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## The role of inflammasome in Alzheimer's disease

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### Abstract

Alzheimer's disease (AD) is a chronic, progressive and irreversible neurodegenerative disease with clinical characteristics of memory loss, dementia and cognitive impairment. Although the pathophysiological mechanism is not fully understood, inflammation has been shown to play a critical role in the pathogenesis of AD. Inflammation in the central nervous system (CNS) is characterized by the activation of glial cells and release of proinflammatory cytokines and chemokines. Accumulating evidence demonstrates that inflammasomes, which cleaves precursors of interleukin-1 $\beta$  (IL-1 $\beta$ ) and IL-18 to generate their active forms, play an important role in the inflammatory response in the CNS and in AD pathogenesis. Therefore, modulating inflammasome complex assembly and activation could be a potential strategy for suppressing inflammation in the CNS. This review aims to provide insight into the role of inflammasomes in the CNS, with respect to the pathogenesis of AD, and may provide possible clues for devising novel therapeutic strategies.

### Keywords

Alzheimer's disease; Inflammasome; Inflammation; Astrocytes; Microglia; Interleukins

### 1. Introduction

Alzheimer's disease (AD) is an irreversible neurodegenerative disease, characterized by extracellular deposition of A $\beta$  beta (A $\beta$ ) plaques and intracellular accumulation of hyper-

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phosphorylated tau protein and NFTs (neurofibrillary tangles) (Huang and Mucke, 2012). Studies in AD patients demonstrate a significant reduction in the size of specific brain regions through the loss of neurons and synapses as compared with healthy subjects (Desikan et al., 2009; Wenk, 2003). The early stages of AD manifest as mild cognitive decline, with progressive memory loss and impairment of other intellectual abilities as the disease progresses, and at the late stages patients suffering from AD alter their personality and lose their bodily functions (Huang and Mucke, 2012). Currently, AD affects more than 38 million people worldwide and every four seconds a new patient is diagnosed with AD (Alzheimer's, 2012). The number of AD patients is expected to double every 20 years and is expected to reach 115 million in 2050 (Martin Prince et al., 2012). As the baby boom generation ages, it is anticipated that the number of AD cases will rise dramatically. However, effective treatment of AD remains a challenge, although numerous hypotheses of possible mechanisms have been put forth.

To date, there are several competing hypotheses that attempt to explain the cause of AD, such as the cholinergic hypothesis, A $\beta$  hypothesis, tau hypothesis and inflammation hypothesis. Cholinergic deficiency is the oldest hypothesis, but has not been widely accepted due to unsuccessful clinical and experimental results (Comim et al., 2012; Nelson et al., 2009). The A $\beta$  and tau hypotheses postulate that A $\beta$  deposits or tau protein abnormalities are the fundamental causes of the disease. Nevertheless, A $\beta$  plaques and neurofibrillary tangles are not sufficient in explaining all the features of AD, since abnormal levels of A $\beta$  plaques have been found among normal healthy elderly individuals (Aizenstein et al., 2008). In addition, highly expressed A $\beta$  or tau protein in animal models of AD do not show significant neurodegenerative changes (Johnston et al., 2011). Furthermore, clinical trials with immune-therapeutics to reduce A $\beta$  level in the brain did not improve cognitive function in AD patients, although evidence suggests clearance of the A $\beta$  plaques (Holmes et al., 2008). These findings suggest that other factors might also be involved in the pathogenesis of AD. Inflammation has been proposed as a key effector of AD, and several features of AD, such as microglial activation, reactive astrocytes and elevated cytokine expression, have all been observed in AD patients, leading to an inflammatory hypothesis. Additionally, epidemiological evidence suggests that non-steroidal anti-inflammatory drugs (NSAIDs) could delay the onset and reduce severity of AD symptoms, although several studies have failed to confirm this (Szekely and Zandi, 2010). More recently, analysis of human brain AD samples has revealed highly expressed inflammatory cytokines during the early stages of AD (Sudduth et al., 2013), and genome-wide studies show an upregulation of inflammatory genes, indicating a potential role of inflammation in the progression of AD (Hollingsworth et al., 2011; Sudduth et al., 2013).

Considering the importance of inflammation in the pathogenesis of AD, several recent review articles have discussed how inflammatory molecules, signaling pathways and processes could be involved in the pathogenesis of AD (Blasko et al., 2000; Johnston et al., 2011; Lau and Yu, 2001; Li et al., 2011; Liu et al., 2013a). In this review, we focus on specific inflammasomes identified in the central nervous system (CNS), in particular the brain, their potential as therapeutics, and the cytokines released by these inflammasomes, with emphasis on environmental factors, i.e. fatty acids, that trigger the generation of cytokines through inflammasomes.

## 2. Inflammatory process in AD

Inflammation is a complex cellular and molecular defense mechanism in response to diverse stimuli, including stress, injury and infection. In the brain, an upregulation in inflammatory signaling (i.e. neuroinflammation) is characterized by the activation of astrocytes and microglia, and the release of proinflammatory mediators. Neuroinflammation is involved in neurodegenerative diseases, such as Parkinson's disease, Huntington's disease, amyotrophic lateral sclerosis, multiple sclerosis, psychiatric, as well as AD (Craft et al., 2006; Pizza et al., 2011; Sudduth et al., 2013; Varnum and Ikezu, 2012). Mounting evidence indicates an upregulation in the inflammatory molecules and activated glial cells surrounding the senile plaques in brains of AD patients and AD transgenic animal models (Bauer et al., 1991; Cagnin et al., 2001; Fillit et al., 1991). Concomitantly, a downregulation in anti-inflammatory molecules is observed in post mortem brains of AD patients (Walker et al., 2009). These findings implicate an involvement of inflammatory responses in the pathogenesis of AD. In the CNS, activated astrocytes and microglia are major sources of inflammatory molecules, such as cytokines, chemokines, neurotransmitters, reactive oxygen species, and nitric oxide (NO) (Tansey et al., 2007). The released cytokines, in particular IL-1 $\beta$ , IL-6 and TNF $\alpha$  (tumor necrosis factor  $\alpha$ ), are the major effectors of the neuroinflammatory signals (Allan and Rothwell, 2003; Rothwell, 1999), and can affect neurophysiologic mechanisms regarding cognition and memory (Gemma and Bickford, 2007; Jankowsky and Patterson, 1999). Cytokines could establish a feedback loop, to activate more astrocytes and microglia and lead to further generation of inflammatory molecules. In addition, the secreted inflammatory molecules also recruit other cells such as monocytes and lymphocytes to cross the blood brain barrier to enhance neuroinflammation in the CNS (Das and Basu, 2008).

### 2.1 Astrocytes and AD

Astrocytes are the most abundant non-neuronal cells in the CNS, constituting about 20–50% of the human brain volume, much higher than microglia (Shimada et al., 2012). Astrocytes have multiple functions, such as regulation of extracellular ions and energy reserves, clearance and metabolism of neurotransmitters, and facilitating the maintenance of normal neuronal functions in the CNS (Dong and Benveniste, 2001; Rossi and Volterra, 2009). Astrocytes can be activated by numerous factors, including pathogens, lipopolysaccharide, oxidative stress (Querfurth and LaFerla, 2010), free saturated fatty acids (Liu et al., 2013a) as well as A $\beta$  (Jana and Pahan, 2010). The activation of astrocytes is believed to last longer than that of microglia, enabling a prolonged engagement of astrocytes in the neuroinflammatory response (Li et al., 2011).

Reactive astrocytes as opposed to quiescent astrocytes, can produce cytokines such as interleukins, TNF $\alpha$  and interferon  $\gamma$  (IFN- $\gamma$ ), etc. (Liu et al., 2013a). They can also generate low amount of A $\beta$ , in addition to neurons, the major producer of A $\beta$  (Blasko et al., 2000; Li et al., 2011). Cytokines, IFN- $\gamma$  along with TNF $\alpha$  or IL-1 $\beta$ , have been demonstrated to induce the generation of A $\beta$  in primary human astrocytes and astrocytoma cells (Blasko et al., 2000). Proinflammatory molecules secreted by reactive astrocytes can elevate the expression of secretases in neurons to enhance the production of A $\beta$  (Tang,

2009; Yu et al., 2009b), and activate microglia to further produce inflammatory factors (Oth et al., 2002).

## 2.2 Microglia and AD

Microglia constitute around 20% of the total glial cells in the brain, and 10% of the cells in the CNS (Lawson et al., 1992; Rubio-Perez and Morillas-Ruiz, 2012). They have various functions in the CNS, including phagocytosis and inducing cytotoxicity (Gehrmann et al., 1995). Microglia can be activated by many factors including A $\beta$ . Activated microglia have phagocytic ability by migrating to damaged cells and clearing debris, and like astrocytes, can also generate inflammatory molecules such as cytokines, chemokines, reactive oxygen and nitrogen species.

Activated microglia colocalize in areas of heavy A $\beta$  concentration (Edison et al., 2008), and preferentially associate with certain types of A $\beta$  plaques, such as fibrillar A $\beta$  (D'Andrea et al., 2004). A $\beta$  binds to microglia and induces phagocytosis or expression of proinflammatory molecules (Ho et al., 2005; Pan et al., 2011), and this activation is dependent on the load of A $\beta$  (Permanne et al., 2002). In an earlier study, upon depletion of microglial cells in animal models the A $\beta$  level increased thereby suggesting microglial cells play a role in the clearance of A $\beta$  by phagocytosis (Broussard et al., 2012; Majumdar et al., 2007). However, more recent work confirmed a detrimental role of microglia in AD that depends upon the degree of activation of the microglial cells. They suggest that with moderate activation by light A $\beta$  concentration microglia have strong phagocytic ability to clear A $\beta$ , while heavily activated microglia by high levels of A $\beta$  increase the production of proinflammatory molecules, such as IL-1 $\beta$  and TNF $\alpha$ , and trigger neuronal damage and coincide with a reduced ability to clear A $\beta$  (Hickman et al., 2008; Johnston et al., 2011). The lower clearance is due to a significant decrease in the expression of A $\beta$  binding scavenger receptors and A $\beta$  degrading enzymes in microglia (Hickman et al., 2008; Johnston et al., 2011). Therefore, the exact function of microglia in AD, whether it is beneficial or detrimental, may depend on the activation state of the microglia.

## 2.3 Neurons and AD

Neurons at around 100 billion, are the core components of the brain (Williams and Herrup, 1988). They are electrically excitable and have diverse functions, including processing and transmitting information through electrical and chemical signals. In AD, a significant loss of neurons in the cerebral cortex and certain subcortical regions have been reported (Annaert and De Strooper, 2002). Both extracellular deposition of amyloid plaques and intracellular aggregation of neurofibrillary tangles are hallmarks of AD. A $\beta$  peptides and hyperphosphorylated tau protein are major components of amyloid plaques and neurofibrillary tangles, respectively (Annaert and De Strooper, 2002). Both are toxic to the cells and are believed to lead to neuronal dysfunction and death (Price and Sisodia, 1994).

Inflammatory processes are thought to play a role in neuronal degeneration (Meng et al., 2013; Morales et al., 2010). In addition to astrocytes and microglia, neurons also contribute to the inflammatory response in the CNS by releasing cytokines, e.g. IL-1 $\beta$  and IL-18. Inflammasomes involve in the maturation of IL-1 $\beta$  and IL-18 are expressed in neurons (de

Rivero Vaccari et al., 2008; Yang-Wei Fann et al., 2013; Zou and Crews, 2012). Trauma, a risk factor for AD, increases inflammasome expression in rat neurons (de Rivero Vaccari et al., 2009; de Rivero Vaccari et al., 2008). Furthermore, postmortem brain tissues from stroke patients show increase protein levels of inflammasomes in primary cortical neurons (Yang-Wei Fann et al., 2013). Currently, the expression levels of inflammasomes in neurons of AD patients are not extensively studied, and much needed insight into the pathogenesis of AD could be gained through further studies in this area.

### 3. Inflammatory cytokine molecules in AD

Elevated levels of inflammatory cytokines, TNF $\alpha$ , IFN $\gamma$ , and interleukins, are found in the brain, in particular near the A $\beta$  plaques, of Alzheimer's patients and AD transgenic animals (Johnston et al., 2011). Cytokines are also detected in the peripheral blood and cerebrospinal fluid of AD patients (Bossu et al., 2008; Di Rosa et al., 2006; Rubio-Perez and Morillas-Ruiz, 2012). Proinflammatory cytokines have multiple functions, which include stimulating or inhibiting cell proliferation, differentiation and apoptosis, as well as inducing inflammation (Rubio-Perez and Morillas-Ruiz, 2012). In the CNS, proinflammatory cytokines induce the release of a number of other proinflammatory molecules from the same or different cell types, amplifying the cytokine effects. Although these studies do not demonstrate whether inflammation is an initiator, contributor or side effect in the pathophysiological changes associated with AD (Li et al., 2011; Sudduth et al., 2013), it (i.e. cytokines) nevertheless is involved in regulating A $\beta$  plaque deposition and BACE1 expression in AD transgenic mice (Yamamoto et al., 2007). Interleukins, in particular IL-1 $\beta$  and IL-18, are upregulated in AD brain, and the overexpression of IL-1 $\beta$  or IL-18 is critical for the onset of the inflammatory process (Rubio-Perez and Morillas-Ruiz, 2012), and both mediate the expression of a vast array of inflammatory genes (Weber et al., 2010). IL-1 $\beta$  and IL-18 are synthesized as inactive precursors, proIL-1 $\beta$  and proIL-18, respectively, and require inflammasomes for their maturation.

#### 3.1 Inflammasome

Inflammasome, the platform for proIL-1 $\beta$  and proIL-18 processing, is an intracellular multiprotein complex. To date, the most well-characterized inflammasomes are NLRP1 (nucleotide-binding oligomerization domain (NOD)-like receptor protein 1), NLRP2, (NOD-like receptor protein 2), NLRP3 (NOD-like receptor protein 3), and NLRC4 (CARD domain-containing protein 4, also called IPAF (ICE-protease activating factor)) inflammasome (Martinon et al., 2009; Minkiewicz et al., 2013). The basic components of inflammasomes include a NOD-like receptor (NLR) that recognizes danger signals or ligands, and procaspase-1 which is central to inflammasome activation. In addition to these basic components, other factors could also be involved in the assembly or activity of inflammasomes, depending on the type of cell and stimulus. They include the adaptor protein, ASC (apoptosis-associated speck-like protein containing a caspase recruitment domain), which is an essential component of the NLRP1, NLRP2 and NLRP3 inflammasomes, but not of the more complex NLRC4 inflammasome (Martinon et al., 2009; Minkiewicz et al., 2013; Schroder and Tschopp, 2010). Under certain but not all conditions ASC (Martinon et al., 2009) or Naip5 (NLR family, apoptosis inhibitory protein 5)

(Lightfield et al., 2011) is required for the activity of the NLRC4 inflammasome. The receptors of the inflammasomes, NLRs, that sense the pathogen associated and damage associated molecules (Martinon et al., 2009), are predominantly cytosolic proteins. They are structurally characterized by three distinct domains: a domain of leucine rich repeats (LRRs) for ligand sensing, a NACHT domain (also called nucleotide-binding oligomerization domain (NOD)) that facilitates oligomerization, and an effector domain which consists of one of three domains, i.e. a pyrin (PYD), CARD (caspase activation and recruitment domains), or BIR (baculovirus IAP repeat) domain (Martinon et al., 2009). In response to danger signals, inflammasomes assemble by self-oligomerizing the NLRs through interactions with the NACHT domain (van de Veerdonk et al., 2011). As for the NLRP1 and NLRP3 inflammasomes, the oligomerized NLRs recruit and interact with the adaptor protein ASC, which in turn recruits the effector protein procaspase-1 that is central to the activation of inflammasomes (Huang et al., 2013). Once procaspase-1 is recruited to assemble the inflammasome, it self-cleaves to form the mature caspase-1. Caspase-1 is the protease that cleaves the precursor of the proinflammatory molecules to form their mature form, such as IL-1 $\beta$  and IL-18 (Schroder and Tschopp, 2010). The activators of the inflammasomes can be divided into two categories; pathogen associated molecular patterns (PAMPs) activate a host-defense reaction, and damage associated molecular patterns (DAMPs) activate a self-defense mechanism in response to danger signals (Salminen et al., 2008). Activators include bacteria, virus, fungus, protozoa, microbial proteins, crystalline urea, RNA, Alum, ATP, potassium efflux, fatty acids, A $\beta$ , and most recently, degraded mitochondrial DNA (Liu et al., 2013a; Mathew et al., 2012; Schmidt and Lenz, 2012). Overall the assembly and activation of inflammasomes are cell type and stimulus specific (Martinon et al., 2009; Schroder and Tschopp, 2010).

### 3.2 Proinflammatory cytokine: Interleukin-1 $\beta$

IL-1 $\beta$  is expressed in many cell types including macrophages, monocytes and neutrophils, and in the CNS IL-1 $\beta$  can be released from astrocytes, microglia and neurons (Dinarello, 2010). IL-1 $\beta$  induces diverse signalings which are cell type specific (Srinivasan et al., 2004), for example, in glial cells IL-1 $\beta$  activates NF $\kappa$ B (nuclear factor kappa B) signaling to upregulate cytokine production. In contrast, in neurons IL-1 $\beta$  activates the MAPK-p38 signaling cascade to increase the secreted APP fragment cleaved by BACE1 to enhance the ability to form A $\beta$  (Salminen et al., 2008; Srinivasan et al., 2004; Sun et al., 2003).

**3.2.1 IL-1 $\beta$  relationship to AD**—IL-1 $\beta$  plays a key role in the onset and development of diverse diseases, including neurodegenerative diseases such as AD (Dinarello, 2010). The IL-1 $\beta$  gene and eight other interleukin 1 family genes form a cytokine gene cluster on chromosome 2 (Webb et al., 1986). Elevated IL-1 $\beta$  has been observed in the serum, cerebrospinal fluid and brain of patients with AD as well as other dementia (Blum-Degen et al., 1995; Cacabelos et al., 1991; Deniz-Naranjo et al., 2008). Elevated levels also have been reported in the cerebrospinal fluid and brain parenchyma of both humans and rodents shortly after traumatic brain injury, the latter is an independent risk factor for AD (Emmanouilidou et al., 2011; Shaftel et al., 2008; Yamasaki et al., 1995). IL-1 $\beta$  can activate other cell types, in particular astrocytes and microglia, to further induce cytokine release (e.g. IL-1 $\beta$ , IL-6 and IL-18), as well as inducible nitric oxide synthase activity to produce the free radical NO,

leading to neurotoxicity (Rossi and Bianchini, 1996; Rubio-Perez and Morillas-Ruiz, 2012). IL-1 $\beta$  secreted from astrocytes has been shown to enhance the production of APP and A $\beta$  from the neurons (Blasko et al., 2000; Bonifati and Kishore, 2007; Li et al., 2011). Additionally, IL-1 $\beta$  has been found to increase the release of S100B from an astrogloma cell line and the plaque-associated activated astrocytes in the primary cortex that promote neuronal APP synthesis (Li et al., 2011; Liu et al., 2005). Injecting IL-1 $\beta$  into the cerebral hemisphere upregulates the levels of APP and amyloidogenesis (Sheng et al., 1996). Furthermore, several studies demonstrate that IL-1 $\beta$  can induce the phosphorylation of the tau protein and mediate the formation of neurofibrillary tangles through the MAPK-p38 pathway (Griffin et al., 2006; Salminen et al., 2008). Blocking IL-1 $\beta$  signaling in the brain of an AD mouse model is able to alter the inflammatory responses of the brain, rescue cognition, attenuate tau pathology, and reduce fibrillar A $\beta$  level (Kitazawa et al., 2011). Conversely, knocking out the IL-1 $\beta$  receptor antagonist in mice increases the neuronal damage induced by A $\beta$  (Craft et al., 2005). These studies implicate a proinflammatory role of IL-1 $\beta$  in the pathogenesis of AD.

**3.2.2 Inflammasomes involved in IL-1 $\beta$  maturation in the CNS**—In the CNS, the production of IL-1 $\beta$  by inflammasomes, specifically NLRP1, NLRP2, NLRP3 and NLRC4, is well-characterized as compared to other interleukins (Minkiewicz et al., 2013; Trendelenburg, 2008). The NLRP1 inflammasome is present in the neurons, astrocytes, oligodendrocytes, and microglia (Abulafia et al., 2009; de Rivero Vaccari et al., 2008; Kummer et al., 2007; Silverman et al., 2009). Spinal cord injury can activate the NLRP1 inflammasome to produce IL-1 $\beta$  in rat spinal cord neurons (de Rivero Vaccari et al., 2008). A subsequent study demonstrated that P2X<sub>7</sub> purinergic receptor is involved in the activation of NLRP1 inflammasome (Silverman et al., 2009). Given that P2X<sub>4</sub> and P2X<sub>7</sub> are the major purinergic P2X receptor subtypes, a study of spinal cord injury in P2X<sub>4</sub> knock-out mice showed a significant reduction in inflammasome activation and proinflammatory cytokine production as compared to wild type (de Rivero Vaccari et al., 2012), supporting the involvement of purinergic receptor P2X<sub>4</sub> in the activation of the NLRP1 inflammasome. Similarly, the P2X<sub>7</sub> purinergic receptor has been shown to activate the NLRP1 inflammasome in primary neurons (Silverman et al., 2009).

ATP, a danger-associated molecular pattern that is released from damaged cells after brain injury, activates the NLRP2 inflammasome, which consists of the NLRP2 receptor, ASC and caspase-1, in human astrocytes (Minkiewicz et al., 2013). The ATP-induced activation of the NLRP2 inflammasome interacts with the ATP-release pannexin 1 channel and ATP-gated P2X<sub>7</sub> receptor leading to the maturation of IL-1 $\beta$  (Minkiewicz et al., 2013).

*In vivo* and cell studies demonstrate that fibrillar A $\beta$  activates the NLRP3 inflammasome which is composed of the NLRP3 receptor, ASC and caspase-1, to produce IL-1 $\beta$  in microglia (Halle et al., 2008). Phagocytosis and subsequent lysosomal damage trigger by A $\beta$  initiate the activation of the NLRP3 inflammasome in the microglia (Halle et al., 2008). In support, a recent study in APP/PS1 mice confirms that the NLRP3 inflammasome contributes to the AD pathology (Heneka et al., 2013). Deficiency of the NLRP3 gene reduces A $\beta$  deposition and plays a protective role on memory and behavior (Heneka et al.,

2013). Similarly, inhibiting the NLRP3 inflammasome reduces the neuritic plaque burden in an AD transgenic mouse model (Shi et al., 2013).

Palmitate, a fatty acid, activates the NLR4 inflammasome in primary astrocytes leading to the release of IL-1 $\beta$  (Liu and Chan, 2014). The adaptor protein ASC is important for the activation of NLR4 inflammasome in astrocytes, while Naip 5 is not (Liu and Chan, 2014). The IL-1 $\beta$  release is upregulated by the astrocytes cultured in palmitate and contributes to a higher production of both BACE1 and A $\beta$  by primary neurons (Liu and Chan, 2014; Liu et al., 2013a; Liu et al., 2013b).

### 3.3 Proinflammatory cytokine: Interleukin-18

Interleukin-18 (IL-18), belonging to the IL-1 superfamily, is constitutively expressed in several cell types, and the active form of IL-18 is generated by cleavage of the precursor, proIL-18. IL-18 has several similarities in their properties with IL-1 $\beta$ , such as, an inactive precursor, is activated by pathogen and danger associated factors, involves inflammasomes, and induces similar signaling events (Das et al., 2008; Huang et al., 2013). However, there are considerable differences between the two cytokines, such as their expression levels, regulation and action (Das et al., 2008; Huang et al., 2013). In normal brain tissue, IL-18 is constitutively and highly expressed, whereas IL-1 $\beta$  is lowly expressed (Culhane et al., 1998). In peripheral immune cells, IL-1 $\beta$  increases rapidly after ischemia, while IL-18 increases much slower (Jander et al., 2002), suggesting differential regulations of the two cytokines. IL-1 $\beta$  activates NF $\kappa$ B and p38 signaling pathways (Griffin et al., 2006; Salminen et al., 2008; Srinivasan et al., 2004) while IL-18 binds to the receptor IL-18R and activates NF $\kappa$ B, STAT3 (Signal transducer and activator of transcription 3) and NFATc4 (Nuclear factor of activated T-cells, cytoplasmic 4) (Suk et al., 2001; Sutinen et al., 2012). Moreover, it activates both Fas and Fas-L promoter activities, and thus has been suggested to be an apoptosis inducer and initiator of atherosclerosis and cardiovascular diseases (Chandrasekar et al., 2006). IL-18 can modulate neuronal excitability (Kanno et al., 2004), and inhibit long term-potential, a form of a neuronal plasticity considered to underlie learning and memory (Curran and O'Connor, 2003).

**3.3.1 IL-18 relationship to AD**—IL-18 is believed to play an important role in various diseases, in particular AD. The IL-18 gene is located in the 11q22.2–22.3 region close to the dopamine receptor D2 locus, near chromosome 11, a chromosomal region of interest in AD as defined by genome studies and suggested as a linkage area for AD pathology in familial AD (Blacker et al., 2003). Moreover, IL-18 promoter polymorphism has been shown to increase the risk of developing sporadic late onset AD in Han Chinese and Italian populations (Bossu et al., 2007; Yu et al., 2009a). In the CNS, IL-18 is produced by astrocytes, microglia and ependymal cells, and has been also detected in neurons (Conti et al., 1999; Ojala et al., 2008; Sugama et al., 2002). The mRNA and protein levels of IL-18 increase significantly in astrocytes, microglia and neurons that co-localized with A $\beta$  plaques in the brains of AD patients (Ojala et al., 2009). IL-18 is elevated significantly in the plasma of mild cognitively impaired and AD patients (Malaguarnera et al., 2006; Ozturk et al., 2007). Moreover, a significant increase in IL-18 is observed in stimulated mononuclear cells and macrophages of peripheral blood from AD patients (Bossu et al., 2008; Di Rosa et al.,



2006), as well as in the blood of patients with ischemic heart disease, type-2 diabetes, and obesity, which are risk factors for AD (Sutinen et al., 2012). These results support that IL-18 is involved in AD disease pathology. Mild AD patients show significant increase in IL-18 level, but slightly lower than moderate AD patients, whereas no significant upregulation is observed in severe AD patients as compared to age-matched control subjects (Motta et al., 2007). This gradual decline in immune response in AD suggests that the immune response, in particular IL-18, could be an initiator of AD pathogenesis. Studies demonstrate that IL-18 triggers an elevation in the protein levels of APP, BACE1, and the N-terminal fragment of presenilin-1 and presenilin enhancer 2 which are components of the  $\gamma$ -secretase complex (Sutinen et al., 2012), suggesting IL-18 accelerates A $\beta$  genesis. Additionally, IL-18 increases the expression of glycogen synthase kinase 3 $\beta$  (GSK-3 $\beta$ ) and cyclin dependent kinase 5 (Cdk5), which are involved in hyperphosphorylation of the tau protein (Ojala et al., 2008). These studies support an important role of IL-18 in AD.

**3.3.2 Inflammasomes involved in IL-18 maturation in the CNS**—Like IL-1 $\beta$ , in most cases, the mature secretable form of IL-18 is generated by caspase-1 through the activation of inflammasome. In contrast, the mature secretable form of IL-18 also can be derived from various extracellular enzymes, including protease 3, serine protease, elastase and cathepsin G (Alboni et al., 2010; Gracie, 2004; Sugawara et al., 2001). However the maturation of IL-18 and IL-1 $\beta$  could be regulated by the same type of inflammasome. For example, down-regulation of NLRP1 in macrophages trigger by *Cordyceps sinensis* mycelium reduces both IL-18 and IL-1 $\beta$  levels (Huang et al., 2013). The release of IL-18 and IL-1 $\beta$  could also be regulated by different inflammasomes even though they are in the same cell type and expose to the same stimuli. A recent report shows that IL-18 and IL-1 $\beta$  are secreted from primed murine dendritic cells in response to *Listeria* protein p60, but inhibiting NLRP3 reduces the production of IL-1 $\beta$ , but not IL-18 secretion (Schmidt and Lenz, 2012). This suggests that the maturation of IL-18 and IL-1 $\beta$  is regulated condition-specifically by different signaling mechanisms upon exposure to similar stimulus.

Spinal cord injury causes IL-18 and IL-1 $\beta$  release from neuronal cells through the activation of the NLRP1 inflammasome, composed of receptor NLRP1, adaptor protein ASC, caspase-1, caspase-11 and X-linked inhibitor of apoptosis protein (de Rivero Vaccari et al., 2008). ASC neutralization reduces the upregulation in IL-18 and IL-1 $\beta$  levels (de Rivero Vaccari et al., 2008). Spinal cord injury elevates extracellular ATP levels during neuroinflammation, which may act on purinergic receptors to trigger the activation of inflammasome (de Rivero Vaccari et al., 2012; Minkiewicz et al., 2013). However, upon further study of purinergic receptor P2X<sub>4</sub> knockout mice with spinal cord injury, the production of IL-1 $\beta$  but not of IL-18 reduces in the neurons as compared with wild-type mice (de Rivero Vaccari et al., 2012). This further supports the differential regulations of IL-18 and IL-1 $\beta$  expression.

In human astrocytes, ATP released from damaged or dying cells after traumatic brain injury activates the NLRP2 inflammasome, leading to the maturation of both IL-1 $\beta$  and IL-18 (Minkiewicz et al., 2013). Inhibiting pannexin 1 by probenecid or Brilliant Blue G significantly reduces the increase levels of IL-1 $\beta$  and IL-18 triggered by ATP (Minkiewicz

et al., 2013), suggesting that the regulation of IL-1 $\beta$  and IL-18 expression in the situation could be similar.

Diverse factors upregulate IL-18 level in the CNS. Kainic acid has been shown to induce IL-18 production from microglia (Jeon et al., 2008). Stress, a risk factor of AD, has been confirmed to promote the synthesis and maturation of IL-18 (Sugama and Conti, 2008). Aging, another risk factor of AD, has been found to activate the NLRP1 inflammasome and upregulate IL-18 and IL-1 $\beta$  levels in the hippocampus of aged mice (Mawhinney et al., 2011). Currently information on the type and composition of the inflammasomes involved in the maturation of IL-18 in the CNS is largely uncharacterized. Further studies are required to reveal the role of IL-18 in AD pathogenesis.

#### 4. Fatty acids involvement in inflammasome activation and pathogenesis of AD

Environmental factors, in particular, saturated fatty acids have been reported to induce the production of proinflammatory cytokines, such as IL-1 $\beta$ , IL-6 and TNF $\alpha$ , the major effectors of neuroinflammatory cascade, in astrocytes and microglia (Gupta et al., 2012; Liu et al., 2013a; Wang et al., 2012). Palmitate, the most common saturated fatty acid in the diet (Guo et al., 2007), triggers the production of cytokines, in particular IL-1 $\beta$ , from many cell types, including astrocytes (Liu et al., 2013a; Luo et al., 2012). Recently we identify that palmitate activates the NLRC4 inflammasome in primary astrocytes to release IL-1 $\beta$ , and ASC participates in the activation of the NLRC4 inflammasome (Liu and Chan, 2014). Reducing NLRC4 or ASC levels in the palmitate (PA)-treated astrocytes significantly reduces IL-1 $\beta$  production (Liu and Chan, 2014). In addition, NLRC4 and ASC levels are upregulated in the brains of AD patients (Liu and Chan, 2014), suggesting a possible role of the NLRC4 inflammasome in AD pathogenesis. In support, palmitate induces IL-1 $\beta$  release from HepG2 (hepatocellular carcinoma cells), and knockdown of NLRC4 inhibits the palmitate induced inflammation and cytokine release (Cho et al., 2013; Luo et al., 2012). Fatty acids can also cause the release of IL-1 $\beta$  from microglia, but the specific inflammasome that regulates this process has not been identified (Wang et al., 2012). These studies highlighted the role of saturated fatty acids in the production of IL-1 $\beta$  by inflammasomes, i.e. NLRC4.

Fatty acids are known to cross the blood-brain barrier through passive diffusion (Dhopeswarkar and Mead, 1973; Smith and Nagura, 2001), and diets high in saturated fats increase brain uptake of fatty acids from the plasma (Karmi et al., 2010; Wang et al., 1994). It is possible that increase uptake of fatty acids by the brain from the plasma (Karmi et al., 2010) could lead to enhance cytokine release and inflammatory response in the brain. Fatty acid metabolism has been suggested as a risk factor for the development of AD (Di Paolo and Kim, 2011; Morris and Tangney, 2010), and epidemiological studies suggest that consumption of saturated fatty acids increases while consumption of unsaturated fatty acids decreases the risk of AD (Scarmeas et al., 2006; Solfrizzi et al., 2005; Takechi et al., 2010). AD brains show high fatty acid content as compared with normal healthy subjects (Roher et al., 2002). Elevated saturated free fatty acids (FFAs) have been found in the plasma of patients with diabetes mellitus, hypertension and obesity, which are risk factors for AD

(Monti et al., 2004; Whitmer et al., 2005). Traumatic brain injury, an independent risk factor for AD (de Rivero Vaccari et al., 2009), induces higher levels of palmitate and stearate in the brain (de Rivero Vaccari et al., 2009; Homayoun et al., 1997; Lipton, 1999). Many *in vivo* studies report that high fat diet induces A $\beta$  deposition and memory deficits in APP transgenic mice (Julien et al., 2010; Levin-Allerhand et al., 2002; Maesako et al., 2012a; Maesako et al., 2012b; Oksman et al., 2006). Concomitantly, our group has demonstrated that palmitate triggers the upregulation of AD-like characteristics, such as elevated BACE1, A $\beta$ , and hyperphosphorylated tau levels in neurons, mediated by conditioned media from astrocytes cultured in palmitate (Liu et al., 2013a; Patil et al., 2007; Patil et al., 2006). A subsequent study shows the cytokines, in particular IL-1 $\beta$ , secreted by the palmitate-treated astrocytes, contribute to the upregulation of BACE1 and A $\beta$ 42 levels in the primary neurons (Liu and Chan, 2014; Liu et al., 2013a). However, palmitate does not directly induce the AD-like changes in neurons (Patil and Chan, 2005; Patil et al., 2006), which could be attributed to the lower ability of neurons to take up and metabolize long chain fatty acids (Blazquez et al., 2000; Qin et al., 2010). In contrast, astrocytes readily metabolize fatty acids, and furthermore peripheral administration of fatty acids is found to accumulate primarily in astrocytes (Bernoud et al., 1998; Morand et al., 1979).

Palmitate and its metabolites, such as ceramides, regulate many gene expression and immunological pathways (Ajuwon and Spurlock, 2005; Haughey et al., 2010; He et al., 2010; Jana and Pahan, 2010; Katsel et al., 2007; Little et al., 2012; Puglielli et al., 2003; Saddoughi and Ogretmen, 2013). Serine palmitoyltransferase (SPT), the rate-limiting enzyme that synthesizes ceramide, is upregulated in palmitate-treated astrocytes. Downregulating SPT to decrease ceramide levels mitigates the increase in IL-1 $\beta$  and TNF $\alpha$  that are secreted by the astrocytes (Liu et al., 2013a; Patil et al., 2007), suggesting that ceramide is involved in the cytokine production. Neutralizing TNF $\alpha$  and IL-1 $\beta$  in the conditioned media significantly reduce the upregulation of BACE1 in the neurons (Liu et al., 2013a). Moreover, our group and others have found the levels of SPT are upregulated in human AD brains as compared to age-matched controls (Geekiyana and Chan, 2011; He et al., 2010). Our group observe that inhibiting SPT decreases A $\beta$ 42 and tau hyperphosphorylation in an AD mouse model (Geekiyana et al., 2013). Taken together, the aforementioned studies suggest an involvement of saturated fatty acids in the neuroinflammation mediated by glial cells. Due to the quantity, location, and function of the astrocytes, they are likely major players in the induction of inflammation in the CNS generated by saturated fatty acids.

## 5. Therapeutic for inflammasome-mediated inflammation

Interleukins, in particular, IL-1 $\beta$  and IL-18, have been implicated in neuronal damage in chronic and progressive neuronal diseases, such as AD, Parkinson's disease, Huntington's disease, and Amyotrophic lateral sclerosis. Several approaches, described below, have been evaluated for their ability to reduce the deleterious effects of excessive inflammatory cytokine production and signaling. Secreted extracellular domains of soluble IL-1 receptor and IL-1 receptor antagonist (IL-1Ra) have been shown to mitigate IL-1 $\beta$  signaling by binding IL-1 $\beta$  (Chakraborty et al., 2010; Eisenberg et al., 1990). Similarly, the IL-1 receptor II acts as a decoy receptor to inhibit IL-1 $\beta$  signaling (Colotta et al., 1993). Constitutively

expressing IL-18 binding protein, which binds IL-18, prevents the binding of IL-18 to its receptor and its downstream signaling (Kim et al., 2003). Consequently, therapeutic approaches have been developed to mimic the actions of these intrinsic factors. To date, blocking IL-1 signaling is the most effective therapy in many auto-inflammatory disorders such as familial cold autoinflammatory syndrome, Muckle-Wells syndrome, chronic infantile neurologic cutaneous and articular syndrome, and TNF-receptor associated periodic syndrome (Moll and Kuemmerle-Deschner, 2013). Cytokine traps such as rIL-1Ra, which recognizes IL-1 $\beta$ , are being tested in clinical trials for several autoinflammatory disorders (Hawkins et al., 2003; Hoffman et al., 2004). In support, inhibitors of IL-1 are effective in treating cryopyrin associated periodic syndromes (CAPS) and gout (Lachmann et al., 2011; So et al., 2013). Although inhibitors of IL-1 $\beta$  have been used to systemically treat inflammation (Hoffman et al., 2004), this type of therapy has not been reported for treating CNS inflammation. Non-steroidal anti-inflammatory drugs (NSAIDs), used to suppress inflammation (Hayden et al., 2007; Lee et al., 2010), appear to reduce the risk of developing AD after 2–3 years of treatment based on cognitive decline if taken prior to age 65 (Broussard et al., 2012; Hayden et al., 2007). This is controversial, since there are other reports of their increasing the risk of developing AD (Breitner et al., 2009). The diverse outcomes could be the result of different conditions, such as the age of the patients at treatment, dose of the drugs, duration of the treatment, and the severity of the neurodegeneration (Lee et al., 2010). Nevertheless there are promising NSAIDs and combination of NSAIDs currently in clinical trial for AD (Galimberti and Scarpini, 2011).

Inflammasome is required for the maturation of IL-1 $\beta$  and IL-18. Probenecid, an inhibitor of pannexin 1, has been shown to significantly inhibit the expression and activation of the NLRP2 inflammasome, and the maturation of both IL-1 $\beta$  and IL-18 in human astrocytes induced by ATP (Minkiewicz et al., 2013). Probenecid has also been demonstrated to reduce the activation of the NLRP1 inflammasome, and improve the learning performance in age-related cognitive decline (Mawhinney et al., 2011). In addition, Brilliant Blue G (BBG), a P2X<sub>7</sub> receptor antagonist, inhibits ATP-induced activation of the NLRP2 inflammasome in human astrocytes (Minkiewicz et al., 2013). P2X<sub>4</sub> knock-out mice has been shown to decrease the level of IL-1 $\beta$  and to have impaired inflammasome signaling (de Rivero Vaccari et al., 2012). These reports support that probenecid and BBG have potential therapeutic value, and that pannexin 1, P2X<sub>4</sub> and P2X<sub>7</sub> receptors could be potential therapeutic targets.

Inflammasome is composed of a receptor such as NLRP1, NLRP2, NLRP3 and NLRC4, and caspase-1, and sometimes ASC. Therefore, targeting any of these components, such as the receptor, caspase-1 or ASC, could prevent inflammasome assembly and activation, and reduce IL-1 $\beta$  and IL-18 generation. In support, knockout of NLRP3 and caspase-1 have been shown to suppress amyloidogenesis and neuropathology, and improve cognition in AD transgenic mice (Heneka et al., 2013). Consequently, designing agents that control the activation or assembly of inflammasome could provide a promising approach to tackling neuroinflammation. Reportedly a neutralizing antibody against NLRP1 is able to reduce cytokine levels after thromboembolic stroke in mice, thereby reducing the detrimental consequences of post-ischemic inflammation (Abulafia et al., 2009). The administration of anti-ASC neutralizing antibodies in spinal cord injury in rats reduces caspase-1 activation and IL-1 $\beta$  and IL-18 levels, leading to better recovery after spinal cord trauma (de Rivero

Vaccari et al., 2008). A tetrapeptide recognition sequence has been uncovered for caspase-1, WEHD (Trp-Glu-His-Asp), which can be inhibited by a compound, N-Acetyl-WEHD-aldehyde (Chakraborty et al., 2010; Garcia-Calvo et al., 1998). Thus far clinical trials of this promising application have not been reported. Therapies for treating inflammasome-mediated inflammation have not been fully investigated and would be an interesting area for future exploration.

## 6. Conclusion

To date, studies on the role of inflammasomes in regulating the maturation of interleukins have enhanced our understanding of inflammation in diseases, including AD. Blocking or neutralizing IL-1 $\beta$  reduces cognitive impairment and decreases AD-like pathological changes in AD mouse models (Gonzalez et al., 2009; Kitazawa et al., 2011). Similarly, knockout of IL-1 $\beta$  receptor antagonist in mice increases the neuronal damage induced by A $\beta$  (Craft et al., 2005). These findings support a role of inflammasome in the pathogenesis of AD, however, the detail mechanism of inflammasome assembly and activation in the CNS is still emerging. AD is a complex neurodegenerative disease with an unclear etiology and thus far, there are no proven treatments to prevent or slow its progression. Evidence suggests that environmental factors, e.g. saturated fatty acids, can initiate the activation of glial cells to release inflammatory molecules, including IL-1 $\beta$  and IL-18 (Liu and Chan, 2014; Liu et al., 2013a; Wang et al., 2012). IL-1 $\beta$  and IL-18 can further induce the activation of glial cells in the CNS to secrete more inflammatory molecules, creating a feedback loop to sustain the production of proinflammatory molecules and result in increased amyloidogenesis and neurofibrillary tangles (Figure 1). Therefore, studying interleukin generation through the inhibition of inflammasomes assembly and activation could prove crucial to gaining a better understanding of the role of inflammasomes in AD, and providing novel therapeutics for AD.

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## List of abbreviations

<b>AD</b>	alzheimer's disease
<b>ASC</b>	apoptosis-associated speck-like protein containing a caspase recruitment domain
<b>A<math>\beta</math></b>	A $\beta$ beta
<b>BACE1</b>	beta site A $\beta$ beta precursor protein cleaving enzyme 1
<b>Cdk5</b>	cyclin dependent kinase 5
<b>CNS</b>	central nervous system
<b>DAMPs</b>	damage associated molecular patterns
<b>GSK-3<math>\beta</math></b>	glycogen synthase kinase 3 $\beta$

<b>HepG2</b>	hepatocellular carcinoma cells
<b>IFN-<math>\gamma</math></b>	interferon $\gamma$
<b>IL-1<math>\beta</math></b>	interleukin-1 $\beta$
<b>IL-18</b>	interleukin-18
<b>IL-1RL1</b>	IL-1 receptor-like 1 (also called ST2)
<b>IL-1RAcp</b>	IL-1 receptor accessory protein
<b>Naip5</b>	NLR family, apoptosis inhibitory protein 5
<b>NF<math>\kappa</math>B</b>	nuclear factor kappa B
<b>NFTs</b>	neurofibrillary tangles
<b>PYD</b>	pyrin domain
<b>BIR</b>	baculovirus IAP repeat
<b>NLRP1</b>	nucleotide-binding oligomerization domain (NOD)-like receptor (NLR) protein 1
<b>NLRP2</b>	(NOD-like receptor protein 2)
<b>NLRP3</b>	NOD-like receptor protein 3
<b>CARD domain</b>	Caspase activation and recruitment domains
<b>NLRC4</b>	CARD domain-containing protein 4
<b>IPAF</b>	ICE-protease activating factor
<b>NSAIDs</b>	non-steroidal anti-inflammatory drugs
<b>TNF<math>\alpha</math></b>	tumor necrosis factor- $\alpha$
<b>NFATc4</b>	nuclear factor of activated T-cells, cytoplasmic 4
<b>NO</b>	nitric oxide
<b>PA</b>	palmitate
<b>SPT</b>	serine palmitoyltransferase
<b>PAMPs</b>	pathogen associated molecular patterns
<b>STAT3</b>	Signal transducer and activator of transcription 3

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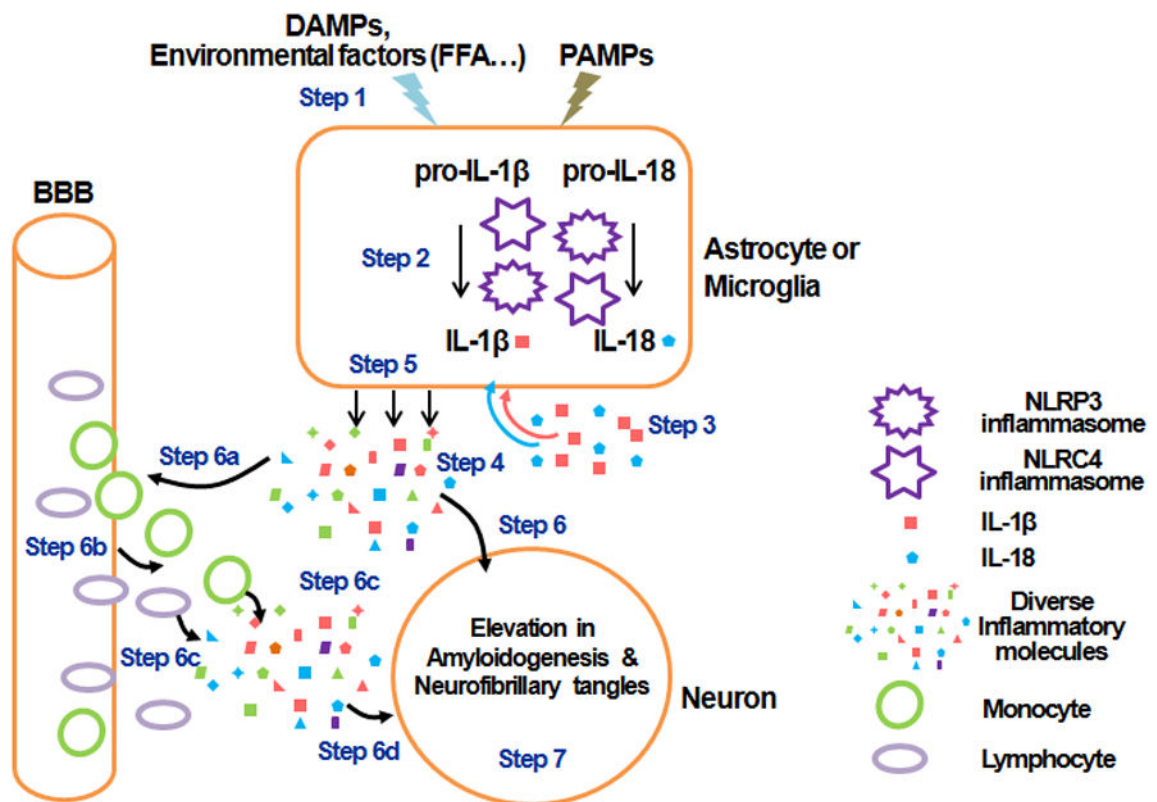
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- Inflammasomes involved in IL-1 $\beta$  and IL-18 maturation contribute to AD progression
- Environmental factors, i.e. fatty acids, can trigger the activation of inflammasome
- Modulating inflammasome activation could be a potential strategy for treating AD



**Figure 1. Scheme of inflammasome-mediated inflammatory response**

Damage associated molecular patterns (DAMPs), environmental factors such as elevated free fatty acids (FFA), or pathogen associated molecular patterns (PAMPs) initiate the activation of inflammasomes in astrocytes or microglial cells to release inflammatory molecules, IL-1 $\beta$  and IL-18 (Step1–3). IL-1 $\beta$  and IL-18 can further activate more astrocytes or microglial cells in the CNS to secrete more diverse inflammatory molecules (Step4, 5). The inflammatory molecules induce the elevation of amyloidogenesis and neurofibrillary tangles in neurons (Step6, 7). Concomitantly, the diverse inflammatory molecules also recruit other cells such as monocytes and lymphocytes to cross the blood brain barrier (BBB) and to release more diverse inflammatory molecules, resulting in an increase in amyloidogenesis and neurofibrillary tangles in the neurons (Step6a–7).