



The Complexity of Microglial Interactions With Innate and Adaptive Immune Cells in Alzheimer's Disease

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In the naïve mouse brain, microglia and astrocytes are the most abundant immune cells; however, there is a complexity of other immune cells present including monocytes, neutrophils, and lymphocytic cells, such as natural killer (NK) cells, T cells, and B cells. In Alzheimer's disease (AD), there is high inflammation, reactive microglia, and astrocytes, leaky blood–brain barrier, the buildup of amyloid-beta ($A\beta$) plaques, and neurofibrillary tangles which attract infiltrating peripheral immune cells that are interacting with the resident microglia. Limited studies have analyzed how these infiltrating immune cells contribute to the neuropathology of AD and even fewer have analyzed their interactions with the resident microglia. Understanding the complexity and dynamics of how these immune cells interact in AD will be important for identifying new and novel therapeutic targets. Thus, this review will focus on discussing our current understanding of how macrophages, neutrophils, NK cells, T cells, and B cells, alongside astrocytes, are altered in AD and what this means for the disorder, as well as how these cells are affected relative to the resident microglia.

Keywords: Alzheimer's disease, microglia, astrocytes, macrophages, neutrophils, T cells, B cells, NK cells

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INTRODUCTION

Alzheimer's disease (AD) is the most prevalent neurodegenerative disease with core neuropathological features including the amassing of amyloid-beta ($A\beta$) plaques and neurofibrillary tangles (NFT) along with neuroinflammation and cognitive decline (Lane et al., 2018; DeTure and Dickson, 2019). Until more recently, the focus on AD has been the treatment of $A\beta$ plaques through directly modulating the plaques; however, these treatments have not cured the underlying neuropathology of the disease (Ceyzeriat et al., 2020). This had led investigators to pursue other avenues, including a newer hallmark feature of neuroinflammation, where increases have been observed throughout the brain in individuals with AD (Bradburn et al., 2019; Hampel et al., 2020). Exactly how this neuroinflammation is accumulating remains to be fully understood, but two glial cells, microglia and astrocytes, both of which are involved inflammation, have come forward as potential major contributors (Webers et al., 2020). Microglia specifically have become the main focus of the neuroinflammation alongside their phagocytic capacity (Hemonnot et al., 2019); however, this focus tends to widely overlook infiltrating peripheral immune cells. It is well known that peripheral immune cells can contribute to neuroinflammation once they enter

the brain (Rezai-Zadeh et al., 2009; Prinz and Priller, 2017). Even in the wild type (WT) mouse brain, a complexity of immune cells have been noted, including monocytes, neutrophils, natural killer (NK) cells, T cells, and B cells, with microglia and astrocytes as the most abundant cells in the central nervous system (CNS) (Mrdjen et al., 2018). A “leaky” blood–brain barrier (BBB) has been observed in individuals with AD (Montagne et al., 2015, 2017; Sagare et al., 2015; van de Haar et al., 2016). Through the use of magnetic resonance imaging (MRI), increases in BBB leakage in the gray matter, hippocampus, and cortex has been shown to be significantly correlated with a decrease in cognitive function (Montagne et al., 2015; van de Haar et al., 2016; Nation et al., 2019). There was an increase in soluble platelet-derived growth factor receptor β , a marker of injury to pericytes, in the cerebral spinal fluid in these individuals, as well as in Tg2576 mice, indicating a breakdown in the BBB (Montagne et al., 2015; Sagare et al., 2015). Through the breakdown of the BBB in AD pathology, cells can infiltrate the brain and affect the pathology; these cells include peripheral macrophages, neutrophils, and T cells in the brains of both humans and mice, and NK cells and B cells in the mouse brain (Montagne et al., 2017; Mrdjen et al., 2018; Gate et al., 2020). However, how these cells interact with microglia in AD warrants further investigation. Here we will discuss current literature regarding microglial interaction with peripheral immune cells in the context of the different models of AD (Table 1) and the questions that remain unanswered.

Microglia in AD

Microglia are the main immune cells for the CNS. They are responsible for monitoring and maintaining the homeostatic environment through surveillance. Once microglia receive “alert” signals, they will react through the morphological altering of their cytoskeleton to respond to pathological events (Dionisio-Santos et al., 2019; Wyatt-Johnson and Brewster, 2020). Based on genome-wide associated studies, roughly 80% of gene changes that occur in AD are found within microglia, implicating them as critical in AD pathology (Marsh et al., 2016; Srinivasan et al., 2016; Zhang et al., 2016; Hansen et al., 2018). However, the exact role of microglia during the progression of AD remains unknown.

The magnitude to which microglia contribute to the AD brain has been up for debate. Microglia have also been observed alongside NFT in both human AD and in animal models (Zilka et al., 2012). Recent studies have suggested that microglia may be contributing to the buildup and accumulation of A β plaques (Baik et al., 2016; Spangenberg et al., 2019; Friker et al., 2020). Initially, alongside their contribution to inflammation, microglia will phagocytose A β clusters in the AD environment, which can lead to a buildup of intracellular A β (Baik et al., 2016). Early studies showed that A β can accumulate inside of microglia, as A β has been observed to be difficult for cells to degrade (Frackowiak et al., 1992). In the presence of an apoptotic-associated speck-like protein, microglia become unable to destroy A β and undergo pyroptotic death (Friker et al., 2020). This can add to the increase of A β plaques through the accrual of dead microglia (Baik et al., 2016). Furthermore, it has been shown that there is an increase in

RIPK1, a marker for necroptosis, co-localizing with ionized calcium binding adaptor molecule 1 (IBA1) + cells in both individuals with AD and APP/PS1 mice (Ofengeim et al., 2017). When RIPK1 is inhibited, there is a decrease in the A β plaque load, reductions in tumor necrosis factor alpha (TNF- α) and interleukin (IL)-1 β , and improved behavioral performance in reversal learning in the APP/PS1 mice (Ofengeim et al., 2017). This further supports the hypothesis that microglial death may be playing a role in the increased plaque formation. In APP/PS1/Trem2^{-/-} mice, the A β plaques are much more defused throughout the brain, with reduced microglial engulfment of the A β plaques (Jay et al., 2017). In models of neuronal injury, Trigger receptors expressed on myeloid cells 2 (Trem2) depletion in microglia limits their ability to migrate toward injured neurons (Mazaheri et al., 2017). Therefore, Trem2 may have a larger role in microglial migration toward A β plaques and their ability to contain them. Similar observations have been observed after depletion of microglia in APP/PS1 mice (Unger et al., 2018). These reports support the idea that microglia are aiding in the accumulation of the plaque formations found in AD.

Profile analysis of microglia at different stages of AD has also given more insight into their role in disease. Early in AD, microglia tend to be associated with increased activation and inflammation, with later phagocytic dysfunction (Hansen et al., 2018; Nordengen et al., 2019). A similar subset of microglia was observed in 18-month-old geriatric mice and 4-month-old APP/PS1 mice; these microglia were notably different from those in WT 4-month-old adult mice (Mrdjen et al., 2018). Microglia in both the geriatric WT and APP/PS1 mice had increased expression of the phagocytic associated markers, CD11c and CD14 (Mrdjen et al., 2018). Specifically, CD11c + microglia were located around A β plaques. Moreover, this subset of microglia, termed “disease-associated microglia” (DAMs), had increased expression of CD44 and CD86, and the inhibitory ligand programmed death-ligand 1 (PD-L1) as well as major histocompatibility complex class II (MHCII) molecules, whereas the homeostatic markers, MerTK, CX3C chemokine receptor 1, and Siglec-H were decreased (Mrdjen et al., 2018). A similar subset of microglia, which also had increases in Trem2 and apolipoprotein E (APOE), was observed in the 5xFAD mouse model of AD (Keren-Shaul et al., 2017). Interestingly, both of these genes have been determined to be AD risk factors (Jay et al., 2017; Keren-Shaul et al., 2017; Mrdjen et al., 2018; Shi and Holtzman, 2018; Zhou et al., 2018). The exact mechanism(s) by which DAMs may affect AD pathology and whether these cells are the result of pathological changes or contribute directly to the pathology requires further investigation. As the data suggest, microglia are not entirely responsible for all of the pathology of AD; this is evident by alterations present in other peripheral immune cells in the brain.

Astrocytes and Microglia in AD

Astrocytes are the other most well-known glial cell. They are involved in the homeostasis and maintenance of neurons and they do this through controlling the BBB and regulating neuronal communication (Matias et al., 2019). *In vitro* work has also

TABLE 1 | An overview of the pathological changes of different pre-clinical models of AD (Hsiao et al., 1996; Hsia et al., 1999; Dudal et al., 2004; Jankowsky et al., 2004; Grootendorst et al., 2005; Radde et al., 2006; Minkeviciene et al., 2009; Tai et al., 2011; Yue et al., 2011; Jawhar et al., 2012; Kamphuis et al., 2012; Cohen et al., 2013; Granger et al., 2016; Myers and McGonigle, 2019).

Models of AD	Genetic background	Mutations		Development (months)			
		APP	PSEN1	Plaque	Tangles	Gliosis	Memory deficit
5xFAD	Mixed: C57BL/6 and SJL	Swedish: K670N/M671L Florida: I716V London: V717I	M146L L286V	1.5	ND	1	4
	C57BL/6J			2	ND	2	3
APP/PS1	C57BL/6J	Swedish: K595N/M596L	L166P	3–4	NMT	1.5	7
APP _{swe} /PSEN1 Δ E9	C57BL/6J	Swedish: K670N/M671L	deltaE9	9	NMT	6	12
PSAPP	C57BL/6J* or Mixed: C57BL/6 and C3H	Swedish: K670N/M671L	M146L	6	6	6	3
PDGF-APP _{swe} , Ind J9	C57BL/6J	Swedish: K670N/M671L Indiana: V717F		21–25	NMT	ND	ND
PDGF-APP _{swe} , Ind J20	C57BL/6J	Swedish: K670N/M671L Indiana: V717F		7–8	Absent	6	4
Tg2576	C57BI/SJL	Swedish: K670N/M671L		11–13	Absent	10–16	6–12
TgCRND8	C57BI/6	Swedish: K670N/M671L Indiana: V717F		3	Absent	4	3
TgF344	Fischer 344	Swedish: K670N/M671L	deltaE9	6	16	6	15
Models of AD	Strain background	Other mutations		Plaque	Tangles	Gliosis	Memory deficits
APOE3-TR	C57BL/6J	ApoE replaced with human APOE3		10 (few)	ND	ND	ND
APOE4-TR	C57BL/6J	ApoE replaced with human APOE4		10 (few)	ND	ND	4–5
rTg4510	Mixed: 129S6 and FVB	Human tau MAPT ^{P301L}		Absent	2.5	2.5	2.5

app, amyloid precursor protein; *psen1*, presenilin 1; ND, no data; NMT, no mature tangles. *PSAPP mice bred on C57BL/6J have a seizure phenotype.

shown that astrocytes contribute to the production of pro-inflammatory cytokines (Matias et al., 2019). Crosstalk between astrocytes and microglia have been observed to induce microglial motility, microglia and astrocyte accumulation, and potentially both of their phagocytic capabilities (Fakhoury, 2018; Vainchtein and Molofsky, 2020). This cross-talk is thought to be through three main mechanisms: the communication between both cells directly, synchrony; both cell responding to injury and aiding in each other's response; and a relay effect, where one cell receives the signal and "relays" it onto the other cell (Vainchtein and Molofsky, 2020). Through RNA sequencing, APOE has been found to be upregulated in astrocytes as well as microglia (Grubman et al., 2019). Similar to microglia, astrocytes have also been found to be in a disease-associated state with an elevation in glial fibrillary acidic protein (GFAP) levels and increases in genes related to inflammatory signaling and response to toxic compounds (Habib et al., 2020). These toxic-reactive astrocytes, termed A1, have been shown to be induced by activated microglia through pro-inflammatory cytokines, and have been observed to be increased in the pre-frontal cortex in individuals with AD (Liddel et al., 2017). It has also been observed that astrocytes require TNF- α to become A1, which has been shown to be released from microglia, macrophages, and neutrophils (Tecchio et al., 2014; Liddel et al., 2017). In AD tissue, the loss of myelination of neurons was followed by an increase in GFAP expression (Han et al., 2019). This increase in GFAP was also found in close proximity to NFTs in both human AD and animal models (Fakhoury, 2018), with the highest expression of GFAP occurring in later tau pathology tissue (Gomez-Arboledas et al.,

2018). The depletion of tau in rTg4510 mice reduced the number of GFAP-, CD3-, and CD4-positive cells and prevented the loss of the BBB (Blair et al., 2015). In the TgF344 rat model of AD, there is an increase in astrocytes that produce a co-agonist of the N-methyl-D-aspartate receptor (NMDAR) on neurons, which could aid in their excitotoxicity and death (Balu et al., 2019). In GFAP[±]/APP_{swe}/PSEN1 Δ E9 mice, astrocytic levels are decreased with no change in plaque formation at 15 months and an increased number of IBA1 + cells (Kamphuis et al., 2015). This implicates astrocytes as a potential contributor to neuronal changes in AD pathology and microglial activation. More studies are necessary to further explore how microglia and astrocytes may work together to aid in the AD environment.

Macrophages and Microglia in AD

Macrophages are the chief phagocytes in the periphery (Patel et al., 2017; Ma et al., 2019). The main job of macrophages is to phagocytose pathogens and apoptotic cells. They are also very critical antigen-presenting cells (APCs) (Ma et al., 2019). Macrophages present peptide antigens via MHC I and II molecules. They also release signals to stimulate the migration of other cells to injured or infected tissue (Ma et al., 2019). Recently, a subset of macrophages has been found in the brain called brain associated macrophages (BAMs) (Mrdjen et al., 2018; Villacampa and Heneka, 2018). BAMs are located within the CNS in the border region areas, including the pia mater, perivascular space, choroid plexus, and dura mater and are replaced by circulating monocytes (Park et al., 2017; Mrdjen et al., 2018; Rajan et al., 2020). Through the use of mass cytometry, Mrdjen et al. (2018)

determined that these cells have an atypical surface receptor profile, with a high expression of MHCII but neither Ly6C, a myeloid marker, nor Siglec-H, a microglia marker (Mrdjen et al., 2018). They also found different expression profiles of the BAMS leading them into three different subsets based on their levels of CD38 and MHCII; these subsets were found in different regions of the brain. The dura mater had the highest expression of MHCII+ CD38- cells, while the pia mater and perivascular space had higher levels of MHCII+ CD38 + cells followed by those that were MHCII-CD38 + ; the choroid plexus had a mixture of all three subclasses (Mrdjen et al., 2018). The finding of BAMS and their unique and heterogeneous expression has been demonstrated in non-pathological and pathological conditions in mice, with similar findings observed in human brain tissue (Jordao et al., 2019; Rajan et al., 2020).

In PDGF-APP^{Sw}, Ind J20 mice, a genetic increase in perivascular BAMS, through the deletion of scavenger receptor class B, enhanced the development of A β deposits and exacerbated the cognitive deficits (Thanopoulou et al., 2010). Similar results were found in the TgCRND8 model of AD mice (Hawkes and McLaurin, 2009). When perivascular BAMS were stimulated, this led to a decrease in vascular levels of A β , whereas depletion of the BAMS led to an increase in A β (Hawkes and McLaurin, 2009). In 3-month-old Tg2576 mice, the number of BAMS was not altered compared to WT controls. However, when BAMS in the perivascular space were depleted with clodronate, that led to cerebrovascular dysfunction that was induced by A β (Park et al., 2017). Thus, this implicates BAMS in AD pathology.

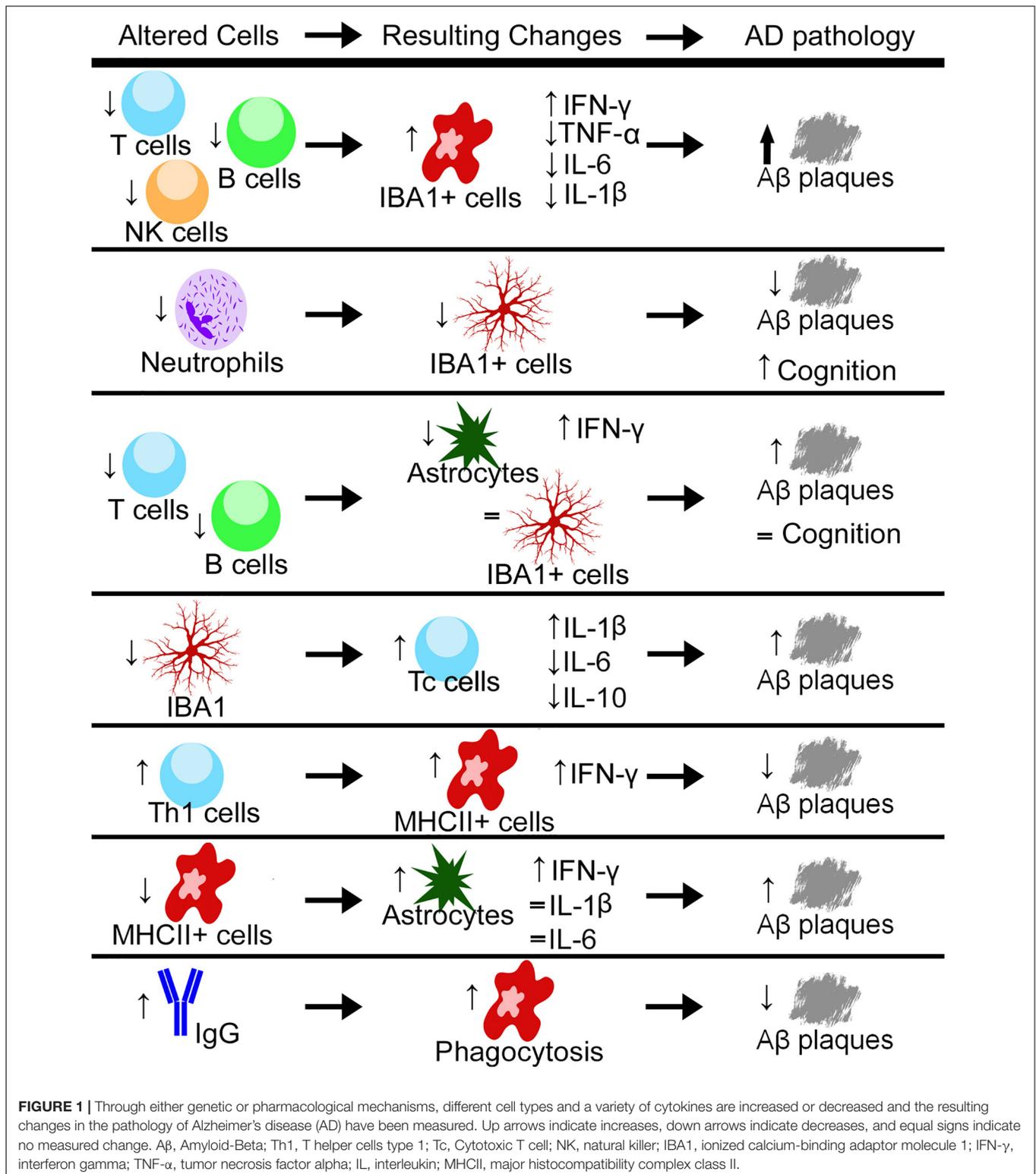
Currently, most studies focus on either microglia or macrophages or collectively analyze both of them together in AD. A reason for this is the similarity in phenotype between the two cell types, making it difficult to fully differentially distinguish them (Bennett et al., 2016; Jordao et al., 2019). However, recent advances in profiling microglia and macrophages have made it easier to look at each population individually (Bennett et al., 2016). For example, initial studies, using electron microscopy, analyzing both microglia and macrophages have shown that macrophages are more effective at eliminating A β plaques (Wisniewski et al., 1991). Moreover, the stimulation of toll-like receptor 9 (TLR9) by its ligand, CpG oligodeoxynucleotides (CpGODNs), caused an increase in the phagocytosis of A β plaques thought to be initially a microglial response. However, CpGODNs cannot easily penetrate the BBB, supporting the hypothesis that it is a macrophage response and not directly a microglial response (Scholtzova et al., 2009). During aging, microglia and macrophages have a directly opposite response (Rawji et al., 2016; Park et al., 2017; Friker et al., 2020); macrophages tend to produce less inflammatory cytokines in response to stimuli, whereas microglia exhibit increased production (Rawji et al., 2016). There has also been much debate about whether microglia or macrophages are associated with the formation of amyloid plaques. Knocking out macrophage chemokine receptor CCR2 in the Tg2576 mice, showed less accumulation of both microglia and macrophages (El Khoury et al., 2007), yet more recent studies have shown that it is specifically microglial cells—and not infiltrating macrophages—that are surrounding the plaques (Reed-Geaghan et al., 2020).

Together, these data suggest that in the AD brain, the microglia are potentially more impactful in the pathology compared to the infiltrating macrophages. More studies are necessary to differentiate how both cells are interacting in AD.

Neutrophils and Microglia in AD

Alongside macrophages, neutrophils are also responsible for phagocytosis but are mainly found in the bloodstream (Rosales, 2018). Typically, neutrophils are one of the first responders to injury or infection. Once they arrive, neutrophils recruit other immune cells to the site of injury and phagocytose microbes. Neutrophils can also increase their response in reaction to multiple signals, which can be beneficial, or in the case of chronic inflammatory disease, can be detrimental (Mayadas et al., 2014; Rosales, 2018; Wang, 2018). In healthy elderly subjects, there is reduced activity of neutrophils present in the blood compared to young adults (Vida et al., 2017). AD patients have an even greater reduction in neutrophil function in the later stages of disease; however, in the early stages of AD, there are more neutrophils present compared to age-matched healthy controls (Vida et al., 2017). The glutathione and oxidized glutathione ratio, an indicator of oxidative stress in cells, has been shown to be increased in the neutrophils from patients in the early stages of AD compared to age-matched healthy subjects; this ratio was further increased in late stages of severe AD and coupled with the increased release of pro-inflammatory cytokines (Vida et al., 2017). However, there was a decrease in anti-inflammatory cytokine production (Vida et al., 2017), together suggesting that neutrophils are altered in AD pathology.

Notably, neutrophils have been found in the brain during homeostatic conditions (Mrdjen et al., 2018). They can cross the BBB through the interactions between lymphocyte function-associated antigen 1 (LFA-1) found on circulating neutrophils and its receptor intercellular adhesion molecule 1 (ICAM-1). When LFA-1 is knocked out genetically (Itgal^{-/-}) or blocked by an antibody, neutrophils are unable to enter the brain (Zenaro et al., 2015). LFA-1-deficient 3xTg mice or normal 3xTg mice treated with an anti-LFA-1 mAb showed reduced cognitive impairments in the Y-maze and reduced accumulation of IBA1 + cells in the hippocampus along with reduced levels of A β plaques (Figure 1; Zenaro et al., 2015). Most frequently, neutrophils were found through live imaging that they traveled via diapedesis to areas with A β deposits and accumulated around A β plaques (Baik et al., 2014; Zenaro et al., 2015). Although A β plaques cannot recruit neutrophils directly, it has been shown that microglia in the AD pathology can recruit neutrophils to the brain (Baik et al., 2014; Park et al., 2019). *In vitro* and *in vivo* studies have suggested that the recruitment of neutrophils may lead to increased production of pro-inflammatory cytokines by microglia (Park et al., 2019). As prior studies have suggested, it is also likely that microglia are involved in the recruitment of neutrophils to A β plaques (Zenaro et al., 2015). Based on these reports, it has been proposed that neutrophils first engulf the A β plaques, and then microglia phagocytose the neutrophils. Further investigation is necessary to understand the relationship between microglia and neutrophils, and their engulfment of A β plaques.



NK Cells and Microglia in AD

NK cells are a subpopulation of cytotoxic lymphocytes that is a part of the innate immune response. NK cells can respond quickly upon activation, and they can kill cells without pre-stimulation

from other cells (Abel et al., 2018). NK cells initiate apoptosis in other cells through the release of cytotoxic granules; most commonly, granzyme B (Solana et al., 2018). They can also release inflammatory cytokines, including interferon-gamma (IFN- γ),

that stimulate other immune cells, especially macrophages, and contribute to chronic inflammation which is detrimental to the brain (Radde et al., 2006; Prinz and Priller, 2017). They can secrete immunosuppressive cytokines (Solana et al., 2018). During aging in healthy individuals, the cytotoxic activity of circulating NK cells is impaired relative to young adults (Vida et al., 2017). In a report studying individuals with mild AD, there was no change in NK cell activation capacity when comparing the expression of CD107a, a marker for granular release, and levels of granzyme B and IFN- γ (Le Page et al., 2015); however, it is not known how these cells may be altered in individuals with severe AD. In Rag2^{-/-}/Il2r γ ^{-/-} knockout mice crossed with 5xFAD mice, NK cells, T cells, and B cells do not develop and these mice have increased A β levels (Marsh et al., 2016). These mice also have elevations in the numbers of IBA1 + cells with an altered morphology, from ramified to bushy and amoeboid (Marsh et al., 2016). However, whether these effects are directly related to the lack of NK cells or the loss of the three lymphocyte subpopulations remains to be investigated.

In T- and B cell-deficient (but NK cell-sufficient) Rag2^{-/-} mice crossed with PSAPP mice, there is a decrease in the number of A β plaques (Spani et al., 2015). However, unlike that observed with the 5xFAD/Rag2^{-/-}/Il2r γ ^{-/-} mice, there was no change in intracellular IBA1 levels, but the number of astrocytes was decreased (Marsh et al., 2016; Mittal et al., 2019). There was also no recovery in performance in novel object recognition in the PSAPP/Rag2^{-/-} mice compared to the PSAPP mice (Figure 1; Spani et al., 2015). Together, these results indicate that NK cells may be more involved in IBA1 activation compared to T- and B cells (these cells are discussed below); however, this warrants further investigation with Il2r γ ^{-/-} deficient mice.

T Cells and Microglia in AD

T cells are a type of lymphocyte that responds to immune challenges as a part of the adaptive immune system (Mundt et al., 2019). There are three major types of T cells: cytotoxic (Tc; CD8+), helper (Th; CD4+), and regulatory (Treg; CD4+) (Schettters et al., 2017; Mundt et al., 2019). Tc cells recognize peptide antigens presented by MHC I molecules. These are found on all nucleated cells in the body, including those in the CNS. Upon antigen recognition, they release cytotoxic granules that kill the recognized cells (Mundt et al., 2019; Gate et al., 2020). Th cells are activated by peptides presented by MHC II molecules on professional APCs (macrophages, dendritic cells, and B cells) and then release specific cytokines depending on the Th cell subtype: Th1 cells produce pro-inflammatory cytokines that stimulate macrophages and cytotoxic T cells; Th2 cells secrete anti-inflammatory cytokines that activate B cells; Th17 cells produce the pro-inflammatory cytokine IL-17 and cytokines/chemokines that recruit neutrophils and macrophages. There are other Th cell subtypes (e.g., Th9, Neurath and Kaplan, 2017), but they will not be described here. Treg cells work to suppress immune cells when they are no longer needed or to maintain immune homeostasis (Schettters et al., 2017; Rostami et al., 2020). During aging, it has been observed that there is an increase in the overall number of T cells in the brain (Mrdjen et al., 2018). This has also been observed in both APP/PS1 and 5xFAD models of AD (Keren-Shaul et al., 2017; Mrdjen et al., 2018; Unger et al., 2018). In human cerebral

spinal fluid and brain tissue of individuals with AD, an increased number of Tc and Treg cells have also been found, respectively (Ciccocioppo et al., 2019; Gate et al., 2020). Tc cells have also been found to be significantly correlated with the increase in NFT and tau pathology in human AD tissue (Merlini et al., 2018).

Although most studies are using both male and female mice grouped together, one found differences in the number of T cells between APP/PS1 males and females, with females having overall more T cells with specific increases in Th and Tc cells (Unger et al., 2018). They also found that when microglia were ablated with PLX5622, there was an overall increase in the number of Tc cells in the APP/PS1 mice (Figure 1). This depletion did not have a gender effect (Unger et al., 2018). In 5xFAD mice, intracerebroventricular injection of A β -specific Th1 cells decreased plaque load and resulted in an increase in MHC II + cells. These IBA1+/MHCII+ cells were less ramified in appearance, with shorter branch length. They also had an increase in A β found inside these cells compared to those that were MHCII- (Mittal et al., 2019), indicating a more phagocytic phenotype induced in IBA1 + /MHCII + cells by the Th1 cells (Figure 1). When MHCII knockout mice were crossed with 5xFAD mice, there was a recovery in GFAP levels when compared with 5xFAD mice at 3 months of age. At 6 months of age, the 5xFAD/MHCII- mice displayed an overall increase in IL-1 β , IL-6, and IFN- γ compared to both 5xFAD and WT mice (Mittal et al., 2019). When A β -specific Th1 cells were injected into the 5xFAD/MHCII- mice, there was no variation in overall plaque load, but the levels of IFN- γ were increased; however, these levels did not reach those found in the 5xFAD mice (Mittal et al., 2019). Together, these data indicate that MHCII + /IBA1 + cells are activated by Th1 cells, potentially indicating a cooperative role between Th cells and microglia/macrophages.

There have also been multiple studies that have implicated the gut microbiota as a factor in affecting the infiltration of T cells into the brain (Minter et al., 2017; Dodiya et al., 2019, 2020). When APP_{SWE}/PSEN1 Δ _{E9} mice were given combinatorial antibiotics (ABX) that alter the gut microbiome, this led to attenuated A β plaque formation as well as increased Treg cells in the brain and blood (Minter et al., 2017). This treatment also reduced the number of plaque-localized microglia and astrocytes, as well as altered the morphology of the microglia, increasing the process length to a more ramified morphology in both APP_{SWE}/PS1 Δ _{E9} mice (Minter et al., 2017; Dodiya et al., 2019). However, these effects were not observed in female APP_{SWE}/PS1 Δ _{E9} mice (Dodiya et al., 2019), indicating the importance of understanding both male and female immune responses. Further work is needed to distinguish between the roles of the different types of T cells in AD and the cognitive consequences of the adoptive transfer of the T cells into the AD mouse brain and to understand AD development in a T cell-deficient environment.

B Cells and Microglia in AD

B cells are involved in the adaptive immune response by producing antibodies that coat target cells, allowing for easier engulfment by macrophages (Tarlington, 2019). B cells can release pro-/anti-inflammatory cytokines that may contribute to chronic inflammation (Nikolajczyk, 2010). As aging occurs,

B cells become less able to produce antibodies, and in chronic inflammation, progenitor B cells are more likely to undergo programmed cell death (Ratliff et al., 2015; Hagen and Derudder, 2020). In both moderate and severe AD, there is an overall decrease in B cells (Bulati et al., 2015). In 5xFAD mice, a whole-brain analysis showed an increase in the number of mature B cells compared to WT (Keren-Shaul et al., 2017), whereas in female APOE4-TR mice, there was a decrease in the expression of B cells in the hippocampus (Zhang et al., 2019). It has also been observed that B cells secrete antibodies that are A β -specific (Schenk et al., 1999). Here, immunization against A β reduced plaques and astrocytic accumulation (Schenk et al., 1999). However, when this was tried in human clinical trials, the immunization caused a T cell response leading to life-threatening complications (Cao and Zheng, 2018). In 5xFAD mice, elevated levels of IgG are present, and with IgG localizing in and around IBA1 + cells (Marsh et al., 2016). These increased levels are absent in 5xFAD/Rag2^{-/-}/Il2r γ ^{-/-} mice (Marsh et al., 2016). Exposure to preimmune IgG elicits a phagocytic response by BV-2 microglia to engulf A β (Marsh et al., 2016). When preimmune IgG is injected into the brain in both the 5xFAD and APP/PS1 mice, there is a reduction in A β levels (Figure 1; Sudduth et al., 2013; Marsh et al., 2016). Also, WT mouse bone marrow transplanted into 5xFAD/Rag2^{-/-}/Il2r γ ^{-/-} mice resulted in an increase in brain IgG levels and a concurrent reduction in plaque volume (Marsh et al., 2016). In APOE3-TR mice, there is no change in the number of B cells harvested from the blood; however, there is an increase in IgG levels across the neocortex, entorhinal cortex, hippocampus, thalamus, and cerebellum (Zhang et al., 2019). This increase in IgG staining was mostly localized to microglia and not astrocytes (Zhang et al., 2019). However, early studies showed B cells harvested from individuals with AD were more hyperactivated to NFT and A β plaques (Kingsley et al., 1988; Xu and Gaskin, 1997). These B cells that reacted with NFT also secreted antibodies that reacted highly with astrocytes (Kingsley et al., 1988). This difference between the secreted antibodies reacting with astrocytes and also co-localizing with microglia could be due to the varied response of B cells in humans with AD compared to animal models of AD. The increase in IgG that is observed, could also be infiltrating from the disrupted BBB. Studies using a drug, 3K3A-APC, that protects cerebrovascular endothelium cells from damage, showed reduced IgG levels in the cortex of 5xFAD mice (Williams et al., 2012; Lazic et al., 2019). It should also be noted that 3K3A-APC

inhibits BACE1 in neurons and treatment of 5xFAD mice with this drug decreases A β levels, as well as reduces the number of microglia and astrocytes (Lazic et al., 2019). Taken together, these data indicate a relationship between B cell production of IgG, IgG infiltration, and phagocytosis, but exactly how IgG is altering microglia requires more investigation.

CONCLUDING REMARKS

In AD, current studies focus on modulating A β plaque formation and altering the local neuroinflammation; however, other important contributors can consume A β plaques and induce neuroinflammation—the peripheral immune cells. Investigations on AD have analyzed microglia either directly or indirectly with macrophages (Hawkes and McLaurin, 2009; Thanopoulou et al., 2010; Park et al., 2017; Mrdjen et al., 2018; Reed-Geaghan et al., 2020), neutrophils (Baik et al., 2014; Zenaro et al., 2015; Vida et al., 2017; Park et al., 2019), NK cells (Marsh et al., 2016; Vida et al., 2017), T cells (Mrdjen et al., 2018; Unger et al., 2018; Ciccocioppo et al., 2019; Mittal et al., 2019; Gate et al., 2020), and B cells (Sudduth et al., 2013; Marsh et al., 2016; Zhang et al., 2019). However, more questions remain in order to increase our understanding of how these cell types may be interacting and contributing to AD pathology as well as the associated cognitive decline. It also remains elusive if the peripheral immune cells must enter the brain to affect the AD pathology. Future studies are necessary to understand these relationships, in terms of how the development of AD occurs and how the pathology continues to become exacerbated, to determine the best course of treatment for AD patients.

AUTHOR CONTRIBUTIONS

SW-J performed the literature search and wrote the manuscript. RB critically revised the work. Both authors contributed to the idea for the article.

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Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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