

RAGE potentiates A β -induced perturbation of neuronal function in transgenic mice

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Receptor for Advanced Glycation Endproducts (RAGE), a multiligand receptor in the immunoglobulin superfamily, functions as a signal-transducing cell surface acceptor for amyloid-beta peptide (A β). In view of increased neuronal expression of RAGE in Alzheimer's disease, a murine model was developed to assess the impact of RAGE in an A β -rich environment, employing transgenics (Tgs) with targeted neuronal overexpression of RAGE and mutant amyloid precursor protein (APP). Double Tgs (mutant APP (mAPP)/RAGE) displayed early abnormalities in spatial learning/memory, accompanied by altered activation of markers of synaptic plasticity and exaggerated neuropathologic findings, before such changes were found in mAPP mice. In contrast, Tg mice bearing a dominant-negative RAGE construct targeted to neurons crossed with mAPP animals displayed preservation of spatial learning/memory and diminished neuropathologic changes. These data indicate that RAGE is a cofactor for A β -induced neuronal perturbation in a model of Alzheimer's-type pathology, and suggest its potential as a therapeutic target to ameliorate cellular dysfunction.

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Introduction

Recent studies have focused attention on amyloid- β peptide (A β) as a key pathogenic entity in neuronal dysfunction and neuropathologic changes of Alzheimer's disease (AD). The steady accumulation of A β over years in AD may result in progressive displacement of essential neuronal structures. However, early in the disease process, when levels of deposited A β are low, mechanisms amplifying and focusing the effects of soluble neurotoxic A β assemblies on vulnerable cellular targets could contribute importantly to cellular dysfunction. Several mechanisms could potentially target A β to cellular elements. In this regard, cell surface-binding sites are logical to consider for multiple reasons: their capacity to concentrate A β at the plasma membrane, where it could directly damage membranes; the possibility that they could function as receptors which engage in intracellular signaling mechanisms; and, their ability to trigger endocytosis, potentially concentrating toxic species in the endolysosomal pathway where disruption of lysosomal integrity could induce severe cellular damage (Yang *et al*, 1999). As might be expected for a pleiotropic peptide such as A β , many cell surface interaction sites have been reported (Snow *et al*, 1995; Paresce *et al*, 1996; Yan *et al*, 1996; Wang *et al*, 2000). We have focused our attention on Receptor for Advanced Glycation Endproducts (RAGE) for several reasons; its enhanced expression in an A β -rich environment (Yan *et al*, 1995; Lue *et al*, 2001; Deane *et al*, 2003); the effectiveness with which it recruits and activates signal transduction mechanisms; and its presence on critical cells (neurons, microglia, and cells of the vessel wall) (Schmidt *et al*, 2001).

A key test of the possible involvement of RAGE (as well as other candidate cell surface binding sites/receptors) in cellular dysfunction associated with A β involves analysis of transgenic (Tg) models in which mutant human amyloid precursor protein (mAPP) and/or presenilins are overexpressed (Duff *et al*, 1996; Hsiao *et al*, 1996; Holcomb *et al*, 1998; Mucke *et al*, 2000). Since many of these Tg models display only subtle changes in neuronal function and neuropathology, especially at early time points, we began our analysis by developing Tg mice with neuronal overexpression of wild-type (wt) RAGE. Our results demonstrate that double Tg mice, with targeted neuronal expression of transgenes for mAPP and RAGE, display functional and pathologic evidence of neuronal perturbation at early times (3–4 months of age), before the steep rise in cerebral A β and plaque formation occur. At later times (14–18 months), activation of microglia/

astrocytes was significantly increased in double Tgs, compared with single Tg mAPP mice. Consistent with these data, experiments in which Tg mice bearing a dominant-negative form of RAGE (DN-RAGE) targeted to neurons were crossed with Tg mAPP animals (to generate Tg mAPP/DN-RAGE mice) showed preservation of spatial learning/memory and diminished neuropathologic changes compared with Tg mAPP mice. Our data support the possibility that blockade of neuronal RAGE might have cytoprotective effects, especially with respect to preserving neuronal function early in the disease process.

Results

Generation of Tg RAGE and Tg mAPP/RAGE mice

RAGE is a multiligand receptor which binds A β in the nanomolar range (apparent K_d 's of 50–100 nM; Yan *et al*, 1996). *In vitro*, interaction of ligands with RAGE-bearing neuronal-like cell lines induces cell stress, characterized by nuclear translocation of NF- κ B, and, eventually, loss of cell viability. In view of increased levels of RAGE in AD brain, including neurons (Yan *et al*, 1996; Lue *et al*, 2001), we sought to develop a model system in which neuronal expression of RAGE would be exaggerated so that consequences of receptor–ligand engagement could be assessed in an A β -rich environment. A transgene bearing full-length wt human RAGE driven by the platelet-derived growth factor (PDGF) B-chain promoter was constructed and used to generate Tg mice, termed Tg RAGE (or Tg wtRAGE), which were backcrossed into the C57BL/6 strain eight times. Southern blotting from two offspring of the same founder demonstrated the presence of the transgene (Supplementary Figure S1a, lanes 1 and 2). This founder was used to establish a line of Tg animals, which has shown normal gross development, reproductive fitness and lifespan (two other Tg RAGE founders were generated, and one of these was used to develop a second line of Tg RAGE mice). Further characterization of the line shown in panel S1a demonstrated increased expression of RAGE transcripts in total RNA harvested from the cerebral cortex of Tg animals compared with nonTg littermates (Supplementary Figure S1b, lanes 1–4). Western blotting of cortical homogenates confirmed the increase in RAGE expression observed in Tg mice, compared with nonTg littermates (Supplementary Figure S1c, lanes 1 and 2). Immunostaining of neocortex from a Tg mouse demonstrated enhanced expression of RAGE antigen, compared with nonTg littermates (Supplementary Figure S1d).

In order to test the effect of neuronal RAGE in an A β -rich environment, Tg RAGE mice were crossed with animals expressing a mutant form of human APP (Tg mAPP) also driven by PDGF B-chain promoter, and in the C57BL/6 background (Mucke *et al*, 2000). The Tg mAPP model was well suited to our strategy of determining whether over-expression of RAGE might enhance/accelerate pathologic changes in the brain, since these animals have been characterized with respect to neuropathologic, biochemical, and physiologic end points (Hsia *et al*, 1999; Mucke *et al*, 2000). The cross of Tg RAGE with Tg mAPP mice produced four genotypes, in the expected Mendelian ratios, single Tgs (Tg RAGE, Tg mAPP), double Tgs (Tg mAPP/RAGE), and nonTg littermate controls (Figure 1A).

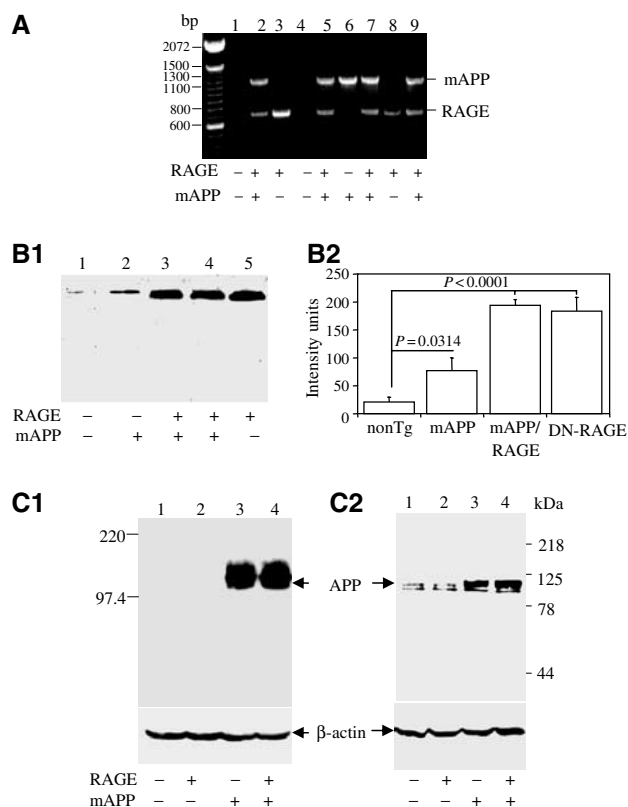


Figure 1 Identification and characterization of Tg mAPP/RAGE mice. (A) PCR analysis of tail DNA from one litter (+, RAGE or mAPP transgene present; -, transgene absent). (B) Western analysis of RAGE expression in the cerebral cortex of the four genotypes of mice. Homogenates of cerebral cortex (100 μ g) were subjected to SDS-PAGE (reduced; 10%), followed by immunoblotting with anti-RAGE IgG (6.5 μ g/ml) (B1). Densitometric analysis of gels is shown in (B2) (nonTg, $n=7$; Tg RAGE, $n=5$; Tg mAPP, $n=7$; Tg mAPP/RAGE, $n=4$). Results are means \pm s.e. (C) Western analysis of APP expression in the cerebral cortex of the four genotypes of mice. Homogenates of cortical protein were prepared as above and subjected to SDS-PAGE (reduced; 7.5%), followed by immunoblotting with either antibody 6E10 (C1; 1 μ g/ml) or 369W (C2; 1:2000 dilution).

Characterization of transgene expression in Tg mAPP/RAGE mice

Expression of RAGE in each of the four genotypes was studied by immunoblotting of homogenates of cerebral cortex with an antibody that recognized both transgene-derived human and endogenous murine RAGE (Figure 1B). Lowest levels of RAGE were observed in nonTg littermates (lane 1), whereas Tg mAPP animals displayed an increase in RAGE antigen (lane 2). This is consistent with enhanced RAGE expression observed in other situations in which the cellular microenvironment is enriched for RAGE ligands, such as AD brain and diabetic vasculature (Lue *et al*, 2001; Schmidt *et al*, 2001; Deane *et al*, 2003). Tg RAGE mice had higher levels of RAGE antigen (Figure 1B1, lane 5) than Tg mAPP animals (Figure 1B1, lane 2), and double Tgs had RAGE levels comparable to Tg RAGE animals (Figure 1B1, lane 4). Levels of RAGE antigen were \sim 3-fold greater in Tg mAPP/RAGE mice than in Tg mAPP mice. These results were obtained with animals at 3–4 months of age, and similar differences in RAGE expression were observed at 14–18 months (not shown).

Expression of the mAPP transgene was also studied in homogenates of cerebral cortex. Immunoblotting with an antibody selective for human APP revealed comparable expression of the transgene in Tg mAPP and Tg mAPP/RAGE (Figure 1C1, lanes 3–4). As expected, no material immunoreactive with murine monoclonal antibody 6E10 (targeted to human A β , residues 1–16) was observed in brain extracts from Tg RAGE or nonTg littermates (Figure 1C1, lanes 1–2). In order to assess the changes in total APP, both endogenous murine and Tg human APP, studies were performed with antibody 369W. Immunoblotting demonstrated similar lower levels of APP in nonTg littermates and Tg RAGE mice (Figure 1C2, lanes 1 and 2), and similar higher levels in Tg mAPP and Tg mAPP/RAGE (Figure 1C, lanes 3 and 4). These results were obtained using animals at 3–4 months of age, and levels of APP remained similar up to the oldest age group analyzed (14–18 months).

Cellular stress in Tg mAPP/RAGE mice: activation of NF- κ B and inflammation

NF- κ B activation has been identified in AD brain (Yan *et al*, 1996; Kaltschmidt *et al*, 1997). Based on the hypothesis that a

receptor for A β should magnify cell stress at early stages of A β accumulation, nuclear binding activity for NF- κ B was studied by gel shift analysis using nuclear extracts from cerebral cortex with a consensus ³²P-labeled NF- κ B probe. At 3–4 months of age, Tg mAPP/RAGE mice displayed a strong gel shift band compared with a virtually undetectable signal in nonTg littermates and single Tgs (Figure 2A1). The data shown in Figure 2A2 are representative of results obtained in mice and densitometry of autoradiograms from this larger group confirmed the striking increase in nuclear translocation of NF- κ B in double Tgs.

In view of the association of NF- κ B activation with inflammation, we considered the possibility that neurons in double Tg mice might incite an inflammatory response, even as early as 3–4 months. Thus, we compared microgliosis and astroglytosis in Tg mAPP/RAGE and Tg mAPP animals. Microglia were visualized immunohistochemically with antibodies to CD11b (complement receptor 3) or phosphotyrosine, and astrocytes were visualized with antibody to glial fibrillary acidic protein (GFAP). The results of image analysis of these sections are shown in the histogram in Figure 2, panel B, and representative micrographs are shown in panels c–f (animals

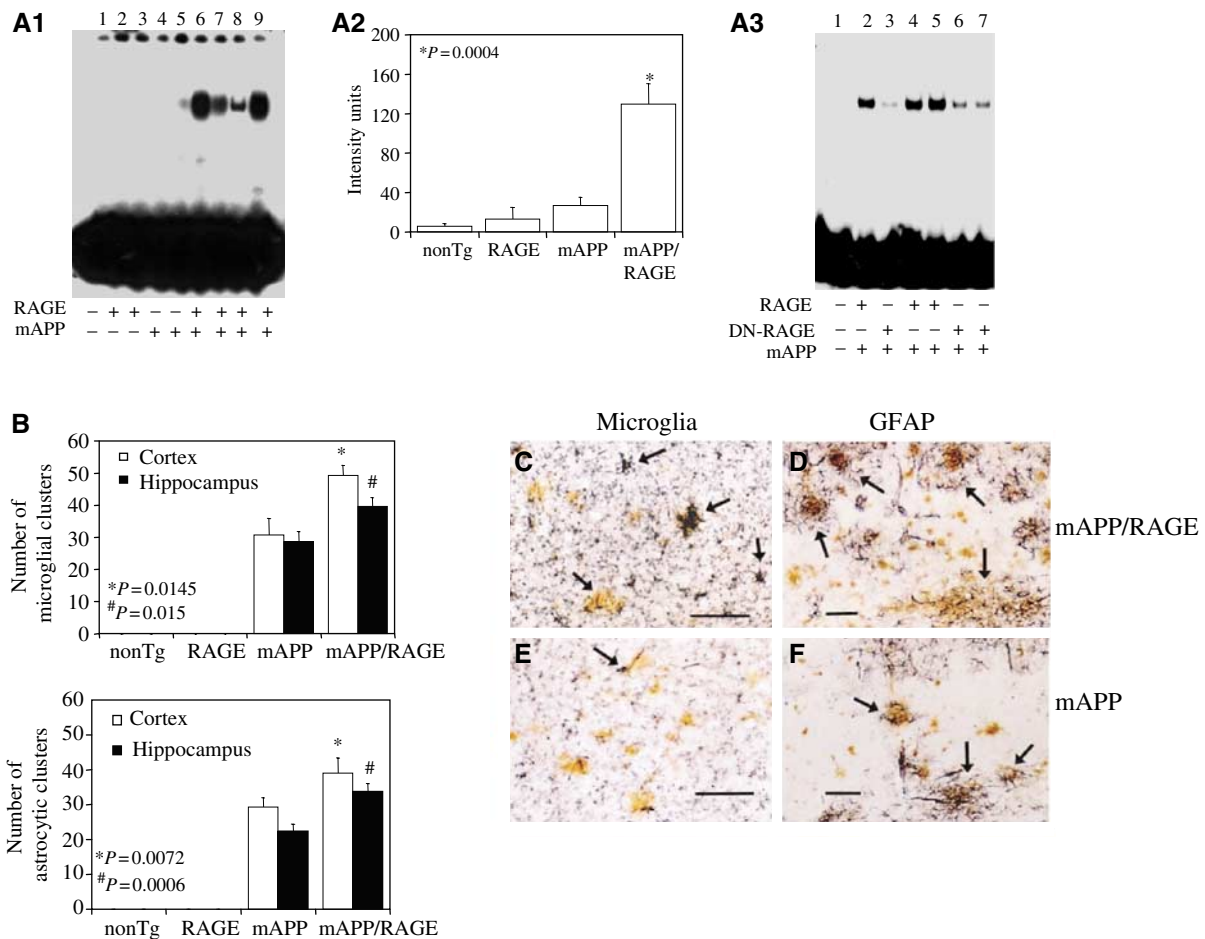


Figure 2 Activation of NF- κ B and inflammation in Tg mAPP/RAGE mice. (A) Nuclear translocation of NF- κ B was assessed in nuclear extracts from cerebral cortex (10 μ g total protein/lane) of mice (age 3–4 months) with ³²P-labeled consensus NF- κ B probe (A1). The adjacent panel displays image analysis of results from a larger group of mice (4–6 mice/genotype) (A2). Panel A3 displays results of NF- κ B gel shift experiments with Tg mAPP/RAGE, Tg mAPP/DN-RAGE, and nonTg littermates. Lane 1 shows migration of free probe. (B) Microgliosis and astroglytosis in brains of Tg mAPP/RAGE mice compared with the other groups. Results of image analysis are shown at 14–18 months in the cerebral cortex and hippocampus (4–6 mice/genotype). (C–F) Representative sections from Tg mAPP/RAGE (C, D) and Tg mAPP (E, F) mice at age 14–18 months to demonstrate microglia (C, E) and astrocytes (D, F). The arrows denote cells reactive with the indicated markers. Scale bar = 10 μ m. Multiple images similar to those in (C–F) were used to construct the histogram shown in panel (B).

were 14–18 months). Analysis of Tg mAPP/RAGE mice at 3–4 and 7–8 months of age did not reveal significantly increased microgliosis or astrocytosis compared with Tg mAPP, Tg RAGE or nonTg littermates (not shown). However, at 14–18 months, Tg mAPP/RAGE mice had significantly more plaque-associated reactive microglia and astrocytes ($P \leq 0.015$; Figure 2B). It is possible that the latter reflect an exaggerated secondary inflammatory response to deposited A β .

Perturbation of neuronal function in Tg mAPP/RAGE mice

In view of accentuated cell stress in brains of Tg mAPP/RAGE mice, as reflected by NF- κ B activation at early times, we sought to determine whether this would be reflected in impaired spatial learning/memory in the radial arm water maze (Diamond *et al*, 1999). At 3–4 months of age, nonTg littermates showed strong learning and memory capacity (Figure 3A). Similar results were obtained with Tg RAGE,

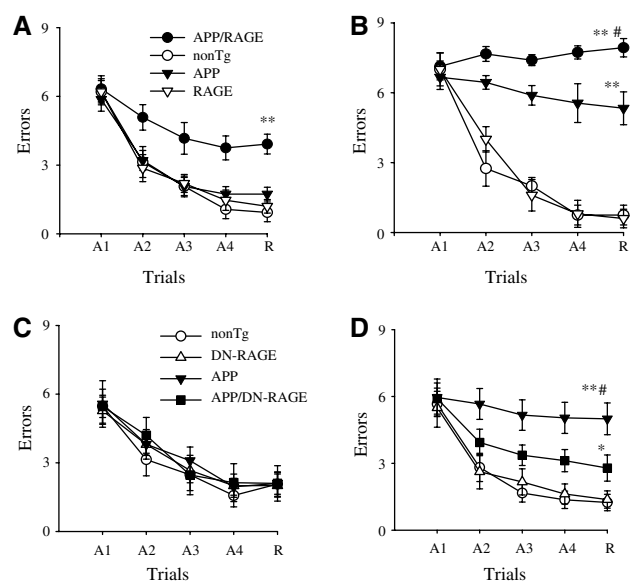


Figure 3 Functional neuronal deficits: spatial learning and memory in Tg mAPP/RAGE and Tg mAPP/DN-RAGE mice. (A, B) Spatial learning and memory was tested in the radial arm water maze at 3–4 (A) and 5–6 months of age (B) in mice of the indicated genotypes (in (A), $n = 7$ –8 male mice/genotype; in (B), $n = 5$ male mice/genotype): Tg mAPP/RAGE (APP/RAGE), nonTg littermate (nonTg), Tg mAPP (APP), and Tg RAGE (RAGE). (A1–A4) denote the acquisition trials, and R denotes the retention trial. In panel A, $**P < 0.01$ Tg mAPP/RAGE compared with nonTg mice (by repeated-measure ANOVA followed by Fisher's protected least significant difference for *post hoc* comparisons in this and the following graphs). In panel B, $**P < 0.01$ Tg mAPP/RAGE and Tg mAPP mice compared with nonTg mice; $*P < 0.01$ Tg mAPP/RAGE compared with Tg mAPP mice. ANOVA revealed a main age effect in Tg mAPP/RAGE mice and Tg mAPP mice ($P < 0.05$ for both), but not in nonTg and DN-RAGE mice ($P > 0.05$ for both). (C, D) Effect of DN-RAGE transgene on spatial learning and memory in Tg mice at 3–4 months ($n = 5$ –8 male mice/genotype (C)) and 5–6 months ($n = 6$ –9 male mice/genotype (D)). The following genotypes were tested: nonTg littermate (nonTg), Tg DN-RAGE (DN-RAGE), Tg mAPP (APP), and Tg mAPP/DN-RAGE (APP/DN-RAGE). In panel D, $**P < 0.01$ Tg mAPP mice compared with nonTg mice; $*P < 0.01$ Tg mAPP mice compared with Tg mAPP/DN-RAGE mice, and $*P < 0.05$ Tg mAPP/DN-RAGE mice compared with nonTg littermates. ANOVA revealed a main age effect in Tg mAPP mice ($P < 0.05$) but not in Tg mAPP/DN-RAGE, DN-RAGE, and nonTg mice ($P > 0.05$).

as well as Tg mAPP mice (Figure 3A). In contrast, Tg mAPP/RAGE mice averaged about four errors by trials 4 or 5, indicative of impaired spatial memory for platform location between trials, as well as during the 30 min delay before trial 5 (Figure 3A). To determine whether combined overexpression of RAGE and mAPP impaired vision, motor coordination or motivation, we also tested mice with a visible platform task. The four genotypes showed no difference in their speed of swimming (Supplementary Figure S2a1) or in the time required to reach the platform in this version of the task (Supplementary Figure S2a2). At 5–6 months, the deficit in learning/memory in Tg mAPP/RAGE mice was worse (≈ 8 errors by trial 4 or 5) than in younger double Tg animals (Figure 3B). A deficit, though less severe, was also observed in age-matched Tg mAPP mice (≈ 6 errors by trials 4 or 5), whereas single Tg RAGE mice and nonTg littermates showed intact spatial memory (Figure 3B). Spatial memory impairment persisted in older double Tg animals (12–13 months; Supplementary Figure S3); their performance was worse than single Tg mAPP mice, Tg RAGE mice, and nonTg littermates.

These results indicate an early perturbation of learning and memory, consequent to overexpression of RAGE and mAPP. Since cognitive abnormalities in AD are thought to be linked to synaptic dysfunction (Selkoe, 2002), we examined synaptic transmission under basal conditions and during long-term potentiation (LTP). Consistent with previous observations (Hsia *et al*, 1999), field-excitatory post-synaptic potential (fEPSPs) in the CA1 stratum radiatum revealed impaired basal synaptic transmission (BST) in young (3–5 months) double Tg mice ($\approx 24\%$ decrease in slope of the input–output curve) and Tg mAPP littermates ($\approx 26\%$ decrease) (Figure 4A). BST was further reduced in aged (8–10 months) Tg mAPP/RAGE (64% decrease) and Tg mAPP animals ($\approx 58\%$ decrease) (Figure 4A). Both younger and older Tg mice overexpressing RAGE alone did not show any significant difference in BST, compared with nonTg littermates (Figure 4A). CA1 LTP was normal in young (3–5 months) Tg mice (all genotypes) and nonTg littermates (Figure 4B). However, LTP was most severely affected in 8–10-month-old double Tg mice compared with the other groups (Figure 4C). At 60 min after LTP induction, we observed a $140 \pm 9\%$ increase in potentiation in Tg mAPP/RAGE mice, whereas nonTg littermates displayed a $202 \pm 13\%$ increase. Single Tg RAGE and Tg mAPP showed levels of LTP comparable to nonTg animals (188 ± 20 and $203 \pm 17\%$, respectively). These findings are consistent with the hypothesis that RAGE-A β interaction alters synaptic plasticity. In addition, since BST is affected both in single Tg mAPP and double Tg mice, but LTP is reduced only in the double Tgs, we infer that overexpression of RAGE in an A β -rich environment further perturbs synaptic function in the older mice, impairing plasticity.

To further evaluate the cellular basis of the apparent learning/memory deficit, we tested the concept that impaired plasticity in older (14–18 months) Tg mAPP/RAGE mice might be due to insufficient release and/or entry of Ca $^{2+}$ during the tetanus. CA1-LTP appears to be due to Ca $^{2+}$ entrance through postsynaptic NMDA receptors following glutamate release during high-frequency stimulation of the presynaptic cell. Thus, an increase of the tetanus strength should release more neurotransmitter, leading to an increase in the amount of Ca $^{2+}$ conveyed through NMDA receptors producing normal potentiation. Similar manipulations have

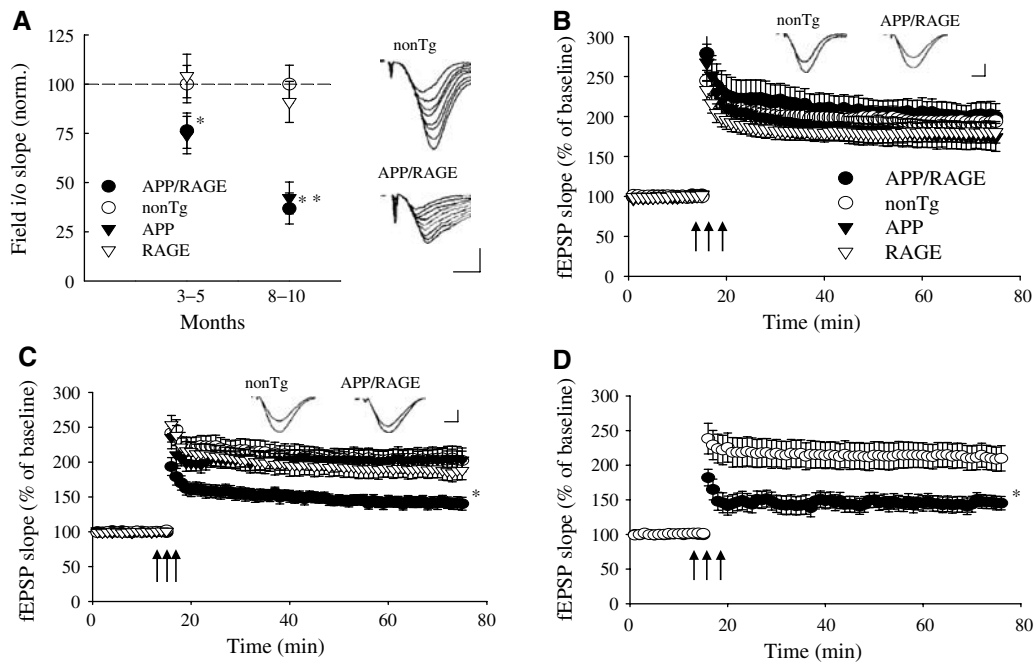


Figure 4 Functional neuronal deficits: decreased LTP in Tg mAPP/RAGE mice. **(A)** BST was reduced in the CA1 region of slices from 3–5-month-old Tg mAPP/RAGE (APP/RAGE) mice ($*P < 0.05$) and Tg mAPP (APP) littermates compared with nonTg littermates (male animals were used throughout in electrophysiologic studies). A greater reduction in BST was observed in 10-month-old Tg mAPP/RAGE mice and Tg mAPP littermates ($**P < 0.01$). Single Tg RAGE (RAGE) mice showed normal values of BST at both ages. The responsiveness of CA1 cells to increasing afferent fiber stimulation (slope of input–output (i/o) relation) was measured to assess BST strength. Results from 3–5-month-old animals were based on recordings in 16 slices from five male Tg mAPP/RAGE mice, 19 slices from six Tg mAPP mice, 16 slices from six Tg RAGE mice, and 19 slices from seven nonTg mice. Results from 8–10-month-old mice were based on recordings in 12 slices from five Tg mAPP/RAGE mice; 10 slices from four Tg mAPP mice; 15 slices from six Tg RAGE; and 12 slices from six nonTg mice. Insets show representative fEPSP for a nonTg mouse and a Tg mAPP/RAGE mouse, illustrating that far higher stimulation strengths are required to elicit synaptic responses in 10-month-old Tg mAPP/RAGE mice. Scale: 0.2 mV; 5 ms. **(B)** LTP was normal in the CA1 region of slices from 3–5-month-old Tg mice (all genotypes) and their nonTg littermates. Theta-burst stimulation indicated by the three arrows was delivered after recording a 15-min baseline ($n = 16$ slices from five Tg mAPP/RAGE mice, $n = 19$ slices from six male Tg mAPP mice, $n = 16$ slices from six male Tg RAGE mice, and $n = 19$ slices from seven male nonTg mice). Insets show traces taken 1 min before and 60 min after LTP induction on 3–5-month-old nonTg and Tg mAPP/RAGE mice. Scale: 0.1 mV (nonTg), 0.05 mV (Tg mAPP/RAGE mice); 2.5 ms. **(C)** LTP was reduced in the CA1 region of slices from 8–10-month-old double Tg mAPP/RAGE mice, whereas Tg mAPP, Tg RAGE, and nonTg mice showed normal potentiation ($n = 12$ slices from five Tg mAPP/RAGE mice; $n = 10$ slices from four Tg mAPP mice; $n = 15$ slices from six Tg RAGE mice; and $n = 12$ slices from six nonTg mice; $*P < 0.05$). Insets show traces taken 1 min before and 60 min after LTP induction on 8–10-month-old nonTg and Tg mAPP/RAGE mice. Scale: 0.1 mV (nonTg), 0.03 mV (Tg mAPP/RAGE mice); 2.5 ms. **(D)** LTP was still reduced in double Tg mAPP/RAGE mice after enhancement of the theta-burst strength ($n = 7$ slices from four Tg mAPP/RAGE mice; and $n = 8$ slices from four nonTg mice; $*P < 0.05$).

previously been used to address this issue (Usdin *et al*, 1999). Slices from 8–10-month-old Tg mAPP/RAGE mice were subjected to enhanced tetanus consisting of a double number of bursts (see Materials and methods). Under these conditions, the amount of LTP was not increased (Tg mAPP/RAGE, $145 \pm 9\%$; nonTg, $209 \pm 18\%$; Figure 4D). These results suggest that impaired LTP in Tg mAPP/RAGE mice was not likely due to insufficient release and/or entry of Ca^{2+} , but, rather, due to dysfunction at steps occurring after Ca^{2+} entrance.

Neuropathologic changes in Tg mAPP/RAGE mice

Diminished density of cholinergic fibers and synapses is associated with AD-like pathology (Masliah *et al*, 1989; Geula, 1998). We sought to determine if such changes might occur in Tg mAPP/RAGE mice by semiquantitatively analyzing acetylcholinesterase (AChE) activity-positive neurites (which predominately identifies cholinergic neurites), using a histochemical method, and synaptophysin (a marker of presynaptic terminals) by immunostaining. Based on the results of the above behavioral studies, we predicted that changes in these neuropathologic properties should be

detectable by 3–4 months of age in the double Tgs, and should be progressive. AChE activity was determined histochemically and reported as the area occupied by positively staining neurites in the subiculum (Sb), CA1 and entorhinal cortex (EC). At the 3–4-month time point, the area occupied by AChE-positive neurites in the subiculum was decreased in Tg mAPP/RAGE mice ($P < 0.001$) compared with single Tgs and nonTg littermates (Figure 5A1). This decrease in AChE-positive fibers in double Tgs appeared to be progressive at the 14–18-month time point (Figure 5A2). At this older age, there was also a statistically significant decrease in AChE-positive fibers in Tg mAPP mice, though not in Tg RAGE or nonTg littermates, indicating that loss of cholinergic fibers was not simply a function of age. A representative example of AChE histochemical-stained sections of subiculum from each genotype of mice used for image analysis is shown in Figure 5B. Similar results, with respect to the density of AChE neurites, were obtained in the EC and CA1 (not shown).

We also assessed the density of presynaptic terminals by determining synaptophysin immunoreactivity. Analysis of presynaptic terminals of defined signal intensity, which

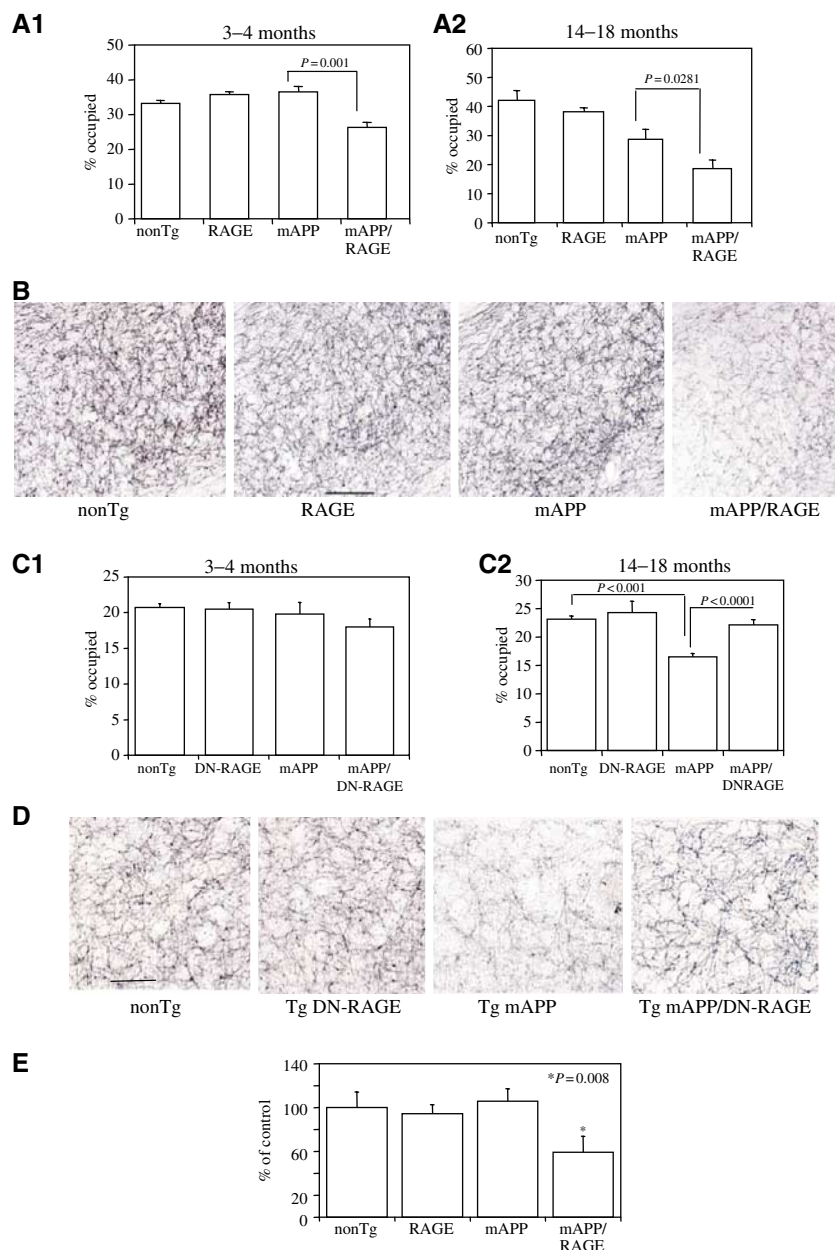


Figure 5 Characterization of Tg mAPP/RAGE and Tg mAPP/DN-RAGE mice: effect on AChE activity. (**A1**, **A2**) AChE activity was determined histochemically in the subiculum of mice of the indicated genotype at 3–4 (**A1**; $n = 4$ –9/group) and 14–18 months (**A2**; $n = 5$ –12/group) of age: nonTg littermate (nonTg), Tg RAGE (RAGE), Tg mAPP (mAPP), and Tg mAPP/RAGE (mAPP/RAGE). The bar graph shows the results of image analysis. (**B**) Representative images from AChE histochemical staining from the subiculum of mice of the indicated genotype from the experiment in **A1**. Scale bar = 10 μ m. (**C1**, **C2**) AChE activity was determined as above in the indicated mice (nonTg, Tg DN-RAGE (DN-RAGE), Tg mAPP (mAPP), and Tg mAPP/DN-RAGE (mAPP/DN-RAGE)) at 3–4 months (**C1**; $n = 7$ –12/group) and 14–18 months of age (**C2**; $n = 7$ –12/group). (**D**) Representative images of AChE staining for the indicated mice from the experiment in **C2**. Scale bar = 10 μ m. (**E**) Area occupied by synaptophysin immunoreactivity in the stratum lacunosum moleculare of the hippocampus in mice of the indicated genotype at 3–4 months of age (mean \pm s.e. are shown).

reacted with anti-synaptophysin antibody in the stratum lacunosum moleculare of the hippocampus, demonstrated a decrease in the area occupied by these structures in samples from Tg mAPP/RAGE mice, compared with single Tgs (Tg mAPP, Tg RAGE) and nonTg littermates, at 3–4 months of age (Figure 5E). Taken together, these data indicate that neuropathologic changes accompany altered neuronal function at both early and later time points, and appear to be accelerated in Tg mAPP/RAGE mice.

Analysis of Tg mAPP/DN-RAGE mice

In view of subtle behavioral and neuropathologic changes in Tg mAPP mice, we favored starting our analysis of the role of RAGE in A β -mediated neuronal stress by overexpressing the receptor and determining if pathologic changes would become more exaggerated. Next, we sought to determine whether RAGE was functioning solely as a cell surface-binding site for A β , possibly increasing the local levels of A β to interact with other sites, or whether it had a more direct role

in transducing a signal important for A β -induced cellular perturbation. For these studies, we employed Tg mice overexpressing a DN-RAGE also under control of the PDGF B-chain promoter (Tg DN-RAGE) backcrossed seven times into the C57BL/6 strain. DN-RAGE comprises a truncated form of the receptor with intact extracellular and membrane-spanning portions, but a deleted cytosolic tail. In previous studies, we have found that expression of DN-RAGE, even in cells expressing wt RAGE, blocks RAGE-dependent cellular activation (Hofmann *et al*, 1999). Tg DN-RAGE animals did not manifest abnormalities with respect to reproductive fitness, development, basic neurologic function or longevity. Southern analysis of tail DNA from two offspring derived from one founder demonstrated the presence of the transgene (Supplementary Figure S1a, lanes 3 and 4). The level of expression of DN-RAGE, at the mRNA (Supplementary Figure S1b, lanes 6 and 8) and antigen levels (Supplementary Figure S1c (lane 3) and d (DN-RAGE)), appeared to be similar to that of the wtRAGE transgene in the line of Tg RAGE mice used in the above studies (see Supplementary Figure S1). DN-RAGE antigen is seen to migrate more rapidly on SDS-PAGE than wtRAGE (Supplementary Figure S1c), in view of deletion of the cytosolic tail in the dominant-negative variant. Tg DN-RAGE mice were crossed with Tg mAPP animals to produce Tg mAPP/DN-RAGE animals, as well as single Tgs (Tg mAPP, Tg DN-RAGE) and nonTg littermates.

First, activation of NF- κ B was analyzed in nuclear extracts of cerebral cortex of animals expressing the DN-RAGE transgene at 3–4 months of age (Figure 2A3). In the electrophoretic mobility shift assay for NF- κ B, Tg mAPP/DN-RAGE animals (Figure 2A3, lanes 3, 6, and 7) displayed a striking reduction in intensity of the gel shift band compared with Tg mAPP/RAGE mice (Figure 2A3, lanes 2, 4, and 5). These data were consistent with decreased evidence of cell stress in brains of Tg mAPP/DN-RAGE mice compared with Tg mAPP/RAGE mice, and led us to analyze the effect of introducing the DN-RAGE transgene on spatial memory/learning (Figure 3C and D). At 3–4 months of age, Tg mAPP/DN-RAGE, Tg DN-RAGE, Tg mAPP, and nonTg littermates all showed strong learning and memory capacity (they averaged about two errors by trials 4 or 5; Figure 3C). However, at 5–6 months of age, when Tg mAPP showed a significant deficit in spatial memory (about five errors by trials 4 and 5), Tg mAPP/DN-RAGE animals made fewer errors in the same trials (about three errors by trials 4 and 5) (Figure 3D). All groups performed similarly during the visible platform trials with respect to swim speed and time required to reach platform (not shown). Thus, compared with Tg mAPP/RAGE animals, there was preservation of spatial learning/memory at the 3–4-month time point in Tg mAPP/DN-RAGE mice. Furthermore, at 5–6 months of age, neurologic function in Tg mAPP/DN-RAGE mice was better maintained compared with Tg mAPP. Neuropathologic changes, as evaluated by the area occupied by AChE-positive fibers in the subiculum (Figure 5C and D), were virtually completely prevented in Tg mAPP/DN-RAGE mice, compared with Tg mAPP animals, at both 3–4 (Figure 6C1) and 14–18 months of age (Figure 5C2). Representative micrographs of AChE histochemical-stained sections are shown in Figure 5D. In contrast, there was accelerated loss of AChE-positive fibers in Tg mAPP/RAGE mice compared with Tg mAPP animals at these

same time points (Figure 5A1–2). Taken together, these data are consistent with the concept that RAGE functions as a signal transduction receptor mediating neuronal dysfunction in the A β -rich environment provided by the mAPP transgene.

Markers of synaptic plasticity in Tg mAPP/RAGE mice

In view of these findings, we explored the possible mechanisms underlying accelerated neuronal dysfunction in Tg mAPP/RAGE mice and apparent protection in Tg mAPP/DN-RAGE animals, compared with findings in Tg mAPP mice, at the level of markers of synaptic plasticity. We therefore analyzed phosphorylation of CREB and activation of mitogen-activated protein kinases (MAP kinases) and calcium/calmodulin-dependent protein kinase II (CAMKII) in hippocampal extracts. Immunoblotting with antibody to total CREB showed no differences in 3–4-month-old mice from each of the genotypes (Figure 6A, lower panel). However, labeling of the same blot with anti-phospho-CREB (Ser-133) antibody revealed significantly greater levels of phospho-CREB in Tg mAPP/RAGE mice, than in single Tgs (Tg RAGE, Tg mAPP) and nonTg littermates (Figure 6A, lower panel). Furthermore, intensity of the phospho-CREB band was reduced to the background in Tg mAPP/DN-RAGE mice (Figure 6A, lower panel). Densitometric analysis of data from a larger group of animals ($n = 5\text{--}8/\text{group}$) demonstrated the reproducibility of these observations (Figure 6A, upper panel).

Activation of MAP kinases is closely associated with synaptic plasticity (especially extracellular signal-regulated kinases, Erk1/2) and cell stress (p38 and jnk) (Kyriakis and Avruch, 2001; Sweatt, 2001). Furthermore, in previous studies, we have found that activation of MAP kinases occurs in RAGE-bearing tumor cells exposed to ligands of the receptor *in vitro* (Taguchi *et al*, 2000). Here, we focused our attention on animals at 3–4 months of age, because of impaired spatial learning/memory at this time, using antibodies to phosphorylated forms of MAP kinases and hippocampal extracts to estimate activation of the related kinases. Phosphorylation of p38 occurred selectively in Tg mAPP/RAGE mice (Figure 6B, lower panel; there was no change in the level of total p38), compared with the other groups, with ≈ 3.5 -fold increased intensity of the immunoreactive band ($P = 0.0016$; Figure 6B, upper panel). As in the experiments analyzing phospho-CREB (above), levels of phospho-p38 were reduced to the background in Tg mAPP/DN-RAGE mice, compared with Tg mAPP/RAGE animals (Figure 6B). In terms of Erk1/2, the phosphorylated forms of Erk1/2 (p44/42) were also increased in Tg mAPP/RAGE mice, compared with other groups (and diminished in Tg mAPP/DN-RAGE mice) (Figure 6C, lower panel). Densitometric analysis of gels (focusing on phospho-Erk1) confirmed this impression and indicated that it achieved statistical significance based on analysis of samples from 5–8 animals/group (Figure 6C, upper panel). However, p-Erk1/2 was also increased in Tg mAPP (though this did not achieve statistical significance), and there was a somewhat more variable inter-animal baseline level of p-Erk in nonTg controls. In contrast, similar studies with anti-phospho-JNK IgG showed no change in p-JNK immunoreactive material in samples from the Tg mAPP/RAGE animals, compared with Tg RAGE, Tg mAPP, and nonTg littermates (data not shown).

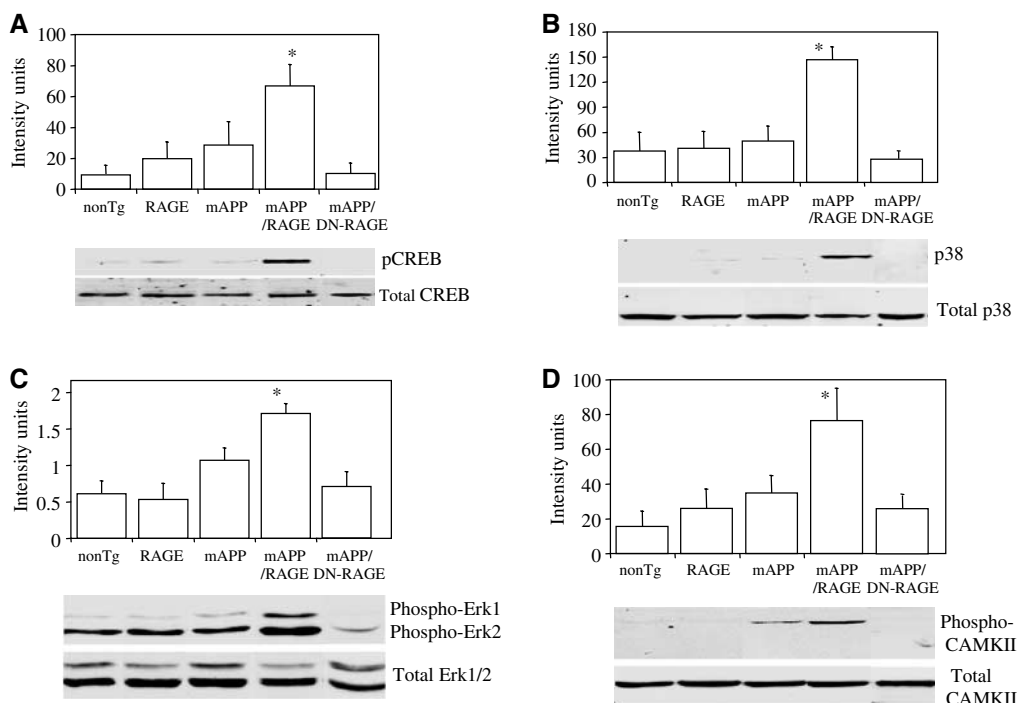


Figure 6 Phosphorylation/activation of CREB (A), MAP kinases (B, C), and CAMKII (D) in Tg mice. (A) Protein extracts were obtained from hippocampii of mice of the indicated genotypes (Tg mAPP, Tg (wt)RAGE, Tg mAPP/RAGE, Tg mAPP/DN-RAGE, nonTg) at 3–4 months of age and subjected to immunoblotting with antibody to total CREB or phospho (p)-CREB. The upper panel displays quantitation of the results with phospho-CREB by image analysis ($n = 5–8$ /group) and the lower panel shows representative immunoblots. (B, C) The experiment was performed as in (A), except that antibodies for immunoblotting were anti-total p38 and anti-phospho-p38 IgG (B), and anti-total Erk1/2 and anti-phospho-Erk1/2 IgG (C). The upper panel displays image analysis of results with the phospho antibodies and $n = 5–8$ /group (in the case of Erk, image analysis was performed on p44). In panel D, the experiment was performed with antibodies to total and phospho-CAMKII (CAMKII) and the animals were 14–18 months of age (image analysis shows results of phospho-CAMKII from 5–13 animals/group).

CAMKII is an important contributor to synaptic plasticity, and, shortly after its activation, becomes autophosphorylated at threonine 286 (Miller and Kennedy, 1986). Using an antibody specific for phospho-CAMKII, levels of the phosphorylated form were found to be increased in hippocampal extracts from Tg mAPP/RAGE mice compared with Tg mAPP animals at 14–18 months of age (Figure 6D, lower panel), whereas they were unchanged at the earlier time point, 3–4 months (not shown). Densitometric analysis (Figure 6D, upper panel) of autoradiograms from a larger group of mice ($n = 5–13$ animals/group) confirmed this impression, and demonstrated that the p-CAMKII band was $\approx 2–3$ -fold more intense in the double Tg mAPP/RAGE mice than in Tg mAPP animals ($P = 0.014$). In contrast, p-CAMKII remained at background levels in Tg mAPP/DN-RAGE animals. Total CAMK II antigen was similar in each of the groups of mice (Figure 6D, lower panel).

Discussion

Previous studies have linked RAGE to evolving immune/inflammatory responses, ranging from colitis to atherosclerosis (Schmidt *et al*, 2001). *In vitro*, interaction of A β with RAGE-bearing neuronal-like cells leads to activation of NF- κ B and expression of macrophage-colony stimulating factor and interleukin-6, suggesting a possible role for neuronal RAGE in inflammation (Yan *et al*, 1997; Wang *et al*, 2000). However, despite strong activation of NF- κ B in Tg mAPP/RAGE mice by 3–4 months of age, evidence of neuroinflammation, based on

microgliosis and astrocytosis, was not significantly increased until much later, 14–18 months. In contrast, more subtle indications of neuronal dysfunction (impaired spatial learning/memory) were the earliest detectable functional defects in Tg mAPP/RAGE mice, and occurred at early times (3–4 months), when overall A β levels are low and there are few plaques (Mucke *et al*, 2000).

There were several findings temporally associated with altered spatial learning/memory in young Tg mAPP/RAGE mice. These findings included decreased area occupied by neurites positive for AChE activity in the subiculum, as well as EC and CA1; decreased area occupied by synaptophysin-positive presynaptic terminals in the striatum lacunosum moleculare of the hippocampus; and altered activation of markers of synaptic plasticity. In each case, our data are consistent with the concept that increased cellular RAGE renders neurons more vulnerable to A β -induced dysfunction. Furthermore, decreased density of presynaptic terminals (synaptophysin immunoreactivity) and cholinergic fibers (the predominant population of AChE-positive neurites) are consistent with pathologic findings in AD. Increased phosphorylation of CREB, CAMKII and the MAP kinases (p38, Erk1/2) might not have been expected to be associated with decreased hippocampal function (spatial learning/memory). However, strongly increased basal activation of these kinases and CREB in Tg mAPP/RAGE mice probably has very different functional consequences than transient activation in wt animals associated with stimuli that accompany physiologic processes (such as LTP or fear conditioning) (Kyriakis and

Avruch, 2001; Sweatt, 2001). The capacity of RAGE to activate multiple signaling pathways (in addition to those analyzed in the current experiments), including recruitment of NADPH oxidase, p21^{ras} (Wautier *et al*, 2001), phosphatidylinositol 3-kinase, Rho GTPases (cdc42 and rac), and the Jak/stat pathway (Miller and Kennedy, 1986; Lander *et al*, 1997; Deora *et al*, 1998; Huttunen *et al*, 1999; Taguchi *et al*, 2000; Huang *et al*, 2001), indicates that receptor-mediated neuronal stimulation could exert effects on diverse cellular functions.

The role of RAGE as a signal transduction receptor mediating the effects of an A β -rich environment is highlighted by our experiments in Tg mAPP/DN-RAGE mice. Although the latter animals expressed about the same levels of RAGE as Tg mAPP/(wt)RAGE animals and the receptor encoded by the DN-RAGE transgene is fully competent with respect to ligand binding, signal transduction from the truncated DN-RAGE variant is blocked (Yan *et al*, 2000). Tg mAPP/DN-RAGE mice displayed decreased activation of NF- κ B and apparent preservation of spatial learning/memory at early times. In addition, there was apparent sparing of AChE-positive neuronal fibers in the subiculum of Tg mAPP/DN-RAGE mice, compared with Tg mAPP animals. These data provide strong support for the concept that RAGE functions as a signaling receptor, rather than solely as a site tethering ligands to the cell surface. Detailed mechanisms linking occupancy of RAGE to NF- κ B activation, diminished neuronal/synaptic function, and neuropathologic changes remain to be elucidated. However, in previous studies, we have found that ligand engagement of RAGE results in activation of MAP kinases and NF- κ B. RAGE-dependent activation of MAP kinases appears to proceed via an oxidant-sensitive mechanism involving p21^{ras} (Lander *et al*, 1997). Links between I κ B phosphorylation and activation of protein kinase A suggest a further mechanism potentially joining the NF- κ B and the cAMP pathways, the latter closely linked to synaptic plasticity (Zhong *et al*, 1997).

The results of our studies add to a growing body of evidence in murine models, suggesting that even before the amyloid deposition in the brain A β can disrupt neuronal

integrity and functions, including synaptic plasticity (Hsia *et al*, 1999; Buttini *et al*, 2002). The current experiments demonstrate that RAGE can act as a cofactor critically magnifying the effects of A β on neuronal function.

Materials and methods

Generation of Tg mice and characterization of transgene expression

Selective overexpression of human RAGE in neurons was achieved using the PDGF B chain promoter, in view of its previous success in driving overexpression of mAPP (Mucke *et al*, 2000). A transgene driving expression of human RAGE or DN-RAGE was similarly placed under control of PDGF B chain promoter. Further description of these Tg mice is provided in Supplementary data.

Immunocytochemical, histochemical and immunoblotting analyses are described in Supplementary data.

NF- κ B activation in Tg mice

For gel shift analysis, nuclear extracts (total protein, 10 μ g/lane) prepared from mouse cerebral cortex were incubated with ³²P-labeled consensus NF- κ B probe and subjected to PAGE (5%) as described (Yan *et al*, 2000).

Behavioral and electrophysiological studies

Behavioral testing and electrophysiological studies were performed as described (Petroni *et al*, 2003; Trinchese *et al*, 2004). Details of these experimental methods are described in Supplementary data.

Statistical analysis

Statistical analyses using STATVIEW were performed with a two-way ANOVA for repeated measure, followed by Fisher's protected least significant difference for *post hoc* comparisons. Results were expressed as mean \pm standard error mean (s.e.m.).

Supplementary data

Supplementary data are available at *The EMBO Journal* Online.

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