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Cellular/Molecular

The Loss of c-Jun N-Terminal Protein Kinase Activity Prevents the Amyloidogenic Cleavage of Amyloid Precursor Protein and the Formation of Amyloid Plaques In Vivo

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Phosphorylation plays a central role in the dynamic regulation of the processing of the amyloid precursor protein (APP) and the production of amyloid-β (Aβ), one of the clinically most important factors that determine the onset of Alzheimer’s disease (AD). This has led to the hypothesis that aberrant Aβ production associated with AD results from regulatory defects in signal transduction. However, conflicting findings have raised a debate over the identity of the signaling pathway that controls APP metabolism. Here, we demonstrate that activation of the c-Jun N-terminal protein kinase (JNK) is essential for mediating the apoptotic response of neurons to Aβ. Furthermore, we discovered that the functional loss of JNK signaling in neurons significantly decreased the number of amyloid plaques present in the brain of mice carrying familial AD-linked mutant genes. This correlated with a reduction in Aβ production. Biochemical analyses indicate that the phosphorylation of APP at threonine 668 by JNK is required for γ-mediated cleavage of the C-terminal fragment of APP produced by β-secretase. Overall, this study provides genetic evidence that JNK signaling is required for the formation of amyloid plaques in vivo. Therefore, inhibition of increased JNK activity associated with aging or with a pathological condition constitutes a potential strategy for the treatment of AD.

Introduction

Alzheimer’s disease (AD) is a neurodegenerative disorder characterized by the presence of extracellular amyloid plaques in the brain formed by the progressive deposition of amyloid-β (Aβ). Aβ is a 4 kDa peptide that derives from the sequential cleavage of the amyloid precursor protein (APP) by β- and γ-secretases (Mattson, 2004). Mutations in genes associated with familial forms of AD (FAD) result in elevated production of Aβ that precedes disease pathology (Hardy, 1997; Bertram et al., 2010). These findings have led to the idea that the accumulation of Aβ is a primary cause of AD. Aβ neurotoxicity is mediated, at least in part, by enhancing the levels of reactive oxygen species (ROS) (Behl et al., 1994). Furthermore, a positive-feedback loop by which oxidative stress produced following the interaction of Aβ with the receptor for advanced glycation endproducts (RAGE) increases RAGE expression, thereby enhancing Aβ neurotoxicity, has been suggested (Yan et al., 1996).

Activation of the proapoptotic c-Jun N-terminal protein kinase (JNK) signaling pathway constitutes one possible mechanism by which ROS produced by the accumulation of Aβ impairs synaptic function and leads to neuronal loss. This is mostly demonstrated by the observation that Aβ-induced neuronal death is significantly reduced in cortical neurons lacking JNK expression (Morishima et al., 2001). JNK is a member of the mitogen-activated protein kinase (MAPK) family that phosphorlates and activates transcription factors of the AP-1 (activator protein-1) family, including c-Jun and ATF2, in response to various stresses (Davis, 2000). Analogous to other MAPKs, JNK is activated by phosphorylation at threonine (Thr) and tyrosine (Tyr) residues by two dual-specificity MAPK kinases (MKK4 and MKK7) (Wang et al., 2007a).

Consistent with a role of JNK signaling in mediating Aβ toxicity (Morishima et al., 2001), recent studies have demonstrated the neuroprotective effect of inhibiting JNK activity in rodent AD models (Braithwaite et al., 2010; Ramin et al., 2011). Furthermore, JNK-mediated APP phosphorylation at Thr (T) 668 has been proposed as a mechanism to prevent APP degradation and increase Aβ production in a neuroglioma cell line (Colombo et al., 2009). The importance of this finding is highlighted by evidence that T668 phosphorylation is increased in human AD brain (Lee et al., 2003). However, the demonstration that the cyclin-dependent kinase 5 (Cdk5) is responsible for phosphorylating APP at T668 in naturally degenerating CAD cell, a CNS-derived neuronal cell line used for studying neuronal cell biology and pathology, has challenged the idea that JNK is implicated in con-
trolling APP metabolism under pathophysiological conditions (Muresan and Muresan, 2007). In contrast, this study provided evidence that JNK phosphorylates APP to control its transport in distal neurites under normal conditions.

Conflicting findings regarding the role of JNK signaling in regulating APP processing may be attributed to differences in experimental conditions and emphasize the importance of using animal models to identify physiologically relevant regulatory mechanisms. Therefore, we decided to test the effect of the loss of JNK activity in the brain of mice that overexpress FAD-linked mutant genes to elucidate the requirement of JNK signaling in AD pathology.

Materials and Methods

Mice. The mouse strains were maintained in a pathogen-free facility at the University of Manchester. All animal procedures were performed under license in accordance with the United Kingdom Home Office Animals (Scientific Procedures) Act (1986) and institutional guidelines.

Genotype determination of mice and tissues. Offspring carrying flox alleles, creERT2, APP, and PS1 transgenes were identified by PCR on tail and brain tissue DNA, as previously described (Jankowsky et al., 2004; Wang et al., 2007b).

Tissue culture. Primary cultures of cortical neurons were prepared from the cerebral cortices of 17-d-old embryos (E17) and cultured on poly-ornithine precoated six well plates in Neurobasal media containing B27 supplement, 1% penicillin/streptomycin, and 1% glutamine. After 30 min with zVAD (10 mM; Sigma-Aldrich), or for 1 h with a neutralizing anti-RAGE antibody (Santa Cruz; sc-8230). Immune complexes were detected with a secondary antibody conjugated to fluorescein (green). DNA was stained with DAPI (blue). Scale bar, 10 μm.

Death assays. Cell survival was quantified by 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) assay. Caspase activity was measured by spectrofluorometer using the DEVD-AMC caspase 3-specific fluorogenic substrate (Alexis Biochemicals).

Histological and immunohistochemical analyses. Mice were anesthetized and perfused with 0.9% saline, followed by 4% PFA. The 30 μm cryosections of brain were blocked in PBS containing 10% goat serum and 0.1% Triton X-100 for 1 h at room temperature before being incubated overnight at 4°C with primary antibodies to MKK4 (1:100; BD Biosciences Pharmingen), MKK7 (1:100; Cell Signaling), or Aβ42 (1:500; Alpha Diagnostic). The following day, the slides were rinsed in PBS and incubated at room temperature for 1 h with secondary biotinylated anti-rabbit antibody. The slides were processed using the ABC detection kit (Vector Laboratories). The presence of the antigens was revealed using the DAB (diaminobenzidine) (brown) peroxidase substrate kit (Vector Laboratories).

Electron microscopy. Brains were fixed with 2.5% PFA and 0.1% glutaraldehyde in phosphate buffer, and then dissected and small blocks of
the cerebral cortex, hippocampus, and entorhinal cortex were postfixed with 1% osmium tetroxide, dehydrated in methanol, and embedded in Epon. Semithin sections were stained with toluidine blue, and selected ultrathin sections, with uranyl acetate and lead citrate.

**Results**

**The deletion of mkk4 and mkk7 protects cortical neurons from Aβ-induced apoptosis**

Mice lacking mkk4 or mkk7 die before birth (Wang et al., 2007a). To circumvent this early lethality and thereby permit the study of the physiological function of JNK signaling in AD pathology, we developed novel mouse models in which specific exons of the mkk4 and mkk7 genes were flanked by LoxP sites [referred to as the flox (fl) allele] (Wang et al., 2007b; A. Hübner and R. J. Davis, unpublished observation). These sites, which do not interfere with the normal expression of the genes, constitute a binding domain for the DNA recombinase Cre. Homozygous mkk4fl and/or mkk7fl mice were mated with a transgenic line expressing Cre fused to a mutated form of the ligand binding domain of the estrogen receptor (ER<sup>T2</sup>) (Hayashi and McMahon, 2002). The ER<sup>T2</sup> moiety ensures the cytoplasmic sequestration of Cre. Cortical neurons were prepared from the embryos and cultured for 7 d before being incubated with 4-OHT to trigger the nuclear translocation of Cre where it specifically recombines the fl alleles.

Immunoblot analysis confirmed that the inactivation of the mkk4 and mkk7 genes only occurred in homozygous flox neurons carrying the creER<sup>T2</sup> transgene (Fig. 1A). The loss of MKK4 and/or MKK7 in matured neurons did not cause any obvious change in cell morphology (Fig. 1B, E). In subsequent experiments, cortical neurons expressing or not Cre and treated with 4-OHT will be referred to as (−/−) and (+/+), respectively.

To determine the functional consequence of the loss of MKK4 and/or MKK7 expression, the viability of mkk4/7<sup>−/+</sup>, mkk4<sup>−/−</sup>, mkk7<sup>−/−</sup>, and mkk4/7<sup>−/−</sup> neurons treated with Aβ42 was examined. Bright-field microscopy indicated that mkk4/7<sup>−/+</sup> neurons underwent extensive cell death after Aβ42 stimulation. The loss of axon integrity, as demonstrated by the disappearance of MAP2 staining, correlated with the detection of fragmented nuclei characteristic of apoptotic cells (Fig. 1B, E). Consistently, caspase 3 activity was increased in neurons treated with Aβ42 (Fig. 1C). Aβ42-induced neuronal death was prevented by incubating the cells with zVAD, a well characterized caspase inhibitor (Fig. 1D). Similarly, Aβ42 induced the apoptotic death of mkk4<sup>−/−</sup> and mkk7<sup>−/−</sup> neurons (Fig. 1E). In contrast, mkk4/7<sup>−/−</sup> neurons remained viable (Fig. 1D, E) and did not display increased caspase 3 activity (Fig. 1C) following Aβ42 stimulation.

Consistent with the requirement of JNK to mediate Aβ-induced apoptosis (Morishima et al., 2001), Aβ42 treatment increased JNK activity in mkk4/7<sup>−/+</sup> neurons with a maximum at 1 h (Fig. 2A). This was prevented following the loss of MKK4 and MKK7 (Fig. 2A) and significantly blocked in neurons pretreated with the antioxidant NAC (Fig. 2B). Impaired JNK activity in mkk4/7<sup>−/−</sup> neurons correlated with low level of c-Jun expression under both basal and stimulated conditions, supporting the evidence that JNK increases the transcriptional activity of c-Jun and
the ability of c-Jun to control its own promoter (Davis, 2000) (Fig. 2C). Furthermore, immunoblot analysis confirmed the proteolytic cleavage of caspase 3 in mkk4/mkk7fl/fl, but not in mkk4/mkk7+/−, neurons stimulated with Aβ42 (Fig. 2C). In addition, mkk4 and mkk7 gene deletion prevented Aβ42-mediated increased RAGE expression (Fig. 2D). The functional importance of this finding is demonstrated by the ability of a neutralizing RAGE antibody to block JNK activation (Fig. 2E) and increased caspase 3 activity (Fig. 2F) in mkk4/mkk7+/− neurons incubated with Aβ42.

Overall, these studies provide genetic evidence that MKK4 and MKK7 are both required to mediate the apoptotic response of neurons by activating JNK following Aβ–RAGE interaction and the production of ROS.

Inactivation of MKK4 and MKK7 in the adult forebrain
To determine the physiological significance of the loss of active JNK in AD pathology, we created a mouse model in which the expression of MKK4 and MKK7 could be specifically abolished in neurons in the adult forebrain. This was achieved by crossing the mkk4/mkk7fl/fl mice with a transgenic line expressing CreER<sup>T2</sup> under the control of the Ca<sup>2+</sup>/calmodulin-dependent protein kinase II (CaMKII) gene promoter (Erdmann et al., 2007). One-month-old mkk4/mkk7fl/fl littermates carrying or not the CaMKII-creER<sup>T2</sup> transgene were injected intraperitoneally every day for 5 d with a nontoxic amount of tamoxifen (200 µg) to induce Cre activation and the neuronal specific deletion of the flox alleles.

Immunostaining of brain sections with antibodies to MKK4 and MKK7 showed that MKK4 was highly expressed in the cerebral cortex, while MKK7 was mostly detected in the CA2 and CA3 regions of the hippocampus (Fig. 3A). The inactivation of the mkk4 and mkk7 genes only occurred in the brain of mkk4/mkk7fl/fl mice expressing Cre and injected with tamoxifen (Fig. 3A). This was confirmed by immunoblot analysis of brain extracts (Fig. 3B). Similar expression of MKK4 and MKK7 in the cerebellum of mkk4/mkk7fl/fl and (CaMKII-creER<sup>T2</sup>)mkk4/mkk7fl/fl mice injected with tamoxifen demonstrated the selective ablation of the proteins in specific areas of the brain (Fig. 3A). JNK activity was decreased by ∼80% following the deletion of the mkk4 and mkk7 genes (Fig. 3C). Consistent with our previous results in cell cultures (Fig. 2), this resulted in impaired c-Jun expression (Fig. 3B). In subsequent experiments, mkk4/mkk7fl/fl and (CaMKII-creER<sup>T2</sup>)mkk4/mkk7fl/fl mice injected with tamoxifen will be referred to as mkk4/mkk7<sup>−/−</sup> and mkk4/mkk7<sup>+/−</sup> animals, respectively.

The loss of MKK4 and MKK7 prevents amyloid plaque formation
To establish the requirement of JNK signaling in the onset and in the progression of AD, we examined the effect of the loss of MKK4 and MKK7 in the brain of mice that overexpress APP. To determine the physiological significance of the loss of active JNK in AD pathology, we created a mouse model in which the expression of MKK4 and MKK7 could be specifically abolished in neurons in the adult forebrain. This was achieved by crossing the mkk4/mkk7fl/fl mice with a transgenic line expressing CreER<sup>T2</sup> under the control of the Ca<sup>2+</sup>/calmodulin-dependent protein kinase II (CaMKII) gene promoter (Erdmann et al., 2007). One-month-old mkk4/mkk7fl/fl littermates carrying or not the CaMKII-creER<sup>T2</sup> transgene were injected intraperitoneally every day for 5 d with a nontoxic amount of tamoxifen (200 µg) to induce Cre activation and the neuronal specific deletion of the flox alleles.

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In contrast, the number of plaques was greatly reduced in the absence of MKK4 and MKK7 (Fig. 4A, B). In fact, plaques that did develop in the brain of the mkk4/mkk7<sup>−/−</sup> mice carrying the APPswe/PS1ΔE9 transgene were found in areas positive for MKK4 and MKK7 expression and JNK phosphorylation. This was expected since the deletion of β genes following tamofoxifen-induced Cre does not occur with 100% efficiency. Consistent with a reduction in Aβ deposition, the level of Aβ42 was significantly decreased following the loss of MKK4 and MKK7 (Fig. 4C). Furthermore, MKK4/7-deficient brains displayed a marked decrease of the number of dystrophic neurite clusters and only a few isolated neurites with residual bodies (Fig. 4D). Together, these results demonstrate that active JNK is required for the formation of amyloid plaques in vivo.

The loss of MKK4 and MKK7 prevents the phosphorylation of APP at T668
To understand the mechanism by which JNK signaling increases Aβ42 production, we investigated the effect of the loss of MKK4 and MKK7 expression on APP phosphorylation at T668. As expected, expression of the APPswe/PS1ΔE9 transgene in mkk4/mkk7<sup>−/−</sup> mice resulted in amyloid deposition as soon as 3 months of age with a progressive increase in plaque number at 5 months (Fig. 4A, B). The amount of Aβ42 produced in the brain increased with a similar kinetics (Fig. 4C). Electron microscopy confirmed the presence of a large number of clustered dystrophic neurites filled with altered mitochondria and residual bodies around amyloid deposits characteristic of AD pathology (Fig. 4D).

Figure 3. Inactivation of MKK4 and MKK7 in the adult forebrain. One-month-old mkk4/mkk7fl/fl mice carrying (+) or not (−) the CaMKII-creER<sup>T2</sup> transgene were injected with tamoxifen to induce Cre-mediated recombination of the flox alleles. A, One month later, brain sections were analyzed by immunohistochemistry using antibodies against MKK4 or MKK7. Scale bar, 10 µm. HC, Hippocampus; DG, dentate gyrus. B, Protein lysates were analyzed by immunoblot with specific antibodies. The data are representative of two independent experiments. C, Endogenous JNK activity was measured by protein kinase assay. Radioactivity incorporated in the GST-c-Jun substrate was quantified by Phosphoimager. Values are expressed as fold of maximum. The data correspond to the mean ± SD (N = 2 animals).
significant decrease in JNK activity (Fig. 5Aii). This prevented the phosphorylation of APP at T668 (Fig. 5Aii). The low level of phospho (P)–T668–APP correlated with the accumulation of the full-length (FL) protein as well as a C-terminal fragment (CTF) produced after the cleavage of APP by α or β secretases (Fig. 5Aii). Further analyses using an antibody raised against amino acids 1–17 of human Aβ (6E10) confirmed that mkk4 and mkk7 gene deletion increased CTFβ (Fig. 5Aii). The effect of mkk4 and mkk7 gene deletion on APP processing was tested in cortical neurons expressing the APPswe/PS1dE9 transgene, after 14 d in culture. Consistent with our in vivo data, the loss of MKK4 and MKK7 following Cre activation prevented APP phosphorylation at T668, causing an accumulation of APP and CTFβ (Fig. 5B). A similar phenotypic abnormality has been observed in cortical neurons defective in PS1 or expressing loss-of-function variants (Naruse et al., 1998). Therefore, we hypothesized that mkk4 and mkk7 gene deletion increased APP level by preventing γ-mediated cleavage of CTF. To test this hypothesis, we examined the effect of DAPT, a known inhibitor of γ-secretase. The results show that incubation of mkk4/7+/− neurons with DAPT increased APP and CTF levels (Fig. 5Ci), as well as cell survival (Fig. 5Cii), to a similar extent as that associated with the loss of MKK4 and MKK7. Consistent with the observation that neurons lacking MKK4 and MKK7 are less sensitive to the expression of the APPswe/PS1dE9 transgene than wild-type cells, the level of Aβ42 was significantly lower in mkk4/7−/− than in mkk4/7+/− neurons (Fig. 5Ciii). In addition, we found that the brain of 5-month-old mkk4/7+/− mice and mkk4/7−/− neurons cultured for 18 d displayed higher levels of APP and CTF compared with wild-type (wt or +/+ ) samples (Fig. 6Ai,Bi). Consistently, the loss of MKK4 and MKK7 prevented increased production of Aβ42 in aging brains and neurons (Fig. 6Ai,Bii).

Together, these results provide strong evidence that increased JNK activity downstream of MKK4 and MKK7 is required for the amyloidogenic cleavage of mutant, but also endogenous, APP and Aβ42 production.

**Discussion**

In this study, we demonstrated that the loss of active JNK associated with the absence of both MKK4 and MKK7 prevents neurons against Aβ-induced toxicity. Furthermore, we discovered that JNK signaling is required for amyloid plaque formation in vivo. Together, these results provide the first genetic demonstration that activation of JNK triggers the cascade of event that leads to AD pathology.

Phosphorylation plays a central role in the dynamic regulation of APP processing (Gandy and Greengard, 1994). This has led to the hypothesis that aberrant Aβ production associated with AD results from regulatory defects in signal transduction. Our results support this idea by providing a genetic link between amyloid plaque formation and increased JNK activity in neurons via the amplification of a positive-feedback loop associated with the phosphorylation of APP at T668 (Fig. 7). However, a previous study showed that, while JNK phosphorylates APP to control its transport in distal neurites under normal conditions, Cdk5 is responsible for phosphorylating APP in naturally degenerating CAD cells overexpressing or not APP (Muresan and Muresan, 2007). Furthermore, the role of T668 phosphorylation in promoting the amyloidogenic cleavage of APP (Lee et al., 2003; Vingtdeux et al., 2005) has been disputed by evidence that a knock-in mouse model in which T668 was replaced by an alanine residue displayed normal level of Aβ (Sano et al., 2006). Similarly, there is an ongoing debate over the role of phospho-T668 in regulating APP metabolism as a consequence of modulating its ability to interact with its binding partners. For example, Pin1 is a peptidyl-prolyl isomerase that binds the phosphorylated T668 motif of APP, leading to increased Aβ production (Akiyama et al., 2005). However, a role of Pin1 in promoting the nonamyloidogenic processing of APP has also been described previously (Pastorino et al., 2006). Similarly, there is an ongoing debate over the role of phospho-T668 in regulating APP metabolism as a consequence of modulating its ability to interact with its binding partners.
Overall, these discrepancies may be attributed to differences in experimental conditions (wild-type vs mutant APP; endogenous vs ectopic expression; neurons in culture vs animal model). For example, T668 phosphorylation prevents /H9253-mediated cleavage of wild-type APP (Feyt et al., 2007), while APPswe mutant requires to be phosphorylated at T668 before being cleaved by /H9253-secretase (Vingtdeux et al., 2005). In contrast, the Swedish mutation that enhances the affinity of APP for /H9252-secretase may bypass the requirement of T668 phosphorylation, which has been proposed to facilitate the cleavage of APP by /H9252-secretase in primary cortical neurons (Lee et al., 2003). Consequently, the functional disruption of JNK signaling may affect /H9252-mediated cleavage of wild-type APP, but not APPswe mutant. This is consistent with the detection of CTFβ in the brain of APPswe/PS1dE9mkk4/7br mice. Together, these apparent conflicting findings support the hypothesis that APP metabolism is regulated by distinct signaling mechanisms during normal aging and in AD condition in which mutant or wild-type forms of APP are often expressed above physiological levels.

Therefore, although we show that JNK signaling is implicated in the processing of endogenous APP, it will be important to determine the requirement of MKK4 and MKK7 in regulating the cleavage of overexpressed wild-type APP. This will help to clarify the specific requirement of T668 phosphorylation in controlling β cleavage of APP. Together, this information will enable us to predict (1) the usefulness of JNK inhibitors to treat late-onset AD patients that do not display the Swedish mutation and (2) take advantage of the biological activity of MKK4 and MKK7 to control APP metabolism in AD conditions.
Increased JNK activity induces amyloid plaque formation and neuronal demise. Under normal conditions, Aβ is a relatively minor product in the brain because the nonamyloidogenic processing of APP by α- and γ-secretases prevails. α-secretase-mediated cleavage of APP generates secreted (s) APP and γ-secretase. Processing of CTFβ by γ-secretase produces p3 and an APP intracellular domain (AICD). However, defects in Aβ/γ cleavage, associated with aging or a pathological condition, lead to a rise in basal Aβ level. Aβ interacts with RAGE to produce ROS and activates JNK via MKK4 and MKK7. Active JNK phosphorylates AIP1 transcription factors to regulate gene expression (i), and APP at T668 to enhance the amyloidogenic cleavage of APP (ii). CTFβ generated by β-secretase is processed by γ-secretase to produce a phosphorylated form of AICD, which can translocate to the nucleus (Chang et al., 2006), and more Aβ, which further stimulates JNK activity. Sustained high-level JNK activity in the brain induces caspase activation (iii) causing synaptic dysfunction (D’Amelio et al., 2011) and/or neuronal death. A key therapeutic strategy may be to interfere with the amplification of positive-feedback loops that shift the balance toward the amyloidogenic pathway leading to increased Aβ42 production and AD pathology, sec, secretase.

References


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