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How does diabetes accelerate Alzheimer disease pathology?

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Abstract

Diabetes and Alzheimer disease (AD)—two age-related diseases—are both increasing in prevalence, and numerous studies have demonstrated that patients with diabetes have an increased risk of developing AD compared with healthy individuals. The underlying biological mechanisms that link the development of diabetes with AD are not fully understood. Abnormal protein processing, abnormalities in insulin signaling, dysregulated glucose metabolism, oxidative stress, the formation of advanced glycation end products, and the activation of inflammatory pathways are features common to both diseases. Hypercholesterolemia is another factor that has received attention, owing to its potential association with diabetes and AD. This Review summarizes the mechanistic pathways that might link diabetes and AD. An understanding of this complex interaction is necessary for the development of novel drug therapies and lifestyle guidelines aimed at the treatment and/or prevention of these diseases.

Introduction

The population of the developed world is aging, and the incidence of age-related metabolic and neurodegenerative diseases is increasing. In the US, diabetes and Alzheimer disease (AD)—two high-morbidity age-related diseases—affect \approx 23.6 and \approx 5.3 million people, respectively. These figures are projected to rise considerably. The Centers for Disease Control and Prevention predict that >29 million people in the US will be affected by diabetes by 2050, while the Alzheimer's Association forecasts that by this date, 11–16 million Americans will have AD. Numerous studies report that patients with diabetes have an increased risk of developing AD compared with healthy individuals.^{1,2} In fact, a study of the Mayo Clinic Alzheimer Disease Patient Registry revealed that 80% of patients with AD exhibited either impairments in glucose tolerance or frank diabetes.³

Diabetes is a complex metabolic disorder characterized by hyperglycemia and associated with microvascular and macrovascular complications, including retinopathy, nephropathy, neuropathy and cardiovascular disease.⁴ An association between diabetes and these complications is well established, although the impact of diabetes on the CNS—particularly in relation to cognitive dysfunction—is not understood in detail. This interaction has,

Competing Interests

Author contributions

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however, received increasing attention over the past decade. 5-10% of patients with diabetes in the US have type 1 diabetes, which is associated with hyperglycemia and insulin deficiency. The severity of cognitive dysfunction experienced by individuals with this type of diabetes is affected by age of onset of diabetes, degree of glycemic control, and duration of diabetes.¹ Type 2 diabetes is the most common form of this disease, accounting for $\approx 90-$ 95% of cases of diabetes in the US, and is characterized by hyperinsulinemia and insulin resistance. Obesity, hypertension, hypercholesterolemia and hyperlipidemia are all associated with type 2 diabetes. Early cognitive changes in learning and memory, mental flexibility and mental speed are also associated with this form of the disease (Box 1).^{5,6}

Box 1

Cognitive processes affected by diabetes

Type 1 diabetes

- Information processing
- Psychomotor efficiency
- Attention
- Visuoconstruction
- Memory
- Visual-motor skills
- Visual-spatial skills
- Motor speed
- Vocabulary
- General intelligence
- Motor strength

Type 2 diabetes

- Psychomotor speed
- Frontal lobe and executive function
- Attention
- Verbal fluency
- Memory
- Processing speed
- Complex psychomotor function

AD accounts for 50–70% of all dementia cases and is characterized by cognitive deficits⁷ as well as several neuropathological markers, which include extracellular senile plaques and intracellular neurofibrillary tangles (NFTs). Familial AD is a rare form of dementia and is caused by autosomal dominant mutations in one or more of the genes encoding the amyloid precursor protein (APP), presenilin 1 or presenilin 2 (the latter two proteins form the catalytic core of γ -secretase).⁸ By contrast, late-onset AD might be caused by environmental and/or lifestyle factors.⁹ Interestingly, late-onset AD is characterized not only by the neuropathological markers mentioned above, but also by vascular lesions, and

hyperglycemia, hyperinsulinemia, insulin resistance, glucose intolerance, adiposity, atherosclerosis and hypertension are all independent risk factors for AD.¹⁰

Owing to the profound socioeconomic impact of diabetes and AD, an understanding of the mechanisms that might link these two diseases together is imperative. Oxidative stress, the formation of advanced glycation end products (AGEs), and impairments in CNS insulin signaling^{11–14} are all associated with diabetes and AD. Furthermore, dysregulation of cellular processes—such as glucose metabolism, apolipoprotein E (ApoE) processing, cholesterol metabolism, mitochondrial activity, calcium homeostasis, and second messenger signaling—is thought to contribute to both diseases.^{3,14–16} This Review discusses the potential biological mechanisms that could underlie how diabetes might accelerate the progression of AD, and highlights possible points of intervention that future therapies could exploit to prevent or delay the progression of AD in patients with this metabolic condition.

Epidemiological studies

Multiple population-based studies have shown that patients with diabetes exhibit an increased risk of developing AD compared with nondiabetic individuals matched for age and sex;^{1–3} however, other studies have failed to find a link between these two conditions.^{17–19} The reason why some studies but not others have identified such a link might reflect differences in study participants (in terms of age, ethnicity and sex) and/or study designs between investigations. In studies that indicated a positive association between the two diseases, obesity, dyslipidemia, and high blood pressure were all identified as potential risk factors for both diabetes and AD.^{3,15} An analysis of nine high-quality studies demonstrated that individuals with probable type 2 diabetes have nearly a twofold higher risk of AD than individuals without diabetes (Table 1).²

Diabetes and progression of AD

Protein processing

Many neurodegenerative disorders are characterized by abnormal protein processing, and two prominent pathological features of AD-both of which result from abnormal protein processing—are amyloid- β (A β) plaques and NFTs. A β is a 39–42 amino acid peptide generated by the proteolytic cleavage of APP by β -secretase and γ -secretase.²⁰ Compared with shorter variants of A β , the 42 amino acid form of this peptide—A β_{42} —has an increased propensity to aggregate, and accumulates as extracellular amyloid deposits; that is, senile plaques.^{21,22} As mentioned above, genetic analyses have shown that mutations in one or more of the genes encoding APP, presenilin 1 or presenilin 2 cause some cases of familial AD; however, the familial form of this disease is rare, and $\approx 90\%$ of known AD cases are sporadic.^{9,10} The amyloid cascade hypothesis states that the neurodegeneration associated with AD is caused by the processing of APP via the amyloidogenic pathway,^{20,22} and now the production of senile plaques is widely considered to affect neuronal activity by impairment of synaptic function and induction of cell death.^{23,24} However, impaired synaptic function might precede the formation of plaques.²⁵ The accumulation of Aβderived diffusible ligands (ADDL), which are soluble AB oligomers, has been shown to lead to functional deficits before plaque formation.²⁶ As impaired insulin signaling, which is associated with type 2 diabetes is known to affect the expression and metabolism of $A\beta$,⁴ diabetes might, therefore, exacerbate the production of AB, synaptic impairment and neuronal cell death in patients with AD.

A major component of NFTs is hyperphosphorylated tau.²⁷ Tau is a soluble microtubuleassociated protein that is expressed in mature neurons and is localized within axons, and maintains the stability of neuronal microtubules. In AD, tau is abnormally phosphorylated

and aggregates in cell bodies and proximal dendrites.²⁸ Cleaved tau is detected in the brains of patients with AD as well as in neurons treated *in vitro* with A β .^{29–31} Cleaved tau has been shown to induce apoptosis of cortical neurons *in vitro*,³⁰ and when expressed in transgenic animals, the cleaved form of this protein is associated with a reduction in spatial memory.³² Several proteases, including caspases (proteases associated with the proteasome) and calpains, are implicated in tau cleavage.³³

In mouse models of diabetes, increased tau phosphorylation is evident in animals with either type 1 or type 2 diabetes compared with control wild-type mice.³⁴ Tau phosphorylation and cleavage is considerably more pronounced in mice with type 2 diabetes than in mice with the type 1 form of this disease,^{34,35} although the change in phosphorylation status in the latter positively correlates with impairments in learning and memory.^{34,36–38} Considering that an increase in tau phosphorylation is observed in postmortem brain samples from patients with type 2 diabetes, patients with this form of the disease might also experience cognitive deficits as a result of dysfunctional glucose metabolism.³⁹

Glycosylation of *O*-linked β -*N*-acetylglucosamine (GlcNAcylation) is a normally occurring post-translational protein modification. Studies show a reduction in GlcNAcylation under diabetic conditions, indicating that GlcNAcylation might attenuate abnormal tau phosphorylation and decrease tau-induced neuronal death.⁴⁰

Insulin and glycogen synthase kinase

Insulin deficiency associated with type 1 diabetes contributes to the cognitive deficits observed in patients with this form of the disease.¹ Furthermore, both spatial learning and hippocampal long-term potentiation (LTP) are attenuated following induction of experimental type 1 diabetes in rats with a single intravenous injection of streptozotocin.^{1,41} These changes in spatial learning and LTP can be prevented by insulin treatment (Figure 1).⁴¹

Type 2 diabetes is also associated with cognitive impairment.⁴² As mentioned above, this form of diabetes is characterized by insulin resistance, hyperinsulinemia and impaired insulin signaling. Insulin receptors are expressed throughout the CNS; however, the function of these receptors in the brain is not fully understood. Insulin receptors might be involved in the regulation of synaptic activity and, hence, might affect cognitive processes.⁴³ Neurodegeneration and cognitive impairment in type 2 diabetes and AD could be caused, in part, by impairments in insulin receptor signaling.⁴⁴ In fact, decreases in the sensitivity of such receptors are known to affect the expression and metabolism of A β and tau,⁴ and impaired insulin receptor activity and hyperinsulinemia are observed in patients with AD and in animal models of this disease.⁴⁵ In addition, dysfunction of insulin receptor signaling is associated with impairments in ADDL clearance.⁴⁶

Insulin-degrading enzyme (IDE) is required for both insulin and A β degradation in neurons and microglia. Elevated insulin levels in type 2 diabetes induce A β accumulation through competition between insulin and A β for IDE.⁴⁷ Insulin and insulin receptor densities are, however, decreased in patients with AD compared with age-matched healthy controls.^{12,45} Nevertheless, insulin sensitizers might increase insulin signaling and decrease the levels of insulin available to compete with A β for degradation by IDE. Treatment of mice with ADlike symptoms with an insulin sensitizer reduces A β_{42} levels and improves memory.⁴⁸ Although a larger study is needed to confirm the findings, patients with early AD who were treated with the insulin sensitizer rosiglitazone failed to demonstrate a decline in A β_{42} levels, despite demonstrating better cognitive performances during treatment than before treatment.⁴⁹

Insulin regulates tau phosphorylation *in vitro*^{50,51} and *in vivo*,^{52,53} and increases the rate of NFT development. In fact, insulin transiently increases tau phosphorylation in primary cortical neurons,⁵¹ and hyperinsulinemia results in tau hyperphosphorylation in rat brains.⁵⁴ Furthermore, insulin receptor substrate 2 knockout mice demonstrate typical pathological signs of type 2 diabetes and have an increased number of NFTs in hippocampal neurons compared with control wild-type mice.⁵³ Thus, impaired insulin signaling could increase tau phosphorylation and cleavage.³⁴

Insulin receptor signaling leads to the activation of two major signaling pathways, the mitogen-activated protein kinase (MAPK) pathway and the Akt signaling pathway. MAPK signaling is a required component of cell differentiation, cell proliferation and cell death,⁵⁵ whereas Akt signaling is involved in the regulation of cell growth, cell proliferation, protein synthesis (via the mammalian target of rapamycin signaling pathway) and cell survival (through the inhibition of several pro-apoptotic agents).^{56,57} Both the MAPK and Akt pathways are implicated in AD pathogenesis. MAPK expression is increased in the brains of patients with AD compared with healthy individuals. Moreover, MAPK immunoreactivity is greater in postmortem brain samples from patients with AD than in such samples from healthy controls.⁵⁸ The expression of this protein kinase is positively associated with Aβ plaques and NFTs. Indeed, MAPK co-localizes with NFTs in hippocampal and cortical regions in AD brains.^{58,59} Studies indicate that MAPK signaling is involved in neuroinflammation, tau phosphorylation and synaptic plasticity.⁵⁹ For example, in transgenic mice with hyperphosphorylated tau, aggregated tau co-localizes with MAPK.⁶⁰

Akt signaling induces the inhibition of glycogen synthase kinase- 3β (GSK- 3β)^{44,61,62} (Figure 1), which phosphorylates and, hence, inactivates glycogen synthase, a key enzyme in glycogenesis. Thus, under normal conditions, insulin signaling via the insulin receptor leads to GSK- 3β inactivation, whereas insulin resistance leads to GSK- 3β dephosphorylation and activation.^{36,61} The regulation of GSK- 3β in the hippocampus and cortex changes in response to changes in glucose and insulin concentrations,⁶³ and in type 2 diabetes an increase in GSK- 3β activity might lead to insulin resistance by reducing glucose clearance.⁶² Increased GSK- 3β activation might also lead to an elevation in A β production (resulting from a GSK- 3β -mediated increase in presenilin 1 activity)⁶⁴ and an increase in tau phosphorylation associated with NFT formation (Figure 1).⁶¹ By contrast, inhibition of GSK- 3β attenuates APP processing and inhibits hyperphosphorylated tau-associated neurodegeneration in cell-culture and animal models of AD.^{64,65}

Abnormal glucose metabolism

Glucose metabolism and insulin signaling are important for normal brain function. Imaging studies have revealed that patients with AD and individuals at risk of developing this disease typically have reductions in glucose metabolism in temporal and parietal brain regions.⁶⁶ Of note, reductions in glucose metabolism have been observed in the hippocampus of patients with AD.⁶⁷ Moreover, compared with healthy individuals, patients with AD might also have increased fasting plasma insulin levels and/or a decreased cerebrospinal fluid (CSF)-toplasma insulin ratio. Intravenous administration of insulin—while maintaining normal blood glucose levels—or glucose facilitates cognitive functioning in patients with AD and in healthy older adults.⁶⁸ This finding indicates that normal glucose metabolism is required for the performance of cognitive functions, and that impairments in glucose metabolism might contribute to cognitive dysfunction. The negative effect of impaired glucose metabolism on cognitive functioning might be caused, in part, by the formation of AGEs, an increase in oxidative stress and, subsequently, an increase in local inflammation within the brain.

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Advanced glycation end products—Hyperglycemia can lead to a number of pathophysiological processes, including oxidative stress and AGE formation that can cause brain damage.⁶⁹ In fact, both abnormal glucose metabolism and oxidative stress contribute to the formation of AGEs. These substances comprise a heterogeneous group of molecules formed by irreversible, non-enzymatic reactions between sugars and the free amino groups of proteins, lipids and nucleic acids. Auto-oxidation of glucose leads to the formation of oxygen radicals, which are intermediates in the AGE pathway and the predominate source of endogenous AGEs (Figure 2).⁷⁰ The formation and accumulation of AGEs occurs during normal aging; however, these processes are exacerbated in patients with diabetes,^{71,72} and the binding of AGE to its receptor (receptor for AGEs or RAGE) induces a series of biological processes that cause further diabetic complications.⁷¹

AGE immunoreactivity is present in both Aβ plaques and NFTs in patients with AD.⁷³ Furthermore, hippocampal neurons from patients with this neurodegenerative disease contain Aβ-positive, AGE-positive and RAGE-positive granules.^{73,74} Whether the modification of Aβ and tau by AGEs is a primary or secondary event in AD is a controversial topic. Nevertheless, AGEs are widely accepted to be active participants in the progression of AD, since AGE-induced glycation of Aβ and tau protein has been shown to cause Aβ aggregation and the formation of NFTs, respectively.⁷⁵ Moreover, diabetic mice with cognitive impairments exhibit increased RAGE expression in neurons and glia compared with wild-type control mice,⁷⁶ and in one clinical study, AGE immunostaining was increased in postmortem brain slices from patients with AD and diabetes compared with nondiabetic patients with AD.⁷⁷ Another study has failed, however, to detect a difference in AGE immunostaining in NFTs and senile plaques between patients with diabetes and agematched control individuals.⁷⁸

Oxidative stress—Abnormal glucose metabolism can increase the production of free radicals such as reactive oxygen species (ROS) and reactive nitrogen species (RNS). This overproduction of free radicals can exhaust a cell's antioxidant capacity and lead to a condition known as oxidative stress, which is a hallmark of both type 1 and type 2 diabetes and a contributing factor to diabetic neuropathy.^{79,80} ROS-induced and RNS-induced protein and/or lipid peroxidation result in cell damage that can lead to cell death, and are increased in patients with diabetes or AD compared with healthy controls.^{81,82} Brain and CSF levels of lipid peroxidation biomarkers—including malondialdehyde and 4-hydroxynonenal—are higher in individuals with AD than in healthy people.¹¹ Furthermore, levels of oxidized proteins are increased in the frontal and parietal lobes and in the hippocampus of patients with mild cognitive impairment compared with healthy controls, ⁸³

Oxidative stress and lipid peroxidation seem to be able to induce A β accumulation: studies in a mouse model of AD have demonstrated that brain lipid peroxidation increases before A β levels increase,⁸⁴ and that the onset of A β deposition is associated with an increase in the level of RNS.⁸⁵ Further evidence supporting this hypothesis has been obtained from studies of a mouse model of AD in which mutations in the genes encoding APP and pre-senilin 1 cause an elevation in A β_{42} production. In these animals, lipid and protein peroxidation are evident at disease onset.⁸⁶ In another transgenic animal model of AD, in which mice develop A β plaques, NFTs and cognitive defects, a decrease in antioxidant capacity and an increase in lipid peroxidation were noted before the development of AD pathology.⁸⁷ Oxidative stress seems to affect APP either directly, by increasing APP levels, or indirectly, by modulating APP processing, and both mechanisms could increase levels of A β . Studies in transgenic mice and postmortem brain tissue from patients with AD also suggest that an increase in A β production leads to a rise in the production of ROS. Evidence indicates that insertion of A β into the mitochondrial membrane can disrupt the mitochondrial electron

transport chain,⁸⁸ markedly attenuating cellular energy production and increasing the formation of ROS and, thus, driving a feedforward process involving ROS and A β . This feedforward process might also result from microglia activation by A β protofibrils: these fibrils can induce a robust inflammatory response and the local release of neurotoxins or neurotoxic cytokines from activated microglia.⁸⁹

Oxidative stress increases both the activation of inflammatory pathways (Figure 2)¹¹ and the release of inflammatory mediators such as C-reactive protein and interleukin 6 into the circulation.⁹⁰ Local inflammation initiated by activated microglia and reactive astrocytes surrounding extracellular A β plaques can lead to activation of the complement cascade and neuronal cell damage.⁹¹ Cytokines and chemokines, including interleukins, tumor necrosis factor and macrophage inflammatory protein 1 α , are all upregulated in patients with AD.⁹¹ Furthermore, transgenic mice exhibiting A β plaques and cognitive defects have higher levels of cytokines and chemokines than have wild-type control animals.⁹² Observational epidemiological studies have reported that anti-inflammatory drugs are associated with a decreased risk of AD;⁹³ however, experimental trials in which patients with AD were treated with anti-inflammatory drugs failed to demonstrate any beneficial effects of taking these agents.^{94,95}

Cholesterol, APOE and metabolic syndrome

Dyslipidemia and hypercholesterolemia both contribute to the pathology of diabetes and are also independent risk factors for AD. Apolipoprotein E (APOE), which is expressed predominantly in the liver and brain, participates in the transport of cholesterol and lipoproteins within the circulatory system and can, therefore, markedly affect blood lipid levels. In fact, humans with an APOE deficiency exhibit an increase in the level of plasma cholesterol,⁹⁶ and *ApoE* knockout mice have been shown to have elevated blood cholesterol levels compared with wild-type control mice, even when the knockout mice have been on a strict low-fat, low-cholesterol diet.⁹⁷ Four alleles of *APOE* exist. In the general population, 77% of individuals have an *APOE* ε 3 allele (the most common *APOE* variant), while 15% possess an ε 4 variant. By contrast, 40% of patients with AD possess an ε 4 allele, and individuals with one ε 4 allele are 3–4-fold more likely to develop AD than individuals who are not ε 4 carriers.

The risk of AD associated with the *APOE* ε 4 allele might be exacerbated by diabetes, as patients with diabetes who are ε 4 carriers are twofold more likely to develop AD than individuals who harbor the ε 4 allele but are not diabetic.⁹⁸ Results from one study have indicated that diabetes only increases the risk of AD in individuals who have the ε 4 allele;¹⁸ however, other investigations have failed to demonstrate a correlation between *APOE* ε 4 status, diabetes and AD.^{99,100}

How *APOE4* carrier status might confer an increased risk of developing AD is unclear. APOE has, however, been implicated in the clearance of $A\beta$,¹⁰¹ and APOE ϵ 4 and $A\beta$ aggregates have been shown to synergistically induce neurodegeneration in the brains of mice with AD-like symptoms.¹⁰² Evidence indicates that APOE ϵ 4 also affects the processing of tau: for example, overexpression of APOE ϵ 4 in transgenic mice seems to promote neuronal tau phosphorylation.¹⁰³ APOE receptors, specifically the low-density lipoprotein receptor (LDLR)-related protein, are known to interact with APP, increasing its endocytic trafficking and amyloidogenic processing.¹⁰⁴

As highlighted above, hypercholesterolemia is associated with an increased risk of type 2 diabetes, and patients with this form of the disease have an increased risk of cognitive decline compared with healthy individuals.¹⁰⁵ In fact, hypercholesterolemia is present in 70% of patients diagnosed with diabetes and 77% of patients with this condition who go

from the circulation to the brain, and by decreasing the expression of IDE and LDLR-related protein, both of which are involved in the clearance of A β from the brain to the circulation.¹⁰⁸ Furthermore, cholesterol, cholesterol oxidase, and APOE have all been shown to co-localize with A β in fibrillar plaques in transgenic mouse models of AD,¹⁰⁹ and cholesterol and oxidized cholesterol (oxysterols) have also been shown to accumulate in the dense core of A β plaques,¹¹⁰ indicating that cholesterol might be involved in the formation of senile plaques. In the brain, cholesterol is oxidized to 24-hydroxycholesterol (24-HC) by 24-hydroxylase, and 24-HC levels in plasma and CSF reflect neuronal cholesterol synthesis. In addition, 27-HC is one of the main oxysterols found in the circulation, and the ratio of 24-HC and 27-HC is altered in AD.¹¹¹ In fact, the level of 27-HC in the brain increases in this neurodegenerative disease¹¹¹ and this rise might, subsequently, elevate the production of A β .¹¹² Thus, statins that cross the blood–brain barrier could reduce the risk of AD,¹¹³ owing to their ability to reduce the levels of neuronal cholesterol, tau phosphorylation and amyloid formation.^{114,115}

Mitochondrial dysfunction

Diabetes and AD are associated with deficits in mitochondrial activity.¹⁶ Mitochondria are essential for ATP synthesis and for maintaining calcium homeostasis, which is required for normal neuronal function.¹¹⁶ The calcium hypothesis states that dysregulation of calcium homeostasis is a central process in normal aging of the brain and in age-related diseases.^{117,118} Excessive calcium uptake by mitochondria leads to an increase in ROS production, inhibition of ATP synthesis, release of cytochrome c, and a sudden increase in inner membrane permeability termed mitochondrial permeability transition.¹¹⁹ Mitochondrial dysfunction triggers neuronal degeneration and cell death, and is thought to contribute to AD pathophysiology. The role mitochondrial dysfunction has in AD is not fully understood, but APP is known to be associated with the outer mitochondrial membrane,¹²⁰ and β -secretase and A β , which inhibits cytochrome oxidase in the presence of copper, are present in mitochondria, indicating that AB could negatively affect mitochondrial electron transport.¹²¹ Indeed, dysfunction of mitochondrial electron transport proteins and a decrease in cytochrome oxidase activity are both associated with AD.¹²² Furthermore, neurons affected by AD pathology exhibit an overall decrease in mitochondrial mass, an increase in cytoplasmic mitochondrial DNA, and an increase in cytochrome oxidase in lipofuscin-containing vacuoles.123

Dysregulation of calcium homeostasis seems to occur as a result of a decrease in mitochondrial function. Brain tissue from patients with AD shows an increase in the concentration of calcium, and intracellular calcium levels are higher in neurons containing NFTs than in neurons from healthy control patients.¹²⁴ Furthermore, levels of calcium-dependent proteases are increased in neurons containing NFTs compared with neurons without NFTs,¹²⁵ and neurons that are vulnerable to degeneration exhibit an increase in the levels of calcium–calmodulin-dependent protein kinase II.¹²⁶ Transglutaminase, a calcium-activated enzyme that induces crosslinking of tau molecules, is also increased in patients with AD compared with controls,¹²⁷ and cells exposed to agents that induce calcium influx show elevations in A β production.¹²⁸ Thus, an increase in levels of intracellular calcium might be involved in enhancement of APP processing and A β production.

Type 1 and type 2 diabetes are associated with an increase in intracellular calcium levels (Figure 2).¹²⁹ Abnormal calcium homeostasis is common in patients with diabetes and has been shown to occur in animal models of this disease.¹²⁹ Hormonal regulation of

intracellular calcium might also be abnormal in diabetes. In type 2 diabetes, high levels of intracellular calcium in pancreatic β -cells and alterations in membrane cation pumps might contribute to impaired insulin secretion.¹³⁰ Insulin deficiency is associated with calcium overload, which affects metabolic function by activating calcium-dependent protein kinases, phosphatases, proteases, phospholipases and lysosomal enzymes.¹³¹ In animal models of type 1 diabetes, neuronal mitochondria exhibit a decrease in antioxidant capacity owing to a low content of coenzyme Q9, which is a result of oxidative stress.¹³² Inherited defects in mitochondrial DNA are known to cause an insulin-deficient form of diabetes mellitus that resembles type 1 diabetes. In patients with type 2 diabetes, the activity of mitochondrial oxidative enzymes is lower than in age-matched controls; however, patients with this form of diabetes are obese, and obesity is associated with smaller mitochondria and reduced bioenergetic capacity than lean controls.¹³³

Animal models of AD and diabetes

Several studies have demonstrated that the induction of diabetes in mouse models of AD leads to an acceleration of AD neuropathology. For example, the induction of type 1 diabetes-by means of an intraperitoneal injection of streptozotocin-in transgenic mice prone to tau pathology is associated with an increase in levels of hyperphosphorylated and insoluble tau.¹³⁴ Furthermore, results from a study conducted by Jolivalt et al. indicate that hypoinsulinemia increases AD pathology in transgenic mice with AD and experimental type 1 diabetes. Rises in the level of A β_{42} the number of immunoreactive A β plaques, GSK-3 β activity, and the degree of tau phosphorylation-all contributing factors to neurodegeneration and neuronal loss in AD-were evident in the transgenic mice after induction of experimental diabetes.¹³⁵ By contrast, induction of type 2 diabetes did not increase A β levels in the brains of transgenic mice that develop AD pathology, although the development of diabetes did cause the early onset of cognitive dysfunction.¹³⁶ Taken together, these results suggest that diabetes-associated cerebral amyloid angiopathy, but not A β accumulation, is responsible for cognitive dysfunction, at least in the early stages of the disease. Owing to the fact that the mice with experimental type 2 diabetes were leptin deficient, and that leptin is involved in synaptic function and affects cognition and behavior, ¹³⁷ the absence of leptin signaling could be responsible for the observed cognitive phenotype. More studies are needed to fully elucidate the link between diabetes and AD.

Conclusions

The work presented here highlights the overlap and the many of the points of intersection that exist between the molecular mechanisms underlying diabetes and AD, and indicates how diabetes could exacerbate AD pathology. Hyperglycemia and hypoglycemia in the CNS result in the dysregulation of multiple extracellular and intracellular signaling cascades, which in turn could lead to decreases in neuronal and synaptic function and, ultimately, to an increase in neuronal loss. An understanding of how each molecular pathway intersects and affects the others is essential for the development of future drug intervention strategies for AD.

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Key points

- Alzheimer disease (AD) and diabetes are both associated with enormous and increasing socioeconomic effects
- Diabetes affects the processing of amyloid-β and tau, and might increase the rate of formation of senile plaques and neurofibrillary tangles, the main neuropathological hallmarks of AD
- Hyperinsulinemia is associated with amyloid-β accumulation and regulates tau phosphorylation
- Oxidative stress activates inflammatory pathways and, hence, might exacerbate AD neuropathology
- Mitochondrial dysfunction is associated with both diabetes and AD, and leads to intracellular calcium dysregulation and abnormal processing of the amyloid precursor protein
- Induction of diabetes exacerbates AD neuropathology in mouse models of this neurodegenerative disease

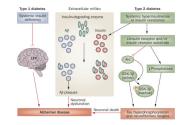


Figure 1.

Altered insulin signaling in diabetes might contribute to Alzheimer disease pathophysiology. In type 1 diabetes, insulin deficiency attenuates LTP and might lead to deficits in spatial learning and memory. In type 2 diabetes, insulin resistance leads to both A β plaque formation and tau hyperphosphorylation. During hyperinsulinemia, insulin and A β compete for insulin-degrading enzyme, leading to A β accumulation and plaque formation. A decrease in insulin receptor signaling leads to inhibition of Akt and dephosphorylation (activation) of GSK-3 β , and results in tau hyperphosphorylation. Abbreviations: A β , amyloid- β ; GSK-3 β , glycogen synthase kinase 3 β ; LTP, long-term potentiation; P, phosphate.

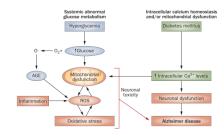


Figure 2.

Pathological mechanisms associated with diabetes might cause AD. Mitochondrial dysfunction, oxidative stress and dysregulated calcium homeostasis are all associated with diabetes and might be contributory factors to the development of AD. Glucose auto-oxidation can lead to AGE formation and, as a result, oxidative stress, which is associated with mitochondrial dysfunction. Oxidative stress combined with an increase in intracellular calcium result in a feedforward cycle of continued mitochondrial damage that can cause neuronal death and, hence contribute to AD pathology. Abbreviations: AD, Alzheimer disease; AGE, advanced glycation end product; ROS, reactive oxygen species.

Table 1

Diabetes increases the risk of developing Alzheimer disease

Reference	Patients (patients with diabetes/total number of patients)	Relative risk [*]
Ott et al. (1999) ¹³⁸	692/6,370	1.9 (95% CI 1.2–3.1)
Brayne et al. (1998) ¹³⁹	25/376 [‡]	OR 1.4 (95% CI 1.1–17.0)
Yoshitake et al. (1995) ¹⁴⁰	70/828	2.2 (95% CI 1.0-4.9)
Peila <i>et al.</i> (2002) ⁹⁸	900/2,574 [‡]	1.7 (95% CI 1.0–2.8)
MacKnight <i>et al.</i> (2002) ¹⁹	503/5,574‡	1.2 (95% CI 0.8–1.8)
Xu et al. (2004) ¹⁴¹	114/1,301	HR 1.3 (95% CI 0.8–1.9)
Leibson <i>et al.</i> (1997) ¹⁴²	1455/75,000 [‡]	SMR 1.6 (95% CI 1.3-2.0)
Luchsinger et al. (2005) ¹⁴³	231/1,138‡	HR 2.4 (95% CI 1.8–3.2)
Arvanitakis et al. (2004) ¹⁴⁴	27/824 [‡]	HR 1.7 (95% CI 1.1–2.5)

Patients with probable type 2 diabetes have nearly a twofold higher risk of AD than individuals without diabetes.

* Relative risk unless otherwise stated.

 ‡ Data represents number of patients at follow-up, all other data represent patient numbers at baseline.

Abbreviations: HR, hazard ratio; OR, odds ratio; SMR, standard morbidity ratio.