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# The "Quad-partite" Synapse: Microglia-Synapse Interactions in the Developing and Mature CNS

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#### Abstract

Microglia are the resident immune cells and phagocytes of our central nervous system (CNS). While most work has focused on the rapid and robust responses of microglia during CNS disease and injury, emerging evidence suggests that these mysterious cells have important roles at CNS synapses in the healthy, intact CNS. Groundbreaking live imaging studies in the anesthetized, adult mouse demonstrated that microglia processes dynamically survey their environment and interact with other brain cells including neurons and astrocytes. More recent imaging studies have revealed that microglia dynamically interact with synapses where they appear to serve as "synaptic sensors," responding to changes in neural activity and neurotransmitter release. In the following review, we discuss the most recent work demonstrating that microglia play active roles at developing and mature synapses. We first discuss the important imaging studies that have led us to better understand the physical relationship between microglia and synapses in the healthy brain. Following this discussion, we review known molecular mechanisms and functional consequences of microglia-synapse interactions in the developing and mature CNS. Our current knowledge sheds new light on the critical functions of these mysterious cells in synapse development and function in the healthy CNS, but has also incited several new and interesting questions that remain to be explored. We discuss these open questions, and how the most recent findings in the healthy CNS may be related to pathologies associated with abnormal and/or loss of neural circuits.

#### Keywords

Glia; Pruning; Plasticity; Neurotransmission; Cortex; Hippocampus; Retinogeniculate

### Introduction

Nearly a century has passed since the concept of microglia was originally described in the healthy mammalian CNS by Pio del Rio Hortega (del Rio-Hortega 1932). Surprisingly, there is still very little known regarding the role of these mysterious cells in the normal, healthy CNS. A vast majority of studies have focused on the role of microglia in the context of disease, in which they display an extraordinary ability to respond rapidly and perform a broad range of functions such as shielding injury sites, phagocytosing cellular material, and

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releasing inflammatory signals to initiate and/or propagate the immune response. While these rapid and robust responses to CNS injury and disease make microglia interesting candidates for developing diagnostic and therapeutic strategies, these qualities also make microglia very difficult to study in the context of the healthy brain. Culture models such as dissociated cells and brain slices can mimic injury resulting in 'activation' of microglia, a shift toward a graded cellular state most associated with injury and characterized by morphological characteristics more closely resembling peripheral macrophages (amoeboid, rod-like, etc.) as well as functional changes such as increased phagocytic capacity, chemotaxis, proliferation, and expression of pro-inflammatory molecules (Kettenmann et al. 2011; Ransohoff and Perry 2009). Thus, while *in vitro* studies are valuable for understanding some of the basic cellular biology, *in vivo* strategies must also be utilized to understand the function of these cells in the context of the healthy, intact CNS.

In 2000, a transgenic mouse was engineered that enabled visualization of microglia by EGFP (CX3CR1+/EGFP)(Jung et al. 2000). These mice, combined with new technology to image the brain in a live, anesthetized mouse by 2-photon microscopy, resulted in groundbreaking findings in 2005. Two studies used a thin-skulled transcranial approach and demonstrated that, in contrast to their name, 'resting' microglia in the healthy, adult cerebral cortex continuously survey their surrounding extracellular environment (Davalos et al. 2005; Nimmerjahn et al. 2005). Time-lapse imaging revealed that microglia rapidly extend and retract their processes on the order of minutes while their cell bodies remain stationary. As a result, within a few hours, microglia processes have the capacity to sample the entire brain parenchyma and physically interact with other cortical cells including astrocytes and neurons (Nimmerjahn et al. 2005). These key findings beg the question, what is the function of microglia in the healthy brain? The following review will focus on the most recent studies that suggest that microglia play important roles at synapses in the developing and mature CNS and that synaptic activity may concomitantly modulate microglia function.

## I. Imaging activity-dependent interactions between microglia and synaptic circuits

The pioneering imaging studies conducted in 2005 laid a foundation for more work focused on the interactions between microglia and synapses in response to spontaneous and sensory experience-driven changes in neuronal activity. The following sections will discuss key imaging studies demonstrating that microglia have the capacity to rapidly respond to changes in neurotransmitters and physically interact with synapses in an activity-dependent manner.

#### Imaging the effects of neurotransmission on microglial dynamics

*In vitro* preparations of microglia demonstrate that microglia have the capacity to express receptors for and respond to neurotransmitters such as acetylcholine, gamma-aminobutyric acid (GABA), glutamate, and purinergics including adenosine-5'-triphosphate (ATP). Such studies have demonstrated that neurotransmitters can affect microglia *in vitro* in numerous ways, including changes in membrane potential, intracellular calcium, cytokine release, and overall cellular motility, reviewed in (Biber et al. 2007; Kettenmann et al. 2011; Pocock and

Kettenmann 2007). Since these original studies, several live imaging studies have demonstrated the capacity of microglia to rapidly respond and change their dynamics in response to neurotransmitters.

Of the many neurotransmitters known to elicit an effect, in vitro and in vivo preparations alike have demonstrated that ATP is one of the most potent signals to elicit a microglial response (Farber and Kettenmann 2006; Inoue et al. 2007). In 2005, Davalos et al. demonstrated that purinergic signaling increased basal microglia motility in vivo. In the anesthetized mouse, the authors applied either apyrase, which hydrolyzes ATP and ADP, or pharmacological blockers of gap junction hemichannels (carbenoxolone or flufenamic acid) to the cortical surface followed by live imaging (Davalos et al. 2005). In all instances, disrupting ATP-dependent signaling resulted in reduced basal velocity of microglial processes. More recently, using mice harboring a deletion in the purinergic receptor P2Y<sub>12</sub> or  $A2_A$ , it has been demonstrated *in vitro* and *in vivo* that purinergic signaling through these receptors can mediate microglial process extension and chemotaxis or retraction, respectively, in response to local nucleotide application or lipopolysaccharide (LPS)induced activation (Haynes et al. 2006; Orr et al. 2009). However, it remains to be determined whether purinergic signaling can regulate microglia under more basal, physiological conditions in the absence of local nucleotide or LPS application. Interestingly, a more recent live imaging studying using ex vivo retinal explants revealed that glutamatergic transmission increased microglia motility indirectly via increased ATP release upon neuronal excitation, data suggesting a role for ATP in regulating microglia dynamics under more basal conditions (Fontainhas et al. 2011).

Similar to purinergic and glutamatergic signaling, inhibitory neurotransmission has also been implicated in regulating microglial dynamics. In the same *ex vivo* study in the retina described above, which demonstrated effects of glutamate and ATP, the inhibitory neurotransmitter, GABA, decreased microglia process motility and overall velocity, while the GABAA receptor antagonist, bicuculline, increased motility and velocity. These data are similar to the *in vivo* work by Nimmerjahn et al. (2005) that showed a similar increase in motility upon bicuculline application; however, overall velocity was unchanged, data that may reflect differences between *ex vivo* and *in vivo* preparations and/or region analyzed (retina vs. cortex).

Overall, the studies discussed above suggest that increased excitatory neurotransmission and purinergic signaling can cause an increase in microglial process motility, while inhibitory neurotransmission may decrease this motility. However, one recently published study by Grinberg et al. (2011) used an *in vitro* hippocampal slice preparation to assess microglial dynamics in response to spreading depression and have results which suggest a seemingly opposite result. When long term potentiation (LTP) was induced by increasing cyclic adenosine monophosphate (cAMP), enhanced neural transmission induced a decrease in microglial movement (Grinberg et al. 2011). Conversely, blockade of activity with tetrodotoxin (TTX) resulted in increased movement, and restoration of activity with glutamate or ATP subsequently decreased movement. However, these results may reflect the *in vitro* nature of this sample preparation, regional differences in neurotransmission (e.g., hippocampus vs. retina or cortex), regional microglial heterogeneity, and/or differences in

stimulation paradigms (e.g., cAMP or TTX  $\pm$  glutamate or ATP vs. direct application of neurotransmitters or antagonists).

Despite discrepancies, it is clear that microglia are responding to changes in neurotransmission and that neurotransmitters can have direct and rapid effects on the overall dynamics of these cells. Future *in vivo* work will be required to further dissect the nature of microglia responses to excitatory and inhibitory neurotransmission as well as identify whether these effects have regional specificity.

#### Imaging physical interactions between microglia and synapses

The imaging studies discussed above suggest that microglia can rapidly change their dynamics in response to neurotransmission. In order to better understand how microglia may be physically interacting with synapses in paradigms known to change neural activity and induce synaptic remodeling, recent work has taken advantage of 2-photon live imaging in the anesthetized mouse, in which microglia and neurons are fluorescently labeled (Tremblay 2012). The first imaging study to characterize physical interactions between microglia and synaptic elements in vivo used 2-photon time-lapse imaging and transgenic mice expressing EGFP in both microglia and neurons (Iba-1-EGFP/Thy1-EGFP M line) (Feng et al. 2000; Hirasawa et al. 2005; Wake et al. 2009). This study revealed that microglial processes briefly (~5 min) contacted synaptic elements in layers II-III of the somatosensory and visual cortices at a rate of ~1 structure per hour and that microglial processes appeared to enlarge once presynaptic terminals were contacted. Following these baseline measurements, the authors investigated how changes in neural activity might affect physical interactions between microglia and synaptic elements. Using 3 independent methods to decrease neural activity (enucleation, retinal TTX administration, or lowered body temperature), the authors observed that microglial processes retracted and made fewer contacts with less active presynaptic terminals within the cortex. These data are in discordance with previously published work demonstrating no change in microglial motility in response to direct cortical application of TTX in vivo (Nimmerjahn et al. 2005) or increased microglial movement in response to TTX administration in an *in vitro* slice preparation (Grinberg et al. 2011). However, these discrepancies may easily be explained by method of TTX administration, regional differences, and/or in vivo vs. in vitro preparations. To further assess interactions between microglia and synapses, Wake et al. (2009) induced ischemic injury by photochemical occlusion of the middle cerebral artery (MCA) and demonstrated that, under these conditions, microglia contact with presynaptic structures was prolonged and was often accompanied or followed by a disappearance of these structures. These data suggest a potential function for microglia in the removal of synapses undergoing, albeit injuryinduced, remodeling.

In order to address whether microglia were indeed playing a role at remodeling synapses under more physiological conditions, Tremblay et al. published a recent study assessing *in vivo* changes in microglia-synapse interactions in the developing primary mouse visual cortex (V1)(Tremblay et al. 2010). This study was able to attain better spatial resolution than previous studies by combining high resolution 3-D serial electron microscopy (3-D serial EM) with 2-photon, transcranial imaging. In addition, in contrast to Wake et al. (2009),

Tremblay et al. (2010) were better able to distinguish processes belonging to microglia versus neurons by using a transgenic mouse line expressing EGFP in microglia and YFP in neurons (CX3CR1+EGFP/Thy-1 YFP). This study utilized the critical period in the mouse visual system to assess sensory experience-induced changes in microglia-synapse interactions (Gordon and Stryker 1996; Tremblay et al. 2010). During this particular period in visual cortex development (postnatal day 21-30 (P21-30) in the mouse), several aspects of visual perception develop (e.g., direction selectivity, ocular dominance, etc.) and are associated with changes in spine dynamics, size, and number (Bence and Levelt 2005; Hooks and Chen 2007; Majewska and Sur 2003). During the peak of the critical period (P28), high resolution imaging demonstrated that microglia in layer II of V1 normally contacted spines, synaptic terminals and synaptic clefts (Tremblay et al. 2010). Importantly, this landmark study revealed for the first time that spines often changed size upon microglia contact, suggesting that microglia may be key regulators of structural spine plasticity. Consistent with this idea, the authors noted that spines which changed size upon microglial contact tended to be smaller and, during later imaging sessions, were often eliminated. These data raise the intriguing possibility that microglia may mediate the elimination of spines in response to sensory experience and suggest one potential mechanism could be phagocytosis.

The idea that glial cells could be phagocytosing circuits undergoing active remodeling is substantiated by earlier work in the *Drosophila* mushroom body and mammalian neuromuscular junction (NMJ) (Freeman 2006; Mallat et al. 2005). In developing *Drosophila*, glial cells engulf the axons of mushroom body  $\gamma$  neurons undergoing anatomical pruning. In addition, in the mammalian system, Schwann cells phagocytose axonal arbor remnants (i.e., "axosomes") undergoing pruning at the developing NMJ (Bishop et al. 2004). Consistent with these data, glia have been shown to engulf destabilized synaptic boutons and presynaptic debris at the developing Drosophila NMJ (Fuentes-Medel et al. 2009). However, except for early work suggesting that microglia and astrocytes could be engulfing axonal debris during large-scale cortical pruning (Berbel and Innocenti 1988), very little was known regarding glial-mediated phagocytosis of mammalian CNS synapses undergoing local, small-scale remodeling. Thus, to further assess physical interactions between microglia and remodeling CNS synapses, Tremblay et al. used a dark adaptation paradigm in which juvenile mice were placed in the dark for 6 days during the visual system critical period (Tremblay et al. 2010). In some cases, mice were then re-exposed to light for 2 days, a paradigm known to elicit synaptic remodeling in V1 (Mower et al. 1983; Philpot et al. 2001; Tropea et al. 2010; Viegi et al. 2002). Although microglia in V1 contacted synapses in a similar manner in dark-adapted mice  $\pm$  light exposure as compared to normal, light exposed mice, the authors provided exciting new evidence that microglia had more phagocytic inclusions upon dark adaptation + light exposure, which resembled pre- and postsynaptic elements. Consistent with these data, a more recent study by the same group demonstrated an age-dependent increase in phagocytic inclusions resembling pre- and postsynaptic elements in either V1 or primary auditory cortex (A1) in association with agedependent loss of vision or hearing, respectively (Tremblay et al. 2012). Furthermore, another recent study used stimulated emission depletion (STED) microscopy and postembedding EM to assess microglia-mediated phagocytosis of synapses in the developing

hippocampus (Paolicelli et al. 2011), data that further suggest that microglia may engulf remodeling synaptic elements.

The data described above suggest that microglia are phagocytosing synaptic elements and that this interaction may underlie synaptic remodeling in response to changes in sensory experience. In order to more directly address whether microglia are phagocytosing synapses undergoing activity-dependent synaptic remodeling, we recently (Schafer et al. 2012) assessed phagocytosis in a system in which activity-dependent synaptic remodeling has been very well characterized: the postnatal retinogeniculate system. In this system, retinal ganglion cells (RGCs) form synapses on relay neurons residing in the dorsal lateral geniculate nucleus (dLGN) of the thalamus and these synapses are known to undergo activity-dependent remodeling during a very small developmental window (see Section III for details regarding retinogeniculate remodeling) (Guido 2008; Hong and Chen 2011; Huberman 2007; Sretavan and Shatz 1986). Using this system, we developed a high throughput in vivo phagocytic assay to demonstrate that microglia phagocytose presynaptic RGC inputs during a peak period in retinogeniculate synaptic remodeling (P5 in mouse), data that were validated by several different light and ultrastructural assays (Schafer et al. 2012). By EM, this phagocytosed material had characteristic features of presynaptic terminal machinery such as 40 nm vesicles; however, engulfment of axonal arbors cannot be excluded. Unlike previous work, this study observed little evidence suggesting engulfment of postsynaptic machinery, which may be due to regional differences, particularly given that RGCs synapse on the cell body and proximal dendrites of postnatal dLGN relay neurons (Guido 2008). We further demonstrated that engulfment and synaptic remodeling were temporally correlated such that as remodeling was nearly complete, engulfment of RGC inputs was also dramatically reduced.

In addition to assessing developmental regulation, we developed an *in vivo* assay to test whether microglia synapse-interactions are affected by activity-dependent synaptic competition. Activity-dependent competition was established by injecting one eye with either TTX to silence firing or forskolin, a cAMP analog, to increase firing (Dunn et al. 2006; Stellwagen and Shatz 2002; Stellwagen et al. 1999). The other eye was injected with vehicle. After pharmacologically decreasing (TTX) or increasing (forskolin) firing in one eye, data revealed that microglia preferentially engulfed inputs originating from the 'weaker' or less active eye (Schafer et al. 2012). Importantly, these particular pharmacological manipulations are known to disrupt normal synaptic remodeling such that inputs from the 'weaker' or less active eye lose territory and inputs from the 'stronger' or more active eye gain territory (Cook et al. 1999; Del Rio and Feller 2006; Huberman et al. 2008; Penn et al. 1998; Shatz 1990; Shatz and Stryker 1988; Stellwagen and Shatz 2002). Thus, these data suggest that microglia are dynamic sensors during activity-dependent synaptic remodeling and may be actively engulfing synapses destined for elimination.

Taken together, these provocative studies demonstrate that microglia, indeed, interact with and engulf synaptic elements and that microglia-synapse interactions are dependent upon neural activity and sensory experience. Importantly, these studies raise new and interesting questions including: 1) What are the molecular mechanisms underlying microglia-synapse

interactions? and 2) What are the physiological consequences of these interactions at remodeling synapses?

## II. Microglia-synapse interactions in the developing CNS: Molecular mechanisms and functional consequences

The mature nervous system is characterized by a remarkably precise neural circuitry. However, in the developing nervous system, this circuitry is far less precise as neurons during early development form exuberant transient synapses. In a process termed synaptic pruning, many synapses are eliminated while the remaining synapses are maintained and strengthened. While it is clear that neural activity regulates pruning as well as synapse maturation (Hua and Smith 2004; Huberman et al. 2008; Katz and Shatz 1996; O'Leary and McLaughlin 2005; Sanes and Lichtman 1999; Torborg and Feller 2005), little is known regarding cellular and molecular mediators. Interestingly, several key papers published in the last decade have identified a critical role for molecules traditionally associated with immune function (MHC class I molecules and receptors, complement components and receptors, and neuronal pentraxins) as modulators of developmental synaptic pruning (Bjartmar et al. 2006; Boulanger 2009; Corriveau et al. 1998; Datwani et al. 2009; Goddard et al. 2007; Huh et al. 2000; Schafer and Stevens 2010; Stevens et al. 2007; Syken et al. 2006). Taken together with the emerging evidence that activity regulates microglia at synaptic sites (Schafer et al. 2012; Tremblay et al. 2010; Wake et al. 2009), it was hypothesized that microglia, the resident CNS immune cells, may be cellular mediators of activity-dependent synaptic pruning and maturation. To determine whether microgliasynapse interactions in the developing brain have physiological consequences and dissect molecular mechanisms underlying these interactions, recent studies have utilized mice harboring deletions of genes specific to microglia in the context of the healthy developing CNS (Paolicelli et al. 2011; Pascual et al. 2011; Roumier et al. 2004; Roumier et al. 2008; Schafer et al. 2012).

#### A role for microglia in developmental synaptic pruning

One molecular pathway that has been proposed as a potential mediator of microglia-synapse interactions and developmental synaptic pruning is the classical complement cascade (Alexander et al. 2012; Schafer and Stevens 2010; Stevens et al. 2007). Molecules belonging to the classical complement cascade, C1q and C3, are localized to synaptic compartments and mediate synaptic pruning in the developing retinogeniculate system (Stevens et al. 2007). In the innate immune system, C1q and/or C3 bind cellular material inducing its removal by several different mechanisms including phagocytic pathways (Gasque 2004; Lambris and Tsokos 1986; van Lookeren Campagne et al. 2007). Thus, an intriguing hypothesis has been proposed that complement proteins bind synapses, which are subsequently eliminated by the primary CNS phagocyte, microglia. Recently, we demonstrated *in vivo* that complement component C3, enriched in synaptic compartments, and its receptor, complement receptor 3 (CR3), expressed on the surface of microglia, mediated engulfment of presynaptic terminals in the developing retinogeniculate system during a peak period in synaptic remodeling (P5 mouse dLGN) (Schafer et al. 2012).

of complement components (C1q and C3) and CR3-dependent engulfment of neurites by microglia. (Linnartz et al. 2012). These data are consistent with the idea that complement proteins may be 'tagging' synapses for removal by complement-receptor expressing microglia (Figure 1, panel B). However, it remains to be determined in vivo whether complement components are regulated by activity and localized to synapses destined for elimination by microglia-mediated engulfment. However, because engulfment of RGC inputs was reduced by 50% in C3 and CR3 KO mice, data suggest that other phagocytic pathways must be involved (Schafer et al. 2012). Interesting candidates may be proteins belonging to the "find-me", "eat-me," and "don't-eat-me" signaling pathways traditionally associated with engulfment of apoptotic cells, for review see (Elward and Gasque 2003; Griffiths et al. 2009; Grimsley and Ravichandran 2003; Nagata et al. 2010; Ravichandran 2011). For example, the Drosophila Draper (CED-1 in C. elegans and MEGF10 in mammals) and CED-6 (GULP in mammals) pathways have been demonstrated to be involved in the removal of axonal debris during developmental axon pruning and following axonal injury (Awasaki et al. 2006; Fuentes-Medel et al. 2009; Logan et al. 2012; MacDonald et al. 2006; Ziegenfuss et al. 2008). It remains to be determined whether these pathways may also be involved in the mammalian system.

Given that C3 and CR3 KO mice had a deficit in engulfment of remodeling RGC inputs, we directly addressed the role of microglia in developmental synaptic pruning by assessing C3 and CR3 KO mice for pruning defects in the developing retinogeniculate system. Early in development, RGCs in the retina project to and form exuberant, transient synaptic connections within the early dLGN of the thalamus (Guido 2008; Hong and Chen 2011; Huberman 2007; Sretavan and Shatz 1986). Within the first week of rodent postnatal development, RGC synaptic inputs compete for territory, resulting in the elimination of many transient synaptic connections and the maintenance and strengthening of those synapses that remain. This competition can occur between synapses originating from the same eye (monocular) as well as between synapses originating from different eyes (binocular). One gross assessment for deficits in retinogeniculate pruning is eye specific segregation, in which inputs from both eyes compete for territory, ultimately resulting in the termination of ipsilateral and contralateral synaptic inputs in distinct non-overlapping domains in the mature dLGN (Godement et al. 1984; Guido 2008; Huberman et al. 2008; Jaubert-Miazza et al. 2005; Sretavan and Shatz 1986; Ziburkus and Guido 2006). Using this system, pharmacological (minocycline) or more specific genetic (C3 or CR3 KO) disruptions in microglia function resulted in deficits in eye specific segregation as well as an increased density of structurally intact synapses (Schafer et al. 2012; Stevens et al. 2007). Importantly, these effects were sustained into adulthood and, in the case of the CR3 KO, could be specifically attributed to microglia in the context of the healthy developing brain. One question that remains unanswered is whether microglia-dependent pruning effects represent an active or passive process. That is, while C3/CR3-mediated engulfment and pruning data are temporally correlated, it remains to be determined whether microglia mediate pruning by actively engulfing intact terminals destined for elimination via C3/CR3 signaling or whether C3/CR3-mediated engulfment represents a 'clean-up' process independent of C3/CR3-dependent pruning. In addition, future work should aim to assess whether these effects are specific to the retinogeniculate system or have broader

implications throughout the CNS and whether other immune pathways previously identified to play a role in retinogeniculate pruning and plasticity throughout the CNS (e.g., MHC class I molecules) may interact with complement and/or microglia to mediate synaptic pruning (Corriveau et al. 1998; Datwani et al. 2009; Goddard et al. 2007; Huh et al. 2000).

#### A role for microglia in synapse maturation in the developing CNS

Consistent with microglia having a broader role in synapse development, recent work in the developing cortex and hippocampus has suggested a role for microglia in remodeling and/or maturation of synaptic circuits (Paolicelli et al. 2011; Roumier et al. 2004; Roumier et al. 2008; Tremblay et al. 2010). For example, Paolicelli et al. (2011) demonstrated a role for the fractalkine receptor (CX3CR1), expressed specifically on the surface of microglia in the developing CNS (Harrison et al. 1998), in hippocampal synapse development. In the context of disease, CX3CR1 has the capacity to modulate microglia number, activation, and recruitment to sites of injury by binding its ligand fractalkine (CX3CL1), expressed by injured neurons (Cardona et al. 2006; Jung et al. 2000). In the healthy, developing hippocampus, *Cx3cr1*<sup>KO</sup> mice exhibited an increase in spine density and PSD-95 immunoreactivity, enhanced hippocampal long-term depression (LTD), and decreased duration of and latency to pentylenetetrazol (PTZ)-induced seizure response, characteristics associated with less mature synapses and possibly associated with abnormal pruning (Paolicelli et al. 2011). However, postnatal (P15) *Cx3cr1<sup>KO</sup>* mice had no deficits in engulfment of postsynaptic elements. In contrast, the authors demonstrated that fewer microglia were present in the postnatal Cx3cr1KO hippocampus as compared to age-matched wild-type controls. Importantly, abnormalities in microglia number and synapse density in postnatal Cx3cr1<sup>KO</sup> mice returned to normal levels by adulthood. Thus, in the context of the developing CNS, it is most likely that fractalkine signaling regulates microglia activation, number, and/or recruitment to synaptic sites in the early postnatal brain (Figure 1, panels A&B). It remains to be determined if microglia play any sustained functional role at these synapses. In contrast to Paolicelli et al. (2011), another recent study demonstrated different defects in microglia and synaptic function in the mature (3 month old) hippocampus of  $Cx3cr1^{KO}$  mice (i.e., increased microglia numbers and reduced LTP: see Section III for more details) raising the possibility that fractalkine signaling may play different roles in the developing and mature brain (Figure 1, panels A-C) (Rogers et al. 2011).

Consistent with microglia playing a role in synapse maturation in the hippocampus, earlier *in vitro* studies demonstrated that maturation of hippocampal synapses was altered in mice harboring a mutation in KARAP/DAP12 (DAP12<sup>KI</sup>), a transmembrane receptor expressed by microglia from birth and a known regulator of macrophage activation in the immune system (Hamerman et al. 2005; Roumier et al. 2004; Roumier et al. 2008; Tomasello et al. 2000; Tomasello et al. 1998; Turnbull et al. 2005). Acute hippocampal slices prepared from P22 DAP12<sup>KI</sup> mutant mice had increased NR2B-containing N-Methyl-D-aspartate (NMDA) receptors as assessed by ifenprodil sensitivity and increased 2-amino-3-(5-methyl-3-oxo-1,2-oxazol-4-yl) propanoic acid (AMPA) receptor calcium permeability, characteristics indicative of less mature synapses (Roumier et al. 2004). In a later study, acute hippocampal slices prepared from P18-P25 DAP12<sup>KI</sup> mice had increased AMPA/NMDA ratios and, while most synapses appeared normal by EM, there was an observed increase in perforated

synapses in the stratum radiatum of CA1 (Roumier et al. 2008). In order to address whether this effect was a prenatal developmental defect, the authors prepared near-pure dissociated hippocampal neuron cultures from P0 DAP12<sup>KI</sup> mice and from mice subjected to *in utero* inflammation with LPS. After 14 days in culture (DIV 14), functional calcium imaging revealed significantly reduced calcium fluctuations in the presence of the AMPAR blocker 6-cyano-7 nitroquinoxline-2 (CNQX) in cultures prepared from P0 mutant and LPS-treated mice as compared to wild-type (WT) or vehicle-treated mice. In addition, there was an increase in immunohistochemical colocalization of pre- and postsynaptic markers in cultures prepared from DAP12<sup>KI</sup> mice. Because DAP12 expression is specific to microglia and cultures were presumably void of microglia, these data suggest that the physiological and structural deficits are the result of a developmental defect in synapse maturation that originates prenatally (Figure 1, panel A) (Roumier et al. 2008). However, because the oldest age at which synapse function was assessed was P25 and most experiments were performed *in vitro*, it remains to be determined whether deficits in DAP12<sup>KI</sup> hippocampal synapse maturation represent a developmental delay and are later recovered.

These new insights offer significant advancements in our understanding of molecular mechanisms and functional consequences of microglia-synapse interactions in the developing brain and have provoked interest in identifying more microglia-related molecular pathways and functions throughout the CNS. Indeed, identification of such mechanisms and functions will have tremendous impact on the way we think about synapse development and plasticity and may also greatly increase our understanding of diseases associated with abnormal brain wiring (e.g., autism, schizophrenia, etc.) and/or synaptic degeneration (e.g., Alzheimer's disease, Parkinson's, etc.) (See Section IV for discussion of disease relevance).

#### III. Microglia-synapse interactions in the mature CNS: Molecular

#### mechanisms and functional consequences

The previously discussed studies suggest that interactions between microglia and synapses play an important role in the pruning and/or maturation of synaptic connectivity in the developing CNS. Could microglia play a role in the plasticity and function of synaptic circuits in the mature CNS? In addition to roles at developing synapses, several studies have now described potential functions for microglia at synapses in the healthy, adult CNS, such as regulating LTP, synaptic scaling, and basal glutamatergic and GABAergic transmission (Figure 1, panel C) (Ben Achour and Pascual 2010; Bessis et al. 2007; Kettenmann et al. 2011). The following section will review several key findings.

#### Microglia-dependent synaptic plasticity in the mature CNS

While studies in the developing brain have demonstrated that physical interactions between microglia and synaptic elements can regulate plasticity, most of the work in the mature CNS has suggested that microglia can function to modulate the plasticity of neural circuits by paracrine signaling. *In vitro* studies have demonstrated that when either cultured microglia or conditioned media from cultured microglia was added to cortical slice cultures, NMDA-receptor mediated excitatory postsynaptic currents (EPSCs) were larger in amplitude and longer in duration (Moriguchi et al. 2003). Work in hippocampal neuron cultures confirmed

this finding and demonstrated that microglia conditioned media also enhanced LTP induction (Hayashi et al. 2006). The same study identified glycine and L-serine secreted by microglia as molecular mediators of this effect. Consistent with a role for microglia in hippocampal LTP, a recent study found that LTP induction was reduced in organotypic hippocampal slices prepared from adult (3 month)  $Cx3cr1^{KO}$  mice as compared to WT littermates (Rogers et al. 2011). Importantly, these *in vitro* deficits were coincident with *in vivo* impairments in learning and memory as assessed by Morris water maze and contextual and cued fear conditioning. Because these mice also had impaired adult neurogenesis *in vivo*, it remains to be determined whether CX3CR1 signaling in microglia has a direct or indirect effect on synapse function. Interestingly, unlike postnatal hippocampus, which was characterized by a transient decrease in microglia density (Paolicelli et al. 2011). These data raise the intriguing possibility that fractalkine signaling could have differential effects on microglia depending on the context (Figure 1, panels A-C).

In addition to synaptic plasticity associated with LTP, microglia have also been suggested to play a role in synaptic scaling, a homeostatic mechanism that promotes long-term stability of neural networks (Stellwagen and Malenka 2006; Turrigiano 2008; Turrigiano and Nelson 2004). Using cultured hippocampal neurons, glial-derived TNF $\alpha$  was shown to be necessary for synaptic scaling in excitatory and inhibitory neurons. Because both astrocytes and microglia produce TNF $\alpha$ , the relative contribution of these two cell types to *in vitro* and *in vivo* synaptic scaling is an interesting and important area for future investigation.

#### Microglia-mediated effects on basal synaptic transmission

Besides synaptic plasticity, it has been suggested that microglia could regulate basal glutamatergic and GABAergic synaptic transmission (Coull et al. 2005; Pascual et al. 2011; Tsuda et al. 2003). For example, when acute organotypic hippocampal slices were exposed to the pro-inflammatory molecule LPS, microglia became more 'activated' and AMPA receptor-mediated spontaneous EPSC frequency was increased in CA1 neurons (Pascual et al. 2011). Pascual et al. (2011) then demonstrated that this effect was attenuated in slices prepared from PU.1 null mice that lack several cells types belonging to the lymphoid and myeloid lineages (McKercher et al. 1996; Scott et al. 1994), data suggesting that the effect was specific to microglia in the context of the CNS. The authors further demonstrated that the microglia-specific effect was, most likely, indirect and suggested that LPS induces microglia to release ATP that binds P2Y1 receptors on astrocytes. Astrocytes subsequently mediate an increase in excitatory transmission via a metabotropic glutamate receptor 5 (mGluR5)-dependent mechanism. While this study was performed *in vitro* and may be more disease relevant as LPS is a proinflammatory agent thought to more closely mimic inflammation, it raises the intriguing question as to whether microglia have the capacity to regulate basal glutamatergic transmission under less pathological conditions in vivo.

In addition to affecting basal glutamatergic signaling, microglia have the capacity to modulate GABAergic synaptic transmission; however, this effect has only been demonstrated in the context of injury (Coull et al. 2005; Tsuda et al. 2003). Briefly, upon injury, ATP is released, which, in turn, stimulates microglia to release brain-derived

neurotrophic factor (BDNF). Microglia-derived BDNF induces a depolarizing shift in the anion reversal potential that results in an inversion of current polarity activated by GABA. Excitatory transmission via GABA receptor activation results in hyperactivity and increased allodynia. It remains to be determined whether similar microglia-mediated mechanisms may be involved in modulating GABAergic neurotransmission in the healthy CNS.

#### IV. Microglia-synapse interactions in the healthy CNS: Disease Relevance

The studies discussed in previous sections demonstrate roles for microglia in the remodeling and maturation of synaptic circuits in the developing CNS as well as in contributing to basal transmission and plasticity in the adult. In addition to elucidating mechanisms of synapse development and function in the healthy brain, these studies have important implications for understanding mechanisms underlying synapse pathology in disease. In the following section, we will briefly touch on some key studies suggesting a function for microglia at diseased synapses.

Microglial pathological functions have been traditionally associated with diseases in which they are known to perform several functions, ranging from removal of debris and shielding injury sites to initiating and propagating immune responses (Hanisch and Kettenmann 2007; Kreutzberg 1996; Prinz et al. 2011; Ransohoff and Perry 2009). In the context of models of neuropathic pain, one of the most intriguing synaptic functions is their role in the reversal of GABA currents resulting in pain hypersensitivity (discussed in previous section). In neurodegenerative diseases of the CNS (e.g., Multiple Sclerosis, Alzheimer's disease, Huntington's disease, glaucoma, etc.), abnormal synapse function and loss are increasingly being recognized as hallmarks of early stages of disease progression (Alexander et al. 2012; Coleman et al. 2004; Mandolesi et al. 2010; Milnerwood and Raymond 2010). While there are instances in which synapse loss associated with neurodegeneration (i.e., Prion disease) is a neuron-autonomous event (Perry and O'Connor 2010; Siskova et al. 2009), data in other contexts suggest that microglia are contributing to early synapse loss and/or dysfunction. (Alexander et al. 2012; Alexander et al. 2008; Beggs and Salter 2010; Rosen and Stevens 2010; Schafer and Stevens 2010). For example, in a mouse model of tauopathy (P301S), in vivo hippocampal synapse loss and microglial activation were coincident and occurred as early as 3 months of age, while significant atrophy is not observed until 9-12 months of age (Yoshiyama et al. 2007). Thus, data place microglia at the right time and place to contribute to early synapse loss and/or dysfunction. In addition, microglia are thought to be involved in synaptic stripping following axotomy (Cullheim and Thams 2007; Kreutzberg 1993; Perry and O'Connor 2010; Trapp et al. 2007). These experiments were first performed in the context of the injured facial nerve where microglial processes were shown to intercalate between pre- and postsynaptic elements within the facial nerve nucleus following injury, which effectively "stripped" away presynaptic terminals from their postsynaptic targets within the facial nerve nucleus (Blinzinger and Kreutzberg 1968). Interestingly, a recent study in the aging brain demonstrated an increase in phagocytic inclusions within microglia resembling pre- and postsynaptic elements that was concommittant with age-dependent loss of vision and hearing independent of any significant neuronal cell loss (Tremblay et al. 2012). Future work will be necessary to better define the role of microglia at dysfunctional

synapses associated with neurodegenerative disease and aging and to determine whether these cells, indeed, contribute to early synapse loss.

In addition to synaptic dysfunction and loss associated with neurodegenerative disorders and acute neuronal injury, particularly timely and relevant is the potential contribution of microglia to synaptic abnormalities associated with psychiatric disorders such as autism spectrum disorder (ASD), obsessive compulsive disorder (OCD), and schizophrenia (Chen et al. 2010; Hashimoto 2008; Havik et al. 2011; Monji et al. 2009a; Morgan et al. 2010; Pardo et al. 2005; Vargas et al. 2005). Several groups have published interesting studies demonstrating that early infection, a risk factor for many psychiatric disorders, can result in abnormalities in synapses, microglia, and/or behavior, particularly after a second immune or stress challenge (Bilbo 2010; Bilbo et al. 2006; Bitanihirwe et al. 2010; Ito et al. 2010; Shi et al. 2003). One of the most relevant and exciting new studies that implicates microglia in the abnormal brain wiring associated with ASD is a very recent study by Derecki et al. (2012). Using a genetic model of Rett syndrome, the Mecp2-null, that has behavioral and synaptic phenotypes resembling those associated with ASD (Chen et al. 2001; Guy et al. 2001), Derecki et al. demonstrated that replenishing an Mecp2-null with wild-type microglia resulted in the attenuation of several behavioral and physiological deficits (e.g., body weight, breathing rate, locomotion, etc.). Furthermore, the beneficial effects of wild-type microglia on Mecp2-null mice were diminished when phagocytic activity was blocked pharmacologically with annexin-V. However, it remains unclear precisely how abnormal phagocytic activity may be contributing to the phenotype and whether and how microglia may be contributing to synaptic abnormalities associated with ASD. Interestingly, one in vitro study using dissociated hippocampal neuron cultures demonstrated that cells treated with conditioned media isolated from Mecp2-null mice had delayed and abnormal dendritic morphology, signs of microtubule disruption and damage to postsynaptic glutamatergic components due to toxic levels of glutamate released by the microglia (Maezawa and Jin 2010). Given that deficits in synaptic circuit development are emerging as important underlying correlates of behavioral outcomes (Belmonte et al. 2004; LeBlanc and Fagiolini 2011; Melom and Littleton 2011; Rubenstein and Merzenich 2003; Waites and Garner 2011) and the new data that microglia are participating in synaptic pruning via phagocytic immune pathways (Schafer et al. 2012), it is a highly speculative yet provocative hypothesis that microglia are contributing to ASD symptoms, in part, through aberrant neural-immune signaling at developing synapses.

#### V. Summary and Remaining Questions

In summary, the recent attention focused on the role of microglia in the healthy brain has elicited several exciting new findings suggesting that microglia play dynamic roles at developing and mature synapses (see Table I). The interactions between microglia and synapses are dependent upon direct, physical contact as well as signaling via soluble factors, and some molecular mediators of these contact-dependent and independent interactions have been identified. On a functional level, it is now clear that microglia interact with and/or engulf synaptic elements in a manner dependent upon neural activity, mediate synaptic pruning in at least one region of the developing CNS, regulate synapse maturation, and modulate plasticity (LTP and synaptic scaling) and basal transmission in the mature CNS.

As a result of these important first studies, several questions have arisen and remain unanswered in the field. First, in the context of the developing CNS, while it is clear that complement-dependent phagocytic signaling is one pathway underlying physical interactions between microglia and remodeling synapses, other, yet to be identified, pathways must also be involved. Second, at this point, data demonstrate that phagocytosis of synaptic elements and pruning are temporally correlated. It remains to be determined whether engulfment of synaptic elements is, indeed, an underlying mechanism of plasticity and an active process by which microglia selectively engulf intact synapses destined for elimination. Furthermore, while current imaging data in the cortex and hippocampus suggest a role for microglia in activity-dependent synaptic remodeling or maturation (Paolicelli et al. 2011; Tremblay et al. 2010), it is still unclear whether microglia are necessary for bona fide synaptic pruning in these other brain regions, and if so, what are the underlying mechanisms? In the context of the mature adult CNS, while several molecular pathways have been identified to contribute to synaptic plasticity and basal transmission in slice and cultured cell preparations, in vivo contributions are still relatively unclear with only a few behavioral correlates. In addition, in the mature CNS, it is unknown whether and how microglia-specific pathways may interact with one another to affect neurotransmission and plasticity and whether many of these molecular pathways have similar or different functions in the context of microglia in the developing brain (Figure 1, panels A-C).

Imperative to our understanding of these mysterious cells is the use of *in vivo* strategies. Microglia are highly reactive cells that respond within minutes to manipulation (Davalos et al. 2005; Hanisch and Kettenmann 2007; Kreutzberg 1996; Nimmerjahn et al. 2005; Ransohoff and Perry 2009). Thus, while in vitro preparations (e.g., isolated cells, acute slice, slice culture, etc.) are important and necessary strategies for dissecting mechanism and function, the physiological relevance and molecular mechanisms identified must be confirmed *in vivo*. Future *in vivo* studies using a combinatorial approach of live imaging and molecular biology, including the use of newly derived cre lines (Parkhurst 2011), should significantly advance our understanding of the function of microglia at synapses and the molecular mechanisms underlying these interactions. Given that several prevalent psychiatric and developmental disorders (e.g., schizophrenia, obsessive compulsive disorder, autism, etc.) have recently been linked to deficits in synapse development and/or function, as well as a growing body of evidence suggesting abnormalities in microglia (Derecki et al. 2012; Maezawa and Jin 2010; Monji et al. 2009b; Morgan et al. 2010; Steiner et al. 2008; Vargas et al. 2005; Yang and Lu 2011), understanding functions and molecular pathways underlying microglia-synapse interactions in the healthy brain becomes imperative. This knowledge of molecules and function is important for advancing our understanding of basic biological mechanisms as well as for the promise of developing novel diagnostic and therapeutic strategies associated with CNS disease and injury.

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#### Figure 1. Models of microglia-synapse interactions in the CNS

A. In the embryonic and early postnatal brain, microglia are of an amoeboid morphology resembling 'activated' cells associated with disease and injury. During this stage, they are actively dividing and recruited to regions throughout the CNS. Fractalkine (purple circles), which may be released by neurons, is proposed to act on fractalkine receptors (purple squares) expressed by microglia to regulate activation, cell number, and/or recruitment to synaptic-enriched regions. DAP12 (orange squares) expressed on the surface of microglia is also thought to affect synapse function during this period, perhaps, by regulating the 'activation' state of microglia. B. In the postnatal brain, during the first 3 weeks of postnatal life, microglia, which are still 'activated' but now have processes, participate in synaptic pruning. Along with the fractalkine receptor (purple squares) and DAP12 (orange squares), microglia express high levels of complement receptor 3 (CR3, red squares) on their surface. We propose that complement component C3 (red stars) is tagging synapses for removal and have demonstrated that synapses are engulfed by phagocytic microglia in a complementdependent manner. Disruptions in any of these processes results in deficits in synaptic pruning or maturation. C. In the adult brain, evidence suggests that microglia are releasing soluble factors (grey and black circles) such as BDNF, TNFa, glycine, L-serine etc., which affect basal neurotransmission and synaptic plasticity (i.e., LTP) via direct action on neurons or indirectly via astrocytes (orange). In addition, fractalkine signaling via soluble fractalkine (purple circles), most likely released by neurons, and the fractalkine receptor (purple squares), expressed by microglia, modulates microglia-synapse interactions to affect LTP and behavior in the mature CNS.

Overview of synapse-related microg	lia functions				
Microglia function	Age of analysis	Molecules mediating function	Method of study	Region studied	References
Activity-dependent synaptic pruning	First postnatal week	C3, CR3, C1q	<i>in vivo</i> : Engulfment assay, EM, anatomical tracing, array tomography	Retinogeniculate system	Schafer et al. 2012
Phagocytic removal of neurites	Cultures prepared from embryonic mice	C3, CR3, C1q, Sialic acids	<i>in vitro:</i> Stalidase treatment of dissociated neuron cultures, IHC, co-culture of neurons and microglia	Hippocampus	Linnartz et al. 2012
Regulation of synapse maturation	Second and third postnatal weeks	CX3CL1, CX3CR1	<i>in vivo</i> : IHC, EM, and chemical seizure induction ex <i>vivo</i> : Physiology in acute slice	Hippocampus	Paolicelli et al. 2011
Regulation of synapse maturation	Third postnatal week	KARAP/DAP12	ex vivo: Physiology in acute slice, calcium imaging in vitro: Physiology in dissociated neuron cultures, IHC	Hippocampus	Roumier et al. 2004, Roumier et al. 2008
Activity-dependent structural remodeling	Adulthood	Unknown	<i>in vivo</i> : Transcranial imaging, EM, calcium imaging, alteration of sensory experience, ischemic injury	Somatosensory and visual cortex	Wake et al. 2009
Sensory experience driven structural remodeling	Third and fourth postnatal weeks	Unknown	<i>in vivo</i> : Transcranial imaging, EM, IHC. alteration of sensory experience	Visual cortex	Tremblay et al. 2010
Age-dependent structural remodeling	Adulthood	Unknown	<i>in vivo</i> : EM, behavioral studies, IHC	Visual and auditory cortex	Tremblay et al. 2012
Extracellular surveillance	Adulthood	Unknown	<i>in vivo</i> : Transcranial imaging, pharmacology, targeted injury	Cerebral cortex	Davalos et al. 2005. Nimmerjahn et al. 2005
Enhancement of NMDA receptor mediated responses, LTP	Adulthood	Glycine, L-serine	<i>in vitro</i> : Physiology in organotypic slice and dissociated neuron cultures, preparation of microglia conditioned medium	Cortex, Hippocampus	Moriguchi et al. 2003, Hayaslii et al. 2006
Regulation of LTP, learning and memory	Adulthood	CX3CL1, CX3CR1	<i>in vivo</i> : Behavioral assays, pharmacology ex <i>vivo</i> : Physiology in acute slice	Hippocampus	Rogers et al. 2011
Synaptic scaling	Adulthood	TNFci	ex vivo: Physiology in acute slice in vitro: Physiology in dissociated neuron cultures	Hippocampus	Stellwagen & Malenka 2006
Modulation of GABAergic transmission	Adulthood	BDNF, ATP	<i>in vivo</i> : Pharmacology, behavioral assays for allodynia ex <i>vivo</i> : Physiology in acute slice	Spinal cord	Tsuda et al. 2003, Coull et al. 2005

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Table I

Microglia function	Age of analysis	Molecules mediating function	Method of study	Region studied	References
Modulation of basal glutamatergic signaling	Adulthood	ATP	ex vivo: Physiology in acute slice, pharmacology in vitro: Physiology in organotypic slice and dissociated neuron cultures	Hippocampus	Pascual et al. 2011
Unknown	Third postnatal week to adulthood	ATP, Glutarmate, GABA	ex vivo.Time lapse imaging in explants, pharmacology, physiology	Retina	Fontainhas et al. 2011
Unknown	Cultures prepared from postnatal mice	ATP, Glutamate	in vitro: Live imaging in slice cultures, pharmacology, physiology	Hippocampus	Grinberg et al. 2011

Summary of studies discussed in this review. IHC- Immunohistochemistry, EM- Electron Microscopy, CR3- Complement receptor 3, CX3CL1-Fractalkine, CX3CR1- Fractalkine receptor, TNFa- Tumor necrosis factor-alpha, BDNF- Brain derived neurotrophic factor, ATP- Adenosine triphosphate, GABA- Gamma-aminobutyric acid

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