# **Chapter 7 Developmental Neuronal Elimination**

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 **Abstract** Microglia, the brain's innate immune cell type, are cells of mesodermal origin that populate the central nervous system (CNS) during early development. Their functions which are best characterized in the developing CNS are related to programmed cell death (PCD), a physiological process that massively affects neural cell lineages and contributes to brain morphogenesis and neuronal network maturation. Although relatively scarce before advanced developmental stages, microglia can remove dead cells in an effective manner due to their migratory and phagocytic behavior. Recent studies indicate that microglia do not only scavenge cell corpses, but also eliminate neural progenitors cells and trigger or induce PCD in different types of developing neurons. Conversely, microglia were also found to promote the neuronal survival by their release of trophic factors. In this chapter we shall discuss the functional involvement of microglia in the loss of neural cells during normal development and review the mechanisms and cell signalling that underlie microglial regulation of PCD and elimination of dead cells.

 **Keywords** Microglia • Neural progenitors • Apoptotic neural cells • Programmed cell death • Survival • Physiology • Development • Phagocytosis • Trophic factors

#### **Bullet Points**

- Microglial cells remove apoptotic neural cells by phagocytosis in different central nervous system (CNS) regions from early to advanced development stages.
- The contribution of microglia to the elimination of apoptotic neural cells varies depending on the species, developmental stage, and CNS region.

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- Secreted "find-me" and exposed "eat-me" signals comprise the central elements of apoptotic cell removal by microglial cells.
- Microglia not only scavenge cell corpses, but also eliminate neural progenitors and trigger or induce programmed cell death in different types of developing neurons.
- Microglia can also promote the survival of some neurons during development through the release of trophic factors.

#### **7.1 Introduction**

 Microglia are the resident macrophages of the central nervous system (CNS). Since the first studies by del Rio-Hortega, it has generally been accepted that ramified microglia are differentiated cells in the CNS that arise from precursor cells of mesodermal origin with ameboid morphology, which are called ameboid microglia and are present in the developing CNS (Cuadros et al. 1992; del Rio-Hortega [1932](#page-12-0); Cuadros and Navascués  $1998$ ; Chan et al.  $2007$ ; Prinz and Mildner  $2011$ ; Prinz et al.  $2011$ ). It is now established that most if not all microglia originate from yolk sac- derived primitive myeloid progenitors that seed the CNS during early developmental stages (Alliot et al. [1999](#page-11-0); Herbomel et al. 2001; Chan et al. 2007; Ginhoux et al. 2010, 2013; Mizutani et al. 2012; Prinz et al. [2011](#page-15-0); Schulz et al. [2012](#page-16-0); Kierdorf et al. 2013). During development, microglia infiltrate all CNS regions through extensive cell migration and proliferation. They progressively differentiate into mature ramified cells and become spread throughout the adult CNS. Ramified microglia appear to continuously scan the extracellular environment by extending and retracting their cell processes. Contact inhibition between microglial cells leads to a characteristic spatial arrangement in which microglial cells occupy non-overlapping territories (Davalos et al. [2005](#page-12-0); Nimmerjahn et al. 2005). Microglia are highly plastic cells that strongly react to any lesion or pathologies affecting the CNS, and they play key roles in neu-roinflammatory reactions (Hanisch and Kettenmann [2007](#page-13-0); Saijo and Glass 2011).

 Since the discovery of microglial cells, their biology has been widely considered with relation to programmed cell death (PCD) taking place in the CNS during normal development. Histological studies have addressed temporal and spatial relationships between the spread of microglial cells and PCD in the CNS of developing vertebrates (Perry et al. [1985](#page-15-0); Ferrer et al. [1990](#page-12-0); Ashwell 1990, 1991; Perry and Gordon [1991](#page-15-0); Marín-Teva et al. [1999](#page-14-0); Peri and Nusslein-Volhard [2008](#page-15-0); Calderó et al. 2009; Rigato et al. 2011). Microglial phagocytosis of dying cells has been described in various regions of the developing CNS (Ferrer et al. 1990; Marín-Teva et al. [1999](#page-14-0); Dalmau et al. 2003; Peri and Nusslein-Volhard 2008). Besides their debris clearance function, microglial cells actively contribute to cell–cell interac-tions that trigger or ensure the execution of PCD (Mallat et al. [2005](#page-14-0); Bessis et al. 2007; Marín-Teva et al. [2011](#page-14-0)). Lately, microglial cells were also shown to promote neuronal survival in the developing CNS (Ueno et al. [2013](#page-17-0)).

 In this chapter, we shall review current knowledge on the role of microglial cells in neuronal elimination/survival during normal development.

## **7.2 Programmed Cell Death of Neurons in the Developing CNS: Where, When, and Which**

 The large-scale loss of neural cells occurs as a normal and essential stage in CNS maturation during the embryonic and postnatal development of many vertebrate species (Oppenheim [1991](#page-15-0); Kuan et al. [2000](#page-17-0); Yuan and Yankner 2000; Buss et al. [2006 \)](#page-11-0). Cell death affects neuronal and glial cell lineages at different stages of development. The loss of neural cells contributes to the morphological sculpting of the developing CNS and is one of the regressive events involved in the remodelling and functional adaptation of neuronal networks (Oppenheim [1991](#page-15-0); Roth and D'Sa 2001). Although a variety of forms of cell death have been reported depending on morphological and biochemical criteria (Edinger and Thompson 2004; Lockshin and Zakeri [2002](#page-14-0)), this physiological PCD appears to occur by two major pathways: apoptosis, recognized by cell rounding, DNA fragmentation, externalization of phosphatidyl serine (PS), caspase activation, and the absence of inflammatory reaction; and autophagy, characterized by the presence of large vacuoles and the fact that cells can be rescued from death until very late in the process. The boundary between apoptosis and autophagy is not sharply defined, and a complex interplay between the two forms of cell death has been described (reviewed in Booth et al. 2013). However, apoptosis is the dominant type of PCD during normal CNS development (Clarke 1990). The major effect of PCD on projecting neurons during development is well-documented. Thus, during the period in which neuronal connections are established, up to 50 % of numerous types of differentiated neurons undergo cell death (Oppenheim  $1991$ ; Raff et al.  $1993$ ). This neuronal death is thought to be in part triggered by the shortage of neurotrophic factors (neural growth factor, NGF; brain-derived neurotrophic factor, BDNF; etc.) released by the target cells innervated by these neurons (Barde 1989; Oppenheim 1989; Raff et al. 1993; Snider [1994](#page-16-0)). Even though less documented, PCD is also observed earlier in development, before synaptogenesis, in populations of undifferentiated cells such as the proliferating neuroepithelial cells and newly postmitotic neuroblasts, where it participates to specific functions (e.g., morphogenic sculpting of the early CNS) and involves various regulatory mechanisms (Kuan et al. [2000](#page-14-0); de la Rosa and de Pablo 2000; Yeo and Gautier 2004; Valenciano et al. [2009](#page-17-0)). It is generally considered that the PCD of neural cells during development requires relatively conserved molecular pathways. These include "proapoptotic" genes of the Bcl-2 family, the apoptosome (cytochrome c, Apaf-1, caspase-9) and downstream caspases (e.g., caspase-3), which lead to the formation of apoptotic bodies that are rapidly phagocytosed and digested by different types of "professional" and "non-professional" phagocytes. Some studies on the in vivo role of caspases in the normal PCD of developing neurons have shown that caspase activation is involved in most cases of neuronal PCD, but is only necessary for the PCD of immature neurons or neuronal precursors at early developmental stages, when neurogenesis is ongoing (Kuan et al. 2000; Oppenheim et al. [2008](#page-15-0) ). Genetic deletion or pharmacological inhibition of caspases was shown to prevent this early type of PCD in the CNS, while being ineffective at

preventing the normal PCD of postmitotic neurons at later developmental stages during the establishment of synaptic connections (Oppenheim et al. 2001, 2008; Yaginuma et al. 2001; Boya et al. 2008). In the absence of caspases, these postmitotic neurons undergo quantitatively normal amounts of PCD by a different caspaseindependent pathway that exhibits signs of autophagy (Oppenheim et al. 2008). Recent studies have also reported that the authophagic machinery provides the energy required for proper cell corpse removal and further degradation of apoptotic cells during the neurogenesis period (Mellén et al. [2008 ;](#page-14-0) Boya et al. [2008](#page-11-0) ; Aburto et al. 2012).

## **7.3 Scavenger Role of Microglia in Different CNS Regions from Early to Advanced Development**

 Cell apoptosis is marked by DNA fragmentation and caspase-triggered cleavage of the cellular proteome (Nicholson [1999](#page-15-0)), which lead to cell shrinkage but spare the integrity of the plasma membrane up to an advanced stage of the death process. Swift elimination of apoptotic cells by tissue phagocytes is important to prevent secondary cell necrosis involving plasma membrane disruption and leakage of intracellular compounds in the extracellular space, which may be responsible for inflammatory reactions or autoimmune diseases (Nagata et al. 2010). Microglial cells remove apoptotic neural cells by phagocytosis in different CNS regions from early (Sorokin et al. [1992](#page-16-0); Cuadros et al. 1991, 1993; Herbomel et al. 1999, 2001; Lichanska and Hume [2000](#page-14-0)) to more advanced development stages (Hume et al. 1983; Ashwell et al. [1989](#page-11-0); Ashwell [1990](#page-11-0), [1991](#page-11-0); Pearson et al. 1993; Perry et al. 1985; Ferrer et al. [1990](#page-12-0); Perry and Gordon [1991](#page-15-0); Thanos 1991; Moujahid et al. 1996; Egensperger et al. [1996 ;](#page-12-0) Marín-Teva et al. [1999](#page-14-0) ; Upender and Naegele [1999](#page-17-0) ; Rakic and Zecevic [2000](#page-16-0); Dalmau et al. [2003](#page-12-0); Peri and Nusslein-Volhard 2008; Calderó et al. [2009 \)](#page-11-0). In the adult mouse CNS, phagocytic microglia ensure the elimination of dying neuroblasts that derive from neural stem cells in the hippocampal neuro-genic niche (Sierra et al. 2010) (see Chap. [10](http://dx.doi.org/10.1007/978-1-4939-1429-6_10)).

 The contribution of microglia to the elimination of apoptotic neural cells appears to vary depending on the species, developmental stage, and CNS region studied (Dalmau et al. 2003; Calderó et al. [2009](#page-11-0); Sierra et al. 2013). For example, in the cerebral cortex, subcortical white matter, and hippocampus of the in vivo perinatal rat brain, microglial cells engulf virtually all cells undergoing PCD (Dalmau et al. [2003 \)](#page-12-0). However, low microglial cell density may be limiting for scavenging activity, especially at the earliest stages of CNS development, when microglial cells remain scarce. During early development of the CNS, in which the phagocytic capacities of microglia appear to be overwhelmed by the number of dying cells, other nonprofessional phagocytes, such as neighboring neuroepithelial cells, neuroblasts, retinal Müller cells, cerebellar Bergmann glia, and spinal cord astrocytes also contribute to the elimination of apoptotic cells or bodies (O'Connor and Wyttenbach 1974; García-Porrero and Ojeda 1979; Kálmán 1989; Cuadros et al. [1991](#page-12-0); Egensperger et al. 1996; Marín-Teva et al. [1999](#page-14-0); Parnaik et al. [2000](#page-15-0); Mellén et al. [2008](#page-14-0)).

## **7.4 How Do Microglia Find and Selectively Engulf Dead Cells in the Developing CNS?**

 The molecular signalling that triggers microglial phagocytosis during normal development remains a key question to be solved, but this issue clearly benefits from the recent discovery of signals that are generated by apoptotic cells and sensed by phagocytes.

Secreted "find-me," exposed "eat-me," and disabled "don't-eat-me" signals comprise the central elements of apoptotic cell removal by professional phagocytes (Savill and Fadok  $2000$ ; Lauber et al.  $2004$ ; Ravichandran  $2010$ ). Soluble forms of lysophosphatidylcholine (LPC), as well as the chemokine and adhesion molecule CX3CL1 (also known as fractalkine) and the nucleotides ATP and UTP, are known to act as "find-me" signals that are released by apoptotic cells and attract phagocytes (Lauber et al. 2004; Ravichandran 2010). LPC has an attractive chemotactic effect on human monocytic cells and macrophages (Lauber et al. 2003). Moreover, it is released from apoptotic cells through the caspase-3-mediated activation of calcium-independent phospholipase  $A_2$  and is recognized by the phagocyte G-protein- coupled receptor G2A (Lauber et al. [2003 ;](#page-14-0) Peter et al. [2008 \)](#page-15-0). On the other hand, fractalkine was found to be released before the loss of plasma membrane integrity by apoptotic lymphocytes via a caspase-regulated mechanism and to have an attractive effect on macrophages expressing the fractalkine receptor CX3CR1, as inferred from in vitro and in vivo studies carried out in humans and mice (Truman et al. [2008](#page-17-0)). Microglial cells are known to be the only CNS cells that express CX3CR1 (Jung et al.  $2000$ ). Interestingly, fractalkine released from cultured neurons damaged by glutamatergic excitotoxicity promotes the phagocytosis of cell debris by microglial cells (Noda et al. [2011](#page-15-0) ). However, the actual role of G2A or CX3CR1 in the elimination of neural cell corpses during development has not yet been established. Finally, it was reported that the caspase-dependent release of ATP and UTP during the early stages of apoptosis in thymocytes acts as a "find-me" signal to promote phagocytic clearance by human monocytes (Elliott et al. [2009](#page-12-0)). Nucleotides could possibly act as "find-me" signals in the normal developing CNS, but this hypothesis needs to be experimentally tested. In particular, extracellular ATP/ADP has chemotactic effects on microglial cells by binding to  $P2Y_{12}$  purinergic receptors (Davalos et al. 2005; Nimmerjahn et al. 2005; Haynes et al. [2006](#page-13-0) ; Ohsawa and Kohsaka [2011 \)](#page-15-0). Costimulation of microglial P2Y receptors and adenosine (A1 type) receptors appears to be required for microglial chemotaxis towards ATP. This cell response is prevented in purified microglia, obtained by shaking mixed glial cell cultures, which are derived from the brain of newborn mice deficient in the expression of CD39, an ectonucleotidase that degrades nucleotides to nucleosides (Farber et al. [2008](#page-12-0)).

 Once recruited in the vicinity of apoptotic cells, phagocytes can selectively recognize the dead cells as prey to be engulfed due to their expression of "eat-me" signals. In mammals, the best-characterized "eat-me" signal is PS, a cell membrane component that translocates from the inner to outer leaflet of the plasma membrane during the apoptotic process (Fadok et al. 2000). Externalized PS can stimulate phagocytosis of apoptotic cells through direct binding to phagocyte receptors such as T cell immunoglobulin- and mucin-domain containing molecule 4 (Tim-4) (Miyanishi et al. [2007](#page-14-0)), brain-specific angiogenesis inhibitor-1 (BAI1) (Park et al.  $2007$  or stabilin-2 (Park et al.  $2008$ ). PS recognition by phagocytes can also involve several "bridging" molecules, binding to both PS and phagocyte receptors, which are released in extracellular fluid (Erwig and Henson  $2008$ ). Among these, milk fat globule epidermal growth factor-8 (MFG-E8) and C3bi can bind to PS and are then recognized by integrins expressed by macrophages/microglia such as vitronectin receptor  $(\alpha \nu \beta 3)$  and complement-receptor-3 (CR3/CD11b), respectively (Hanayama et al. 2002; Mevorach et al. [1998](#page-14-0); Savill and Fadok [2000](#page-16-0)). The bridging molecules and receptors responsible for PS-mediated clearance of dead cells (see Sierra et al.  $2013$  for a review) are not yet clearly defined in the developing CNS, but the capacity of purified microglial cells to engulf PS-coated cells or par-ticles is well-documented (Witting et al. 2000; Konduru et al. [2009](#page-14-0); Neher et al. [2011](#page-15-0); Liu et al. [2013](#page-14-0)).

 Another possible "eat-me" signal is the endogenous cellular ligand for the triggering receptor expressed on myeloid cells 2 (TREM2). TREM2 is specifically expressed by microglial cells, as demonstrated by in vivo immunocytochemical studies in different regions of the mouse brain (Hsieh et al. [2009 \)](#page-13-0). Its ligand is upregulated on apoptotic neurons, mediates signal transduction by association with DNAX adaptor protein-12 (DAP12) on microglia, and promotes the phagocytosis of dying neurons in cell cultures (Takahashi et al. [2005](#page-16-0); Hsieh et al. 2009; Neumann et al. [2009 \)](#page-15-0). Finally, another in vivo and in vitro study demonstrated that activation of  $P2Y_6$  receptors in microglial cells by UTP/UDP released from damaged neurons triggers the clearance of dying cells by phagocytosis (Koizumi et al. [2007](#page-13-0) ). Hence, in addition to PS, the ligands for TREM2 and UTP/UDP may be considered as putative "eat-me" signals promoting microglial phagocytosis during development.

#### **7.5 PCD and Microglial Cell Distribution in the Developing CNS**

 Besides its role in the elimination of dead cells, apoptotic cell-to-phagocyte signalling clearly fulfills the expected criteria for a mechanism that underlies microglia distribution in the developing CNS. Consistent with the recent demonstration that dying cells release chemotactic "find-me" signals, the codetection of microglia and dying cells provided evidence that local PCD may account for the entry, local spread, or transient clustering of macrophages/microglia in different CNS regions of embryonic vertebrates, including the retina (Santos et al. [2008](#page-16-0) ), spinal cord (Rigato et al. [2011](#page-16-0) ), choroid plexus (Swinnen et al. [2013 \)](#page-16-0), cerebellum, or cerebral cortex (Ashwell 1990, 1991). However, microglia infiltration in cell layers of the developing CNS is not always correlated with the occurrence of PCD (Marín-Teva et al. [1999 \)](#page-14-0), and it involves different molecular cues unrelated to the signalling generated by dying cells. In particular, developing neural cells can produce chemoattractants that target macrophages/microglia, such as chemokines (Ransohoff [2009 \)](#page-16-0), macrophage colony-stimulating factor 1 (M-CSF), or vascular endothelial growth factor A (VEGF-A) (Breier et al. [1992](#page-11-0); Nandi et al. [2012](#page-14-0); Lelli et al. 2013). In zebra fish embryos, M-CSF receptor expressed by mononuclear phagocytes is required for the early infiltration of yolk-sac-derived phagocytes in the CNS (Herbomel et al. [2001](#page-13-0)), whereas VEGF receptor 1 signalling promotes early postnatal infiltration of microglia in deep layers of the mouse cerebral cortex (Lelli et al. [2013 \)](#page-14-0). Furthermore, microglia appear to use radially oriented processes of glial or neural progenitors cells as a substrate for cell migration in the retina (Sánchez-López et al. 2004), spinal cord (Rigato et al. 2011), and cerebral cortex (Swinnen et al. [2013](#page-16-0) ). Strikingly, ameboid microglial cells migrate in embryonic quail retina following well-defined routes that are not altered when they pass close to regions of abundant cell death, and they only engulf apoptotic bodies encountered along these routes (Marín-Teva et al. [1999](#page-14-0)). This observation suggests that cell guiding cues unrelated to PCD may outcompete "find-me" signals emitted by dead cells. Alternatively, the capacity of microglial cells to sense "find-me" (and/or "eat-me") signals may vary according to their localization in the developing CNS. In connection with this possibility, the heterogeneity of microglial cell phenotypes during normal development is well-documented (Hristova et al. 2010; Verney et al. 2010; Scheffel et al. [2012 ;](#page-16-0) Arnoux et al. [2013](#page-11-0) ). Moreover, recent studies have emphasized that the functional implications of microglial recruitment are not limited to the scavenging of dead cells. Microglia fulfill other roles during development, including the remodelling of developing synapses (Hoshiko et al. 2012; Tremblay et al. 2010; Paolicelli et al.  $2011$ ; Schafer et al.  $2012$ ), homeostatic regulation of neuronal firing (Ji et al.  $2013$ ) (Chap. [9\)](http://dx.doi.org/10.1007/978-1-4939-1429-6_9), stimulation of angiogenesis (Checchin et al.  $2006$ ; Fantin et al. [2010 \)](#page-12-0) (Chap. [8](http://dx.doi.org/10.1007/978-1-4939-1429-6_8)), and the induction or prevention of developmental cell death, which is discussed below.

## **7.6 Switching Programmed Cell Death On and Off in the Developing CNS: A Dual Role for Microglia**

 Various studies have demonstrated that microglial cells not only have a scavenger role during development, but can also trigger or promote PCD in different types of developing neurons. The mechanisms involved in the induction of PCD by microglial cells are diverse and can be classified into two groups: phagocytosis-unrelated PCD induction; and PCD triggering/execution involving phagocytosis, designated as "engulfment-promoted PCD."

 The former group relies in part on the capacity of microglia to produce neuronal growth factors such as neurotrophins (Mallat et al. 1989; Elkabes et al. 1996). Frade and Barde (1998) showed that nerve growth factor (NGF) is produced in vivo during early chick embryo development by primitive macrophages and microglial cells, which are present in the vitreous body and scattered within the retinal neuroepithelium between embryonic day 3 (E3) and E6. In addition, in vitro experiments in cultured eye cups demonstrated that macrophage/microglial production of NGF triggers PCD in retinal progenitor cells expressing neurotrophin receptor p75 at E3 (Fig.  $7.1a$ ). Another type of phagocytosis-independent involvement of macrophages in PCD was observed in the embryonic rat spinal cord, in which motoneurons undergo PCD from E15. Experiments in cultured explants of ventral horns of rat embryo spinal cord showed that embryonic motoneurons acquire the competence to undergo PCD between E12 and E13, 2 days before the onset of cell death (Sedel et al. [2004 \)](#page-16-0). This neuronal commitment to a death fate appears to be driven by tumor necrosis factor  $\alpha$  (TNF- $\alpha$ ) released from primitive macrophages that invade the surrounding somitic mesenchyme at these developmental stages (Fig. 7.1b). Macrophage-derived TNF- $\alpha$  signals through the TNF- $\alpha$ receptor 1 expressed by embryonic motoneurons (Sedel et al. 2004). Subsequent in vivo studies provided evidence that these somitic macrophages eventually infiltrate the spinal cord and therefore contribute to microglial development (Rigato et al.  $2011$ ).

 Engulfment-promoted PCD was observed in postmitotic neurons in the cerebellum of 3 day-old (P3) mice (Marín-Teva et al. [2004](#page-14-0) ); at this developmental stage, Purkinje cells initiate a death program marked by caspase-3 activation. In vivo observations in the P3 mouse cerebellum and in vitro experimental studies in P3 cerebellar slices showed that the majority of these neurons are engulfed by microglial cells that actively prevent abortion of the death program in the engulfed cells (Marín-Teva et al. [2004](#page-14-0)). In the developing mouse CNS, microglia express all of the genes encoding multimeric Nox1- and Nox2-dependent NADPH oxidases that catalyze the formation of superoxide ions (Bedard and Krause 2007; Chéret et al. 2008). Activation of microglial NADPH oxidases is detectable at early postnatal stages and is required for microglia-promoted Purkinje cell death (Fig.  $7.1c_1$  $7.1c_1$ ) (Marín-Teva et al. [2004 ;](#page-14-0) Lelli et al. [2013 \)](#page-14-0). This mechanism is reminiscent of the innate immune response in which bacteria or fungi ingested by neutrophils or macrophages are neutralized by reactive oxygen species derived from the activation of phagocyte Nox2-dependent NADPH-oxidase (Bedard and Krause 2007; Lam et al. [2010](#page-14-0)). It is noteworthy that the involvement of engulfing cells in the execution of developmental cell death is not limited to vertebrates (Mallat et al. [2005](#page-14-0) ). It is well documented

**Fig. 7.1** (continued) MiCs promote neuron death by the engulfment of differentiated neurons (DNs) undergoing early reversible steps of cell death  $(c_1)$  and healthy neural progenitor cells  $(c_2)$ . ( **c1** ). In the developing cerebellum (Marín-Teva et al. [2004](#page-14-0) ) and hippocampus (Wakselman et al. [2008 \)](#page-17-0), DNs in early stages of cell death express activated caspase-3 and show putative "eat-me" signals on their surface. These signals are recognized by the  $\beta$ 2-integrin CD11b and the immunoreceptor DAP12 on the MiC surface, thereby triggering neuron engulfment and the progression of late stages of cell death, which are promoted by microglial-derived apoptotic effectors such as superoxide ions  $(O_2^-)$ .  $(c_2)$ . MiCs colonize proliferative zones in the developing cerebral cortex and phagocytose postmitotic neural progenitors (NPs) that show no sign of cell death, thereby contributing to regulate the size of the NP pool (Cunningham et al. [2013](#page-12-0)). (**d**) In the developing cerebral cortex, MiCs recruited in the subcortical white matter limit the extent of PCD in DNs of the cortical layer V through production of insulin-like growth factor 1 (IGF-1) (Ueno et al. [2013](#page-17-0) )

<span id="page-8-0"></span>

 **Fig. 7.1** Dual role of microglial cells in promoting either the death or survival of neurons and neural progenitor cells, as demonstrated by studies performed in different parts of the developing central nervous system (CNS). ( **a** ) In the early retina, ameboid microglial cells (MiCs) are involved in the death of some neuroepithelial cells (NCs) expressing the  $p75$  neurotrophin receptor ( $p75<sup>NTR</sup>$ ) by producing nerve growth factor (NGF). The NGF secreted by MiCs remains bound to the cell surface and promotes NC death after  $p75<sup>NTR</sup>$  stimulation (Frade and Barde [1998](#page-13-0)). (**b**) Primitive macrophages (MACs) have a role in committing differentiating motoneurons (MNs) in the developing spinal cord to death (Sedel et al. [2004](#page-16-0) ). These MNs acquire competence to die through signalling via the tumor necrosis factor  $\alpha$  (TNF- $\alpha$ ) released from MACs in adjacent somites. Two days after TNF- $\alpha$  signalling, committed MNs undergo neurotrophic PCD, and cell debris is phagocytosed by ameboid MiCs invading the spinal cord. ( **c** ) In other developing CNS regions,

in the nematode *Caenorhabditis elegans*, in which "death-committed" cells expressing activated CED-3 (homologue to vertebrate caspase-3) can be rescued by inactivation of genes that control the engulfment of cell corpses (Hoeppner et al. 2001; Reddien et al. 2001). Therefore, it was suggested that live Purkinje cells committed to a death fate readily express "eat me" signals, which stimulate engulfing behavior and superoxide production in microglia (Marín-Teva et al. 2004).

 Further insights into this issue came from an in vivo study in newborn mice, which showed that, similar to developing Purkinje cells, hippocampal neurons in contact with microglia undergo PCD through a mechanism requiring the microglial production of superoxide ions (Wakselman et al. [2008](#page-17-0) ). Both microglial superoxide generation and neuronal death were found to be reduced in newborn mice bearing a mutation preventing expression of the CD11b integrin subunit or activity of the DAP12 signalling protein. CD11b/CD18 ( $\alpha$ M $\beta$ 2) functions as an engulfment receptor of prey tagged with the C3bi complement component (Bohana-Kashtan et al. 2004). The involvement of complement components in developmental neuronal death has not yet been established, whereas a recent investigation shows that C3bi-CD11b signalling contributes to microglia-mediated synapse elimination during normal development (Schafer et al. [2012 \)](#page-16-0). DAP12 is a transmembrane-anchored signalling adaptor containing an immunoreceptor tyrosine-based activation motif that transmits signals from CD11b/CD18 and other immunoreceptors (Ivashkiv  $2009$ ). Among these, TREM2 and β3 integrins are expressed by microglia and can directly or indirectly bind to "eat-me" signals, including PS. Whether these receptors contribute to the DAP12-dependent loss of hippocampal neurons (Wakselman et al. [2008 \)](#page-17-0) has not been addressed. However, recent in vitro studies show that a transient PS externalization can occur in neurons cultured in the presence of microglial cells under proinflammatory conditions. Although these PS-tagged neurons remain viable, they are recognized and engulfed by microglia via mechanisms involving the cross-linking of neuronal PS and microglial αvβ3 integrin (vitronectin receptor) by the MFG-E8 bridging protein, and in fact, neuronal death is executed by microglial phagocytosis (Neher et al. 2011; Fricker et al. [2012](#page-13-0)).

 In the developing CNS, the engulfment-mediated loss of cells is not restricted to postmitotic neurons. Thus, a study carried out in vivo, as well as in in vitro cultured slices, revealed that microglia limit the production of cells in the developing cerebral cortex of macaques and rats by phagocytosing neural progenitor cells, mostly during late stages of cortical neurogenesis (Cunningham et al. [2013 \)](#page-12-0). Interestingly, this study found that most neural progenitor cells contacted or engulfed by microglial cells in the cortical proliferative zones show no signs of cell death or apoptosis as defined by the expression of cleaved caspase-3, TUNEL-labelling of fragmented DNA, PS exposure, or nuclear breakdown. The mechanisms by which microglia recognize these healthy progenitor cells as a prey to be engulfed have not been determined. Nevertheless, these findings reveal a new type of engulfment-promoted cell elimination during development, with microglial cells eliminating viable neural progenitor cells through a process unrelated to apoptosis (Fig.  $7.1c_2$ ).

Besides engulfing behavior and the generation of prodeath signals, microglia can produce various neuronal growth factors that may directly support the survival of neurons or the growth of their processes. These functional capacities were documented in culture studies more than 20 years ago (Mallat and Chamak 1994), and their relevance to normal brain development was recently demonstrated in an in vivo and in vitro study of the mouse cerebral cortex (Fig.  $7.1d$ ) (Ueno et al. [2013 \)](#page-17-0). It was observed that, during early postnatal stages (P3–P7), microglial cells recruited in the subcortical white matter limit the extent of PCD in corticospinal and callosal neurons, the cell bodies of which are localized in the cortical cell layer V. This neurotrophic influence is mediated by the microglial production of insulin-like growth factor 1 (IGF-1), as shown in vitro in cocultures of cortical neurons and microglia (Ueno et al. [2013 \)](#page-17-0). The apparently contradictory roles of microglia (elimination of viable cells versus trophic effect on neuron survival) revealed by the two studies performed in the rat and mouse cerebral cortex (Cunningham et al. [2013 ;](#page-12-0) Ueno et al. [2013](#page-17-0), Fig. [7.1c](#page-8-0)., d) are in agreement with the current view that microglia have a dual role on cell death depending on the microenvironment and interac-tions with other cell types (Hanisch and Kettenmann [2007](#page-13-0); Mallat and Chamak 1994; Czeh et al. [2011](#page-12-0)). Microglia analyzed in the above studies act in different microenvironments and at different developmental periods. Thus, the study by Cunningham et al. ( [2013 \)](#page-12-0) reported that microglial phagocytosis of neural precursor cells occurs in the proliferative ventricular zone of E19-P2 rat embryo cerebral cortex, whereas Ueno et al. ( [2013 \)](#page-17-0) observed microglia-promoted survival of differentiated corticospinal neurons, through IGF-1 signaling, in the cortical layer V of the P3–P5 mouse brain. These differences may reflect functional heterogeneity of microglia driven by diverse environmental cues. They may also arise from the selective capacities of target cells to express compounds required for microglial phagocytic or neurotrophic activities, such as "eat-me" signals, the IGF-1 receptor or other signaling components.

#### **7.7 Concluding Remarks**

 Taken together, current evidence indicates that microglia not only eliminate dead cells, but also play an important role in cell–cell interactions that regulate PCD in the developing CNS. Notably, the signalling by which microglia can promote PCD is diversified. Studies of the developing cerebral cortex have revealed that microglia exert a dual influence on local cell production, promoting the elimination of progenitor cells but preventing the loss of differentiated neurons. It is now clear that the role of microglial phagocytosis extends beyond the elimination of dead cells, as it plays a part in the mechanisms that determine the death fate of neural cells during brain development. The engulfment of apoptotic cells was also shown to impact on the ability of the engulfing cells to produce cytokines or lipid mediators (Elliott and Ravichandran 2010). Therefore, microglial phagocytosis could additionally modulate microglial capacity to influence neural cell survival, growth, or differentiation during normal development. This issue warrants further investigation. The variety of mechanisms by which microglia regulate developmental neural cell death, as

<span id="page-11-0"></span>reviewed here, is consistent with the high functional plasticity of microglia, whose behavioral repertoire appears to be location- and time-specific. The physiological effects of microglia on neural cell survival is most likely tuned by the local cell environment or by specific interactions between microglia and target cells, which may change according to the CNS region and developmental time.

 **Acknowledgments** This study was supported by grants from the Spanish Ministry of Science and Innovation (BFU2010-19981) and from Agence Nationale de la Recherche (ANR-06- NEURO-028). The authors thank Richard Davies for improving the English.

**Statement of Interest: None.** 

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