# Inflammatory Cyclooxygenase Activity and PGE<sub>2</sub> Signaling in Models of Alzheimer's Disease

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**Abstract:** The inflammatory response is a fundamental driving force in the pathogenesis of Alzheimer's disease (AD). In the setting of accumulating immunogenic Aß peptide assemblies, microglia, the innate immune cells of the brain generate a non resolving immune response and foil to adequately clear

immune cells of the brain, generate a non-resolving immune response and fail to adequately clear accumulating AB peptides, accelerating neuronal and synaptic injury. Pathological, biomarker, and imaging studies point to a prominent role of the innate immune response in AD development, and the molecular components of this response are beginning to be unraveled. The inflammatory cyclooxygenase-PGE<sub>2</sub> pathway is implicated in pre-clinical development of AD, both in epidemiology of normal aging populations and in transgenic mouse models of Familial AD. The cyclooxygenase-PGE<sub>2</sub> pathway modulates the inflammatory response to accumulating AB peptides through actions of specific E-prostanoid G-protein coupled receptors.

**Keywords:** Alzheimer's disease, microglia, amyloid beta, inflammation, cyclooxgenases, prostaglandin E<sub>2</sub>, EP2 receptor, EP3 receptor, EP4 receptor.

### **INTRODUCTION**

Currently, 1/9 people older than 65 years of age have Alzheimer's disease (AD), and 1/3 of people older than 85 are diagnosed with AD. As the aging population expands, the projected number of AD cases is expected to triple by 2050 [1], with economic costs escalating from \$214 billion in 2014 to \$1.2 trillion in 2050 (Alzheimer's Association Facts 2014). This represents an enormous economic and societal challenge, particularly given the limited efficacy of currently available AD therapeutics. In terms of AD prevention, attention to diet, exercise, and cardiovascular health are likely to help reduce the risk of developing AD. However, identification of specific molecular pathways that could be targeted for prevention in aging and at-risk populations is an urgently needed strategy to help stem this dementia epidemic. Given that the prevalence of AD doubles every 5 years in persons above the age of 65 and given the average human longevity of ~80 years, preventive strategies that could delay the onset of cognitive decline by only 5 years may reduce disease burden by half. Recent data indicate that cyclooxygenase/PGE2/EP receptor signaling may play important roles in preclinical development of AD in both human epidemiology and in mouse models of AD.

### NEUROINFLAMMATION IN AD

Pathological changes in AD consist of amyloid  $\beta$  accumulation, tau phosphorylation, and synaptic and

neuronal loss. These pathological hallmarks develop in the context of a potent and chronic inflammatory response characterized by glial activation, generation of cytokines and chemokines, complement proteins, inflammatory enzymes, and oxidative stress [2, 3]. This chronic inflammatory response is not only injurious to synapses, neurons, and circuits, but is persistent and non-resolving.

Recent studies indicate that the pre-clinical development of AD begins years to decades before initial diagnosis [4]. While the initiating pathological events are not well defined, they are likely to involve synergistic interactions between Aß oligomer-mediated synaptic injury and dysregulated inflammatory responses. Amyloid ß peptide generation and accumulation is an initiating event, and precedes onset of symptoms by years to decades. However, amyloid PET imaging studies suggest that AB peptide accumulation can occur in subjects who do not have evidence of cognitive decline, leading to the hypothesis that AB peptide accumulation is "necessary but not sufficient" for progression to AD [5]. One or more additional factors may be necessary. In that regard, recent GWAS studies of sporadic late-onset AD have identified genes involved in the innate immune response that are expressed in microglia, the resident myeloid cells of the central nervous system (CNS). These microglial genes include the sialic acid-binding immunoglobulin-like lectin CD33 [6-8] and TREM2 [9, 10] which regulate phagocytosis and Aß peptide clearance [11-13], and the complement receptor CR1 [14, 15]. These findings indicate that the microglial immune response may play a pathogenic role in the development of AD.

Microglia have embryonic origins [16, 17] and genetic signatures [18-20] that distinguish them from monocytes and

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other tissue macrophages. Like other tissue macrophages, microglia maintain local homeostasis by clearing toxic proteins and noxious substances and regulating inflammation. However, because microglia reside in the brain, they have additional and unique functions, including the establishment, maintenance, and pruning of synapses that are dynamically regulated by synaptic activity [21-24]. Microglia can also regulate synaptic activity through mechanisms that are beginning to be identified, including secretion of neuroactive and neurotrophic factors [25]. A role for microglia in the development of AD may therefore involve not only perpetuation of dysfunctional innate immune responses to accumulating AB peptides and dying synapses and neurons, as the current literature supports, but may also involve direct effects of microglia on synapse integrity and function.

### NON-STEROIDAL ANTI-INFLAMMATORY DRUGS (NSAIDS) AND PGE<sub>2</sub> IN AD DEVELOPMENT

The primary action of NSAIDs is the enzymatic inhibition of the cyclooxygenases COX-1 and COX-2, cytosolic enzymes that generate PGH<sub>2</sub> from membrane stores of arachidonic acid. PGH2 is the precursor of the prostaglandins PGE<sub>2</sub>, PGD<sub>2</sub>, PGI<sub>2</sub>, and PGF<sub>2a</sub>, and the thromboxane TXA<sub>2</sub>. In epidemiologic studies of cognitively normal aging populations, NSAIDs prevent and delay development of AD [26-31]. Although selected NSAIDs may have cyclooxygenase-independent effects, structurally distinct NSAIDs, including ibuprofen, naproxen, and sulindac all reduce risk of developing AD in large epidemiologic studies [27, 28, 30] suggesting that inflammatory prostaglandin signaling plays an important role in pre-clinical development of AD. Interestingly, the beneficial effects of NSAIDs are restricted to the pre-clinical asymptomatic phase, as NSAIDs or selective COX-2 inhibitors do not help patients with mild cognitive impairment (MCI) or AD [31-36]. NSAIDs however are not a good choice for large scale AD prevention because both toxic as well as beneficial downstream prostaglandin signaling pathways are inhibited and lead to adverse effects, including renal and gastric toxicity and increased risk for vascular disease [37]. Recent studies have identified beneficial prostaglandin signaling pathways downstream of NSAID action, including the vasodilatory prostacyclin PGI<sub>2</sub> receptor and the neuroprotective, anti-inflammatory, and vasodilatory PGE<sub>2</sub> EP4 receptor [38-42]. This major limitation may help explain why in MCI and AD, NSAIDs show no benefit, either because protective downstream prostaglandin pathways are inhibited or disease progression is already too advanced.

Inhibition of COX enzymatic activity by NSAIDs therefore has different consequences depending on timing of AD development, and inhibition of COX-1/COX-2 by non-selective NSAIDs is beneficial in preventing disease in healthy aging individuals but ineffectual once symptoms begin [31, 43]. Given the expanding population of aging individuals and the anticipated rise in AD cases, understanding the molecular mechanisms by which NSAIDs prevent AD has taken on significant urgency. Targeting of toxic inflammatory prostaglandin signaling downstream of COX may potentially slow or prevent progression to AD.

PGE<sub>2</sub>, PGD<sub>2</sub>, PGI<sub>2</sub>, and PGF<sub>2a</sub>, and the thromboxane TXA<sub>2</sub> are lipid signaling molecules that bind and activate specific G-protein-coupled receptors designated EP (for Eprostanoid receptor), DP, IP, FP, and TP, respectively [44]. PGE<sub>2</sub> in particular has generated interest as a potential inflammatory agent in pre-clinical AD, as it was found to be increased 5-fold in cerebrospinal fluid (CSF) of patients with early or probable AD [45, 46], but then declined with disease progression [46]. In parallel, levels of the breakdown product of prostacyclin (PGI<sub>2</sub>), widely considered to be antiinflammatory, were significantly decreased in CSF of probable AD subjects [45]. PGE<sub>2</sub> binds four G-protein coupled receptors (GPCRs) termed E-prostanoid receptors (EP1-4) that have distinct downstream signaling cascades and cellular distributions in brain. In vivo, all four EP receptors are expressed in neurons; microglial expression of EP2, EP3, and EP4 receptors has been confirmed in mouse brain [41, 47, 48]. Activation of EP receptors leads to changes in the production of cAMP and/or phosphoinositol turnover and  $Ca^{2+}$  mobilization. EP2 and EP4 receptors couple positively to G<sub>s</sub> to increase cAMP formation whereas EP3 couples negatively to cAMP through G<sub>i</sub>; EP1 couples to G<sub>q</sub>, and activation results in increased intracellular calcium concentrations.

### MOUSE MODELING OF PREVENTIVE EFFECTS OF NSAIDS

To better understand the inflammatory PGE<sub>2</sub> signaling pathway in the context of Aß peptide accumulation, investigators have studied mouse models of Familial AD, where transgenic mice express mutant forms of the amyloid precursor protein (APP) and/or presenilin 1 (PS1) genes. Microglial activation and elaboration of inflammatory and oxidative stress are well documented in these models, particularly as these mutant APP mice age and accumulate  $A\beta_{42}$  and  $A\beta_{40}$  peptides. Mutant APP models display either loss of synapses or loss of synaptic proteins that are associated with spatial memory deficits, and these have been linked to effects of AB oligomers, which are directly toxic to synapses [49], and to effects of inflammatory mediators like IL1 $\beta$  [50] or TNF $\alpha$  [51]. However, because mutant APP models do not develop significant neuronal loss and tau pathology, two hallmarks of MCI and AD in human subjects, these models are believed to be more reflective of the preclinical or asymptomatic phases of human AD [52]. If considered in this way, the beneficial effects of NSAIDs have been validated in these mutant APP models, where a correlation between NSAID administration and reduction of brain inflammation,  $A\beta$  deposition, and rescue of learning and memory deficits has been established [53-56].

### ROLES OF MICROGLIAL EP RECEPTORS IN MODELS OF AD: OPPOSING EFFECTS OF MICROGLIAL EP2 AND EP4 IN MOUSE AD MODELS

Early studies examining *in vivo* effects of EP2 signaling in innate immunity demonstrated a significant reduction in lipid peroxidation following intracerebroventricular (ICV) administration of lipopolysaccharide (LPS) [57], a canonical inducer of the innate immune response. LPS-dependent

increases in F2-isoprostanes, which are free radicalgenerated isomers of prostaglandin PGF2a, and F4neuroprostanes (isoPs), which are products of neuronal docosohexanoic acid (DHA) oxidation, were significantly suppressed in cerebral cortex of EP2-/- mice [57]. In vitro studies parsing out the cellular specificity of the EP2 oxidative effect demonstrated that microglial EP2 elicited paracrine neurotoxicity in co-cultures of neurons and LPSprimed microglia, and this effect was dependent on increased inducible nitric oxide synthase (iNOS) and COX-2 activity [58]. In vivo, in the APPSwe-PS1AE9 model of AD (APP-PS1), global deletion of EP2 also led to significant decreases in lipid peroxidation in aging APP-PS1 mice [59], suggesting a toxic inflammatory role for the EP2 receptor. Additional in vivo studies demonstrated that global deletion of EP2 reduced expression of oxidative enzymes iNOS and components of the NADPH oxidase complex in the APP-PS1 model of AD, and reduced expression of COX-1, COX-2, iNOS, and many components of the NADPH oxidase complex in a related model of familial amyotrophic lateral sclerosis (ALS) [60].

A major target of the innate immune response in the CNS is the clearance of toxic and misfolded proteins. The accumulation of misfolded or aggregated proteins is a common feature of several chronic neurodegenerative diseases, including AD, Parkinson's disease, Huntington's disease, and ALS. Microglia play a crucial role in the clearance of these toxic protein assemblies [61]. However, with progression of disease, notably in AD models, the healthy phagocytic response of microglia to AB peptides falters, either because microglia become ineffective or because they are overwhelmed by levels of accumulating Aß peptides. In parallel with progression of pathology in AD model mice, microglia also develop a more toxic inflammatory phenotype [62]. This leads to a damaging feedforward cycle, with increasing accumulation of toxic Aß peptide assemblies along with increased elaboration of toxic cytokines. In AD model mice, Aβ signaling through Toll-like receptors 2 and 4 (TLR2 and TLR4) drives downstream activation of NF-KB transcription factors as a central inflammatory pathway in AD [63].

A role for microglial EP2 in inhibiting phagocytosis of Aß fibrils was demonstrated in an ex vivo acute preparation using AD brain sections coated with EP2-/- microglia [64]. Microglia lacking EP2 receptor cleared human Aß peptides more effectively than wild type microglia, and EP2-/microglia were associated with lower paracrine neurotoxicity. In vivo, in the APP-PS1 model, deletion of EP2 resulted in significant reductions in total AB40 and AB42 levels and amyloid plaque deposition, an outcome likely reflecting both a more benign inflammatory milieu and an enhanced clearance of AB peptides [65]. In chimeric APP-PS1 mice subjected to whole body irradiation followed by transplantation of wild type or EP2-/- bone marrow, levels of amyloid plaque were reduced in APP-PS1 mice receiving wild type bone marrow, however EP2-/- bone marrow elicited even larger decreases in cerebral cortical plaque load [66]. A role for EP2 signaling in phagocytosis has been shown in non-CNS models, where myeloid EP2 suppresses phagocytosis of latex beads [67, 68] and bacteria [69-72].

Recent in vivo studies using conditional knockout strategies have further defined the critical toxicity of microglial EP2 signaling in models of AD. Conditional knock down of myeloid EP2 receptor using the Cd11bCre recombinase line, where levels of myeloid EP2 are reduced ~50% [47], had multiple beneficial effects and restored healthy microglial responses to AB peptides [73]. Cellspecific knockout of microglial EP2 increased microglial clearance of AB peptides and suppressed toxic inflammatory gene expression. In addition, unbiased genomic studies of microglia isolated from brain revealed that knockdown of microglial EP2 receptor increased generation and local signaling of insulin-like growth factor 1 (IGF1) in hippocampus in response to AB peptide stimulation [73]. IGF1 promotes synaptogenesis, neurogenesis, and neuroprotection through the PI3K/Akt pathway in brain [74]. Deletion of microglial EP2 also increased expression of members of the PPAR signaling pathway, including RXRy which along with its binding partner PPARy reduces proinflammatory gene expression [75] and enhances clearance of AB peptides [76]. RXR $\gamma$  is also the target of the FDA-approved RXR agonist bexarotene (Targretin) that has been shown in some studies to lower interstitial levels of soluble Aß peptides and to prevent memory deficits in AD model mice [77, 78]. Additional genes suppressed by EP2 signaling in microglia but intimately related to AB peptide clearance included the cholesterol transporter ABCA1 [79] and apolipoprotein E (ApoE) [80], proteins that enhance proteolytic degradation of soluble Aß peptides [81, 82]. In addition, lipoprotein lipase (lpl), which binds AB peptide [83], was also upregulated in hippocampus with EP2 microglial deletion; interestingly, an intronic polymorphism in Lpl is associated with reduced Lpl mRNA, increased amyloid and neurofibrillary tangle densities, and increased prevalence of AD [84]. The upregulation of these genes in microglial EP2 knockout mice suggests that EP2-deficient microglia respond to AB<sub>42</sub> peptides in vivo by inducing antiinflammatory and AB-clearing nuclear hormone receptor signaling genes. Consistent with this beneficial effect of reduced EP2 signaling, conditional deletion of microglial EP2 in the APP-PS1 model prevented synaptic injury and spatial memory deficits [73].

The toxic effects of microglial EP2 contrast significantly with the beneficial anti-inflammatory effects of microglial EP4 in vivo. Previous in vivo studies of innate immune responses to LPS had identified a pronounced antiinflammatory effect of microglial EP4 activation that was associated with reduced nuclear translocation of NF-KB subunits p65 and p50 [41] in myeloid cells. Conversely, following LPS stimulation in vivo, conditional deletion of the EP4 receptor in myeloid cells led to an increase in brain pro-inflammatory gene expression and lipid peroxidation, suggesting that the function of myeloid EP4 is to attenuate and/or terminate innate immune responses. Subsequent studies of AB42-mediated inflammatory responses confirmed the anti-inflammatory nature of microglial EP4 signaling, where in primary microglial cells, EP4 stimulation attenuated levels of AB42-induced inflammatory factors and potentiated phagocytosis of  $A\beta_{42}$  [41]. Unbiased genomic studies showed that EP4 receptor activation broadly opposed A $\beta_{42}$ -driven gene expression changes in microglia, with



Fig. (1). Summary of inflammatory effects of the  $COX/PGE_2/EP$  receptor signaling pathways in mouse mutant APP models. Modeling of EP2 and EP3 (top) and EP4 (bottom) inflammatory signaling in mouse models of AD indicates that EP2 and EP3 receptors enhance inflammatory oxidative stress, pro-inflammatory gene expression and are pro-amyloidogenic. In contrast, EP4 signaling in the setting of A $\beta$ -mediated innate immune responses is anti-inflammatory and enhances A $\beta$  phagocytosis. In preclinical AD, use of NSAIDs is preventive only in normal cognitive aging populations. Later symptomatic stages do not respond, potentially because beneficial PGE<sub>2</sub> signaling pathways such as the EP4 receptor, as well as others including the prostacyclin (IP) receptor, are inhibited along with the toxic EP2 and EP3 pathways.

significant enrichment of targets of IRF1, IRF7, and NF- $\kappa$ B transcription factors [41]. *In vivo*, in APP-PS1 mice deficient for microglial EP4, inflammatory gene expression, oxidative protein modification, and A $\beta$  deposition in brain were significantly increased at early stages of pathology, but not at later stages, suggesting an early anti-inflammatory function of microglial EP4 signaling in the APP-PS1 model. Thus, although both the EP2 and EP4 receptors are positively coupled to cAMP, they appear to have divergent inflammatory functions in models of A $\beta$  peptide mediated neuroinflammation and neurodegeneration. The carboxy terminal cytoplasmic tail of the EP4 receptor is significantly longer than that of the EP2 receptor and can recruit distinct signaling molecules [85], a difference likely to influence downstream signaling of the EP4 versus EP2 receptors [86].

## FUNCTION OF THE EP3 AND EP1 RECEPTORS IN THE INFLAMMATORY RESPONSE IN AD

The EP3 receptor is a central component in the regulation of the febrile response [87, 88]. In classical models of innate immunity, the function of inflammatory EP3 has been explored in a model of bacterial lung infection, where deletion of EP3 resulted in a marked increase in clearance of bacteria, reduced neutrophil ingress, and improved survival [89], a phenotype reminiscent of the pro-phagocytic and antiinflammatory effects of EP2 deletion, as discussed above. In the brain, EP3 has mainly been localized to neurons, however immunocytochemical induction of EP3 expression has been observed in striatal microglia in the setting of an excitotoxic lesion induced by injection of quinolinic acid, a potent ligand at the glutamatergic N-methyl-D-Aspartate (NMDA) receptor [48]. This suggests that although EP3 may not be expressed in microglia under physiological conditions, in the appropriate inflammatory or injury contexts, EP3 might be induced and contribute to innate immune responses.

The function of EP3 signaling has been examined in two models relevant to AD, namely the ICV AB injection model, which elicits a potent and long lasting inflammatory response to Aß peptides [90, 91] and in the APP-PS1 model. In a recent study [92], the induction of pro-inflammatory gene expression, cytokine generation, and lipid peroxidation following ICV AB<sub>42</sub> was significantly abrogated in EP3-/mice, suggesting that microglial EP3 signaling engaged a toxic inflammatory response to Aß peptides. In the APP-PS1 model, deletion of EP3 significantly blunted the induction of proteins capable of increasing oxidative injury, including iNOS, components of the NADPH oxidase complex, and COX-2. Moreover, suggestive of a role in Aß clearance, APP-PS1 mice lacking either one or both EP3 alleles had significantly lower amyloid accumulation. This effect is consistent with the lung infection studies mentioned above, where deletion of EP3 led to enhanced clearance of bacteria [89]. However, it is also possible EP3 deletion impacted on the generation of AB peptide from APP, as prior studies have correlated increased oxidative stress and inflammation with increased expression and activity of B-secretase [93, 94], the first enzyme to cleave APP in the formation of the AB42 peptide. Indeed, loss of EP3 receptor in the APP-PS1 background resulted in decreased ß-secretase expression and

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activity. This would suggest that in the APP-PS1 model, EP3 signaling may both suppress Aß clearance, and by increasing inflammatory oxidative stress, increase the generation of Aß peptide. Interestingly, loss of just one allele of EP3 in the APP-PS1 background had significant effects on Aß peptide levels.

Of the four EP receptors, the role of inflammatory EP1 in AD is less clear. EP1 is highly expressed in neurons [95, 96] and functions in neuronal survival. In models of neuronal injury, including NMDA excitotoxicity and cerebral ischemia, pharmacologic or genetic deletion of EP1 reduces cerebral injury [96-98]. However, EP1 expression is not found in microglia after hypoxia-ischemia, nor does microglial activation induce neurotoxicity in hippocampal slices treated with LPS and IFN $\gamma$ [96]. A lack of effect of inflammatory EP1 would suggest that the toxic function of this receptor is primarily neuronal. Supporting this conclusion are findings that inflammation sensitive neural progenitor cells (NPC) [99] in the subgranular zone of the dentate gyrus are vulnerable to microglial EP2 signaling and NPC EP1 signaling [100].

### CONCLUSION

Aß peptides are highly immunogenic, and generate toxic inflammatory responses that injure synapses. Pre-clinical development of AD begins decades prior to diagnosis, and NSAIDs act during this time period to delay onset and progression to AD. The early accumulation of AB42 peptides, beginning years to decades before cognitive symptoms arise, triggers microglial inflammatory responses that are initially robust, but falter as disease progresses. A summary of findings relevant to inflammatory actions of the PGE2 EP2, EP3, and EP4 receptors in the APP-PS1 model are diagrammed in Fig. (1). The opposing actions of the EP2/EP3 and EP4 receptors highlight the importance of targeting selected EP receptors downstream of COX-1/COX-2, as upstream inhibition of COX-1/COX-2 activity may inhibit beneficial as well as toxic PGE<sub>2</sub> EP receptor signaling. This is a potential future indication, as we await the identification and validation of biomarkers that can reliably predict subjects at risk for AD.

### **CONFLICT OF INTEREST**

The authors confirm that this article content has no conflict of interest.

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