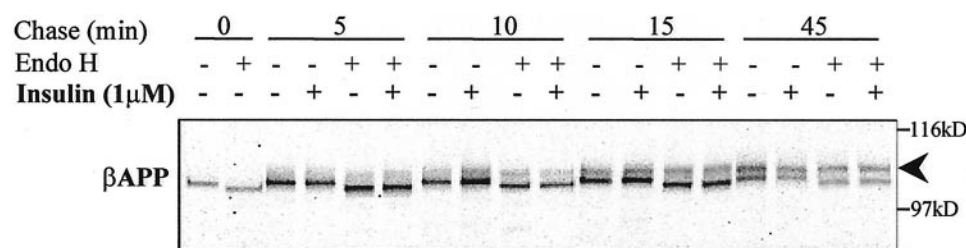


Figure 6. Lack of effect of insulin on β APP trafficking from ER to Golgi. N2a cells were pulse-labeled for 5 min with [35 S]methionine and chased in serum-free medium in the absence or presence of 1 μ M insulin for 5–45 min. β APP was immunoprecipitated by using 369 antibody; one-half of the sample was digested by endoglycosidase-H (*Endo H*). The arrowhead indicates the endoglycosidase-H-resistant β APP species.

secretion of 3 kDa A β _{x-40/42} peptides, mainly composed of A β _{11–40/42} (Gouras et al., 1998), which are resistant to IDE degradation (Qiu et al., 1998; Vekrellis et al., 2000). β APP levels and s β APP β secretion were not affected by insulin, suggesting that insulin may not regulate the β -cleavage of β APP but only β APP/A β trafficking. Elucidation of the potential contribution of the endosomal compartments to the effect of insulin on A β secretion awaits further investigation.

In agreement with previous studies (Qiu et al., 1998; Vekrellis et al., 2000), we have found in using both neuronal cell lines and primary neuronal cultures that IDE is a protease involved in A β degradation and that insulin inhibits A β degradation by competing for IDE. It was reported previously that IDE can be secreted by microglial cell cultures (Qiu et al., 1998), whereas it is cell-associated in primary neurons (Vekrellis et al., 2000). Although a significant amount of IDE is detectable in the medium of primary neurons after 16 hr, this is likely an experimental artifact of the cell culture setting, because IDE

does not have the signal peptide required for targeting into the secretory pathway and therefore should not be secreted. The slow kinetics of IDE release further support this view. Evidence was presented recently that a neutral endopeptidase, but not IDE, was involved in A β degradation when radiolabeled A β was injected into rat brain (Iwata et al., 2000). Thus, IDE-mediated A β degradation may be of less physiological relevance *in vivo*. In addition, proteases other than IDE also may take part in A β degradation in neuronal cultures, as suggested by the incomplete inhibition of A β degradation by insulin, phenantroline, or immunodepletion of IDE in our *in vitro* system (see Fig. 3c). This possibility is supported further by the observation that A β was still degraded when neuronal cultures were treated with phenantroline to inactivate IDE (see Fig. 3d).

We have performed a number of experiments on the signal transduction pathway by which insulin might stimulate the intracellular trafficking of β APP and A β . The effect of insulin was

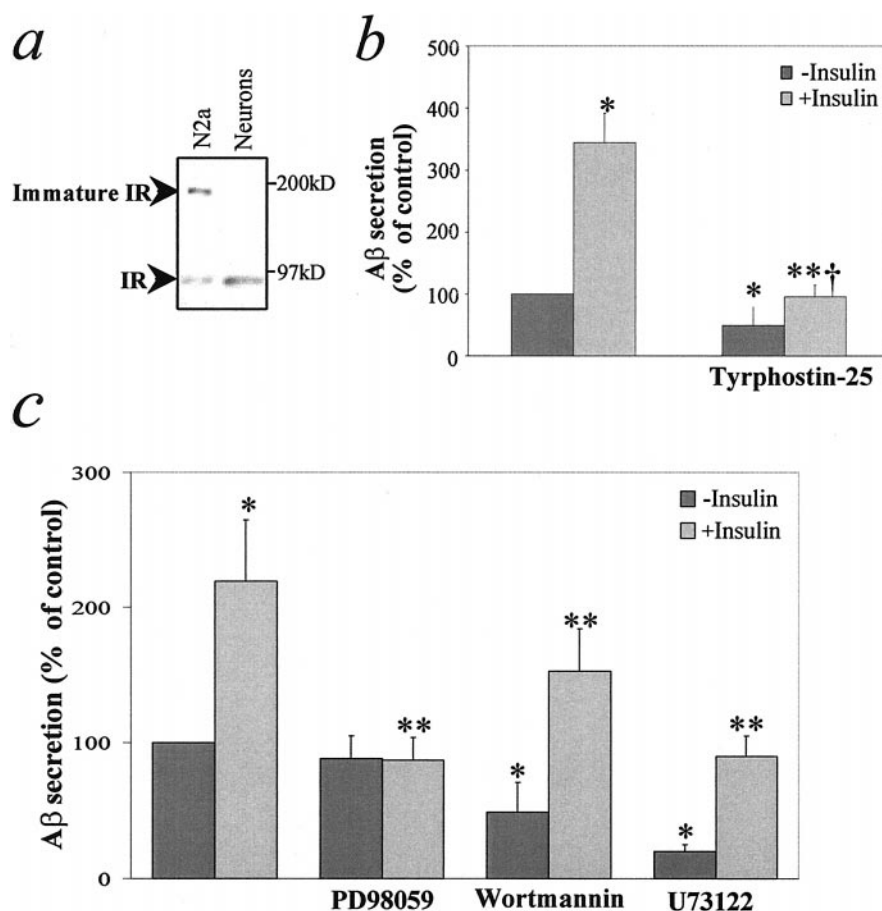


Figure 7. The effect of insulin requires tyrosine kinase activity and is mediated via the MEK/MAP kinase cascade. *a*, Western blot analysis for insulin receptor (IR) in N2a cells and primary neurons. *b*, N2a cells were incubated for 16 hr in serum-free medium in the absence or presence of 1 μ M insulin and/or 25 μ M tyrphostin-25. Data represent means \pm SD; $n = 3$. * $p < 0.05$ with respect to no addition; ** $p < 0.05$ with respect to treatment with insulin alone; †Not significant with respect to tyrphostin-25 alone. *c*, N2a cells were incubated for 4 hr in serum-free medium in the absence or presence of 1 μ M insulin, 25 μ M U73122, 500 nM wortmannin, and/or 10 μ M PD98059. Data represent means \pm SD; $n = 3$. * $p < 0.05$ with respect to no addition; ** $p < 0.05$ with respect to treatment with insulin alone.

abolished by tyrphostin-25 and was not mimicked by several other growth factors, indicating that the activation of the insulin/IGF-1 receptor tyrosine kinase is required for insulin-dependent A β secretion. Many of the present studies were performed by using a concentration of insulin of 1 μ M. Although at this concentration a contribution of IGF-1 receptor could not be excluded, the fact that insulin is effective at a concentration of 50 nM indicates the specificity of its effect. A similar effect of IGF-1 on β APP metabolism is to be expected because activation of IGF-1 receptors triggers the same downstream signaling molecules as those of insulin receptors (Lopaczynski, 1999). In an effort to elucidate the signaling pathway downstream of the insulin receptor, we studied the effect of selective inhibitors of the three known insulin-activated signal transduction cascades. The effect of insulin on A β secretion was abolished by PD98059, a highly selective inhibitor of the activation of the MAP kinase kinase (Alessi et al., 1995; Dudley et al., 1995), indicating that the effect of insulin is mediated by activation of the MAP kinase cascade. In contrast, wortmannin, an inhibitor of PI3-kinase, and U73122, a potent inhibitor of phospholipase C, nonselectively reduced basal- and insulin-stimulated secretion of A β .

Accumulation of A β plaques within the brain is widely believed to initiate the pathological cascade culminating in clinical AD. Generally, it is assumed that secreted A β , and particularly A β 42, plays a crucial role in amyloid neuropathology. Recent evidence suggests the hypothesis that intracellular A β 42 may play an important role in amyloid deposition and neuronal degeneration (for review, see Wilson et al., 1999; Gouras et al., 2000): high intracellular levels of A β 42 are

observed in cell lines expressing FAD mutant PS-1 (Wild-Bode et al., 1997); an insoluble pool of A β 42 increases in a neuronal cell line during aging in culture (Skovronsky et al., 1998); intraneuronal accumulation of A β 42 and extensive neuronal degeneration occur in the absence of A β plaques in transgenic mice expressing an FAD mutant form of PS-1 (Chui et al., 1999). Recent immunohistochemical studies have reported intraneuronal A β 42 accumulation in early (Gouras et al., 2000) and late (Mochizuki et al., 2000) AD. Although direct neurotoxicity of intracellular A β has not been demonstrated, we cannot exclude that intracellular A β 42 may play a direct role in initiating AD neuropathology.

Although basal insulin levels do not appear to be reduced in aging, insulin resistance and impaired insulin release in response to a glucose challenge are age-related phenomena (for review, see Lamberts et al., 1997; Perry, 1999). Several clinical studies suggest that insulin plays an important role in cognitive function and memory (for review, see Wickelgren, 1998; Lovestone, 1999). Recent findings demonstrated insulin receptor upregulation and reduced insulin receptor-mediated tyrosine kinase activity in AD brains (Frolich et al., 1998, 1999). Higher-fasting plasma insulin levels and reduced cerebrospinal fluid-to-plasma ratios of insulin, indicative of insulin resistance, have been described in patients with AD (Craft et al., 1998, 1999). Although recent cross-sectional and prospective population-based studies have indicated that diabetes mellitus is a risk factor for dementia and AD (Yoshitake et al., 1995; Ott et al., 1996; Leibson et al., 1997), the issue is still controversial (Landin et al., 1993; Mortel et al., 1993; Nielson et al., 1996; Heitner and Dickson, 1997). According to the

present data, insulin dramatically reduces intracellular levels of A β 40/42 and increases A β 40/42 secretion in neuroblastoma cells and primary neuronal cultures by promoting β APP trafficking via a known insulin-signaling pathway, suggesting that the insulin/IGF-1 pathway may play a role in AD pathogenesis. However, future studies must assess whether diabetes contributes to AD and, if so, whether such effects are mediated via the action of insulin on β APP/A β metabolism.

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