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**Neural Progenitor Cell-astroglia cross-talk:
involvement of the NF- κ B p50 subunit.**

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Chapter 1.

1.1 Adult neurogenesis

Adult neurogenesis is a process of structural plasticity characterized by *de novo* generation and integration of new neurons into the existing circuitry in the adult central nervous system (CNS) (Ming and Song 2005; Christian et al. 2014). Basal level of neurogenesis occurs in two discrete regions: the subventricular zone (SVZ) in the lateral wall of the lateral ventricle and the subgranular zone (SGZ) of the dentate gyrus (DG) in the hippocampal region (Lois and Alvarez-Buylla 1994; Kempermann and Gage 2000). These two areas, referred to as “canonical” sites of active adult neurogenesis, have been found in all mammalian species studied so far, including primates and humans (Eriksson et al. 1998; Gould et al. 1999; Knoth et al. 2010). The adult neurogenic process comprises a series of sequential developmental steps spanning from proliferation, survival, fate specification, neuronal migration, maturation and functional integration of neuronal progeny. The source of new neurons is the adult neural stem cell (NSC) and their progeny of neural progenitor cells (NPC), which were first isolated from the adult CNS of rodents by Reynolds and Weiss (1992) and Kilpatrick and Bartlett (1993). Their pioneering work on SVZ-derived NSC was pivotal, given that they reported methods for culturing and growing neural stem/progenitor cells (NSPC) in the neurosphere culture assay, where individual cells proliferate and generate clusters of cells in suspension. Additionally, these scientists obtained adhesive cultures of NSPC, grown as monolayers on coated substrates (such as laminin). In these proliferating conditions, basic fibroblast growth factor (bFGF) and epidermal growth factor (EGF) were defined as the primary mitogens used to propagate adult neural progenitors *in vitro* (Reynolds and Weiss 1992; Kilpatrick and Bartlett 1993). These studies established the basic protocols for keeping and maintaining NSPC self-renewal property among many passages, as well as their multipotentiality. Under these conditions the differentiation ability of NSPC is maintained over many passages and can be observed upon growth factor removal, when the molecular machinery drives the differentiation toward the

neural lineages, neurons, astrocytes and oligodendrocytes. Later on, similar protocols were also developed to expand and differentiate NSPC derived from the SGZ of the hippocampus (Gage et al. 1995).

The adult rodent SVZ contains four distinct cell types defined by their morphological features, molecular markers and electrophysiological properties: (1) migrating neuroblasts, type A cells; two distinct populations of NSPC, type B (2) and type C (3) cells; and (4) ependymal ciliated (type E) cells, that separate the SVZ from the ventricular cavity (Lledo et al. 2006). Type B cells are slow cycling cells lining the border between the striatum and the lateral ventricle and expressing the astrocytic marker glial fibrillary acidic protein (GFAP). Type B cells give rise to rapidly dividing intermediate progenitors, type C cells, that lose GFAP expression, and acquire distal-less homeobox-2 (Dlx-2). From type C cells type A cells arise, expressing PSA-NCAM (poly-sialylated neural cell adhesion molecule) and the neuroblast marker doublecortin (DCX) (Alvarez-Buylla and Garcia-Verdugo 2002). Neuroblasts leave the SVZ and move along the rostral migratory stream (RMS) in chains formed by glial cells before terminating into the olfactory bulb (OB), where they differentiate into granule and periglomerular interneurons (Doetsch et al. 1999) (Figure 1). Herein integrated newborn interneurons make local contacts and modulate sensory information. The current hypothesis is that they are involved in changes in the neural representation of an odor, thereby supporting long-term olfactory memory (Sultan et al. 2010).

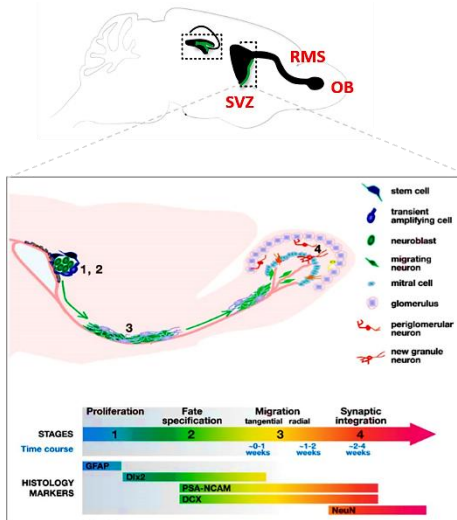


Figure 1. Adult neurogenesis in rodent SVZ/olfactory systems undergoes four developmental stages. A schematic illustration of the adult mouse brain with an enlarged view of the olfactory system. Four developing stages define SVZ neurogenesis. Stage 1. Proliferation: stem cells in the SVZ of the lateral ventricles give rise to transient amplifying cells (type C cells). Stage 2. Fate specification: transient amplifying cells differentiate into immature neurons or neuroblasts (type A cells). Stage 3. Migration: immature neurons migrate in chains through the RMS to the OB. Stage 4. Synaptic integration: immature neurons differentiate into either granule neurons (orange) or periglomerular neurons (red), (Modified from Ming and Song 2005).

Similarly, in the SGZ, stem cells termed type 1 cells, display a radial glial morphology and give rise to intermediate progenitors (type 2a and type 2b cells) which express neuronal specific markers before moving to the dentate granule cell layers where they integrate into the trisynaptic hippocampal circuit (Seri et al. 2001, 2004; Bonaguidi et al. 2012). The detailed developing process of adult hippocampal neurogenesis is described in the following section.

In addition to neurogenesis occurring in the two forebrain regions, emerging evidence arose for a novel neurogenic site along the walls of the third ventricle, in the hypothalamic area. The hypothalamus is a small brain area that surrounds the third ventricle and contains distinct nuclei. It serves as a central homeostatic regulator of numerous physiological and behavioural functions, such as feeding, metabolism, body temperature, thirst, fatigue, aggression, sleep, circadian rhythms, and sexual behaviour. Several studies using 5-bromo-2'-deoxyuridine (BrdU) as well as genetic lineage tracing techniques reported constitutive neurogenesis in the adult hypothalamus of mammals, including rats and mice (Kokoeva et al. 2005; Pierce et al. 2010; Lee et al. 2014). Under unstimulated

conditions, the hypothalamus displays a low rate of neurogenesis and it is suggested that the *de novo* formation of newborn neurons might serve as a compensatory mechanism contributing to homeostatic control of dietary and energy balance (Yon et al. 2013).

Within the adult hypothalamus, cells around the third ventricle form neurospheres, a hallmark of NSC. Cells like tanycytes, ependymocytes, subventricular astrocytes and parenchymal glial cells, all reside near the third ventricle and contribute to the neurogenic potential of the hypothalamus. Hypothalamic tanycytes constitute the main cell type lining the ventral third of the third ventricle, and resemble radial glia. Within the hypothalamus, tanycyte cell bodies are localised in the central and posterior hypothalamus, at the level of median eminence. Morphological studies have mapped and defined two subpopulations of tanycytes according to their position and process projection: i) β -tanycytes, lining the median eminence (ME); ii) α -tanycytes, extending into the arcuate nucleus (ARC) and ventromedial nucleus (VMN), lining the ventricular zone. Robins and colleagues (2013) have reported that α -tanycytes have NSC features and act as neural progenitors in the hypothalamus. It was demonstrated through a series of *in vitro* and *in vivo* experiments in adult mice (using a GLAST::CreER^{T2} mice), that GLAST⁺ (glutamate transporter) α -tanycytes are self-renewing cells, that constitutively give rise to new tanycytes, as well as astrocytes and neurons (Robins et al. 2013) (Figure 2).

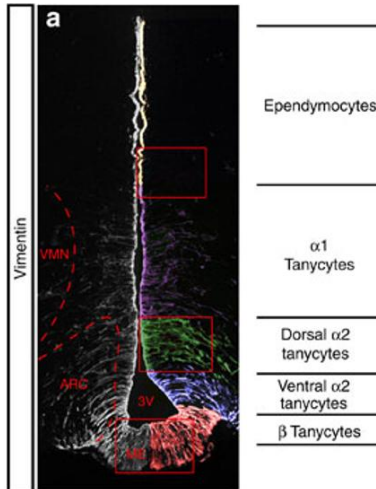


Figure 2. Distinct tanycyte populations line the hypothalamic third ventricle. Left hand panel: coronal section through third ventricle (3V), immunolabelled with Vimentin (intermediate filament marker, expressed by all tanycytes). The position and projection of the Vimentin⁺ process defines different tanycyte subsets: β -tanycytes line the median eminence; $\alpha 2$ -tanycytes reside adjacent to the arcuate nucleus; $\alpha 1$ -tanycytes extend from the level of the ventromedial nucleus to the dorso-medial nucleus. VMN, ventromedial nucleus; ARC, arcuate nucleus; ME, median eminence. (Modified from Robins et al. 2013)

1.1.1 Adult hippocampal neurogenesis

Large efforts have been made in order to understand the function of adult-born neurons both in physio- and pathological conditions. The enormous interest in hippocampal neurogenesis was mainly due to the involvement of this brain region in cognition, and in particular in learning and memory (Aimone et al. 2010). In the hippocampus, neurogenesis is localized within the SGZ in the dentate gyrus, where excitatory granule cells are continuously produced throughout life. The adult SGZ is a narrow band of cells made only of one to three nuclei. The SGZ contains the cell bodies of radial glia-like cells, with processes extending to the densely packed granular cell layer and to the molecular layer (Nicola et al. 2015). The DG has a complex local circuitry, with both inhibitory interneurons and excitatory feedback neurons (mossy cells) constituting the network. The sensory information from specific cortical areas converge to the entorhinal cortex and flows through medial and lateral perforant pathways to the dentate DG, then to CA3 pyramidal cells via mossy fiber axons of granule cells, then to CA1 pyramidal cells via Schaffer collateral projections

of CA3 and back to the entorhinal cortex and subiculum regions, closing the “hippocampal loop” (Aimone et al. 2014).

According to a recent publication by Kempermann and colleagues, adult hippocampal neurogenesis can be divided into four phases, defined as precursor cell phase, early survival phase, postmitotic phase and late survival phase (Kempermann et al. 2015) (Figure 3). The proposed model goes through different developmental stages, identified based on cell morphology and protein marker expression (Kempermann et al. 2004). In general, from a radial glia-like precursor cell, adult neurogenesis progresses throughout three identifiable progenitor stages to a postmitotic maturation phase ending with the formation of a mature granule cell.

In the adult SGZ, four developmental stages define the precursor cell phase of NSPC pool expansion. *In vivo* clonal studies have revealed that radial glia-like precursors are multipotent stem cells capable of repeated self-renewal and generation of both neurons and astrocytes, but not oligodendrocytes, in the adult hippocampus (Bonaguidi et al. 2011). As it is, adult neurogenesis starts with a precursor cell population with morphological and antigenic properties of radial-glia-like cells, named as type 1 cells. The type 1 cells express GFAP, astrocytic marker and nestin. The cell body of type 1 cells is found in the SGZ whereas the processes extends into the molecular layer. Type 1 cells show also electrophysiological properties of astrocytes (Filippov et al. 2003). One transcription factor specifically expressed in SGZ radial glia stem cells is the SRY-related HMG box family member Sox2, which plays a key role in stem cells self-renewal and is critical for the balance between proliferation and differentiation (Julian et al. 2013). The developmental potential of type 1 cells has been demonstrated with tracing experiments (Seri et al. 2004, 2001). Radial glial-like type 1 cells divide asymmetrically, with one daughter cell giving rise to the stem cell and another one giving rise to the intermediate progenitor, or type 2 cells.

Type 2 cells or transient amplifying progenitors, are highly proliferative cells with short processes, that are oriented tangentially to the DG. Type 2 cells come in two subtypes, both nestin positive. One subtype called type-2a cells still express the GFAP, but lack the radial glia morphology, the other cellular phenotype called type-2b, is positive for the immature neuronal marker DCX. Another recently added marker for type 2 cells, is the T-box brain protein 2 (Tbr2), a transcription factor that identifies basal progenitor cells. Tbr2 is required for neurogenesis in the dentate gyrus of developing and adult mice. Tbr2 is critical for the transition from stem cells to intermediate progenitors, as it regulates Sox2 suppression (Hodge et al. 2008). Type 2 cells receive first synaptic GABAergic inputs, and respond to physiological stimuli, such as voluntary wheel running (Kronenberg et al. 2003) or pharmacological stimulation via serotonergic-dependent mechanisms (Encinas et al. 2006) and GABA sets the pace for this regulation (Ge et al. 2007). Kempermann and colleagues propose that the fate choice decision occurs at the level of type 2a cells, given that already in type-2b cells there is expression of transcription factors NeuroD1 and Prox1, involved in neuronal fate choice (Kempermann et al. 2015).

Type 3 cells are DCX positive, but nestin negative. DCX expression extends from a proliferative stage, through cell cycle exit, for a period of two to three weeks (Couillard-Despres et al. 2005). DCX also overlaps with PSA-NCAM expression.

Next phase is the early survival phase. DCX expression still persists until next stage of newly formed neurons expressing the postmitotic marker NeuN and the transient Ca²⁺ binding protein calretinin (Brandt et al. 2003). The majority of cells reaches this stage only three days after the initial division, then these immature neurons are subjected to a selection process, during which they can be eliminated or functionally recruited. Within few more days the number of NeuN⁺ new neurons decreases via apoptosis (Kuhn et al. 2005). The majority of

cells are eliminated before that the functional integration is established in the CA3 region and before receiving dendritic input from the enthorinal cortex, in the molecular layer. Cells that survive retain a vertical morphology with a rounded nucleus and the apical dendrite projecting toward the molecular layer. Then cells send out their axon to target CA3 region, in what is called the mossy fiber path. The main synaptic input to these cells is still GABAergic, and GABA remains excitatory (depolarizing), until sufficient glutamatergic contact has been made (Wang et al. 2000). GABA plays different roles in NPC and their progeny. The response to GABA is depolarizing in NPC and young neurons. In NPC this depolarizing effect is necessary for calcium influx, that in turn promotes the expression of *NeuroD* gene and leads to neuronal differentiation. The absence of a strong GABAergic inhibition in immature neurons represents an important property of these newborn cells. In acute brain slices newborn neurons exhibit enhanced synaptic plasticity, due to lower induction threshold and larger amplitude of long-term potentiation (LTP) of perforant path synaptic inputs compared with mature granule cells. It is suggested that regulation occurs at the survival phase, where only stimuli that are more specific to the hippocampus can affect it as it is for exposure to an enriched environment or to learning and memory hippocampus-dependent tasks, that are known to increase survival at this stage (Gould et al. 1999; Döbrössy et al. 2003).

During the maturation phase, the depolarizing GABA has been shown to be critical for the maturation of adult-born neurons allowing the formation of glutamatergic synapses (Chancey et al. 2013). GABA remains depolarizing in young neurons until two week of age, which corresponds to the time in which glutamatergic spines develop on immature neurons. By 2.5 week of age young neurons have dendrites with spines and receive glutamatergic inputs. Once glutamatergic synaptic connections have been made, the new neurons go through a phase of late survival where increased synaptic plasticity occurs. Inhibitory inputs to adult-born neurons gradually increase with maturation. This

increased inhibition, together with the onset of glutamatergic synapses, sets up a critical period for newborn neurons in memory encoding.

The period of calretinin expression lasts for three to four weeks. Then newly formed cells switch the expression of calretinin to calbindin, that is found in all mature granule neurons (Brandt et al. 2003). At this time point the new neurons still require several additional weeks to become electrophysiologically indistinguishable from older granule neurons.

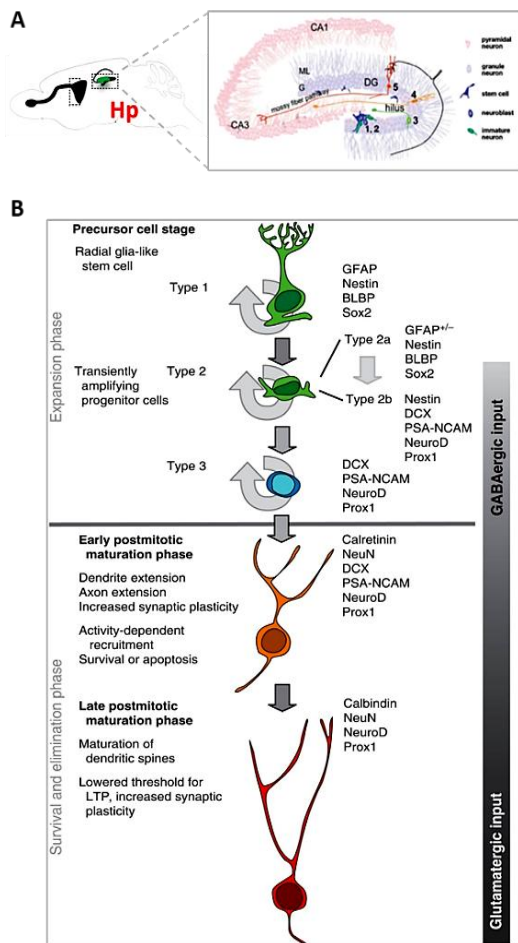


Figure 3. Developmental stages in the course of adult hippocampal neurogenesis. Panel A. Sagittal section view of an adult rodent brain and a detailed representation of the cytoarchitecture of the different layers of the hippocampus (hilus; SGZ; G, granular cell layer; ML molecular layer; area CA3 and CA1) and the hippocampal circuitry of the mossy fiber path connecting axonal projections of immature neurons to CA3 pyramidal cell layer; Schaffer collaterals (not shown) connect CA3 axon projections to CA1, (modified from Ming and Song 2005). Panel B. The precursor cell stage is characterized by cells with radial glia-like morphology (Type 1 cells) and three putative transient amplifying progenitor cells (Type 2a and type 2b cells and type 3 immature neuroblast).

The early postmitotic phase is characterized by calretinin expression. The late postmitotic phase is characterized by terminally differentiated new granule cell, expressing calbindin. (Modified from Kempermann et al., 2015)

An intriguing question is about the functional importance of adult hippocampal neurogenesis. In other words, how newborn neurons contribute to adult brain functions. Many behavioural studies described the correlation between the

levels of adult hippocampal neurogenesis with performances in hippocampus-dependent tasks and thereby put a correlation link between neurogenesis and hippocampus-dependent processes of cognition and emotion. Adult hippocampal neurogenesis is a highly dynamic process, and many extrinsic positive modulators (i.e. enriched environment; physical activity; treatment with certain antidepressants; learning) have been reported to enhance performance in spatial navigation tasks (Morris water maze, MWM) and in tests of anxiety-related behaviour (Kim et al. 2012). On the other hand, physio- and pathological conditions like aging, stress and inflammation, have been reported as negative regulators, and result in decreased performance in tasks of spatial navigation and emotion-related behaviour. Methodological advances in conditional gene targeting improved temporal and tissue specific control of NSPC and neuronal progeny. The primary function that has been mainly associated with adult neurogenesis is pattern separation and hippocampal newborn granule neurons contribute to dentate gyrus pattern separation. Behaviourally, mice in spatial discrimination tasks (i.e. radial-arm maze; two choice discrimination task) must discriminate between two spatially proximate stimuli. Pattern separation is the ability to distinguish similar stimuli, and it is linked to discrimination tasks. It has been demonstrated by several groups that by manipulating the rate of adult neurogenesis, new neurons are critical for making discrimination between close spatial location or similar environments in tests of working memory and long-term memory (Lacar et al. 2014; Vadodaria and Jessberger 2014). In this context, Sahay et al. (2011) reported that by boosting adult hippocampal neurogenesis, utilizing transgenic mice where the proapoptotic gene BAX is conditionally deleted in nestin-expressing NSPC ($iBAX^{nestin}$ mice), enhances discrimination between similar contexts (Sahay et al. 2011). Thus, one of the current hypothesis is that adult hippocampal neurogenesis is critical for discriminating between highly dissimilar contexts, given that discrimination of proximal (in space) choices is selectively impaired without neurogenesis (two-choice discrimination task). Another possible role for adult hippocampal neurogenesis is in encoding time. Young neurons have

been shown to be suited to encode novel information better than the mature neurons (Aimone et al. 2006).

1.1.2 The neurogenic niche

The SGZ, as well the SVZ, form the proper microenvironment that is permissive for neurogenesis to occur. This microenvironment is called neurogenic “niche” (Morrens et al. 2012; Kempermann et al. 2015; Lin and Iacovitti 2015). Neurogenic niches can be defined as functional units within neurogenic areas, consisting of the stem cell pool, their progeny and the immature neurons, and other cellular components, like glial cells, microglia, and endothelial cells. The niche includes as well the extracellular matrix (ECM) and the vasculature and provides a milieu of non-cellular factors, such as secreted molecules and cell-to-cell contacts. All the components in the niche contributes to create a permissive microenvironment that tightly regulate neuronal development of the precursor cells. The local environment in the niche is pivotal in regulating the balance between self-renewal and multipotentiality maintenance for stem cells and for their progeny, and for controlling the different steps of lineage progression until the final commitment.

The architecture within the niche of the SGZ and SVZ differs because it reflects the location differences of the two neurogenic areas (Lin and Iacovitti 2015) (Figure 4). The SVZ is located in proximity to the cerebral ventricles and along the anterolateral ventricle multi-ciliated ependymal cells (E cells) are located. The cilium of B stem cells protrude the E cell wall and senses the cerebrospinal fluid (CSF) in order to gain access to its components, whereas the SGZ does not get in contact with the CSF.

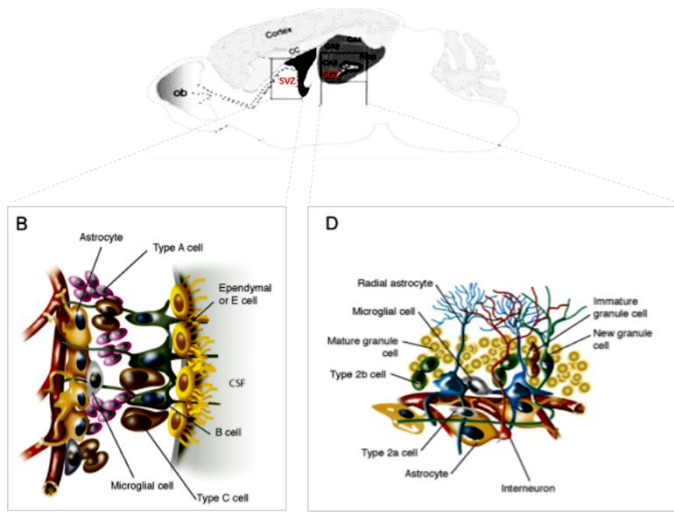


Figure 4.
Schematic representations of the cellular components that made up the canonical neurogenic niches of SVZ (panel B) and SGZ (panel D).
 (modified from Lin and Iacovitti 2015)

A key factor of both neurogenic niches is the vasculature that surrounds the stem cells. For this reason, the niche is often referred also as “neurovascular niche” (Palmer et al. 2000). The vessels of the neurovascular niche lack the astrocytic end feet and the tight junctions that are found in endothelial cells of other tissues. These results in a highly permeable blood-brain barrier in the niche site (Shen et al. 2008). As a consequence, NSPC have access to both local endothelial derived factors and to systemic factors (i.e. vascular endothelial growth factor – VEGF; brain derived neurotrophic factor – BDNF; signaling components like cytokines and chemokines, etc.) (Lin and Iacovitti 2015).

Microglia are present in both SVZ and SGZ niches. They are considered important in regulating neurogenesis in both healthy and injured/diseased brain (Ekdahl 2012). The role of microglia in the niche includes phagocytosis. Sierra and colleagues (2010) showed that newborn cells undergoing death by apoptosis are rapidly cleared via phagocytosis by unchallenged microglia present in the adult SGZ niche (Sierra et al. 2010). In the healthy brain, microglia is also involved in synaptic elimination and pruning by phagocytic engulfment of synapses on mature neurons (Paolicelli et al. 2011) and by this way microglia keep the homeostasis in the presynaptic environment (Platel et

al. 2010). Altogether these functions contribute to the establishment of proper connections for newly formed neurons. Conversely, microglial activation and release of proinflammatory cytokines, upon brain inflammation, has been associated with decline in neurogenesis (Pluchino et al. 2008). Therefore, microglial activation state determines different, even opposite, effects on adult neurogenesis.

Astrocyte cells represent one of the major contributors of the neurogenic niches. Astrocytes have a broad diversity of subtypes, some behaving like stem cells (Seri et al. 2001) and some providing synaptogenic factors, that are critical for local dendritic spine maturation (Sultan et al. 2015). It has been shown that astrocytes also provide neurogenic signals, by instructing neural precursor cells to adopt a neuronal phenotype (Lim and Alvarez-Buylla 1999; Doetsch et al. 1999; Song et al. 2002). Astrocytes of the SGZ are coupled by gap junctions and create what is called the glial syncytium, by which they propagate signals and may regulate activation and differentiation of the stem cell pool. It has been shown that abolishment of gap junctions coupling, via inducible or constitutive depletion of connexin-30 and 43 in radial glia-like cells of the DG (Cx43^{fl/fl}/hGFAP-Cre/Cx30^{-/-} mice), results in a dramatic decrease in the proliferation of precursor cells and in the percentage of newly born granule neurons (Kunze et al. 2009).

ECM molecules (i.e. laminin; heparan sulfate proteoglycans - HSPG; etc.) are also important because they modulate the signals coming from the surrounding parenchyma and blood vessels. Stem cells have laminin receptors (i.e. integrins) on their surface that interact with components of the basal lamina and this interaction helps stem cells to orient in the niche (Erickson and Couchman 2000). Moreover, HSPG of the matrix bind to mitogens like FGF, thereby regulating neurogenesis *via* ECM/growth factors interactions in structures called fractones (Kerever et al. 2007).

Astrocytes also contribute to the production of the ECM. Thrombospondins (TSP) are a family of astrocyte-secreted proteins that are components of the ECM. Previous work suggested that TSP may play a role in hippocampal synaptogenesis. In particular, TSP-1 has been included in the key neurogenic factors released by astrocytes, given that TSP-1 is able to enhance precursor cell proliferation and neuronal differentiation (Hughes et al. 2010; Lu and Kipnis 2010). In the neurogenic niche, astrocytes are as well a source of other important justacrine/paracrine signals (Notch; sonic hedgehog – SHH; bone morphogenetic proteins - BMP), neurotrophic factors (FGF2; BDNF; VEGF) and cytokines (Interleukine-1 β - IL-1 β ; Interleukine-6 - IL-6). An interesting molecule, secreted by astrocytes is the high-mobility group box 1 (HMGB1) protein. HMGB1 is a common proinflammatory cytokine mediating inflammation, neuronal apoptosis and tissue damage. However emerging data suggest that HMGB1 appears to have beneficial effects in brain development, as it is associated with neurogenesis, in NPC survival, proliferation and neuronal differentiation via NF- κ B signaling (Meneghini et al. 2013; Li et al. 2014). Altogether astrocytes may be considered as crucial contributors to neurogenic permissiveness and their contribution in CNS and neurogenesis is deeply discussed in the following section.

1.2 Astroglia in the central nervous system

Astrocytes are a heterogeneous cell population and represent the most abundant cells in the mammalian brain. Astrocytes exist in several morphologically different types all bearing multiple stellate processes. In the morphological classification the following phenotypes are included: i) the protoplasmic astrocytes (localized in brain grey matter), that provide para-arterial influx of nutrients, and para-venous clearance of toxic metabolites, with their perivascular endfeet; ii) the fibrous astrocytes (in brain white matter), involved mainly in the repair of damaged tissue; iii) the radial astrocytes that are the first developing cells from NSC during early embryogenesis and form a scaffold for neuronal migration and, after maturation, became stellate astrocytes. In the mammalian CNS astrocytes undergo differentiation from the NSC pool with the peak of astrogliogenesis occurring during late prenatal/early postnatal developmental (P1 to P10 in rodents), once the bulk of neurogenesis has occurred (Liddelow and Barres 2015). This switch from a largely neurogenic to a gliogenic potential of the NSC pool, reflects the temporal separation in the production of neurons *versus* astroglia. The increase in the number and complexity of glia during evolution suggests a possible function for astrocytes in human cognition (Molofsky et al. 2012). From an evolutionary point of view, both number and complexity of astrocytes increased. Invertebrates CNS contains only 20% of astrocytes, whereas human astrocytes constitute up to 50% of CNS cells, contacting up to 10-times more synapses than their rodent counterparts (Oberheim et al. 2009). Thus, human cortical astrocytes have a more complex branching profile than those of rodents. Given the astrocyte dimension in the human brain, Sloan and Barres (2014) raised the hypothesis that information processing and circuit organization modulated by astroglia may contribute to improved human cognitive abilities. Conversely, human neurodevelopmental disorders may represent an end-product of disrupted astrocyte development (Sloan and Barres 2014).

Originally astrocytes were considered mainly as passive bystanders, playing critical functions in energy metabolism, K^+ buffering and neurotransmitter recycling finalized to keep brain homeostasis. Nowadays, astrocytes cannot be considered anymore only as sustaining cells in the CNS, providing only metabolic and trophic support and passively maintaining neurons (Banker 1980). Great interest grew for the role of astrocytes in regulating neuron-glia interaction, in order to control several processes of brain development, like neurogenesis and differentiation (Barkho et al. 2006; Lim and Alvarez-Buylla 1999; Song et al. 2002), synapses formation (Pfrieger and Barres 1997) neuronal migration/axon guidance (Anton et al. 1997) and neuronal signaling (Fróes et al. 1999), all aspects that provide metabolic and electrophysiological interconnections between neurons and astrocytes. A vast array of data proves the importance of astrocyte-derived soluble factors in several steps of neuronal morphogenesis, ranging from precursor cell proliferation to neuronal differentiation and circuit formation (Gomes et al. 2001). Recently, the metabolic coupling between astrocytes and neurons has been considered to be essential for memory formation. The high metabolic demand for long-term memory formation is possible because astrocytes provide to neurons lactate as a source of energy, that is essential for long-term but not short-term memory formation, for synaptic plasticity and LTP (Suzuki et al. 2011).

Emerging evidence support the concept of regional differences in glia/neuron ratio across the different mammalian brain areas (Herculano-Houzel 2014). This issue highlights how important for brain physiology and function must be the interaction between glial cells and neurons.

1.2.1 Astroglial phenotypes in health and disease

Astrocytes become reactive and undergo reactive astrogliosis/astrocytosis in response to a vast array of CNS insults and the related mediators (such as cytokines and chemokines released from damaged surrounding neurons and active microglia) or in chronic neurological CNS diseases associated with aging, like Parkinson's, Huntington's and Alzheimer's diseases (Sofroniew 2015). In response to tissue damage or inflammation, the category of scar-forming astrocytes form scar borders that segregate the damaged tissue in the CNS. This scar-forming astrocytes demarcate the compartment where damage (trauma, stroke, or infection) has occurred, meanwhile preserving the adjacent tissue. Conversely, in response to mild or diffuse CNS insults, or in tissue that are adjacent to scar borders, astrocytes undergo hypertrophic reactive astrogliosis. The changes that are associated with hypertrophic astrogliosis vary with the nature of the triggering insult, the CNS region involved, the distance from the insult itself, and the presence of co-morbid peripheral infections (Khakh and Sofroniew 2015). Under these circumstances, reactive astrocytes display hypertrophic cell body and nucleus, and the expression of characteristic marker of astrocytes, like GFAP and S100 β is upregulated. Reactive astrocytes are involved in neuroinflammatory processes through the release of proinflammatory molecules and reduced expression of genes involved in neuronal support.

I focused on two specific astrocytic players.

i) *Lipocalin-2* (LCN-2)

LCN-2 is a 24-kDa secreted protein, described first as neutrophil gelatinase-associated lipocalin (NGAL) and also known as 24-kDa superinducible protein (24p3). LCN-2 is a member of the lipocalin family of proteins, known to bind and transport hydrophobic molecules (Flower et al., 2000). LCN-2 acts as a carrier/transporter and binds to two distinct cell-surface receptors: brain type organic cation transporter (BOCT/24p3R), that is expressed constitutively and interestingly is present at high levels in astrocytes; megalin receptor, that is expressed in a wide variety of tissues and cell types, such as ependymal cells; neural progenitors in mice spinal cord; oligodendrocytes, neurons and cultured astrocytes (Lee et al., 2015).

LCN-2 has been implicated in a vast array of cellular processes ranging from cell death/survival (Devireddy et al., 2005), cell migration/invasion in cancer (Yang and Moses, 2009), cell differentiation (Bolognani et al., 2010). In physiological conditions LCN-2 is reported to have high affinity to bind and transport iron, therefore it modulates iron content into cells and this impacts on cell homeostasis (regulating proliferation *versus* apoptosis). Iron-containing (holo-) LCN-2 binds to 24p3R and is internalized, thereby increasing intracellular iron concentration. Instead, when the iron-free (apo-) LCN-2 is internalized upon 24p3R binding, it chelates iron and transfers it in the extracellular milieu, thereby impacting on cellular apoptosis (Devireddy et al. 2005) (Figure 5). As well, LCN-2 can transport iron when complexed with siderophores. This binding confers the ability to LCN-2 to work as part of the acute-phase response in innate immunity.

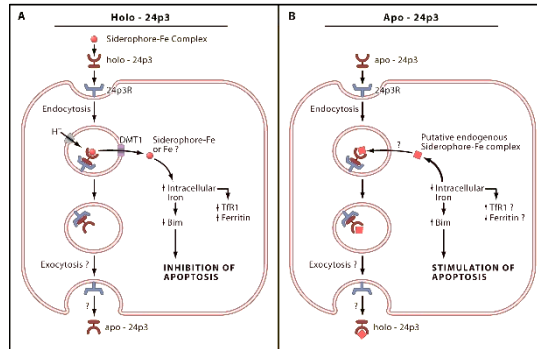


Figure 5. Potential effects of LCN-2 iron binding activity on cellular homeostasis. **A.** Holo-24p3/LCN-2 containing the iron complex of the bacterial siderophore, donates iron to cells via the 24p3 receptor. Internalization of LCN-2 and its receptor leads to the uptake of iron from the siderophore-iron complex. Donation of iron to the cell leads to a decrease in TfR1 (Transferrin receptor protein 1) expression and an increase in ferritin levels. In addition, donation of iron to the cell prevents apoptosis by decreasing the expression of the proapoptotic protein Bim. **B.** Apo-24p3/LCN-2 binds to 24p3R and is internalized into the cell. Devireddy and colleagues (2005) suggest that a putative intracellular mammalian “siderophore iron complex”, called catechol, becomes bound to apo-24p3 to form holo-24p3, which is subsequently released from the cell by exocytosis. Depletion of iron from the cell results in the upregulation of the proapoptotic molecule, Bim, which leads to apoptosis (Richardson 2005).

Initially, LCN-2 was proposed as an acute phase protein also in glial cells, based its role in the regulation of astrocyte death, morphology and migration. LCN-2 can be secreted by reactive astrocytes under inflammatory brain conditions. In astrocytes, released LCN-2 acts as an autocrine mediator of reactive astrocytosis. Following exposure to LCN-2, it was reported that, in parallel to increased GFAP immunoreactivity, there are also phenotypic changes in the number and length of cellular processes of astrocytes (Lee et al., 2015). LCN-2 upregulation also promotes astrocytes migration *via* release of chemokine ligand-10 (CXCL10), chemokine that promotes the migration of other CNS cell types. Altogether LCN-2 upregulation is consistent with the hallmarks of reactive astrocytes.

In the CNS, LCN-2 expression is described to occur in response to injury or in inflammatory conditions. On the other hand, Chia and colleagues (2011) reported the distribution and the expression of LCN-2 in normal rat brain in physiological conditions, with low levels of expression in most brain regions, comprising the hippocampus, and highest levels of expression in olfactory bulbs, brainstem and cerebellum (Chia et al. 2011). They reported that LCN-2⁺ cells co-localize with GFAP⁺ astrocytes of these regions.

In addition, in the CNS, LCN-2 has been suggested to play a modulatory action in cognitive functions (Ferreira et al. 2013), neuronal excitability and anxiety (Mucha et al. 2011) and depression in human beings (Naudé et al. 2013).

Interestingly, LCN-2 null mice (LCN-2^{-/-}) display anxious and depressive-like behaviors, cognitive impairment in spatial learning tasks and in parallel synaptic impairment in hippocampal long-term potentiation (Ferreira et al. 2013). Altogether behavioral studies, performed in order to assess animal anxiety-like behavior (i.e. elevated plus maze; light/dark box; open field test; Morris water maze), associated with hippocampal morphological analysis support the involvement of LCN-2 in emotion and cognition.

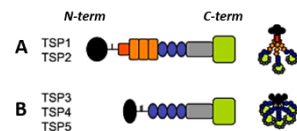
Recently, Mucha and colleagues (2011) reported that both *Lcn2* gene and the encoded protein are upregulated in the hippocampus of male mice upon psychological stress (6h/daily restraint for 3days) and that LCN-2 contributes to stress-induced neuronal plasticity (Mucha et al., 2011). On one hand, they demonstrated through *in vitro* experiments that treating hippocampal neurons with lipocalin-2 (100 ng/ml), as an attempt to mimick stress induced upregulation, caused a decrease of mushroom spines (known as “memory spines”) and a concomitant increase of the thin spines (known as “learning spines”) in neuronal cultures. On the other hand, they observed that disruption of the *Lcn2* gene in mice, caused an increase in the percentage of mushroom spines and their density. These *ex vivo* data correlated also with higher anxious-

like behavior of LCN-2^{-/-} mice upon stress. This scientific report suggested that lipocalin-2 might regulate brain allostasis, and act as a mechanism of cellular adaptation in response to harmful stimuli.

ii) *Thrombospondins (TSPs)*

TSPs are astrocyte-secreted glycoproteins, that participate in a vast array of biological functions as part of the ECM, and regulate cell-cell and cell-matrix interactions. The TSP family consist of two subgroups. Subgroup A includes TSP-1 and TSP-2, whereas subgroup B includes TSP-3/4/5 (Figure 6).

Figure 6. Thrombospondin isoforms and oligomerization states. TSP can be divided into two subgroups. TSP-1 and -2 (subgroup A) form trimers, whereas TSP-3, -4, and -5 (subgroup B) are assembled as pentameric proteins. All TSPs contain a variable number of epidermal growth factor (EGF)-like repeats (blue) that are contiguous with calcium binding repeats (gray) and a globular conserved C-terminal region. A distinctive characteristic of group A TSP is the presence of a pro-collagen (von Willebrand factor, red) homology domain and the three properdin-like repeats (orange). In contrast, the pentameric thrombospondins (TSP-3, -4, and -5) lack the procollagen homology domain and the properdin-like repeats and contain four (instead of three) copies of the EGF-like repeats (Modified from Eroglu et al. 2009)



Recently, it was demonstrated that TSPs promote the formation of new excitatory/gabaergic synapses in the developing CNS. This process is called synaptogenesis and is stimulated by astrocyte-secreted prosynaptogenic signals, such as TSP, and not only controlled by intrinsic neuronal mechanism (Risher and Eroglu 2012). Excitatory synaptogenesis occurs in the mammalian brain after birth, and in rodents this period occurs during the second/third postnatal weeks. TSP-1 and TSP-2 were found to play active roles by promoting the formation of excitatory synapses, and their deficiency resulted in reduced synaptic density during development (Liauw et al. 2008). The increase in synaptogenesis is mediated by the interaction between the TSP EGF-like

domains (common to all TSPs) and the $\alpha\delta$ -1 neuronal receptor (Eroglu et al. 2009).

Considering the critical role of TSPs in developmental synaptogenesis, it was also discovered that these ECM proteins are upregulated/downregulated in response to CNS injuries and in several neuropathological disorders. The role of TSPs was investigated in stroke recovery. It was reported that TSP-1/2 double KO mice exhibited significant reduction in synaptic density, as well as significant deficits in their ability to recover functions. Instead in WT mice TSP-1 and TSP-2 mRNA and protein expression levels increased following stroke and the proteins were found to be co-localized primarily in astrocytes (Liau et al., 2008). These studies suggest that deficiency in TSP-1 and TSP-2 leads to impaired recovery after stroke and that TSPs are beneficial in term of synaptic formation. Altered TSPs expression has been reported in many neurological disorders. Down's syndrome (DS) is the most common genetic form of mental retardation and characterized by a reduced number of dendritic spines. DS astrocytes were found directly involved in the development of spine malformations and reduced synaptic density. This DS neurological feature has been linked to TSP-1 decreased protein expression in astrocytes (Garcia et al. 2010).

Alzheimer's disease (AD) is a progressive neurodegenerative disorder characterized by impaired cognitive function and memory loss due to deposition of senile plaques (Amyloid- β), neurofibrillary tangles, loss of neurons, microgliosis and astrogliosis. Along with these features the clinical stage is characterized by massive synaptic loss that correlates with cognitive decline in AD patients. It has been demonstrated that TSP-1 expression levels are significantly decreased in brain of AD patients and in two AD mouse models (Tg2576 mouse model, expressing A β precursor protein - APP, Swedish mutation K670N; and Tg6799/5xFAD mouse model, overexpressing mutant human APP695 with the Swedish, Florida I716V, and London V717I familial AD mutations, as well as human Presenilin 1 harboring two familial mutations M146L and L286V), (Son et al. 2015). This authors demonstrated that TSP-1

plays a protective role against A β toxicity on dendritic spines. A β ₁₋₄₂ was found able to regulate TSP-1 secretion levels in astrocyte and only TSP-1 was found reduced in astrocyte secretome. Furthermore, co-incubation of TSP-1 with A β ₁₋₄₂-treated hippocampal neurons, was shown to mitigate the dendritic spine loss *via* interaction with the TSP-1 receptor α 2 δ -1 *in vitro*.

Lu and Kipnis (2010) proposed TSP-1 as key regulator of NPC self-renewal and neuronal differentiation *in vivo* and *in vitro* (Lu and Kipnis 2010). The authors showed that deficiency of TSP-1 impairs cell proliferation and neuronal differentiation, both *in vivo* and *in vitro*. First, by quantifying BrdU and Ki67 proliferation markers in the neurogenic SGZ and SVZ areas in WT and TSP-1^{-/-} mice (on a B6129SF1/J background), they reported a significant reduction in proliferating cells in mutant mice. In addition, in the same regions they observed a significant reduction in the number of DCX⁺/BrdU⁺ cells in TSP-1^{-/-} mice compared with the WT counterparts. In the *in vitro* study, the authors confirmed that TSP-1 deficiency impairs cell proliferation and neuronal differentiation of SVZ-derived NPC. For the first time they demonstrated that incubation with exogenous TSP-1 (2 μ g/ml) promotes WT NPC neuronal differentiation and that TSP-1^{-/-} astrocytes are unable to promote neuronal differentiation of NPC. Altogether these observations suggest that TSP-1 is one of the astrocytes-derived molecules required for astrocyte induced neuronal differentiation.

Our group demonstrated that adult hippocampal NPC derived from wild type animals do express the α 2- δ subunit of the voltage-gated calcium channels, and this receptor mediates the proneurogenic effect of antiepileptic drugs, pregabalin and gabapentin (Valente et al., 2012).

1.2.2 Astrocytes as modulators in adult neurogenesis

Astrocytes, as influential components of the neurogenic niches, can contribute to regulation of new neurons formation. An active role for astrocytes and astrocyte-secreted factors in neurogenesis, has been demonstrated in the last two decades. The group of Gage was among the first ones to provide evidence for the active and instructive role of astrocytes in the adult brain (Song et al. 2002). These scientists showed that hippocampal-derived astrocytes from postnatal mice actively regulate neurogenesis, instructing the adult stem cells to a neuronal fate commitment. They examined the fate choice of adult hippocampi-derived neural stem cells *in vitro*, establishing a co-culture system where NSPC were plated onto a feeder layer of primary hippocampal astrocytes from neonatal rodent brain. Furthermore, they showed that even adult hippocampal astrocytes retain the ability to promote neurogenesis from adult NSPC but are less effective compared to neonatal astrocytes. By examining the effects of astrocytes derived from neonatal and adult spinal cord, an area where multipotent stem cells can be isolated, they demonstrated the lower magnitude of neuronal differentiation of adult NSPC in presence of neonatal astrocytes. Conversely, astrocyte derived from adult spinal cord were ineffective in promoting neurogenesis, while were effective in promoting glial over neurogenic fate, data confirmed also by others (Lie et al., 2002). These results suggested for the first time the possibility that astrocytes from different CNS regions may provide regionally specific cues and that astrocytes at different developmental stages may exhibit different effects on NSPC fate choice.

Previously, in a similar work, Lim and Alvarez-Buylla reported that the interaction between NPC derived from adult SVZ and astrocytes is necessary to stimulate neurogenesis (Lim and Alvarez-Buylla, 1999). Given the ultrastructure of SVZ, with astrocytes in close contact with type B cells (slow cycling stem cells, GFAP⁺) and with type C cells (rapidly dividing intermediate progenitors, Dlx-2⁺), these researchers explored the possibility that astrocytes

provide a neurogenic microenvironment for SVZ cells. In their publication they showed that direct contact with astrocytes deriving from cortices supports the proliferation of SVZ precursor cells and the differentiation of these into type A neuroblasts (DCX⁺). They suggested that it is necessary the direct contact between astrocytes and precursor cells for neurogenesis to occur. When they tested conditioned medium from astrocytes, they observed that the medium does not support neurogenesis, suggesting that the soluble factors produced by astrocytes may be necessary but not sufficient.

Later on the work by Song and colleagues was supported by the group of Zhao, who performed gene expression profiling studies on primary astrocytes isolated from different regions of the CNS. The gene expression analysis was performed on astrocytes isolated from regions with high plasticity, such as newborn and adult hippocampal astrocytes and newborn spinal cord astrocytes, shown to promote neuronal differentiation of adult NSPC, and on astrocytes isolated from the non-neurogenic area of adult spinal cord, known to inhibit neuronal differentiation (Barkho et al., 2006). Briefly, they demonstrated that astrocytes isolated from different brain areas have distinct gene expression profiles. They found out that neurogenesis-promoting astrocytes release interleukine-1 β (IL-1 β) and interleukine-6 (IL-6), which are proinflammatory cytokines, that could promote NSPC neuronal differentiation in combination with other factors, such as vascular cell adhesion molecule-1 (VCAM-1) and interferon-induced protein 10 (IP-10). Instead, astrocytes isolated from the spinal cord express more insulin-like growth factor binding protein 6 (IGFBP6, a negative regulator of insulin-like growth factor II signaling) and decorin (proteoglycan that can inhibit transforming growth factor 2, TGF β 2) which negatively regulate neurogenesis. In fact, it was demonstrated that this factors inhibited neuronal differentiation *in vitro* when co-cultured NSPC and neurogenic astrocytes were treated with them. These authors proposed that the combo of factors secreted by astrocytes into the local microenvironment is affecting the neurogenic niche.

Recently, IL-6 was identified to promote neuronal, but not glial differentiation of adult hippocampal NPC when applied directly in culture, by increasing the fraction of class III β -tubulin (TUJ1⁺) cells (Oh et al., 2010). Furthermore, was observed that also the average length of TUJ1⁺ neurites was increased compared to vehicle-treated cells, and the effects observed were abolished when IL-6 was neutralized by anti-IL-6 blocking antibody. This recent study supported the work done by Barkho and colleagues (2006).

While proinflammatory cytokines generally inhibit neurogenesis, the fact that neurogenesis-promoting astrocytes express higher levels of inflammation related proteins than neurogenesis-inhibiting astrocytes, suggests that cytokines and chemokines may have context- and concentration-dependent effects.

Astrocytes have been shown to secrete many positive, as well as negative modulators of hippocampal neurogenesis. Neurogenesis-1 (Ng1) has been identified as an astrocyte-derived soluble factor that positively modulate neuronal differentiation both *in vitro* and *in vivo* (Ueki et al., 2003). Ng1 has been found prominently expressed in neurogenic brain areas and co-localize mainly within astrocytes of the SGZ of the adult brain. The implication of Ng1 in adult neurogenesis was confirmed by *in vitro* experiments on hippocampus-derived NSC cultures. In these experiments, Ng1 neurogenic effect was antagonized with a neutralizing antibody against-Ng1 applied to NSC cultures prior to treatment with the supernatant of hippocampal neurogenic region. Thus this work supported the concept that astrocytes from the hippocampus contribute to the favorable environment for adult neurogenesis.

Among other positive modulators, components related to glial metabolism also appear to be key regulators of adult hippocampal neurogenesis. ATP released from astrocytic metabolism, has been identified as a proliferative factor required for astrocyte-mediated NSC proliferation in the adult hippocampus and *in vitro* (Cao et al., 2013). Similarly, Sultan and colleagues reported that D-serine a secreted astrocyte gliotransmitter abundant in hippocampal astrocytes, is able to increase the number of NSPC *in vitro* and when administered exogenously, it increases both the number of RGL cells and the number of

transient amplifying progenitors in the dentate gyrus of adult mice (Sultan et al., 2013). Additionally the reduction of glutamine levels with fluorocitrate, that selectively inhibits glial metabolism, results in decreased levels of NPC proliferation (Fonnum et al., 1997).

An interesting protein is the neurotrophin brain-derived neurotrophic factor (BDNF), that has been reported to regulate adult hippocampal neurogenesis (Waterhouse et al., 2012). Exogenous BDNF injection into the hippocampus boost NPC proliferation *in vivo* and promotes the differentiation and maturation of adult NPC in culture. This is consistent with evidence that neurogenesis decreases in BDNF knockout mice. SGZ-derived NPC have been demonstrated to express BDNF-TrkB receptor, that is required for the neurogenic and behavioral response to antidepressive treatments, supporting the concept that SGZ-derived NPC are a required component in the amelioration of depression (Li et al., 2008). Studies have demonstrated that hippocampal astrocytes minimally produce BDNF (Rudge et al., 1992). During aging the levels of the neurotrophin does not change, whereas those of the receptor TrkB decreases with increasing age, as reported in the rat hippocampus. This phenomenon might be one of those correlated with aging-related decline in adult hippocampal neurogenesis.

Instead, among the factors that are secreted by astrocytes and promote astroglial differentiation, there are bone morphogenetic proteins (BMP), that alter the fate of neural stem cells from neurogenesis to astrocytogenesis. BMP are members of the transforming growth factor β (TGF β) superfamily, known to mediate the growth, differentiation and survival of different cell types (Dobolyi et al., 2012). First studies reported that BMPs (BMP2/4/5/6) induce astroglial differentiation of SVZ-generated neural progenitors and the concurrent decrease of oligodendroglial and neuronal lineages, and this instructive effect is merely differentiative with no influences on both proliferation and survival of astroglial lineage cells (Gross et al., 1996). Other reports, demonstrated that NSC treatment with low-dose of BMP4 (0.1 ng/ml) or BMP7 (1 ng/ml) stimulates NSC to differentiate exclusively into neurons (TUJ-1⁺ cells), with other cells

remaining undifferentiated, suggesting a concentration-dependent effect of BMP (Chang et al., 2003). Astrocytes secrete ciliary neurotrophic factor (CNTF), that is also important for differentiation of NSC into astrocytic lineage. CNTF alone (20 ng/ml) has a dramatic effect on astrocytic differentiation, with GFAP⁺ cells > 90% of total cells. Interestingly, the combined treatment of the astrogenic/neurogenic BMP with CNTF results in a decrease in astrocytes and an increase in neurons. This demonstrates the context-dependent regulation of BMP, regardless its concentration (Chang et al., 2003).

Furthermore, Ng1 involvement in neurogenic activity can be traced back to its structure (cysteine-rich domains), characteristic of BMP antagonists. Ueki and colleagues (2003) reported that Ng1 antagonize BMP-4 and alters that fate commitment of NSC from gliogenesis to neurogenesis, and suggests that Ng1 may influence hippocampal plasticity by downregulating the activity of BMP (Ueki et al., 2003).

Astrocytes supplement the local microenvironment also with mitotic factors and most notably FGF-2 and VEGF contribute to neurogenesis by enhancing the proliferation of NSC and their progeny both in the SVZ and SGZ. Bernal and colleagues (2011) asked whether a decreased availability of astrocyte-derived growth factors may contribute to the age-related decline in neurogenesis (Bernal and Peterson, 2011). GFAP-positive astrocytes express both VEGF and FGF-2. The authors found out that the gene expression levels of VEGF are significantly reduced in aged DG of rats compared to young one. In DG aging resulted also in the emergence of a phenotype of GFAP-positive astrocytes that did not co-express FGF-2, whereas in the young brain all GFAP-positive astrocytes co-localize with FGF-2. Anyway the overall density of astrocytes immunopositive for GFAP did not change. The authors suggested that the local environment during aging becomes less supportive for neurogenesis. VEGF and its receptor Flk-1 were detected also in Type-1 NSC and the co-expression of the two proteins is prior to neuronal commitment. Once DCX⁺ cells are formed, VEGF expression is lost, but Flk-1-mediated sensitivity to VEGF signaling is still maintained indicating that immature neuroblasts may be influenced by VEGF

signals presented by neighboring astrocytes. Normal aging is accompanied by a decline in CNS function and reduced neurogenesis. Astrocytes might be considered as the major contributors in maintaining the neurogenic niche given that the altered profile of astrocytes in the aged hippocampus negatively affects neurogenesis.

In a recent publication a group of researchers investigated the neurogenic-inducing properties of astrocytes isolated from cortical, hippocampal and midbrain areas, by analyzing the secretome profile of astrocyte conditioned media (ACM) and the effect on neuronal differentiation of human NPC (Cordero-Llana et al., 2011). They cultured human NPC for one and four weeks in control media or in regional-specific ACM. They demonstrated that ACM derived from hippocampus and midbrain is able to promote neuronal differentiation of human NPC, whereas ACM from cortex does not. By using mass spectrometry (MS) they identified clusterin as a protein secreted only in hippocampal and midbrain and not in cortex ACM. Clusterin is an interesting secretory glycoprotein (75-80 kDa), with high expression in astrocytes. It has no effect on the proliferation of NPC, but it promotes both their neuronal differentiation and survival.

Generally, the control of NSPC proliferation and differentiation is influenced by several factors that are expressed by many cell types residing within the niche. The role of astrocytes on cell fate is different, with no clear knowledge of the instructive signals and the molecular pathways involved. Astrocytes are organizers of the neurogenic niche, and the signal they provide to the neurogenic process are a sum of complex microenvironmental cues, and influences of the surrounding cells. Astrocytes by releasing soluble factors, may play both instructive and permissive functions. On one hand astrocytes secreted molecules, may promote one fate at the expenses of another, on the other hand astrocytes cues may inhibit one fate, thereby allowing progenitors to differentiate in the other cell populations.

1.3 NF- κ B signaling in adult neurogenesis

The term NF- κ B (Nuclear Factor kappa-light chain enhancer of activated B cells) refers to a family of transcription factors, firstly discovered in immune B cells to bind to the intronic enhancer of the kappa light chain gene, the κ B site. These proteins are ubiquitously expressed and responsible for regulating the expression of multiple genes whose products are involved in pleiotropic functions, including cell survival/death, cell proliferation and differentiation and immune and inflammatory responses (Grilli and Memo 1999; Kovacs et al., 2004; Vallabhapurapu and Karin 2009). Often NF- κ B signaling appears to play opposite functions, since the final outcome of its activation depends on the cell type involved, the type of disease or injuries and the target genes activated or repressed. NF- κ B signaling was initially studied mainly in the field of immunology, as major regulator of innate and adaptive immunity. In the CNS, NF- κ B is expressed in both neurons and glia. In general, NF- κ B in CNS glial cells, like microglia and astrocytes is mainly neurodestructive due to the overwhelming production of pro-inflammatory cytokines (Mattson and Meffert 2006). The role of NF- κ B in the nervous system has gained interest because of its involvement in neuroprotection/neurodegeneration (Kaltschmidt and Kaltschmidt 2009), synaptic plasticity and neurite outgrowth (Gavalda et al. 2009) and formation of functional dendritic spines (Boersma et al. 2011). On one hand many *in vitro* studies have demonstrated that neurotoxic and pro-apoptotic stimuli including high concentrations of glutamate, amyloid- β or cytokines, cause NF- κ B activation in neuronal cells, resulting in cell death (Uberti et al. 1999; Ryan et al. 2000; Zhang et al. 2005). On the other hand, it has been reported that neuron-restricted ablation of NF- κ B-driven gene expression promotes neurodegeneration (Fridmacher et al. 2003), suggesting a neuroprotective role for NF- κ B. Recent findings suggest that neuronal NF- κ B signaling is neuroprotective via maintaining neuronal survival, synaptogenesis, synapse to nucleus communication (Gutierrez and Davies 2011; Mattson and

Meffert 2006) and its role in regulating neural development and CNS functions such as learning and memory (Salles et al. 2014).

Herein, I will introduce NF- κ B family members and their signaling pathways, focusing on the emerging evidence of a role for NF- κ B in regulating adult hippocampal neurogenesis.

1.3.1 The NF- κ B family of transcription factors

The NF- κ B family of transcription factors is composed of five subunits including p50/p105 (NF- κ B1), p52/p100 (NF- κ B2), p65 (RelA), RelB and c-Rel, which all share a conserved and multifunctional N-terminal Rel homology domain (RHD), that is responsible for homo- and hetero-dimerization of the different subunits. The RHD also contains nuclear localization signal (NLS), masked by binding of inhibitory I κ B protein. Upon I κ B inhibitory protein phosphorylation and degradation, NLS unmasking allows nuclear translocation of dimeric complexes which can then bind regulatory sequences in target genes. According to their structure, NF- κ B family members can be divided into two subfamilies. One includes RelA, RelB and c-Rel, which are generally considered as transcriptionally activators, since they possess a transactivational domain (TAD), while the other subfamily includes p50 and p52 subunits, which are commonly considered as repressors, since lacking the TAD. The I κ B family of inhibitors includes p100, p105, I κ B α , I κ B β , I κ B γ and Bcl-3. These inhibitors proteins are characterized by the presence of ankyrin-repeat motives that mediate their binding with NLS/RHD of NF- κ B subunits. The p105 and p100 C-terminal region, also containing ankyrin repeats, that acts like an auto-inhibitory domain, retaining the precursors of p50 and p52, respectively, into the cytoplasm. (Figure 7)

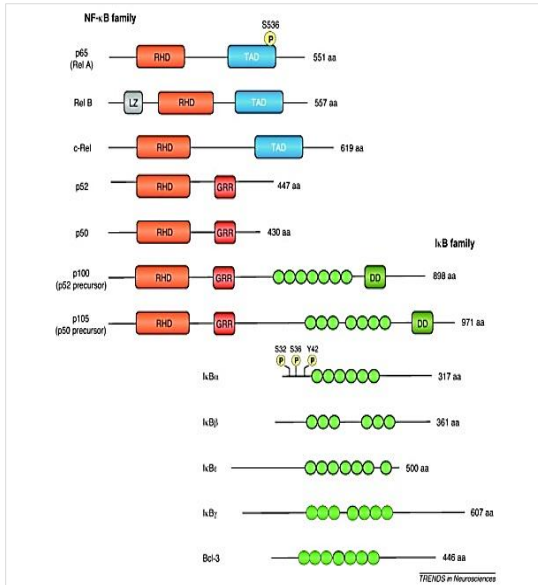


Figure 7. Schematic diagram illustrating the mammalian members of the NF- κ B and I κ B families. NF- κ B family members have in common the RHD, with RelA, RelB and c-Rel additionally possessing a TAD [RelB has a leucine zipper (LZ), while p52 and p50 and their precursors have glycine-rich regions-(GRR); additionally, precursor possess regions with homology to death domains (DD)]. The ankyrin repeats of I κ B members (denoted by green ovals), mediates the RHD binding. Within the I κ B α phosphorylation sites relevant for regulation of NF- κ B activation have been identified (Ser32, Ser36, Tyr42). (Gutierrez and Davies, 2011)

At the cellular level, under unstimulated conditions, NF- κ B dimers are retained in the cytosol through the interaction with a member of the inhibitory I κ B protein family. The I κ B subunit masks NLS in the NF- κ B complex, thereby maintaining it in the inactive state. There is a variety of potential combinations for the NF- κ B complex. In the CNS, the most abundant forms are the p65/p50 heterodimer and the p50 homodimer, while I κ B α is the most common inhibitory protein. A vast array of stimuli are able to trigger NF- κ B activation by two main activating pathways, referred to as the canonical (or classic) and the non-canonical (or alternative) pathways. The canonical NF- κ B signaling pathway is crucial for the activation of innate immunity and inflammation. In response to a several proinflammatory cytokines, including tumor necrosis factor α (TNF α), interleukin-1 β (IL-1 β), and agonists of the toll-like receptor, such as lipopolysaccharide (LPS), the I κ B Kinase (IKK) complex is activated via phosphorylation of consensus motives in the I κ B members (I κ B α , I κ B β , I κ B ϵ and p105). More specifically, phosphorylation on Ser32 and Ser36 residues of

I κ B α by the I κ B Kinase- β (IKK β) catalytic subunit of the IKK complex (composed of IKK β and IKK α , the catalytic subunits and the IKK γ , the regulatory subunit) triggers ubiquitination and proteasome-mediated degradation of the inhibitory protein. Once activated, the NF- κ B p65/p50 dimer translocates into the nucleus, where it binds to the DNA consensus sequence of target genes. The non-canonical NF- κ B signaling pathway is critical for adaptative immunity and is activated by some particular members of the TNF family (i.e. CD40L; Lymphotoxin- β). The non-canonical pathway involves NF- κ B inducing kinase (NIK)-induced phosphorylation of the IKK α subunit within the IKK complex, that in turn phosphorylates the p100 subunit of the heterodimeric complex p100/RelB. Following ubiquitination and proteasome-dependent processing of p100 into p52, the nuclear translocation of the dimer RelB/p52 occurs and regulates target genes expression. A third distinct NF- κ B signaling pathway triggered by DNA damage and oxidative stress, is defined as atypical because it is independent of the IKK complex (Viatour et al. 2005). Although in this pathway proteasome-mediated degradation of I κ B is involved, the upstream kinases and the phosphorylation sites are different and, as of today, not completely understood (Figure 8).

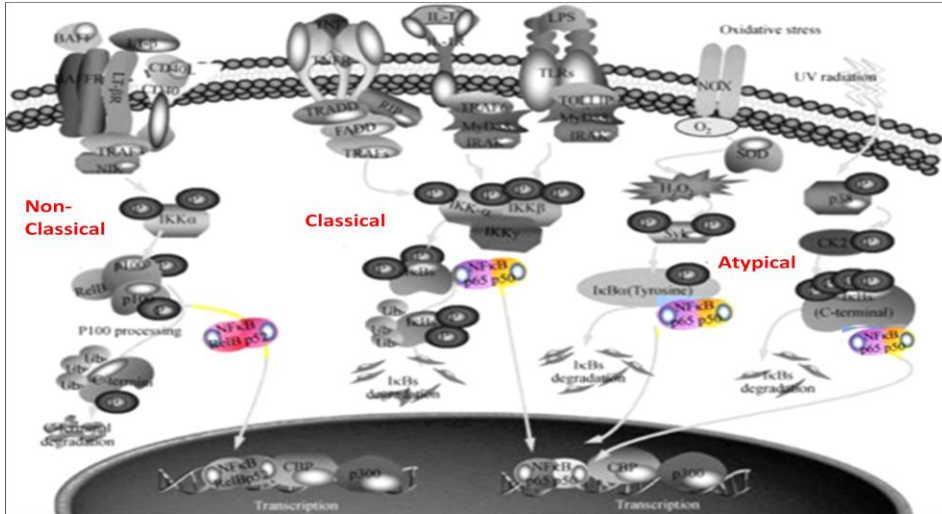


Figure 8. Signaling pathways involved in NF-κB activation. In the canonical pathway, IKK complex activation leads to IκBs phosphorylation, followed by their ubiquitination and degradation by the proteasome, allowing the release of the heterodimer of p65/p50, which then translocates into the nucleus. The non-canonical pathway is dependent on NIK and IKKα, but not on the trimeric IKK complex, and mediates the activation of RelB/p52 complex. In the non-canonical pathway NIK-induced phosphorylation of IKKα leads to the phosphorylation of p100 and proteasome-mediated processing of p100 into p52, allowing the nuclear translocation of RelB/p52 to regulate target gene expression. The atypical or IKK-independent pathway of NF-κB is stimulated by DNA damage (Modified from Zhang and Hu 2012)

1.3.2 NF- κ B signaling in the central nervous system: a role in synaptic plasticity, learning and memory

In adult rodent brain all the DNA-binding subunits of NF- κ B have been detected, with p50/p65 and p50/p50 dimers representing the most abundant forms (Kaltschmidt et al. 1993; Bakalkin et al. 1993; Schmidt-Ullrich et al. 1996). It has been reported that constitutive NF- κ B activity (that is in the nucleus, free of I κ B and capable of binding DNA) has been detected in hippocampal neurons *in vitro*, as well as it occurs *in vivo* in several rodent brain regions, within neurons of neocortex, olfactory bulbs, amygdala, hippocampus and hypothalamus (Bhakar et al. 2002; Schmidt-Ullrich et al. 1996). This piece information supports a physiological role for NF- κ B in maintaining survival or functions of central neurons. In addition, NF- κ B acts also as a signal transducer, not only as a transcriptional regulator, by delivering transient signals from distant active synaptic sites to the nucleus, for example upon glutamatergic stimulation or depolarization (with potassium chloride or kainate) (Wellmann et al. 2000). This supports the concept that the activated transcription factor has a central role in translating short-term synaptic events into long-term gene expression changes (Kaltschmidt and Kaltschmidt 2009).

Several studies based on genetic and pharmacological evidence have shown that NF- κ B is implicated in the regulation of structural changes observed in synaptic connections and in controlling structural plasticity of neural processes (i.e. development, outgrowth, guidance and branching of axons and dendrites) (Gavalda et al. 2009; Salles et al. 2014), and that the signaling pathway is activated by the enhancement of synaptic activity (Meberg et al. 1996; Romano et al. 2006).

Genetic studies associated with a variety of behavioural paradigms are in support for the role of NF- κ B signaling in learning and memory. The availability of transgenic and knock out (KO) mouse lines for specific NF- κ B subunits has been instrumental to research efforts aimed at a better

understanding of their specific role in CNS-related physiological and pathophysiological functions. Mice with a null mutation of p50 (p50^{-/-}) show cognitive defects on novel task acquisitions (Kassed et al. 2002) and reduced short term spatial memory performance, without impairment of long term memory (Denis-Donini et al. 2008). However, the p50 subunit is a component of the transcriptional activator p65/p50 heterodimer, and a repressor in the p50/p50 homodimers, that lack the TAD. Therefore, some of the effects observed in p50^{-/-} mice might be due to the loss of p50 function as repressor as well as activator in combination with p65. Recently, the group of Albensi, found that the lack of the NF-κB p50 subunit led to significant decreases in late long-term potentiation (LTP) and deficits in long-term memory (Oikawa et al., 2012), supporting the idea that p50 may also be required for spatial memory and long term plasticity. The deletion of p65 gene has been shown to be lethal during embryonic development, because of TNFα induced liver degeneration. To overcome this problems, the strategy of p65 deletion in association with concurrent deletion of type I TNF receptors, allowed to create a mouse line (TNFR^{-/-}/p65^{-/-}-double KO mice) which survive postnatally (Alcamo et al., 2001). These mice have normal brain cytoarchitecture but when analysed for memory performance using a spatial version of a radial arm maze, they showed selective learning deficits in spatial information storage (Meffert et al., 2003). These studies point out the role of NF-κB p50 and p65 in memory and hippocampal dependent tasks.

LTP and long-term depression (LTD) are models of synaptic plasticity, believed to be the molecular correlates of memory formation and forgetting, since able to increase or decrease the synaptic efficacy, respectively (de la Fuente et al., 2015; Snow et al., 2014). These paradigms are used to measure synaptic function and along with behavioral test of memory are used to assess hippocampal-dependent alterations in memory. *In vivo*, LTP induction increases the expression and the activation of NF-κB, with both p50 and p65 increase in the hippocampus in response to LTP (Meberg et al. 1996). Instead, the

inhibition of NF- κ B reduces LTP. This suggests that NF- κ B is required for LTP induced synaptic plasticity, and that NF- κ B activation is at the basis of synaptic changes occurring during memory storage. The dendritic spines, site of excitatory and inhibitory synapses on neurons, are dynamic structures with a high turnover during development, whereas throughout life the spine turnover continues at a low rate. The finding that instructive experience leads to spine stabilization, changes in spine density and increase in spine volume during LTP (Alvarez & Sabatini, 2007; Saneyoshi et al., 2010), have linked structural spine plasticity occurring in learning and memory with NF- κ B signaling.

As of today, many target genes being regulated by NF- κ B in neurons have been discovered, and many of these genes are related to memory and plasticity. Those genes can be categorized in several groups. There are genes codifying for molecules involved in cell adhesion and growth, including neural cell adhesion molecule (NCAM) and β 1-integrins, both controlling signals that induce neurite outgrowth during synaptic plasticity. Genes encoding for neurotrophins, such as BDNF, which results upregulated upon NF- κ B activation in astrocytes and neurons and might be connected to cellular growth, differentiation and neurogenesis. Genes involved in cell signaling, such as protein kinase A (PKA). Kaltchmidt and colleagues (2006) reported that NF- κ B regulates the PKA/CREB pathway that in turn controls spatial memory formation and synaptic plasticity (Kaltschmidt et al., 2006). Imielski and colleagues (2012) using a mouse model with neuronal specific ablation of NF- κ B subunits (I κ B/tTA mice - where neuronal NF- κ B ablation can be established by CamKinaseII promoter-driven tetracycline controlled transactivator (tTA) regulating the expression of I κ B in mice, where the transgene expression starts in type 2b DCX⁺ cells) reported that NF- κ B controls the transcription of the forkhead transcription factor (FOXO1) and PKA kinase cascade, involved in axonal outgrowth (Imielski et al., 2012).

In glial cells, there is no constitutive activity of NF- κ B, suggesting that NF- κ B activation might be connected to pathological events. It is suggested that NF- κ B signaling in astrocytes might regulate chemokine-induced infiltration of immune cell into lesioned brain (Kaltschmidt and Kaltschmidt 2009). Astrocytes have beneficial functions during the early stages of injury, whereas sustained activation of astrocytes enhances neuroinflammation. Astroglial-derived mediators, like TNF α , recruit and enhance microglia/macrophage responses. This event is driven by astroglial NF- κ B, which establishes an inflammatory intracellular environment. The group of Zheng and colleagues (2015) reported that in astrocytes, upon amyloid- β stimulation, there is a sustained activation of NF- κ B, that leads to target upregulation and subsequent release of complement factor C3, which in turn, impairs neuronal function in Alzheimer's disease (Lian et al., 2015). Dvorianchikova and colleagues (2009) showed that the transgenic inhibition of astroglial NF- κ B (using GFAP-I κ B α -dominant-negative mice, in which NF- κ B is inactivated in astrocytes through overexpression of a dominant negative form of I κ B α super-repressor under control of GFAP promoter) abrogates the establishment of the inflammatory cellular environment in a model of retinal ischemia-reperfusion injury (Dvorianchikova et al., 2009). The same group reported that the transgenic inhibition of astroglial NF- κ B has also neuroprotective effects in experimental autoimmune encephalomyelitis (Brambilla et al., 2009). Both publications demonstrate that selective inhibition of astroglial NF- κ B significantly reduces the severity of the disease models. NF- κ B role in astrocytes needs further investigation in order to explore the neuroprotective and and/or neurotoxic functions of the pathway in glial cells.

1.3.3 NF- κ B signaling in adult neurogenesis: *in vitro* and *in vivo* supporting data

NF- κ B family members are expressed in zones of active neurogenesis in the adult as well as in the postnatal mouse brain (Denis-Donini et al., 2005), suggesting that the signaling pathway may be actively involved in some of the steps accounting for neurogenesis, such as proliferation, survival, migration and differentiation (Widera et al. 2008; Zhang and Hu 2012). Direct evidence of the presence of NF- κ B in neurogenic zones has been validated by immunofluorescent microscopy (Meneghini et al., 2010). The group of Zhang and colleagues (2012) has shown that NF- κ B signaling is required for NSPC initial differentiation (Zhang et al., 2012). They demonstrated that selective inhibition of canonical NF- κ B subunits and blocking of the signaling pathway (by pharmacological stimuli, such as SN50 - a cell-permeable direct inhibitor for NF- κ B nuclear translocation; or with small hairpin RNA (shRNA) knockdown of NF- κ B subunits; or with NSPC-specific transgene dominant-negative I κ B α), it is possible to attenuate neural differentiation and enhance self-renewal capability of Nestin⁺ and Sox2⁺ NSPC (Zhang et al., 2012). Under physiological conditions, NF- κ B is a crucial regulator of dentate gyrus tissue homeostasis (Imielski et al., 2012), however under pathological conditions, stress and/or aging, over-activation of NF- κ B signaling could impair and reduce neurogenesis by depleting the hippocampal neural stem cell pool (Encinas and Sierra, 2012; Koo et al., 2010).

Many additional studies have implicated NF- κ B signaling in regulating neurogenesis during development and in the adulthood. Several extracellular signals have been identified as being able to affect NSPC via NF- κ B activation. Several cytokines have been shown to regulate neurogenesis by stimulating NF- κ B signaling. IL-1 β is a potent proinflammatory cytokine shown to mediate stress-induced depression via inhibiting adult neurogenesis. Koo and Duman (2008) reported that blockade of the IL-1 β receptor (IL-1R1) by using either an antagonist or IL-1R1 null mice (adult male IL-1R1 null mice bearing a C57BL/6

background) blocks the antineurogenic effect of stress. In the *in vitro* studies they demonstrated that hippocampal neural progenitor cells express IL-1 β receptor-1 (IL-1R1) and that activation of this receptor decreases cell proliferation via the NF- κ B signaling pathway (Koo and Duman, 2008). It has also been reported that IL-1 β exerts also anti-proliferative, anti-neurogenic and pro-gliogenic effect on embryonic rat hippocampal NSPC, which is mediated by IL-1R1 (Green et al., 2012). On the contrary, the cytokine TNF- α was identified to trigger survival, proliferation, and neuronal differentiation from both adult rat and neonatal mice cultured SVZ NSPC, via NF- κ B signaling *in vitro* (Widera et al. 2006; Zhang and Hu 2013). Other data suggest that TNF- α promotes gliogenesis but not neurogenesis (Ricci-Vitiani et al., 2006). A recent report suggests that the mechanism that may contribute to the neuroprotective activity of TNF- α is dependent on the regulation of Ephrin B2 gene by NF- κ B pathway activation (Pozniak et al. 2013). Notwithstanding the negative effect of TNF- α on neuronal survival is dependent on the context, timing and dosage of its activity (Lenzlinger et al. 2001). Increasing interest has been devoted to another cytokine, interleukin-6 (IL-6), that has been shown to induce NSPC to differentiate specifically into glutamate-responsive neurons, as well as into phenotypically different glia types (Mukaino et al., 2008). Another example comes from studied from Wada and colleagues, who demonstrated that vascular endothelial growth factor (VEGF) directly promotes adult NSC survival (Wada et al., 2006). It was also demonstrated that VEGF can influence adult hippocampal neurogenesis and hippocampus-dependent memory (Licht et al., 2011), and the different VEGF phenotypes were blocked by inhibition of the downstream effector NF- κ B. Taken together, these data support the hypothesis that in physiological conditions or in early stages of injuries, transient NF- κ B activation by these cytokines/GF, may be beneficial by promoting neural progenitor cells differentiation for neural repair or maintaining daily neurogenesis. However, chronic activation of NF- κ B would result in the reduction of the pool of NSPC (Koo et al., 2010). These results emphasise the

influence of local environment on the differentiation of NSPC during development,

Many receptors have been identified playing a pivotal role in NF- κ B activation and NPC differentiation, upon ligand binding. Schwartz and colleagues, identified toll like receptors (TLR) as players in adult neurogenesis. TLR2 and TLR4 are found on adult NSPC and have distinct and opposing functions in NPC proliferation and differentiation both *in vitro* and *in vivo*. The activation of TLRs on the NPCs was mediated *via* activation of the NF- κ B signaling pathway (Rolls et al., 2007).

Among the upstream receptors that activate NF- κ B signaling pathway in NPC of adult mouse hippocampal and SVZ neurogenic region, there is the receptor for advanced glycation end-products (RAGE). RAGE ligands, like the alarmin HMGB1, S100 β and AGE-BSA, are able to stimulate both proliferation and neuronal differentiation of SVZ-derived NPC *in vitro* (Meneghini et al., 2010). It was also demonstrated that HMGB1 and A β ₁₋₄₂ oligomers, both involved in Alzheimer's disease pathophysiology (Takata et al., 2003) are able to promote neuronal differentiation of adult hippocampal NPC via activation of the RAGE/NF- κ B axis (Meneghini et al., 2013).

Literature data supports the involvement of NF- κ B signaling pathway in mediating the proneurogenic and antidepressant-like activity of some clinically relevant drugs. Increasing evidence demonstrate that neuroplasticity is disrupted in mood disorders and in animal models of depressive-like behaviour and it can be upregulated by antidepressant drugs (Pittenger and Duman, 2008; Hanson et al., 2011). An intriguing observation is that most antidepressant and environmental interventions (i.e. learning tasks; enriched environment; physical activity) that confer antidepressant like-behavioral effects also stimulate adult neurogenesis (Sahay and Hen, 2007).

Our group recently identified novel positive modulators of hippocampal neurogenesis, as potential new antidepressant drugs, that require NF- κ B

activation for exerting their proneurogenic effects *in vitro* and *in vivo*. The first example is represented by pregabalin (PGB) and gabapentin (GBP) that are clinically relevant anticonvulsant, analgesic and anxiolytic drugs. Our group demonstrated that GBP and PGB produce a concentration dependent increase in the number of mature and immature neurons *in vitro*, acting through the $\alpha 2\text{-}\delta$ subunit of the voltage-gated calcium channels, which was found to be expressed on adult hippocampal NPC (Valente et al., 2012). The activation of NF- κ B is involved in the proneurogenic effect elicited by $\alpha 2\text{-}\delta$ ligands, because the inhibition of both p50 and p65 nuclear translocation of IKK β counteracted drug-mediated effects. Moreover, PGB antidepressant like activity was confirmed *in vivo* in adult mice. The appearance of depressive-like behaviour induced by chronic restraint stress was prevented by administration of the drug, and in parallel promoted adult hippocampal neurogenesis (Valente et al., 2012). Another drug reported to have proneurogenic and antidepressant-like activity *via* modulation of NF- κ B signaling pathway is acetyl-L-carnitine (ALC) (Cuccurazzu et al., 2013). Our group demonstrated that the drug-mediated effect on neuronal differentiation of adult hippocampal NPC *in vitro*, is dependent on the acetylation of p65 subunit and subsequent NF- κ B-mediated upregulation of metabotropic glutamate receptor 2 (mGlu2). In addition, when tested *in vivo*, chronic ALC treatment significantly increased the number of adult born neurons in mouse hippocampi and reverted depressive-like behaviour induced by unpredictable chronic mild stress.

Our group investigated the role of NF- κ B p50 subunit in adult hippocampal neurogenesis by taking advantage of p50^{-/-} mice generated in the laboratory of David Baltimore (Sha et al., 1995). In this genetically modified animal model, we demonstrated a marked deficiency in neurogenesis in the hippocampus, as a result of a defect in the late maturation of newly born neurons (Denis-Donini et al., 2008). Interestingly, no defects have been observed in the rate of NSPC proliferation in the SGZ of p50^{-/-} animals compared to wt mice. However, the survival of neural progenitors after 21 days post BrdU injection was

significantly reduced in mutant compared to wt animals. A detailed phenotypic characterization of newly generated cells revealed no differences in the number of DCX⁺ neuroblasts, but a marked reduction in calretinin post mitotic neurons in the DG of p50^{-/-} animals. Based on this findings, our group proposed that the absence of the p50 subunit of NF-κB may trigger a selective defect in adult neurogenesis progression at the transition phase between neuroblasts and postmitotic neurons.

p50^{-/-} mice were also evaluated at the behavioural level. When tested in the Morris water maze (MWM), wt and p50^{-/-} performed equally in the acquisition test, with normal retrieval of hippocampal-dependent spatial memory in the probe test performed 24h after the last training trial. Conversely, when wt and p50^{-/-} mice were subjected to the place recognition test for evaluating their ability to discriminate familiar versus a novel environment, the two genotypes performed differently. Specifically, p50^{-/-} mice showed a selective impairment in short-term spatial memory. The correlation between neurogenesis defects and deficits in short-term spatial memory performances deserves further investigation.

Many other publications support the importance of NF-κB in synaptic plasticity and memory. The group of Albensi investigated the role of NF-κB in both synaptic plasticity and long term spatial memory in p50^{-/-} mice (Oikawa et al., 2012). They used the same KO mice model to test if the absence of p50 negatively affected *in vivo* MWM performance and *in vitro* LTP response. LTP is an experimental paradigm used to assess hippocampal-dependent alterations and measure synaptic function, and is commonly regarded as the molecular correlate of memory formation. Albensi's group found that the lack of the NF-κB p50 subunit led to a significant decrease in late LTP and in selective alterations in MWM test, with defects in long term spatial memory. Given these contradictory findings, open questions remain concerning the complex contribution of NF-κB p50 subunit in long versus short memory.

The group of Imielski found out that NF-κB ablated mice (IκB/tTA mice) show a strongly increased number of activated astrocytes that is accompanied with a

severe dentate gyrus atrophy. They proposed that this condition may rely on a dual function of NF- κ B in the hippocampus. They suggest a cell autonomous role of NF- κ B in axogenesis, with NF- κ B being necessary for axogenesis and maturation in neuronal progenitors, whereas in mature granule cells NF- κ B may play a distinct role by regulating neuroprotection as well as synaptic transmission. When NF- κ B is inactivated, the dentate gyrus cannot be regenerated by the addition of new-born neurons (Imielski et al., 2012).

Several studies that demonstrated increased hippocampal neurogenesis following hippocampus-dependent tasks, have established a connection between neurogenesis in the adult hippocampus and learning and memory. The deficiency in hippocampal neurogenesis and the selective defect in short term spatial memory performance in adult p50^{-/-} animals, provide evidence that NF- κ B-regulated hippocampal neurogenesis, can influence aspects connected to hippocampal dependent learning.

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Chapter 2.

Outline of the thesis

Neurogenesis occurring during postnatal life is restricted to two selected regions in the adult brain of mammals. The canonical and well-studied neurogenic sites are the subgranular zone (SGZ) in the dentate gyrus of the hippocampus, and the subventricular zone (SVZ) of the lateral ventricles (Ming and Song 2005; Bond et al. 2015; Kempermann et al. 2015). Emerging evidence recently suggested a novel neurogenic site along the walls of the third ventricle in the hypothalamic area, whose features and functions are under investigation (Kokoeva et al., 2005; Robins et al., 2013).

Large interest grew for hippocampal adult neurogenesis due to the involvement of this brain region in cognitive functions like learning and memory, processes relevant for human beings, as well as in emotional behavior and stress response (Aimone et al. 2010; Vadodaria and Jessberger 2014).

Adult hippocampal neurogenesis (ahNG) is a highly dynamic and plastic process strictly connected to environmental changes. Many extrinsic modulators (i.e. learning, enriched environment, physical activity; antidepressants drugs) have been reported to enhance hippocampal neurogenesis and, in parallel, long term potentiation and cognitive performances, reversal of depressive-like behaviour and resilience to stress in mouse models (Hanson et al., 2011). On the other hand, physio- and pathological conditions like aging, stress and inflammation, as well as addictive drugs, have been reported as negative regulators which result in decreased neurogenesis and performance in tasks of spatial navigation and alteration in emotion-related behaviour (Lazarov et al., 2010; Yirmiya and Goshen, 2011; Lucassen et al., 2015). Last, but not least, it is of great interest that hippocampal neurogenesis is deregulated in various neuropsychiatric disorders, such as major depressive disorder and anxiety, schizophrenia, and neurodegenerative diseases (Winner et al., 2011; Nath et al., 2012; Schoenfeld and Cameron, 2015; Miller and Hen, 2015).

Adult neurogenesis occurs in a permissive and instructive microenvironment referred to as the “neurogenic niche” where local environmental cues, both

membrane-associated and released by resident cells, interact with neural/stem progenitor cells (NSPC) and in turn affect their functionality (self-renewal and multipotential properties), their commitment and their survival. The niche is the result of the complex interplay among different cell types (NSPC, precursors, astroglia, microglia, neuroblasts), secreted molecules (i.e. cytokines, growth factors, secreted proteins), vasculature, and extracellular matrix (ECM) components (Lin and Iacovitti 2015). Cell-to-cell communication plays a key role in the homeostatic modulation of the neurogenic process under physiological conditions and also in response to damage in pathological conditions. Among the niche cellular components, astrocytes represent one important contributor which can instruct NSPC proliferation, affect the maintenance of their undifferentiated state, modulate their lineage differentiation through both contact-dependent interactions and secretion of soluble molecules. In addition, astrocytes can also influence maturation, survival and integration of newly generated neurons by providing signals which are critical for synaptogenesis and dendritic spine maturation (Song et al. 2002; Oh et al. 2010; Risher and Eroglu 2012).

In previous publications our lab proved that members of the NF- κ B family of transcription factors are important contributors of signaling pathways in the SGZ neurogenic niche (Denis-Donini et al., 2005) and may be potentially involved in mediating the proneurogenic and antidepressive-like effect of some clinically relevant drugs (i.e. α 2 δ ligands, acetyl-L-carnitine) (Cuccurazzu et al., 2013; Valente et al., 2012) and in the proneurogenic effects of damage-associated signals (i.e. HMGB-1 and A β ₁₋₂) (Meneghini et al., 2013). In the CNS NF- κ B is involved in neuroprotection/neurodegeneration, differentiation, migration, synaptic plasticity, and neurite outgrowth, all functions making this family of transcription factors potential candidates for translating complex signals that regulate the neurogenic process (Grilli and Memo, 1999; Kaltschmidt and Kaltschmidt, 2009; Bortolotto et al., 2014). Within the family, the p50 subunit appears to play a crucial role since p50KO mice display

dramatically reduced adult hippocampal neurogenesis which is in association with short-term memory defects (Denis-Donini et al., 2008).

The main project for this PhD thesis developed from the observation that unlike *in vivo*, hippocampal NPC derived from WT and p50KO adult mice have similar rate of differentiation, proliferation and apoptosis *in vitro*.

Based on these observations, in our first investigation, we exploited the advantages of using primary cell culture systems from WT and p50KO mice to investigate the contribution made by astrocyte conditioned media (ACM) under basal conditions, on adult NPC fate specification. In particular, we wanted to elucidate the involvement of astroglial cells in the defective hippocampal neurogenesis observed *in vivo*. In our *in vitro* experimental context, we analyzed the role of the NF- κ B pathway in the cross-talk between astrocytes and adult hippocampal NPC. For this purpose, we set up primary astroglial cultures, as well NPC cultures, from both WT and p50KO mice and analyzed the phenotypic commitment of WT/p50KO NPC when exposed to WT/p50KO astrocyte-conditioned media. On one hand, we attempted to identify the proneurogenic and/or antineurogenic signals that, under control of NF- κ B p50, may be differentially expressed in astrocytes. On the other hand, we also focused our attention on the possibility that absence of p50 may produce cell-autonomous defects in NPC responsiveness to astrocytic-derived factors.

An additional line of investigation was devoted to explore region-specificity in astrocyte-derived signals that may affect cell fate commitment of hippocampal adult NPC. In particular, we analyzed differences in the pro-neurogenic and -gliogenic potential of hippocampal and hypothalamic astrocytes of both WT and p50KO origin. Additional work was performed with the purpose of understanding fate specification of hypothalamic WT NPC cultures in presence hippocampal ACM.

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Chapter 3.

Astrocyte-mediated neuronal fate specification of adult hippocampal neural progenitors is regulated by NF- κ B p50.

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Running title: Role of NF- κ B p50 in adult neural progenitor-astrocyte cross-talk

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Abstract

In previous work we demonstrated that NF- κ B p50 acts as crucial regulator of adult hippocampal neural progenitor cells (ahNPC). Indeed, NF- κ B p50KO mice are characterized by remarkably reduced hippocampal neurogenesis. As a follow up to that work, herein we show that when cultured *in vitro*, ahNPC from wild type (WT) and p50KO mice are not significantly different in their neurogenic potential. This observation prompted us to investigate cell autonomous and non-cell autonomous consequences of p50 absence on neuronal fate specification of ahNPC. In particular, we focused our attention on astrocytes, known to provide soluble proneurogenic signals, and investigated the influence of WT and p50KO astrocyte conditioned media (ACM) on WT and p50KO ahNPC differentiation. Interestingly, while WT ACM promoted both neuronal and astroglial differentiation, p50KO ACM only supported astroglial differentiation of WT ahNPC. By using a LC-MS/MS approach we identified several proteins which are significantly upregulated in p50KO compared to WT astrocytes. Among them, lipocalin-2 (LCN-2) was recognized as a novel astroglial-derived signal regulating neuronal fate specification of ahNPC. Interestingly, LCN-2 proneurogenic effect is greatly reduced in p50KO NPC. In addition, we demonstrated p50KO NPC unresponsiveness to both neuronal and astroglial fate specification signals from WT and p50KO ACM, and we identified reduced expression of $\alpha 2\delta 1$, a thrombospondin-1 receptor, as one phenotypic change occurring in ahNPC in absence of p50. Altogether our data suggest that reduced hippocampal neurogenesis in p50KO mice *in vivo* is to be ascribed to cell autonomous and, likely, to non-cell autonomous defects in the hippocampal neurogenic niche.

Introduction

Adult neurogenesis occurs in restricted areas of postnatal and adult mammalian brain, namely the subventricular zone (SVZ) in the lateral wall of the lateral ventricles and the subgranular zone (SGZ) in the dentate gyrus (DG) of the hippocampus (Ming and Song, 2005; Bond et al., 2015; Kempermann et al., 2015). Recent studies also suggested an additional neurogenic region in the adult hypothalamus (Kokoeva et al., 2005; Whalley, 2012).

Adult hippocampal neurogenesis (ahNG) has attracted great research interest due to its potential involvement in critical functions including cognition, mood and emotional behavior, stress response (Aimone et al., 2010, 2014; Lacar et al., 2014). The molecular mechanisms that modulate neurogenesis are very complex and deserve extensive investigation which may turn into a better understanding of homeostatic regulation of ahNG but also of neuropsychiatric/neurodegenerative disorders where this process is profoundly deregulated (Winner et al., 2011; Apple et al., 2016).

Within the CNS, NF- κ B pathway activation has been involved in a wide range of functions both under physiological and pathological conditions (Grilli and Memo, 1999; Mattson and Meffert, 2006; Kaltschmidt and Kaltschmidt, 2009). A few years ago our group demonstrated that NF- κ B proteins are expressed at considerable levels in areas of active neurogenesis in the postnatal and adult mouse brain (Denis-Donini et al., 2005). Since then, a vast array of information has been collected on the complex involvement of NF- κ B proteins in different aspects of postnatal neurogenesis. In particular, several extracellular signals and membrane receptors have been identified as being able to affect neural stem cells/neural progenitor cells (NSC/NPC) and their progeny via NF- κ B activation (Wada et al, 2006; Rolls et al., 2007; Widera et al., 2008; Meneghini et al., 2010, 2013).

Since the initial observations of NF- κ B members being present in adult neurogenic areas, our group further explored the role of these regulatory proteins in adult neurogenesis with a specific focus on the NF- κ B1 p50 subunit

(Grilli and Meneghini, 2012; Bortolotto et al., 2014). By taking advantage of p50KO mice (Sha et al., 1995) we demonstrated that absence of p50 can deeply affect the *in vitro* response of adult hippocampal NPC (ahNPC) to several endogenous signals (Meneghini et al., 2013), and to proneurogenic drugs (Valente et al., 2012). *In vivo* we also proved that p50KO mice display a dramatic reduction in adult hippocampal neurogenesis which correlates with a selective defect in hippocampal-dependent short-term memory (Denis-Donini et al., 2008). Interestingly, *in vivo* the proliferation rate of hippocampal NSC/NPC in p50KO mice appeared to be similar to that of WT mice. Similarly, the apoptotic rate in the hippocampal region was not increased in mutant mice compared to their WT counterpart. A detailed phenotypic characterization of newly generated hippocampal cells in p50KO mice suggested that absence of the NF- κ B p50 subunit may trigger a rather selective defect in neuronal differentiation of ahNPC (Denis-Donini et al., 2008).

Homeostasis of adult neurogenesis requires permissive and instructive signals for aNSC/NPC. Several elegant studies have investigated the mechanisms through which local environment in the neurogenic niche may control fate specification of aNSC/NPC (Schänzer et al., 2004; Bonaguidi et al., 2005; Lu and Kipnis, 2010). Among others, a pivotal work by Song et al. (2002) demonstrated that astroglial cells are crucial regulators in the hippocampal niche, where they can instruct stem cells to adopt a neuronal fate (Song et al., 2002). Since the well documented role of NF- κ B in neuronal but also in non-neuronal cells (Brambilla et al., 2009; Lian et al., 2015; de la Fuente et al., 2015), we recently decided to further dissect the role of NF- κ B p50 in the cross-talk between adult neural progenitor cells and astrocytes. Exposure of WT and p50KO hippocampal NPC to conditioned media from WT and p50KO astrocytes was utilized as a strategy to study the potential influence of astroglia on adult NPC fate specification and the cell autonomous or non-cell autonomous role played by NF- κ B p50 signaling in that context.

Materials and Methods

Animals. Wild type (WT; C57BL/6; The Jackson Laboratories) and NF- κ B p50^{-/-} (p50KO; C57BL/6 Nfkb1^{tm1Bal}/J; The Jackson Laboratories) mice were housed under light- and temperature-controlled conditions in high-efficiency particulate air (HEPA)-filtered Thoren units (Thoren Caging Systems) at the University of Piemonte Orientale animal facility. Mice were kept 3-4/cage with *ad libitum* access to water and food. Animal care and handling were performed in accordance with European Community Directive and approved by the local IACUC (Institutional Animal Care and Use Committees).

Isolation and culture of WT and p50KO adult mouse hippocampal neural progenitor cells (NPC). For preparing NPC primary cultures from hippocampi, three adult (3-4 month-old) male WT and p50KO mice were sacrificed and cell suspension was prepared. Briefly, the brains were removed, and hippocampi were isolated and collected in ice-cold PIPES buffer pH 7.4 containing 20 mM PIPES, 25 mM glucose, 0.5 M KCl, 0.12 M NaCl (Sigma-Aldrich), and 100 U/100 μ g/ml Penicillin/Streptomycin solution (Life Technologies). After 5 min at 110 x g centrifugation, the tissue was digested for 40 min at 37°C using the Papain Dissociation System (Worthington DBA). Cell suspension was plated onto 25 cm² cell-culture flask (Thermo-Fisher Scientific) and cultured as floating neurospheres through subsequent passages (Valente et al., 2012). Primary (Passage 1, P1) neurospheres were dissociated after 7-10 days *in vitro* (DIV), whereas P2-P30 neurospheres every 5 DIV. At each passage cells were plated in T25 flask at a density of 12,000 cells/cm² in growing medium: Neurobasal-A medium, supplemented with B27 supplement, 2 mM L-glutamine (Life Technologies), human Epidermal Growth Factor (hEGF, 20 ng/ml; Peprotech), basic Fibroblast Growth Factor (bFGF, 10 ng/ml; Peprotech) and Heparin sodium salt (0.0004 %, Sigma-Aldrich).

Adult hippocampal NPC differentiation. NPC (P5-P30) were used for differentiation experiments. Briefly, neurospheres were dissociated into single cells and plated onto laminin-coated Lab-Tek 8-well permanox chamber slides (Nunc) at a density of 43,750 cells/cm² in NPC differentiation medium (NDM, Neurobasal-A medium supplemented with B27, 2 mM L-glutamine and 100 U/100 µg/ml Penicillin/Streptomycin). NPC were treated in presence of hippocampal astrocyte conditioned media (ACM) derived from WT or NF-κB p50KO glial cultures, or in presence of NDM (referred to as standard, STD medium) for 24 h. WT NPC were treated in presence of 0.01–1 µg/ml recombinant mouse lipocalin-2 (LCN-2, Cell Signaling), 2 µg/ml purified human Thrombospondin-1 (hTSP-1, Amsbio), or corresponding vehicle for 24 h. For RAGE inhibition, 60 min before ACM or NDM treatment, 20 µg/ml neutralizing polyclonal anti-RAGE antibody (α-RAGE Ab; R&D System) was added to NPC.

Primary Astrocyte Cultures. Primary mixed glial cultures were prepared from cortex and hippocampus of neonatal (P1-2) C57BL/6 WT and p50KO mice and grown in DMEM high glucose, 10% FBS, 15 mM HEPES, 2 mM L-glutamine and 100 U/100 µg/ml Penicillin/Streptomycin at a density of 78,000 cells/cm². When cells reached confluence (around 10-12 D.I.V.), proliferation of non-astrocytic cells was blocked by 10 µM cytosine arabinoside (Sigma-Aldrich) for 96 h. Then cultures were switched to NDM. Phenotypic characterization of cultures was performed by immunocytochemistry with antibodies against GFAP (mouse monoclonal, 1:600, Millipore) and CD11b (rat monoclonal; 1:150, Millipore). Contaminating microglial cells were below 3% of total cells, with the remaining cells being GFAP⁺. ACM was collected after 48h in presence of NDM. Briefly, ACM was centrifuged for 10 min at 16,000 x g to eliminate cells and debris and passed through 0.2 µm filter before use. ACM was used diluted 1:2 in fresh NDM.

Immunolocalization studies in differentiated adult hippocampal NPC. NPC were treated in presence of WT or p50KO ACM or STD medium for 24 h. Phenotypic characterization of NPC-derived cells was carried out by immunolocalization for MAP-2 (rabbit polyclonal, 1:600; Millipore), GFAP (mouse monoclonal, 1:600; Millipore), nestin (chicken monoclonal, 1:1,500; Neuromics), NG-2 (rabbit polyclonal, 1:500; Millipore). Secondary antibodies were as follows: Alexa Fluor 555-conjugated goat anti-rabbit (1:1,400; Molecular Probes); Alexa Fluor 488-conjugated goat anti-chicken antibody (1:1,400; Molecular Probes); Alexa Fluor 488-conjugated goat anti-rabbit (1:1,400; Molecular Probes), Alexa Fluor 488-conjugated goat anti-mouse (1:1,600; Molecular Probes). Nuclei were counterstained with 0.8 ng/ml Hoechst (Thermo Fisher scientific) diluted in PBS. In each experiment, 5 fields/well (corresponding to about 150-200 cells/well) were counted with a 60X objective by a Leica DMIRB inverted fluorescence microscope. Cells positive for each marker were counted and their percentage over total viable cells was calculated. All experiments were run in triplicates using different cell preparations and repeated at least three times.

LC-MS/MS analysis and protein quantification. WT and p50KO ACM were subjected to trichloroacetic acid (TCA) precipitation. Briefly, TCA was added to a final 12% (wt/vol) concentration and kept on ice for 2 h, followed by centrifugation (16,000 x g, 10 min). The supernatant was carefully removed, and 1 ml of tetrahydrofuran (pre-cooled in ice) was added to each pellet and vortexing was carried out until the pellet dissolved completely. Proteins were then reduced with dithiothreitol (DTT), alkylated by iodoacetamide, and digested with trypsin at 37°C O/N. Data acquisition was performed with a micro-LC Eksigent Technologies (Dublin, USA) system, with a stationary phase a Halo Fused C18 column (0.5 x 100 mm, 2.7 µm; Eksigent Technologies), interfaced with a Triple-TOF 5600+ system (AB SCIEX) equipped with a DuoSpray Ion source. The mobile phase was a mixture of 0.1% (v/v) formic acid in water (A) and 0.1% (v/v) formic acid in acetonitrile (B),

eluting at a flow-rate of 15.0 mL min⁻¹ at an increasing concentration of solvent B from 2% to 40 % in 30 minutes. An initial data-dependent acquisition (DDA) analysis was performed on the p50KO and WT ACM samples to generate the SWATH-MS (Sequential Window Acquisition of Theoretical mass spectra) spectral library: the mass spectrometer analysis was performed using a mass range of 100–1500 Da (TOF scan with an accumulation time of 0.25 s), followed by a MS/MS product ion scan from 200 to 1250 Da (accumulation time of 5.0 ms) with the abundance threshold set at 30 cps (35 candidate ions can be monitored during every cycle). The samples were then subjected to cyclic data independent analysis (DIA) of the mass spectra, using a 25-Da window: the mass spectrometer was operated such that a 50-ms survey scan (TOF-MS) was performed and subsequent MS/MS experiments were performed on all precursors. These MS/MS experiments were performed in a cyclic manner using an accumulation time of 40 ms per 25-Da swath (36 swaths total) for a total cycle time of 1.5408 s (Venable et al., 2004, Geromanos et al., 2009; Gillet et al., 2012). The ions were fragmented for each MS/MS experiment in the collision cell using the rolling collision energy. The MS data were acquired with Analyst TF 1.7 (AB SCIEX). Three replicates for each sample were subjected to the DIA analysis. The mass spectrometry files were searched using Protein Pilot (AB SCIEX) and Mascot (Matrix Science Inc.). For Protein Pilot the following parameters were used: cysteine alkylation, digestion by trypsin, no special factors and False Discovery Rate at 1%. For Mascot the following parameters were used: the digestion enzyme selected was trypsin, with 3 missed cleavages, a search tolerance of 0.4 Da was specified for the peptide mass tolerance, and 0.4 Da for the MS/MS tolerance. The charges of the peptides to search for, were set to 2+, 3+ and 4+, the search was set on monoisotopic mass and the following modifications were specified for the search: carbamidomethyl cysteins as fixed modification and oxidized methionine as variable modification. The search was conducted using the UniProt Swiss-Prot database of mouse proteins (version 25.03.2015, containing 41741 sequence entries). Dual filtering criteria for protein identification were employed by combining

FDR test from target-decoy database search with a cutoff p -value of 0.05, and protein/peptide confidence above 95% probability, with a minimum of two unique peptides per protein.

The label-free quantification was performed with Skyline (MacCoss Lab Software, University of Washington) by importing the SWATH-MS runs. The library of the identified proteins used for the processing of SWATH data was generated by combining the results of the database search performed with Protein Pilot and Mascot (Manfredi et al., 2016). The quantification was performed by integrating the extracted ion chromatogram of all the unique ions for a given peptide and by using MSstats, a Skyline external tool for statistical analysis as previously described (Choi et al., 2014).

Western blot analysis. Protein extracts from adult NPC cultures were obtained by lysis in 3X vol/vol of ice-cold hypotonic RIPA buffer [50 mM Tris-HCl pH 7.5, 150 mM NaCl, 0.5 mM EDTA pH 8, 1% (vol/vol) Triton X-100, 1% (wt/vol) SDS, 10 mM NaF, 1 mM NaVO₄, 1 mM DTT, protease inhibitor mix (Sigma-Aldrich)] for 30 min on ice. To complete the lysis, incubation at -80°C for 3 min followed by 2 min at 37°C was repeated three times. All lysates were centrifuged at 16,100 x g for 10 min at 4°C and supernatants were collected. Protein concentration was determined by Bradford assay (Sigma-Aldrich) and equivalent protein amounts (25 μ g) were separated by SDS-PAGE gel (6%) and transferred onto nitrocellulose membranes. The membranes were blocked with 5% BSA in TBS (Sigma-Aldrich) buffer for 60 min at RT. Immunoblots were carried out overnight in an antibody solution containing 3% (wt/vol) BSA in TBS-tween 0.1% with the primary antibody against the α 2 δ subunit of Dihydropyridine (DHP)/Thrombospondin-1 receptor (1:500; Sigma-Aldrich). After washing, blots were incubated with peroxidase-conjugated goat anti-mouse antibody (1:10,000; BIO-RAD) for 60 min at RT and immunocomplexes were visualized by the Supersignal West Pico Chemiluminescent substrate (Pierce). Densitometric analysis was performed using the Image Lab software

system (Bio-Rad Laboratories) and each band was normalized to β -actin signal (mouse monoclonal, 1:1,000; Sigma-Aldrich).

Statistical analysis. All experiments were run in triplicates using different cell preparations and repeated at least three times. Data were expressed as mean \pm S.D. and analyzed using Student's *t*-test when only two independent groups were compared, or by one-way analysis of variance (ANOVA) followed by Tuckey's *post-hoc* test when three or more groups were compared. Statistical significance level was set for *p* values < 0.05 .

Results

In vitro WT and p50KO ahNPC do not show differences in neuronal differentiation and survival rates.

Multipotent nestin⁺, sox2⁺ and GFAP⁻ NPC isolated from adult mouse hippocampi can be maintained for several passages in an undifferentiated proliferative state (Cuccurazzu et al., 2013). When grown onto laminin-coated chamberslides, upon removal of growth factors and exposure to a serum-free defined medium (STD medium), NPC stop dividing and spontaneously differentiate toward the neuronal lineage. As previously published, by double immunolabeling for markers of neuronal cells (MAP-2) and undifferentiated progenitors (nestin), the appearance of newly generated MAP-2⁺/nestin⁻ neurons and MAP2⁺/nestin⁺ neuroblasts can be evaluated and quantified (Meneghini et al., 2013). Under these experimental conditions, both WT- and p50KO-derived NPC gave rise to the same percentage of neurons and neuroblasts (MAP-2⁺/nestin⁻ cells: 3.13 ± 0.63 and 3.34 ± 0.84 % for WT and p50KO NPC; MAP2⁺/nestin⁺ cells: 32.22 ± 3.16 and 33.15 ± 2.21 % for WT and p50KO NPC; Figure 1). No significant difference in the subpopulations of undifferentiated neural progenitors (MAP-2⁻/nestin⁺ cells: 12.86 ± 2.29 and 12.29 ± 3.20 % for WT and p50KO NPC; Figure 1) and non-neuronal cells (MAP-2⁻/nestin⁻ cells: 51.79 ± 3.83 and 51.23 ± 4.51 % for WT and p50KO NPC; Figure 1) was present. Interestingly, despite the role of the NF- κ B signaling pathway in the regulation of cell fate, no significant difference in the basal apoptotic rate could also be observed between the two genotypes (11.84 ± 6.52 and 12.40 ± 5.87 % for WT and KO NPC; Figure 1), in agreement with the *in vivo* observation in hippocampi of adult WT and p50KO mice (Denis-Donini et al., 2008). Altogether we concluded that, unlike *in vivo* conditions, absence of NF- κ B p50 did not correlate with altered neuronal differentiation of ahNPC *in vitro*.

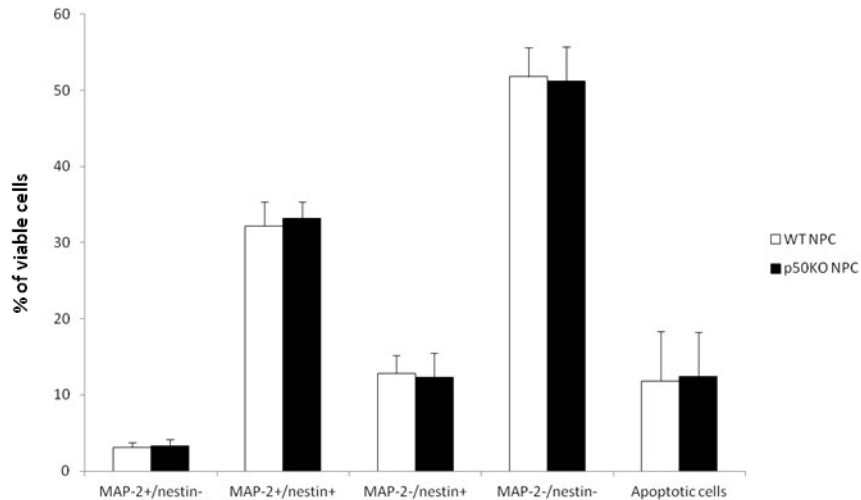


Figure 1. Adult hippocampal NPC derived from WT and p50KO mice show similar *in vitro* differentiation and apoptotic rates. Under differentiating conditions, WT and p50KO hippocampal NPC spontaneously give rise to four cell populations identified by double MAP-2 and nestin immunolabelling: MAP-2⁺/nestin⁻ neurons, MAP-2⁺/nestin⁺ neuroblasts, MAP-2⁻/nestin⁺ undifferentiated progenitors and MAP-2⁻/nestin⁻ cells (NG-2⁺ and GFAP⁺ cells). WT and p50KO hippocampal NPC show similar differentiation rate toward neuronal lineage. The apoptotic rate is not significantly different in WT and p50KO NPC. Data are expressed as mean values \pm S.D. of $n = 4$ experiments, run in triplicates, and analyzed by Student's *t*-test.

Effect of WT- and p50KO-derived ACM on neuronal differentiation of WT ahNPC cultures

The dissonant consequences of NF- κ B p50 absence between *in vitro* and *in vivo* data prompted us to investigate the possibility that non-cell autonomous effects were contributing to remarkably reduced hippocampal neurogenesis in p50KO mice (Denis-Donini et al, 2008). Soluble factors released by astrocytes have been shown to modulate differentiation of NPC (Song et al., 2002). Based on these observations we decided to evaluate the effect of astrocyte-generated

soluble factors on the differentiation rate of ahNPC toward neuronal and non-neuronal lineages. Primary mixed glial cultures were prepared from cortex and hippocampus of neonatal (P1-2) WT and p50KO mice. When cells reached confluence, proliferation of non-astrocytic cells was blocked by 10 μ M cytosine arabinoside so to obtain an astrocytic-enriched cell population with less than 3% of contaminating microglia. Astrocyte-conditioned medium (ACM), diluted 1:2, was added to WT ahNPC cultures. Under these experimental conditions, we tested the effect of WT and p50KO ACM, in comparison with standard differentiation (STD) medium, on NPC differentiation. After 24 h, a significant increase in the percentage of *in vitro* generated neurons and neuroblasts was observed in WT cells exposed to WT-derived ACM, compared to STD medium condition (2.74 ± 0.70 , 5.13 ± 0.88 %, for MAP-2⁺/nestin⁻ cells in STD and WT ACM, $p < 0.001$; 31.12 ± 3.75 , 41.8 ± 3.03 %, for MAP2⁺/nestin⁺ cells in STD and WT ACM, $p < 0.001$; Figure 2 A, B). Under the same *in vitro* experimental conditions, no significant difference was observed between STD medium and p50KO ACM (2.74 ± 0.70 , 2.91 ± 0.64 %, for MAP-2⁺/nestin⁻ cells in STD and p50KO ACM; 31.12 ± 3.75 , 31.66 ± 4.63 % for MAP2⁺/nestin⁺ cells in STD and p50KO ACM; Figure 2 A, B), suggesting that, in absence of NF- κ B p50, primary astrocytes could not produce proneurogenic molecule(s) or, alternatively, did produce antineurogenic molecule(s). To investigate whether ACM could affect cell survival in addition to neuronal differentiation, we also analyzed the apoptotic rate of ahNPC and their progeny. No difference in the percentage of apoptotic cells was observed in the different media conditions (16.37 ± 2.04 , 14.92 ± 1.51 , 14.56 ± 1.46 %, in presence of STD, WT and p50KO ACM, respectively; Figure 2 C). Altogether these data suggested that the absence of NF- κ B p50 may affect the secretory profile of astrocytes, and in particular may reduce their proneurogenic potential on ahNPC, in absence of changes in their effect on the survival rate of NPC and/or their progeny.

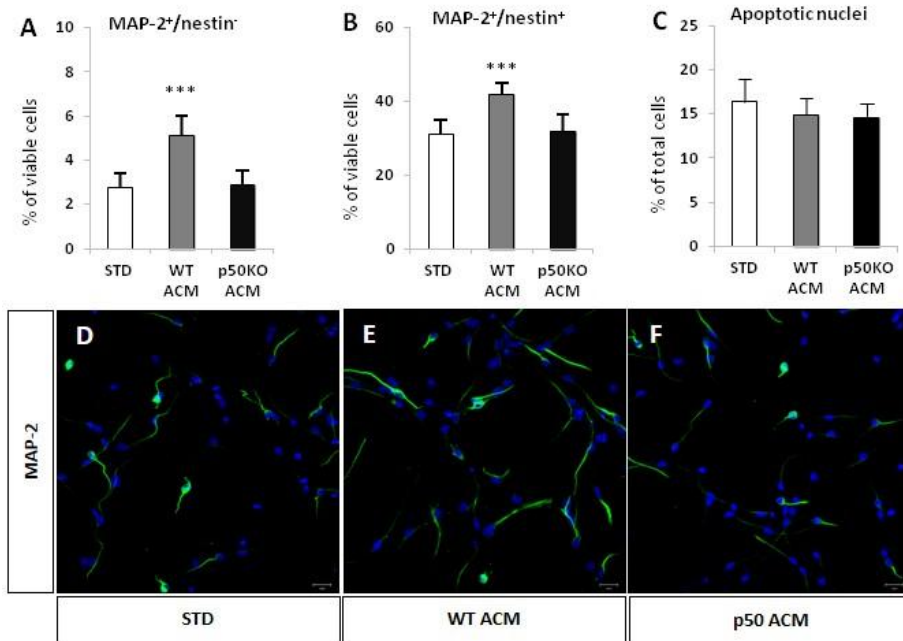


Figure 2. WT, but not p50KO ACM, promote neuronal differentiation of WT ahNPC cultures. *A-B*, Under differentiating conditions, 24h treatment of WT hippocampal NPC with WT ACM significantly increased the percentage of the subpopulations of MAP-2⁺/nestin⁻ neurons (*A*) and MAP-2⁺/nestin⁺ neuroblasts (*B*) compared with standard differentiation medium (STD). Conversely, p50KO ACM was devoid of proneurogenic effect. (*C*) No significant difference was observed in the apoptotic rate of cultures exposed to STD medium, WT and p50KO ACM. *D-F*, Representative confocal microscopy images of MAP2⁺ cells (green) generated from ahNPC after 24h in presence of STD medium (*D*), WT ACM (*E*) and p50KO ACM (*F*). Nuclei were counterstained with Hoechst (blue). Magnification = 400X. Scale bar = 20 μ m. Data are expressed as mean values \pm S.D. of $n = 3$ experiments, run in triplicates and analyzed by one-way ANOVA followed by Tukey's *post hoc* test. *** $p < 0.001$ versus STD.

Both WT- and p50KO-derived ACM significantly promote astroglial differentiation of WT ahNPC.

When exposed to STD medium in absence of growth factors, ahNPC also spontaneously differentiate toward the astrocytic lineage and that commitment is already apparent within 24 h (Meneghini et al., 2013). Interestingly, we observed a significant increase in the percentage of GFAP⁺ cells being generated *in vitro* when WT NPC were exposed to both WT and p50KO ACM compared to STD medium condition (GFAP⁺ cells: 16.0 ± 1.71 , 36.98 ± 6.36 , 44.63 ± 6.70 % in presence of STD, WT and p50KO ACM, respectively; WT ACM vs vehicle, $p < 0.001$, p50KO ACM vs vehicle; $p < 0.001$; Figure 3 A, C-E). These data confirm that primary astrocytes produce soluble factors that promote astrogliogenesis *in vitro*, as previously demonstrated (Barkho et al., 2006). Moreover, they indicate that such soluble activities, unlike proneurogenic ones, are secreted even in absence of NF- κ B p50. Actually, in our experimental setting p50KO ACM was more effective than WT ACM in promoting astrogliogenesis in ahNPC cultures (Student's *t*-test, $p < 0.05$; Figure 3 A). A subpopulation of NG-2⁺ cells is also present both in proliferating and differentiating conditions in our cellular model. NG-2⁺ cells, also known as polydendrocytes, are commonly regarded as oligodendrocyte precursor cells (Nishiyama et al., 2009) but also, although controversial, as multipotential progenitors (Kondo and Raff, 2000). Interestingly, under differentiating conditions, the percentage of NG-2⁺ cells was not different in presence of the different media conditions (NG-2⁺ cells: 23.44 ± 11.35 , 23.50 ± 11.83 , 23.17 ± 9.87 % in STD, WT and p50KO ACM, respectively; Figure 3 B, F-H). Altogether these *in vitro* data suggest that the absence of p50 selectively disrupts the proneurogenic potential of ACM, while astrogliogenic-derived signals are intact and even enhanced. Furthermore, these results support the hypothesis that defective hippocampal neurogenesis in p50KO mice could, at least in part, be ascribed to NF- κ B p50-dependent changes in the secretome of astrocytes.

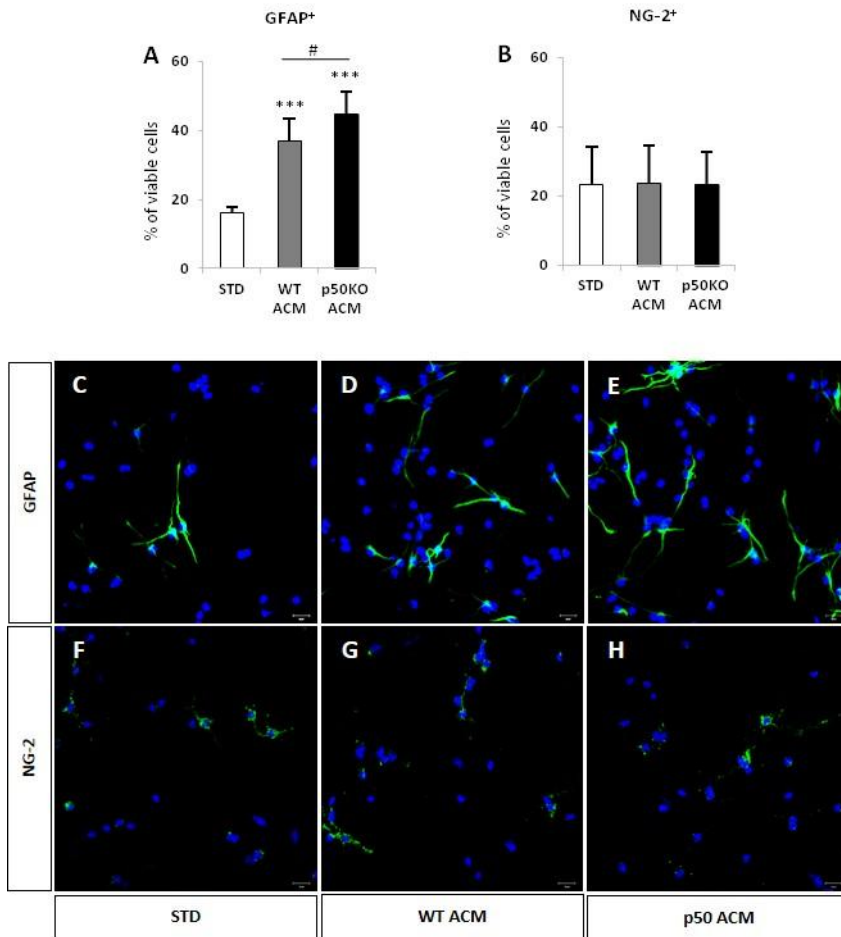


Figure 3. Both WT and p50KO ACM promote astroglial differentiation of WT ahNPC. **A**, 24h treatment of WT ahNPC with WT and p50KO ACM significantly increased the percentage of GFAP⁺ astroglial cells, compared with STD medium. p50KO ACM was more effective on astrogliogenesis than WT ACM. **B**, No significant differences was observed in the percentage of NG-2⁺ cells upon different media conditions. **C-H**, Representative confocal microscopy images of ahNPC cultures differentiated in presence of STD (**C**, **F**), WT ACM (**D**, **G**) and p50KO ACM (**E**, **H**), and immunolabelled for GFAP (**C-E**) and NG-2 (**F-H**). Nuclei were counterstained with Hoechst (blue). Magnification = 400X. Scale bar = 20 μm. Data are expressed as mean values ± S.D. of $n = 3$ experiments, run in triplicates and analyzed by one-way ANOVA followed by Tukey's *post hoc* test. *** $p < 0.001$ versus STD; # $p < 0.05$ versus WT ACM.

p50KO NPC are unresponsive to neuronal and astroglial differentiation signals from both WT- and p50KO astrocytes

We then tested the effect of WT and p50KO ACM on p50KO NPC differentiation toward neuronal and non-neuronal lineages. Surprisingly, neither WT nor p50KO ACM promoted neuronal differentiation of p50KO NPC, and their effect was indistinguishable from that of STD medium (Figure 4 A-B). Specifically, the percentages of neurons (MAP-2⁺/nestin⁻ cells) were 3.04 ± 0.32 , 3.05 ± 0.43 , 3.14 ± 0.59 % in response to STD medium, WT and p50KO ACM treatment, respectively (Figure 4 A). Similarly, the percentage of neuroblasts (MAP2⁺/nestin⁺ cells) were 32.69 ± 2.19 , 31.06 ± 3.54 , 31.81 ± 2.34 % in presence of STD medium, WT and KO ACM, respectively (Fig. 4 B). In addition, unlike WT NPC, p50KO NPC did not even respond to the astrogligenic effects of WT and p50KO ACM (GFAP⁺ cells: 14.25 ± 2.57 , 16.87 ± 3.64 , 17.08 ± 2.14 %, for STD medium, WT and p50KO ACM treatment, respectively; Figure 4 C). Similarly, to what observed in WT NPC, we could not observe any difference in the percentage of NG-2⁺ cells when p50KO NPC were exposed to different media conditions (NG-2⁺ cells: 29.49 ± 5.13 , 28.29 ± 5.48 , 27.38 ± 6.82 %, in STD, WT and p50KO ACM, respectively, Figure 4 D). Finally, also in p50KO cultures apoptotic rates were not different in presence of the different media conditions (*data not shown*). Altogether these data suggested that, in absence of NF-κB p50, cell autonomous changes profoundly affect both neuronal and astroglial differentiation of ahNPC in response to astroglia-derived molecules.

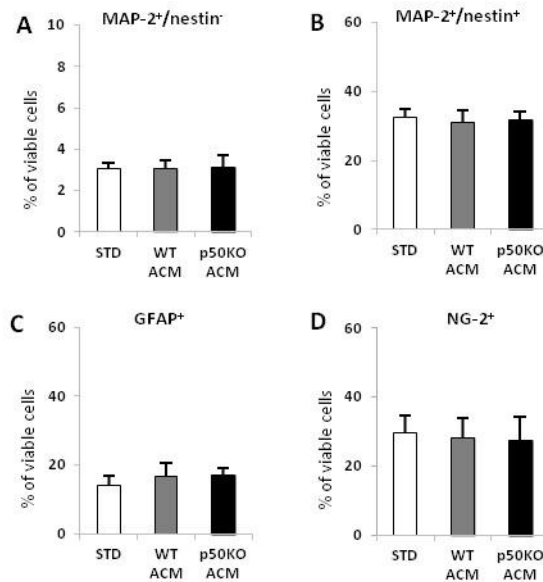


Figure 4. p50KO NPC are unresponsive to neuronal and astroglial differentiation signals from both WT- and p50KO astrocytes.

A-B, Under differentiating conditions, 24h treatment of p50KO hippocampal NPC with WT and p50KO ACM did not affect the percentage of MAP-2⁺/nestin⁻ neurons (**A**) and MAP-2⁺/nestin⁺ neuroblasts (**B**), compared to STD condition. **C-D**, No significant difference was observed in the percentage of GFAP⁺ cells (**C**), and NG-2⁺ cells (**D**) in presence of STD medium, WT and p50KO ACM. Data are expressed as mean values \pm S.D. of $n = 3$ experiments, run in triplicates. Data were analyzed by one-way ANOVA followed by Tukey's *post hoc* test.

The Receptor for Advanced Glycated End-products (RAGE) is not involved in the proneurogenic activity of ACM on WT ahNPC differentiation.

RAGE is functionally expressed by ahNPC (Meneghini et al., 2010) and its activation by astrocytes derived molecules, like HMGB-1 and S100B, results in the promotion of neuronal differentiation of ahNPC (Meneghini et al., 2013). In order to understand if the activation of RAGE could mediate the proneurogenic activity of WT ACM on WT ahNPC we pretreated cultures with a neutralizing

anti-RAGE antibody (α -RAGE Ab, 20 μ g/ml). α -RAGE Ab was not able to counteract the increase of MAP-2⁺ cells induced by WT ACM (MAP-2⁺ cells: 3.56 \pm 0.31, 6.23 \pm 0.71, 5.15 \pm 0.62 %, for STD, WT ACM and WT ACM + α -RAGE Ab respectively, $p < 0.05$; Figure 5 A). These data suggested that proneurogenic soluble factors in WT ACM do not exert their action through RAGE engagement on ahNPC.

p50KO NPC are defective in their response to thrombospondin-1 and display downregulated α 2 δ 1 expression

Thrombospondin-1 (TSP-1) is an astrocyte-derived proneurogenic factor (Lu and Kipnis, 2010). Recently our group demonstrated that α 2 δ 1, a thrombospondin-1 (TSP-1) receptor (Eroglu et al., 2009), is functionally expressed by ahNPC (Valente et al., 2012). We evaluated α 2 δ 1 expression levels in WT and p50KO ahNPC by western blot analysis. Interestingly, the α 2 δ 1 subunit appeared strongly downregulated in p50KO compared with WT NPC (-59.5%; $p < 0.05$, Figure 5 B, C). As expected, recombinant human TSP-1 (2 μ g/ml) promoted an increase in the percentage of newly formed neurons in WT NPC cultures, when compared to vehicle (MAP-2⁺/nestin⁻ cells: 5.90 \pm 1.18 and 3.45 \pm 0.64 % in TSP-1 and vehicle-treated cells; $p < 0.001$ vs vehicle; Figure 5 D). Conversely, TSP-1 treatment was ineffective in p50KO NPC (Figure 5 D). Overall these data confirmed that, in absence of p50, cell autonomous defects may reduce ahNPC responsiveness to astrocyte-derived proneurogenic signals.

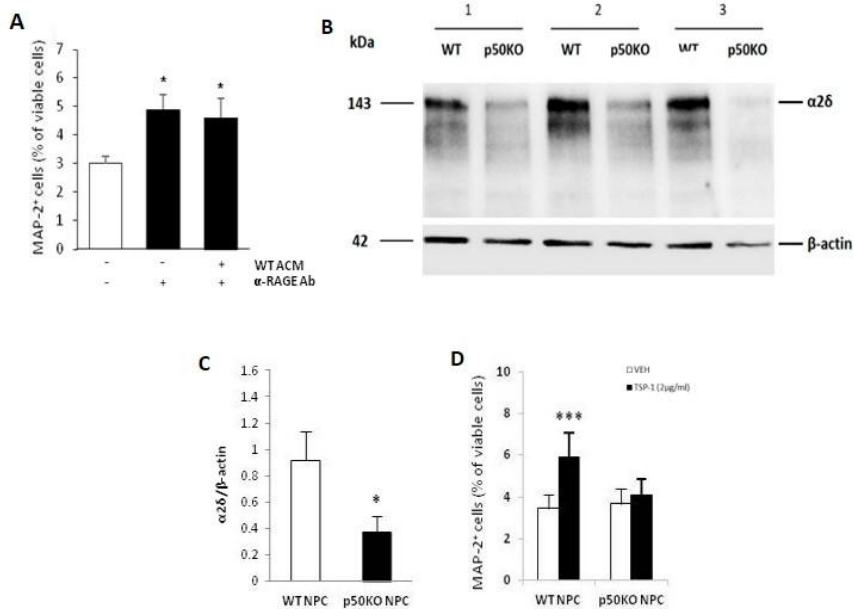


Figure 5. p50KO NPC are defective in their proneurogenic response to thrombospondin-1 and display downregulated expression of $\alpha 2\delta 1$. **A**, Pretreatment with the neutralizing anti-RAGE antibody (20 $\mu\text{g/ml}$) does not counteract the proneurogenic effects of WT ACM on WT ahNPC, as assessed by MAP2⁺ cell counting. **B**, Representative immunoblot analysis of $\alpha 2\delta 1$ subunit and β -actin expression levels in extracts of undifferentiated WT and p50KO ahNPC. Three (1-3) different WT and p50KO cell preparations were collected at different passages and analyzed. **C**, Quantification of $\alpha 2\delta 1$ expression levels in WT and p50KO NPC, normalized by β -actin, demonstrated downregulation in p50KO cultures. **D**, Under differentiating conditions, 24 h treatment of WT NPC with human recombinant TSP-1 (2 $\mu\text{g/ml}$) significantly increased the percentage of MAP-2⁺/nestin⁻ cells, compared to vehicle. No effect was observed in TSP-1-treated p50KO NPC. Data are expressed as mean values \pm S.D. of $n = 4$ experiments, run in triplicates. *** $p < 0.001$ versus vehicle (Student's t -test).

Proteomic analysis of soluble factors released by WT and p50KO ACM

In order to identify proteins which are secreted by WT and p50KO astrocytes, we analyzed ACM samples using a proteomic approach based on a hybrid system that combines the triple quadrupole technology with a time of flight mass spectrometer platform (Andrews et al. 2011). The analysis was performed on peptides produced by tryptic digestion of ACM proteins. Each sample was injected separately into the HPLC columns in a data dependent acquisition mode in order to build the spectra library of proteins identified from the runs. The protein spectra were then subjected to SWATH (Sequential Window Acquisition of Theoretical mass spectra) analysis, that allows complete evaluation of fragmented ions from detectable peptides in biological samples. Through this approach we were able to identify and quantify, as listed in Table 1, four proteins differentially expressed in WT and p50KO ACM, with p values below 0.05. Specifically, the following proteins appeared upregulated in p50KO compared to WT ACM: neutrophil gelatinase-associated lipocalin-2 (NGAL/LCN-2), C-C motif chemokine-2 (CCL2), H-2 class I Histocompatibility antigen, K-K alpha chain (HA1K) and retinoic acid receptor responder 2 (RARRES2). The complete list of proteins identified in WT and p50KO ACM is enclosed as supplementary files.

Table 1. Differentially expressed proteins in WT and p50KO ACM as identified by LC-MS/MS. Peptide number, protein coverage percentage and fold change as KO/WT ratio are provided for each identified protein.

Protein name	Accession number (UniProt Swiss-Prot)	Coverage (%)	Peptides	p -value	Fold change (KO/WT)
H-2 class I histocompatibility antigen, K-K alpha chain	sp P04223 HA1K_MOUSE	11.1%	1	0.0081	3.90
C-C motif chemokine 2 (Monocyte chemoattractant protein 1)	sp P10148 CCL2_MOUSE	22.3%	4	0.0010	3.35
Neutrophil gelatinase-associated lipocalin (NGAL) (Lipocalin-2)	sp P11672 NGAL_MOUSE	31.5%	4	0.0013	2.75
Retinoic acid receptor responder protein 2 (Chemerin)	sp O9DD06 RARR2_MOUSE	17.3%	1	0.0017	3.11

Lipocalin-2 is an astrocyte-derived proneurogenic factor whose activity is strongly reduced in p50KO NPC.

Lipocalin-2 (LCN-2) is 24 kDa iron-related protein whose modulatory role for diverse cell phenotypes in the CNS has recently attracted interest (Jha et al., 2015). Since no data are currently available on the role of LCN-2 on adult neurogenesis we focused our attention on this protein as a contributor to lack of proneurogenic effects elicited by p50KO ACM. We tested recombinant mouse LCN-2 (0.01-1 $\mu\text{g/ml}$) initially on differentiating WT ahNPC. LCN-2 promoted, in a concentration-dependent manner, neuronal differentiation of WT NPC, as assessed by counting the percentages of $\text{MAP2}^+/\text{nestin}^-$ cells in presence of the recombinant protein or vehicle (One-way ANOVA; $p < 0.001$, Fig. 6 A). Maximal proneurogenic effect was elicited in presence of 0.1 $\mu\text{g/ml}$ LCN-2 (% of increase over vehicle-treated cells: $+ 84.11 \pm 7.96$, $p < 0.001$; Figure 6 A). Under the same concentration range, LCN-2 had no effect on cell survival (Figure 6 B). When tested on p50KO NPC LCN-2 was ineffective up to 0.3 $\mu\text{g/ml}$ and only 1 $\mu\text{g/ml}$ of the protein significantly increased the percentage of newly generated neurons (% of increase over vehicle-treated cells: $+ 67.83 \pm 15.73$, $p < 0.001$; Figure 6 A). Altogether, these data suggest, for the first time, that the protein LCN-2 is an astroglial-derived signal which promotes neuronal fate specification of ahNPC and whose activity is strongly reduced in p50KO NPC.

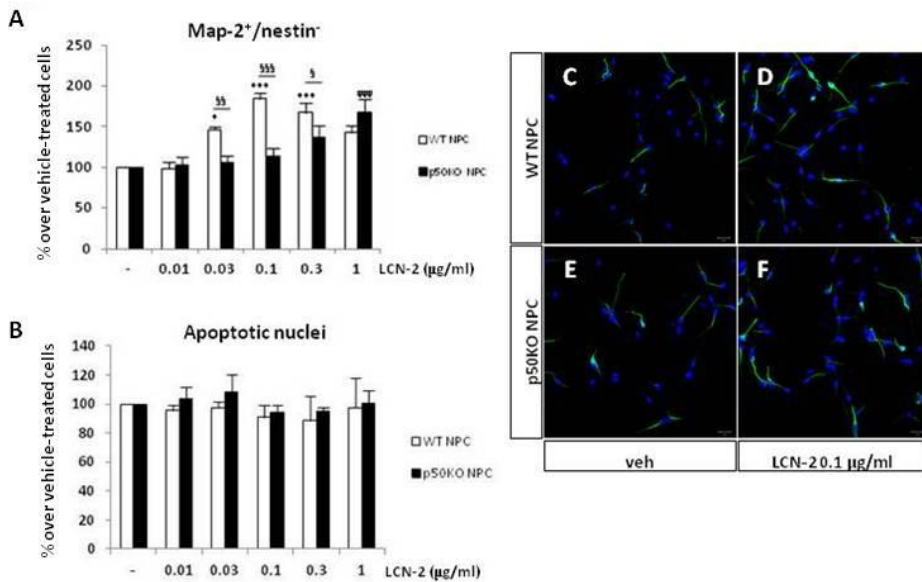


Figure 6. Lipocalin-2 is a proneurogenic signal whose activity is reduced in p50KO NPC. **A**, 24 h treatment with mouse LCN-2 (0.01-1 μg/ml) promoted neuronal differentiation of WT ahNPC in a concentration-dependent manner. Under the same experimental conditions, p50KO NPC responsiveness to LCN-2 is reduced by one order of magnitude. **B**, LCN-2 has no effect on the survival rate of WT and p50KO NPC and their progeny. **C-F**, Representative confocal microscopy images of WT (**C**, **D**) and p50KO (**E**, **F**) ahNPC differentiated in presence of vehicle (**C**, **E**) and 0.1 μg/ml of LCN-2 (**D**, **F**) and immunolabeled for MAP-2 (green). Nuclei were counterstained with Hoechst (blue). Magnification = 400X. Scale bar = 20 μm. Data are expressed % as mean values ± S.D. of n = 3 experiments, run in triplicates, and analyzed by one-way ANOVA, followed by Tukey's *post hoc* test. * $p < 0.05$, *** $p < 0.001$ versus WT vehicle; ### $p < 0.001$ versus p50KO vehicle; § $p < 0.05$, §§ $p < 0.01$, §§§ $p < 0.001$ in WT versus p50KO NPC.

Discussion

Since the initial observation of NF- κ B members being present in adult neurogenic areas, data on the involvement of these regulatory proteins in the regulation of adult neurogenesis have exponentially grown (Grilli and Meneghini, 2012; Bortolotto et al., 2014). A few years ago we proved that p50KO mice display a dramatic reduction in adult hippocampal neurogenesis and, in parallel, a selective defect in hippocampal-dependent short-term memory. A detailed phenotypic characterization of newly generated hippocampal cells in p50KO mice suggested that absence of the NF- κ B p50 subunit may trigger a rather selective defect in neuronal differentiation of ahNPC, in absence of alterations in survival and proliferation rates (Denis-Donini et al., 2008). As a follow up to that previous work, herein we show that when cultured *in vitro*, ahNPC from WT and p50KO mice are not significantly different in their neurogenic potential. Additionally, despite the well-known role of the NF- κ B signaling pathway in the regulation of cell fate, no significant difference in the basal apoptotic rate could also be observed between the two genotypes. Altogether we concluded that, unlike *in vivo* conditions, absence of NF- κ B p50 did not correlate with altered neuronal differentiation of ahNPC cultured alone *in vitro*.

Since the well documented role of NF- κ B in non-neuronal cells (Brambilla et al., 2009; Lian et al., 2015; de la Fuente et al., 2015), we decided to further dissect the role of NF- κ B p50 in the cross-talk between ahNPC and astrocytes. Exposure of WT and p50KO hippocampal NPC to conditioned media from WT and p50KO astrocytes was utilized as a strategy to study the potential influence of astroglia-generated soluble factors on adult NPC fate specification and the cell autonomous or non-cell autonomous role played by NF- κ B p50 signaling in that context. Initially we tested the effects of WT and p50KO astrocyte-conditioned media on WT ahNPC. As previously shown (Song et al., 2002), also in our hands WT ACM promoted neuronal differentiation of ahNPC. Additionally, WT primary astrocytes produced soluble factors that promoted

astrogliogenesis of ahNPC, again in line with published reports (Chang et al., 2003; Barkho et al., 2006). Interestingly, when ahNPC were exposed to p50KO ACM we could not observe proneurogenic effects. Conversely, p50KO ACM-derived astroglial signals appeared intact and even enhanced, compared to WT ACM. Of note, no differences in the survival rate of NPC and/or their progeny could be observed after exposure to WT or p50KO ACM. Altogether these data suggested that the absence of NF- κ B p50 could affect the secretory profile of astrocytes, and in particular reduce their proneurogenic potential on ahNPC. It could be hypothesized that, in absence of NF- κ B p50, primary astrocytes could either be defective in the production of proneurogenic molecule(s) or, alternatively, they could produce antineurogenic molecule(s). Furthermore our data suggested that astrocyte-derived astroglial potential was not affected by NF- κ B p50 absence. Altogether these results supported the initial hypothesis that defective hippocampal neurogenesis in p50KO mice could, at least in part, be ascribed to NF- κ B p50-dependent changes in the secretome of astrocytes. We then tested the effect of WT and p50KO ACM on p50KO NPC differentiation toward neuronal and non neuronal lineages. Surprisingly, neither WT nor p50KO ACM promoted neuronal and astroglial differentiation of p50KO NPC over basal conditions, suggesting that in absence of NF- κ B p50 cell autonomous changes may also occur and affect responsiveness of ahNPC to astrocyte-derived proneurogenic and astroglial signals.

In search for phenotypic changes in p50KO NPC which may explain their reduced responsiveness to astrocyte-proneurogenic signals we took into consideration the possibility that these cells may have lost membrane receptors for astrocyte-derived proneurogenic molecules. RAGE, the Receptor for Advanced Glycation End-products is expressed by adult NPC (Meneghini et al, 2010) and in hippocampal progenitors it mediates the proneurogenic effects of several molecules like HMGB-1 and S100B which can be secreted by astrocytes (Meneghini et al, 2013). Additionally, RAGE is the product of an NF- κ B-regulated gene (Li and Schmidt, 1997). Based on these observations we

initially evaluated the contribution of RAGE engagement on the proneurogenic effects of WT ACM in WT ahNPC, by using a previously characterized neutralizing antibody (Meneghini et al., 2013). Receptor neutralization had no effect on WT ACM proneurogenic effects, suggesting that RAGE activation was not involved in ACM-mediated neuronal differentiation of ahNPC. Thrombospondin-1 (TSP-1) is an established astrocyte-derived proneurogenic factor (Lu and Kipnis 2010). A few years ago, our group demonstrated that $\alpha 2\delta 1$ a thrombospondin-1 (TSP-1) receptor (Eroglu et al., 2009), is functionally expressed by ahNPC (Valente et al., 2012). Interestingly, $\alpha 2\delta 1$ expression levels resulted dramatically reduced in p50KO ahNPC, compared to their WT counterpart. In line with this observation, recombinant human TSP-1 promoted an increase in the percentage of newly formed neurons in WT, but not in p50KO, NPC cultures. Overall these data confirmed that, in absence of p50, cell autonomous defects may reduce ahNPC responsiveness to astrocyte-derived proneurogenic signals, and in particular to TSP-1, via downregulation of its receptor.

We also actively searched for proteins which may be differentially secreted by astrocytes in absence of p50 and whose expression could correlate with the lack of proneurogenic effects of p50KO ACM. LC-MS/MS allowed us to identify several proteins in WT and p50KO ACM. More specifically, four molecules were differentially expressed in WT and p50KO ACM, namely C-C motif chemokine-2 (CCL-2), neutrophil gelatinase-associated lipocalin-2 (NGAL/LCN-2), H-2 class I Histocompatibility antigen, K-K alpha chain (HA1K), and retinoic acid receptor responder 2 (RARRES2, also known as chemerin). Interestingly, absence of NF- κ B p50 is commonly linked to inflammation both in periphery and in brain (Bernal et al., 2014; Rolova et al., 2014) and CCL-2, NGAL/LCN-2 and RARRES2/chemerin all share an established role in inflammation. Indeed, CCL-2 and RARRES2/chemerin are proinflammatory chemokines, while LCN-2 is an acute phase-protein. Moreover, among them, CCL-2 and LCN-2 are both encoded by NF- κ B target

genes and NF- κ B signaling activation lies downstream their receptor activation (Ueda et al., 1994; Bu et al., 2006; Kohda et al., 2014).

We decided to focus our attention on NGAL/lipocalin-2 (LCN-2). It is a 24 kDa iron-related protein which belongs to a family of diverse secreted proteins (Flower, 2000) and whose modulatory role in the CNS has recently attracted interest (Jha et al., 2015), although its function is not completely understood (Ferreira et al., 2015). LCN-2 is secreted by brain astrocytes and regarded as an autocrine promoter of their classical proinflammatory activation (Bi et al., 2013; Jang et al., 2013). Since no information was available on the role of LCN-2 on adult neurogenesis, we decided to concentrate our attention on this protein, whose expression is increased in p50KO compared to WT ACM. To our surprise, LCN-2 promoted, in a concentration-dependent manner, neuronal differentiation of WT NPC, as assessed by counting the percentage of MAP2⁺/nestin⁻ and MAP2⁺/nestin⁺ cells in presence of the recombinant protein or of vehicle. In contrast with previous observations that LCN-2 can promote apoptosis of neurons and astrocytes (Lee et al., 2012; Tong et al., 2013), in our cellular model and within the tested concentration range the protein had no negative effect on the survival of ahNPC and their progeny. Interestingly, under the same experimental conditions, LCN-2 had no or little effect on neuronal differentiation of p50KO NPC, except at the highest concentration. Altogether, these data suggest, for the first time, that LCN-2 is an astroglial-derived signal which promotes neuronal fate specification of ahNPC and that p50KO NPC are less responsive to this protein. This observation points also to the possibility that, like for the TSP-1 receptor $\alpha\delta 1$, downregulation of LCN-2 receptor(s) may also occur in ahNPC and explain their reduced sensitivity (one order of magnitude) to this protein. In such respect increased expression of LCN-2 in conditioned media from p50KO astrocytes may represent a compensatory event to reduced LCN-2 receptor expression or sensitivity. Alternatively, the absence of p50 may disrupt intracellular signaling downstream several membrane receptors, including LCN-2 receptors, an aspect which deserves further investigation. LCN-2 is known to bind two cell-surface proteins. One is the

brain type organic cation transporter (BOCT or 24p3R or LCN2R) which is constitutively expressed at high levels in brain, including in the hippocampus, in neuronal cells and in the choroid plexus (Ip et al, 2011), but it also appears to be upregulated in astrocytes and microglia in EAE and peripheral nerve injury (Berard et al, 2012; Jha et al., 2013). Interestingly, NF- κ B signaling activation has been shown to occur downstream of LCN-2/24p3R (Dizin et al., 2013). The second one is megalin, also known as low density lipoprotein receptor-related protein 2 (Lrp2), a multiligand endocytic receptor which is expressed by many cell types in the CNS. Lrp2 binds about 40 different ligands ranging from lipoprotein and protease inhibitor complex to growth factors and extracellular matrix proteins (Spuch et al., 2012). Increasing evidence has implicated Lrp2, in A β clearance in Alzheimer's Disease (Alvira-Botero et al., 2010). Recently, Lrp2 has also been involved in regulating the behaviour of neural stem and progenitor cells during development and in adulthood, although not in the hippocampus (Auderset et al., 2016). A detailed evaluation of BOCT and Lrp2 expression levels in WT and p50KO ahNPC is certainly needed to complete the picture.

Of course we cannot disregard other differentially expressed proteins as contributors in the lack of response of ahNPC to astrocyte-derived signals. Contradictory data have been gathered on the contribution of CCL-2 to neurogenesis. *In vitro* studies showed that incubation of NPC with CCL-2 increase neuronal differentiation as proven by an increase in the number of Tuj1-positive cells, in absence of effects on cell proliferation and survival (Liu et al, 2007). On the other hand an involvement of CCL-2 in the reduction of neurogenesis associated with cranial irradiation has also been suggested (Lee et al., 2013). Preliminary evidence suggest that exogenously added CCL-2 is also proneurogenic on WT ahNPC cultures (*data not shown*). As far as RARRES2/chemerin, limited information is available on the physiological role of this protein and its receptor (CMKLR1) in the CNS. Targeting CMKLR1:chemerin interaction has been proposed to be beneficial in preventing or treating multiple sclerosis (Graham et al, 2014). Although there

are no data pointing to its involvement in neurogenesis, recently a novel role for chemerin and its CMKLR1 receptor in regulating adipogenesis and osteoblastogenesis of bone marrow-derived precursor cells has been proposed (Muruganandan et al, 2010). Interestingly, chemerin is also an inducer of CCL-2 (Eisinger et al., 2011).

H-2 class I Histocompatibility antigen, K-K alpha chain (HA1K) was also identified by LC-MS/MS in astrocyte-conditioned media and appeared upregulated in p50KO ACM. Of course at this stage we cannot exclude that its presence in media may represent an artefact, but since exosomes bearing MHC Class I molecules have been previously demonstrated in other cell types (Zitvogel et al., 1998) in the future we will also explore the possibility that extracellular microvesicles of astrocytic origin could contribute, with their cargo, to ACM composition.

Last but not least, in our experimental setting p50KO NPC appeared also unresponsive to astroglial astrocyte-derived signals and we could exclude astroglial effects of LCN-2 treatment on ahNPC (*data not shown*). In the future we may also attempt identifying the nature of those astrocyte secreted signals that regulate astroglialogenesis.

In summary, although we cannot underestimate the contribution of other niche cell types which were not investigated (microglia in primis), herein we propose that reduced hippocampal neurogenesis in p50KO mice *in vivo* could be ascribed to complex cell autonomous and to non-cell autonomous defects in the cross-talk between astrocytes and adult hippocampal neural progenitor cells. Moreover, we show for the first time that LCN-2 is an astroglial-derived signal which promotes neuronal fate specification of ahNPC. Finally, these data add further complexity to the growing body of data suggesting the relevance of NF- κ B signaling in the modulation of adult neurogenesis.

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Supplementary material

Table 2. List of WT ACM proteins identified by Mascot and Protein Pilot database search, filtered for protein/peptide confidence above 95%, with a minimum of two unique peptides per protein.

Cov. %	Accession	Protein names	Mass (Da)
42.2	CO3_MOUSE	Complement C3 (HSE-MSF) [Cleaved into: Complement C3 beta chain; C3-beta-c (C3bc); Complement C3 alpha chain; C3a anaphylatoxin; Acylation stimulating protein (ASP) (C3adesArg); Complement C3b alpha' chain; Complement C3c alpha' chain fragment 1; Complement C3dg fragment; Complement C3g fragment; Complement C3d fragment; Complement C3f fragment; Complement C3c alpha' chain fragment 2]	186,484
29.7	CLUS_MOUSE	Clusterin (Apolipoprotein J) (Apo-J) (Clustrin) (Sulfated glycoprotein 2) (SGP-2) [Cleaved into: Clusterin beta chain; Clusterin alpha chain]	51,656
25	NGAL_MOUSE	Neutrophil gelatinase-associated lipocalin (NGAL) (Lipocalin-2) (SV-40-induced 24P3 protein) (Siderocalin LCN2) (p25)	22,875
22.2	APOE_MOUSE	Apolipoprotein E (Apo-E)	35,867
22	CATA_MOUSE	Catalase (EC 1.11.1.6)	59,795
17.3	CH3L1_MOUSE	Chitinase-3-like protein 1 (BRP39 protein) (Breast regression protein 39) (Cartilage glycoprotein 39) (CGP-39) (GP-39)	43,893
15.4	K1C10_MOUSE	Keratin, type I cytoskeletal 10 (56 kDa cyokeratin) (Cytokeratin-10) (CK-10) (Keratin, type I cytoskeletal 59 kDa) (Keratin-10) (K10)	57,770
15.1	FIGL1_MOUSE	Fidgetin-like protein 1 (EC 3.6.4.-)	74,850
12.4	AL1A1_MOUSE	Retinal dehydrogenase 1 (RALDH 1) (RalDH1) (EC 1.2.1.36) (ALDH-E1) (ALHDII) (Aldehyde dehydrogenase family 1 member A1) (Aldehyde dehydrogenase, cytosolic)	54,468
11.4	CYTC_MOUSE	Cystatin-C (Cystatin-3)	15,531
10.8	CCL2_MOUSE	C-C motif chemokine 2 (Monocyte chemoattractant protein 1) (Monocyte chemotactic protein 1) (MCP-1) (Platelet-derived growth factor-inducible protein JE) (Small-inducible cytokine A2)	16,326
10.4	NT5D3_MOUSE	5'-nucleotidase domain-containing protein 3 (EC 3.1.3.-) (GRP94-neighboring nucleotidase)	63,170
8.7	CERU_MOUSE	Ceruloplasmin (EC 1.16.3.1) (Ferroxidase)	121,151
8.2	CXL10_MOUSE	C-X-C motif chemokine 10 (10 kDa interferon gamma-induced protein) (Gamma-IP10) (IP-10) (C7) (Interferon-gamma induced protein CRG-2) (Small-inducible cytokine B10)	10,789
8.1	GRAG_MOUSE	Granzyme G (EC 3.4.21.-) (CTL serine protease 1) (MCSP-1)	27,381
7.3	PAFA_MOUSE	Platelet-activating factor acetylhydrolase (PAF acetylhydrolase) (EC 3.1.1.47) (1-alkyl-2-acetyl-glycerophosphocholine esterase) (2-acetyl-1-alkyl-glycerophosphocholine esterase) (LDL-associated phospholipase A2) (LDL-PLA (2)) (PAF 2-acylhydrolase)	49,258
6.7	K1C17_MOUSE	Keratin, type I cytoskeletal 17 (Cytokeratin-17) (CK-17) (Keratin-17) (K17)	48,162
6.6	SKIT1_MOUSE	Selection and upkeep of intraepithelial T-cells protein 1 (Skint-1) (Immunoglobulin-like and transmembrane domain-containing protein expressed in skin and thymus protein 1)	41,792
6.5	MCF2L_MOUSE	Guanine nucleotide exchange factor DBS (DBL's big sister) (MCF2-transforming sequence-like protein)	129,113
6.3	NAA50_MOUSE	N-alpha-acetyltransferase 50 (EC 2.3.1.-) (N-acetyltransferase NAT13) (NatE catalytic subunit)	19,414
6.2	RUND1_MOUSE	RUN domain-containing protein 1	67,865
6.2	TRRAP_MOUSE	Transformation/transcription domain-associated protein (Tra1 homolog)	291,557
6.1	CBPE_MOUSE	Carboxypeptidase E (CPE) (EC 3.4.17.10) (Carboxypeptidase H) (CPH) (Enkephalin convertase) (Prohormone-processing carboxypeptidase)	53,256

5.8	LIFR_MOUSE	Leukemia inhibitory factor receptor (LIF receptor) (LIF-R) (D-factor/LIF receptor) (CD antigen CD118)	122,574
5.5	SCEL_MOUSE	Sciellin	72,972
5.3	ARAP3_MOUSE	Arf-GAP with Rho-GAP domain, ANK repeat and PH domain-containing protein 3 (Centaurin-delta-3) (Cnt-d3) (Dual specificity Rho- and Arf-GTPase-activating protein 1)	169,741
5.2	LG3BP_MOUSE	Galectin-3-binding protein (Cyp-C-associated protein) (CyCAP) (Lectin galactoside-binding soluble 3-binding protein) (Protein MAMA)	64,491
5.2	ESYT3_MOUSE	Extended synaptotagmin-3 (E-Syt3)	100,068
5	BRE1A_MOUSE	E3 ubiquitin-protein ligase BRE1A (BRE1-A) (EC 6.3.2.-) (RING finger protein 20)	113,520
4.6	SPRC_MOUSE	SPARC (Basement-membrane protein 40) (BM-40) (Osteonectin) (ON) (Secreted protein acidic and rich in cysteine)	34,450
4.6	K1C42_MOUSE	Keratin, type I cytoskeletal 42 (Cytokeratin-42) (CK-42) (Keratin-17n) (Keratin-42) (K42) (Type I keratin Ka22)	50,133
4.6	TDR12_MOUSE	Putative ATP-dependent RNA helicase TDRD12 (EC 3.6.4.13) (ES cell-associated transcript 8 protein) (Tudor domain-containing protein 12)	137,627
4.5	K22E_MOUSE	Keratin, type II cytoskeletal 2 epidermal (Cytokeratin-2e) (CK-2e) (Epithelial keratin-2e) (Keratin-2 epidermis) (Keratin-2e) (K2e) (Type-II keratin Kb2)	70,923
3.9	NAA20_MOUSE	N-alpha-acetyltransferase 20 (EC 2.3.1.88) (Methionine N-acetyltransferase) (N-acetyltransferase 5) (N-terminal acetyltransferase B complex catalytic subunit NAA20) (N-terminal acetyltransferase B complex catalytic subunit NAT5) (NatB complex subunit NAT5) (NatB catalytic subunit)	20,368
3.8	K2C5_MOUSE	Keratin, type II cytoskeletal 5 (Cytokeratin-5) (CK-5) (Keratin-5) (K5) (Type-II keratin Kb5)	61,767
3.4	WDR64_MOUSE	WD repeat-containing protein 64	123,893
3.3	NCAN_MOUSE	Neurocan core protein (Chondroitin sulfate proteoglycan 3)	137,200
3.2	RAD54_HUMAN	DNA repair and recombination protein RAD54-like (EC 3.6.4.-) (RAD54 homolog) (hHR54) (hRAD54)	84,352
3.1	LBP_MOUSE	Lipopolysaccharide-binding protein (LBP)	53,055
3	BAI2_MOUSE	Adhesion G protein-coupled receptor B2 (Brain-specific angiogenesis inhibitor 2)	169,863
2.9	FNIP2_MOUSE	Folliculin-interacting protein 2 (FNIP1-like protein) (O6-methylguanine-induced apoptosis 1 protein)	122,520
2.8	CWC25_MOUSE	Pre-mRNA-splicing factor CWC25 homolog (Coiled-coil domain-containing protein 49) (Spliceosome-associated protein homolog CWC25)	48,811
2.7	FREM1_MOUSE	FRAS1-related extracellular matrix protein 1 (Protein QBRICK)	244,544
2.3	SYIM_MOUSE	Isoleucine--tRNA ligase, mitochondrial (EC 6.1.1.5) (Isoleucyl-tRNA synthetase) (IleRS)	112,804
2.3	ABHD8_MOUSE	Protein ABHD8 (EC 3.-.-) (Alpha/beta hydrolase domain-containing protein 8) (Abhydrolase domain-containing protein 8)	48,229
2.3	K2C79_MOUSE	Keratin, type II cytoskeletal 79 (Cytokeratin-79) (CK-79) (Keratin-79) (K79) (Type-II keratin Kb38)	57,552
2.3	CATZ_MOUSE	Cathepsin Z (EC 3.4.18.1)	33,996
2	FRAS1_MOUSE	Extracellular matrix protein FRAS1	442,369
1.9	ACTBL_MOUSE	Beta-actin-like protein 2 (Kappa-actin)	42,004
1.9	PNCB_MOUSE	Nicotinate phosphoribosyltransferase (NAPRTase) (EC 6.3.4.21) (Nicotinate phosphoribosyltransferase domain-containing protein 1)	58,265
1.7	ZBT49_MOUSE	Zinc finger and BTB domain-containing protein 49 (Zinc finger protein 509)	83,091
1.6	PPARD_MOUSE	Peroxisome proliferator-activated receptor delta (PPAR-delta) (Nuclear hormone receptor 1) (NUC1) (Nuclear receptor subfamily 1 group C member 2) (Peroxisome proliferator-activated receptor beta) (PPAR-beta)	49,715
1.5	5HT2B_MOUSE	5-hydroxytryptamine receptor 2B (5-HT-2B) (5-HT2B) (5-HT-2F) (NP75 protein) (Serotonin receptor 2B)	53,597
1.5	TMPSD_MOUSE	Transmembrane protease serine 13 (EC 3.4.21.-) (Membrane-type mosaic serine protease) (Mosaic serine protease)	59,806
1.5	COBA2_MOUSE	Collagen alpha-2(XI) chain	171,535
1.2	NSD1_MOUSE	Histone-lysine N-methyltransferase, H3 lysine-36 and H4 lysine-20 specific (EC 2.1.1.43) (H3-K36-HMTase) (H4-K20-HMTase) (Nuclear receptor-binding SET domain-containing protein 1) (NR-binding SET domain-containing protein)	284,084

1.2	MKKS_MOUSE	McKusick-Kaufman/Bardet-Biedl syndromes putative chaperonin (Protein Bbs6 homolog)	61,924
1.1	FAT2_MOUSE	Protocadherin Fat 2 (FAT tumor suppressor homolog 2)	480,109
0.9	SSH3_MOUSE	Protein phosphatase Slingshot homolog 3 (EC 3.1.3.16) (EC 3.1.3.48) (SSH-like protein 3) (SSH-3L) (mSSH-3L)	72,227
0.8	TLR4_MOUSE	Toll-like receptor 4 (CD antigen CD284)	95,519
0.7	ATP9B_MOUSE	Probable phospholipid-transporting ATPase IIB (EC 3.6.3.1) (ATPase class II type 9B)	129,017
0.6	RC3H2_MOUSE	Roquin-2 (EC 6.3.2.-) (Membrane-associated nucleic acid-binding protein) (RING finger and CCCH-type zinc finger domain-containing protein 2)	131,295

Table 3. List of p50KO ACM proteins identified by Mascot and Protein Pilot database search, filtered for protein/peptide confidence above 95%, with a minimum of two unique peptides per protein.

Cov (%)	Accession	Protein names	Mass (Da)
48.2	APOE_MOUSE	Apolipoprotein E (Apo-E)	35,867
42.6	SAA3_MOUSE	Serum amyloid A-3 protein	13,774
36.4	CO3_MOUSE	Complement C3 (HSE-MSF) [Cleaved into: Complement C3 beta chain; C3-beta-c (C3bc); Complement C3 alpha chain; C3a anaphylatoxin; Acylation stimulating protein (ASP) (C3adesArg); Complement C3b alpha' chain; Complement C3c alpha' chain fragment 1; Complement C3dg fragment; Complement C3g fragment; Complement C3d fragment; Complement C3f fragment; Complement C3c alpha' chain fragment 2]	186,484
31.5	NGAL_MOUSE	Neutrophil gelatinase-associated lipocalin (NGAL) (Lipocalin-2) (SV-40-induced 24P3 protein) (Siderocalin LCN2) (p25)	22,875
28.1	CLUS_MOUSE	Clusterin (Apolipoprotein J) (Apo-J) (Clustrin) (Sulfated glycoprotein 2) (SGP-2) [Cleaved into: Clusterin beta chain; Clusterin alpha chain]	51,656
25.7	CYTC_MOUSE	Cystatin-C (Cystatin-3)	15,531
23.9	SPA3N_MOUSE	Serine protease inhibitor A3N (Serpins A3N)	46,718
22.3	CCL2_MOUSE	C-C motif chemokine 2 (Monocyte chemoattractant protein 1) (Monocyte chemoattractant protein 1) (MCP-1) (Platelet-derived growth factor-inducible protein JE) (Small-inducible cytokine A2)	16,326
19.7	CBPE_MOUSE	Carboxypeptidase E (CPE) (EC 3.4.17.10) (Carboxypeptidase H) (CPH) (Enkephalin convertase) (Prohormone-processing carboxypeptidase)	53,256
18.1	CERU_MOUSE	Ceruloplasmin (EC 1.16.3.1) (Ferroxidase)	121,151
17.3	RARR2_MOUSE	Retinoic acid receptor responder protein 2 (Chemerin)	18,350
15.9	SCG3_MOUSE	Secretogranin-3 (Secretogranin III) (SgIII)	53,326
15	CFAB_MOUSE	Complement factor B (EC 3.4.21.47) (C3/C5 convertase) [Cleaved into: Complement factor B Ba fragment; Complement factor B Bb fragment]	85,004
12.4	CCL7_MOUSE	C-C motif chemokine 7 (Intercrine/chemokine MARC) (Monocyte chemoattractant protein 3) (Monocyte chemoattractant protein 3) (MCP-3) (Protein FIC) (Small-inducible cytokine A7)	10,999
11.7	HAIK_MOUSE	H-2 class I histocompatibility antigen, K-K alpha chain (H-2K(K))	41,646

9.3	CATD_MOUSE	Cathepsin D (EC 3.4.23.5)	44,954
9.3	KIF5C_MOUSE	Kinesin heavy chain isoform 5C (Kinesin heavy chain neuron-specific 2)	109,275
8.8	TIMP1_MOUSE	Metalloproteinase inhibitor 1 (Collagenase inhibitor 16C8 fibroblast) (Erythroid-potentiating activity) (EPA) (TPA-S1) (TPA-induced protein) (Tissue inhibitor of metalloproteinases 1) (TIMP-1)	22,628
8.2	CXL10_MOUSE	C-X-C motif chemokine 10 (10 kDa interferon gamma-induced protein) (Gamma-IP10) (IP-10) (C7) (Interferon-gamma induced protein CRG-2) (Small-inducible cytokine B10)	10,789
7.9	NAA20_MOUSE	N-alpha-acetyltransferase 20 (EC 2.3.1.88) (Methionine N-acetyltransferase) (N-acetyltransferase 5) (N-terminal acetyltransferase B complex catalytic subunit NAA20) (N-terminal acetyltransferase B complex catalytic subunit NAT5) (NatB complex subunit NAT5) (NatB catalytic subunit)	20,368
7.7	K1C10_MOUSE	Keratin, type I cytoskeletal 10 (56 kDa cytokeratin) (Cytokeratin-10) (CK-10) (Keratin, type I cytoskeletal 59 kDa) (Keratin-10) (K10)	57,770
7.7	CATB_MOUSE	Cathepsin B (EC 3.4.22.1) (Cathepsin B1) [Cleaved into: Cathepsin B light chain; Cathepsin B heavy chain]	37,280
7.6	CAND1_MOUSE	Cullin-associated NEDD8-dissociated protein 1 (Cullin-associated and neddylation-dissociated protein 1) (p120 CAND1)	136,332
7.1	K1C42_MOUSE	Keratin, type I cytoskeletal 42 (Cytokeratin-42) (CK-42) (Keratin-17n) (Keratin-42) (K42) (Type I keratin Ka22)	50,133
6.9	FSTL1_MOUSE	Follistatin-related protein 1 (Follistatin-like protein 1) (TGF-beta-inducible protein TSC-36)	34,554
6.8	PAFA_MOUSE	Platelet-activating factor acetylhydrolase (PAF acetylhydrolase) (EC 3.1.1.47) (1-alkyl-2-acetylgllycerophosphocholine esterase) (2-acetyl-1-alkylglycerophosphocholine esterase) (LDL-associated phospholipase A2) (LDL-PLA(2)) (PAF 2-acylhydrolase)	49,258
6.7	LBP_MOUSE	Lipopolysaccharide-binding protein (LBP)	53,055
6.3	SPRC_MOUSE	SPARC (Basement-membrane protein 40) (BM-40) (Osteonectin) (ON) (Secreted protein acidic and rich in cysteine)	34,450
5.9	CAB39_MOUSE	Calcium-binding protein 39 (MO25alpha) (Protein Mo25)	39,843
5.9	DDX27_MOUSE	Probable ATP-dependent RNA helicase DDX27 (EC 3.6.4.13) (DEAD box protein 27)	85,939
5.6	CTRO_MOUSE	Citron Rho-interacting kinase (CRIK) (EC 2.7.11.1) (Rho-interacting, serine/threonine-protein kinase 21)	235,389
5.5	CATA_MOUSE	Catalase (EC 1.11.1.6)	59,795
5.5	CH3L1_MOUSE	Chitinase-3-like protein 1 (BRP39 protein) (Breast regression protein 39) (Cartilage glycoprotein 39) (CGP-39) (GP-39)	43,893
5.4	MYH7_MOUSE	Myosin-7 (Myosin heavy chain 7) (Myosin heavy chain slow isoform) (MyHC-slow) (Myosin heavy chain, cardiac muscle beta isoform) (MyHC-beta)	222,879
5.1	PLTP_MOUSE	Phospholipid transfer protein (Lipid transfer protein II)	54,453
4.4	KRT35_MOUSE	Keratin, type I cuticular Ha5 (Hair keratin, type I Ha5) (Keratin-35) (K35)	50,530
4.2	ABCA1_MOUSE	ATP-binding cassette sub-family A member 1 (ATP-binding cassette transporter 1) (ABC-1) (ATP-binding cassette 1)	253,912
4.1	K2C79_MOUSE	Keratin, type II cytoskeletal 79 (Cytokeratin-79) (CK-79) (Keratin-79) (K79) (Type-II keratin Kb38)	57,552
4	ACTBL_MOUSE	Beta-actin-like protein 2 (Kappa-actin)	42,004
3.6	NCAN_MOUSE	Neurocan core protein (Chondroitin sulfate proteoglycan 3)	137,200
3.5	FAM3C_MOUSE	Protein FAM3C (Interleukin-like EMT inducer)	24,753

3.3	K22E_MOUSE	Keratin, type II cytoskeletal 2 epidermal (Cytokeratin-2e) (CK-2e) (Epithelial keratin-2e) (Keratin-2 epidermis) (Keratin-2e) (K2e) (Type-II keratin Kb2)	70,923
3.2	PPARD_MOUSE	Peroxisome proliferator-activated receptor delta (PPAR-delta) (Nuclear hormone receptor 1) (NUC1) (Nuclear receptor subfamily 1 group C member 2) (Peroxisome proliferator-activated receptor beta) (PPAR-beta)	49,715
3.1	TMCC1_MOUSE	Transmembrane and coiled-coil domains protein 1	71,642
2.8	CWC25_MOUSE	Pre-mRNA-splicing factor CWC25 homolog (Coiled-coil domain-containing protein 49) (Spliceosome-associated protein homolog CWC25)	48,811
2.7	MOT10_MOUSE	Monocarboxylate transporter 10 (MCT 10) (Aromatic amino acid transporter 1) (Solute carrier family 16 member 10) (T-type amino acid transporter 1)	55,320
2.3	RC3H2_MOUSE	Roquin-2 (EC 6.3.2.-) (Membrane-associated nucleic acid-binding protein) (RING finger and CCCH-type zinc finger domain-containing protein 2)	131,295
2.3	ABHD8_MOUSE	Protein ABHD8 (EC 3.-.-) (Alpha/beta hydrolase domain-containing protein 8) (Abhydrolase domain-containing protein 8)	48,229
2	K1C15_MOUSE	Keratin, type I cytoskeletal 15 (Cytokeratin-15) (CK-15) (Keratin-15) (K15)	49,138
1.9	PNCB_MOUSE	Nicotinate phosphoribosyltransferase (NAPRTase) (EC 6.3.4.21) (Nicotinate phosphoribosyltransferase domain-containing protein 1)	58,265
1.8	LIFR_MOUSE	Leukemia inhibitory factor receptor (LIF receptor) (LIF-R) (D-factor/LIF receptor) (CD antigen CD118)	122,574
1.7	DDX46_MOUSE	Probable ATP-dependent RNA helicase DDX46 (EC 3.6.4.13) (DEAD box protein 46)	117,448
1.5	TMPSD_MOUSE	Transmembrane protease serine 13 (EC 3.4.21.-) (Membrane-type mosaic serine protease) (Mosaic serine protease)	59,806
1.4	CO4A2_MOUSE	Collagen alpha-2(IV) chain [Cleaved into: Canstatin]	167,325
1.2	LG3BP_MOUSE	Galectin-3-binding protein (Cyp-C-associated protein) (CyCAP) (Lectin galactoside-binding soluble 3-binding protein) (Protein MAMA)	64,491
1	AKAP4_MOUSE	A-kinase anchor protein 4 (AKAP-4) (A-kinase anchor protein 82 kDa) (AKAP 82) (mAKAP82) (FSC1) (Major sperm fibrous sheath protein) (Protein kinase A-anchoring protein 4) (PRKA4) (p82)	93,796
0.5	BAI2_MOUSE	Adhesion G protein-coupled receptor B2 (Brain-specific angiogenesis inhibitor 2)	169,863

Table 4. Raw data file of peptides used in SWATH-MS analysis for the identification and quantitation of the 4 differentially expressed proteins listed in Table 1.

Protein	Peptide	Replicate	Precursor Result	Total Area MS1	Total Area Fragment
sp P04223 HA1K_MOUSE	GNEQIFR	SWATH_WTrep3	2471	#N/D	2471
sp P04223 HA1K_MOUSE	GNEQIFR	SWATH_WTrep2	2702	#N/D	2702
sp P04223 HA1K_MOUSE	GNEQIFR	SWATH_WTrep1	2340	#N/D	2340
sp P04223 HA1K_MOUSE	GNEQIFR	SWATH_KOrep_4	10868	#N/D	10868
sp P04223 HA1K_MOUSE	GNEQIFR	SWATH_KOrep_2	10213	#N/D	10213
sp P04223 HA1K_MOUSE	GNEQIFR	SWATH_KOrep_1	11466	#N/D	11466
sp P04223 HA1K_MOUSE	GNEQIFR	SWATH_KOrep_3	9993	#N/D	9993
sp P10148 CCL2_MOUSE	EAVVFTK	SWATH_WTrep3	153687	128204	25483
sp P10148 CCL2_MOUSE	EAVVFTK	SWATH_WTrep2	164498	140179	24319
sp P10148 CCL2_MOUSE	EAVVFTK	SWATH_WTrep1	146597	122183	24414
sp P10148 CCL2_MOUSE	EAVVFTK	SWATH_KOrep_4	661337	554219	107119
sp P10148 CCL2_MOUSE	EAVVFTK	SWATH_KOrep_2	662528	562099	100429
sp P10148 CCL2_MOUSE	EAVVFTK	SWATH_KOrep_1	605459	508588	96871
sp P10148 CCL2_MOUSE	EAVVFTK	SWATH_KOrep_3	583708	492348	91360
sp P10148 CCL2_MOUSE	KEWVQTYIK	SWATH_WTrep3	32001	31099	902
sp P10148 CCL2_MOUSE	KEWVQTYIK	SWATH_WTrep2	20523	19090	1433
sp P10148 CCL2_MOUSE	KEWVQTYIK	SWATH_WTrep1	38266	36820	1446
sp P10148 CCL2_MOUSE	KEWVQTYIK	SWATH_KOrep_4	71301	68320	2981
sp P10148 CCL2_MOUSE	KEWVQTYIK	SWATH_KOrep_2	85960	83370	2591
sp P10148 CCL2_MOUSE	KEWVQTYIK	SWATH_KOrep_1	63109	59543	3566
sp P10148 CCL2_MOUSE	KEWVQTYIK	SWATH_KOrep_3	94789	91994	2794
sp P10148 CCL2_MOUSE	EWVQTYIK	SWATH_WTrep3	155424	128374	27050
sp P10148 CCL2_MOUSE	EWVQTYIK	SWATH_WTrep2	160812	128365	32447
sp P10148 CCL2_MOUSE	EWVQTYIK	SWATH_WTrep1	178413	146207	32206
sp P10148 CCL2_MOUSE	EWVQTYIK	SWATH_KOrep_4	329334	282540	46794
sp P10148 CCL2_MOUSE	EWVQTYIK	SWATH_KOrep_2	329952	279782	50170
sp P10148 CCL2_MOUSE	EWVQTYIK	SWATH_KOrep_1	329134	281025	48109

sp P10148 CCL2_MOUSE	EWVQTYIK	SWATH_KOrep_3	332781	289702	43079
sp P10148 CCL2_MOUSE	SSAPLNVK	SWATH_WTrep3	153153	144243	8909
sp P10148 CCL2_MOUSE	SSAPLNVK	SWATH_WTrep2	160286	150394	9892
sp P10148 CCL2_MOUSE	SSAPLNVK	SWATH_WTrep1	191953	182631	9322
sp P10148 CCL2_MOUSE	SSAPLNVK	SWATH_KOrep_4	516756	476254	40502
sp P10148 CCL2_MOUSE	SSAPLNVK	SWATH_KOrep_2	487764	448050	39714
sp P10148 CCL2_MOUSE	SSAPLNVK	SWATH_KOrep_1	499476	461775	37701
sp P10148 CCL2_MOUSE	SSAPLNVK	SWATH_KOrep_3	459034	422266	36768
sp P11672 NGAL_MOUSE	WYVVGLAGNAVQK	SWATH_WTrep3	135741	120467	15273
sp P11672 NGAL_MOUSE	WYVVGLAGNAVQK	SWATH_WTrep2	199234	183843	15391
sp P11672 NGAL_MOUSE	WYVVGLAGNAVQK	SWATH_WTrep1	171925	155501	16423
sp P11672 NGAL_MOUSE	WYVVGLAGNAVQK	SWATH_KOrep_4	398726	350009	48717
sp P11672 NGAL_MOUSE	WYVVGLAGNAVQK	SWATH_KOrep_2	439424	392870	46555
sp P11672 NGAL_MOUSE	WYVVGLAGNAVQK	SWATH_KOrep_1	421031	371489	49542
sp P11672 NGAL_MOUSE	WYVVGLAGNAVQK	SWATH_KOrep_3	435008	391324	43684
sp P11672 NGAL_MOUSE	TFVPSSR	SWATH_WTrep3	160713	124169	36544
sp P11672 NGAL_MOUSE	TFVPSSR	SWATH_WTrep2	150379	117681	32698
sp P11672 NGAL_MOUSE	TFVPSSR	SWATH_WTrep1	164657	128344	36312
sp P11672 NGAL_MOUSE	TFVPSSR	SWATH_KOrep_4	414944	320077	94868
sp P11672 NGAL_MOUSE	TFVPSSR	SWATH_KOrep_2	423173	326578	96596
sp P11672 NGAL_MOUSE	TFVPSSR	SWATH_KOrep_1	414689	319453	95236
sp P11672 NGAL_MOUSE	TFVPSSR	SWATH_KOrep_3	462786	369185	93601
sp P11672 NGAL_MOUSE	YPQVQSYNVQVATT DYNQFAMVFFR	SWATH_WTrep3	88085	84257	3828
sp P11672 NGAL_MOUSE	YPQVQSYNVQVATT DYNQFAMVFFR	SWATH_WTrep2	86929	84550	2379
sp P11672 NGAL_MOUSE	YPQVQSYNVQVATT DYNQFAMVFFR	SWATH_WTrep1	72685	69959	2727
sp P11672 NGAL_MOUSE	YPQVQSYNVQVATT DYNQFAMVFFR	SWATH_KOrep_4	257914	248498	9416
sp P11672 NGAL_MOUSE	YPQVQSYNVQVATT DYNQFAMVFFR	SWATH_KOrep_2	288924	277626	11298
sp P11672 NGAL_MOUSE	YPQVQSYNVQVATT DYNQFAMVFFR	SWATH_KOrep_1	284240	271784	12456
sp P11672 NGAL_MOUSE	YPQVQSYNVQVATT DYNQFAMVFFR	SWATH_KOrep_3	223409	213629	9780
sp P11672 NGAL_MOUSE	ELSPCLK	SWATH_WTrep3	146315	128375	17940
sp P11672 NGAL_MOUSE	ELSPCLK	SWATH_WTrep2	181324	154967	26357
sp P11672 NGAL_MOUSE	ELSPCLK	SWATH_WTrep1	123902	109046	14856
sp P11672 NGAL_MOUSE	ELSPCLK	SWATH_KOrep_4	399498	352069	47429

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sp P11672 NGAL_MOUSE	ELSPK	SWATH_KOrep_2	399837	352828	47008
sp P11672 NGAL_MOUSE	ELSPK	SWATH_KOrep_1	380892	328979	51913
sp P11672 NGAL_MOUSE	ELSPK	SWATH_KOrep_3	356387	316640	39746
sp Q9DD06 RARR_2_MOUSE	HPPVQLAFQEIGVD R	SWATH_WTrep3	1874	#N/D	1874
sp Q9DD06 RARR_2_MOUSE	HPPVQLAFQEIGVD R	SWATH_WTrep2	1877	#N/D	1877
sp Q9DD06 RARR_2_MOUSE	HPPVQLAFQEIGVD R	SWATH_WTrep1	1707	#N/D	1707
sp Q9DD06 RARR_2_MOUSE	HPPVQLAFQEIGVD R	SWATH_KOrep_4	5360	#N/D	5360
sp Q9DD06 RARR_2_MOUSE	HPPVQLAFQEIGVD R	SWATH_KOrep_2	6580	#N/D	6580
sp Q9DD06 RARR_2_MOUSE	HPPVQLAFQEIGVD R	SWATH_KOrep_1	5245	#N/D	5245
sp Q9DD06 RARR_2_MOUSE	HPPVQLAFQEIGVD R	SWATH_KOrep_3	5373	#N/D	5373

Chapter 4.

Modulatory events within the adult hippocampal neurogenic niche: novel insights into region-specific astrocyte-derived signaling cues and the role played by NF- κ B p50.

[UNPUBLISHED RESULTS]

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Introduction

Active adult neurogenesis is well documented to occur in the adult mammalian brain in two discrete areas, the subventricular zone (SVZ) of the lateral ventricles, and the subgranular zone (SGZ) of the dentate gyrus (Ming and Song, 2005; Bond et al., 2015; Nicola et al., 2015). Growing evidence suggest the presence of another neurogenic area in the hypothalamus, along the walls of the third ventricle (Kokoeva et al., 2005). The neurogenic areas called niches are composed of different cell types and non-cellular components, such as soluble cues (i.e. growth factors, cytokines, hormones) originated both locally and distantly. In the neurogenic niches neural stem/progenitor cells (NSPC) are housed and supported by the different cellular and molecular components, that balance their proliferation and differentiation rates (Lin and Iacovitti, 2015). Astrocytes represent one major contributor of the hippocampal neurogenic niche. Hippocampal-derived astrocytes were demonstrated to promote neurogenesis both via directly interacting with adult hippocampal NSPC and through secreted factors (Song et al., 2002). On the other hand, astrocytes isolated from non-neurogenic areas, such as those isolated from the adult spinal cord were proven ineffective in promoting neurogenesis, while were effective in promoting gliogenesis (Barkho et al., 2006). Overall these data suggested that the astrocytes in adult neurogenic regions contribute to generate an instructive environment for neuronal differentiation of adult NSPC.

Our group reported that the NF- κ B family of transcription factors are expressed at considerable levels in areas of active neurogenesis in the postnatal and adult mouse brain (Denis-Donini et al., 2005). Previous publications from our lab also reported that mice lacking NF- κ B p50 subunit display a dramatic reduction in adult hippocampal neurogenesis associated with short-term memory defects (Denis-Donini et al., 2008). Interestingly, NPC derived from WT and p50KO hippocampi do not display any differences in neuronal differentiation and apoptotic rates *in vitro*. These data suggested that impaired hippocampal neurogenesis observed *in vivo* could also be due to inadequate signaling

between niche cells. In previous work in our laboratory (Cvijetic et al., under submission) we investigated NPC-astroglia communication and, in particular, the effect of soluble factors released by astrocytes derived from mixed culture of cortex/hippocampi on cell fate specification of WT and p50KO adult hippocampal NPC (ahNPC). We demonstrated that WT ACM promote both neuronal and astroglial differentiation of WT NPC, whereas p50KO ACM can only promote astroglial differentiation. Additionally, when p50KO NPC were exposed to WT and p50KO ACM, they showed unresponsiveness to both treatment conditions.

As a follow up of these experiments, in the first part of this study, we addressed the importance of region-specificity in the effect of ACM on ahNPC. More specifically, we compared the proneurogenic and proglial properties of hippocampal and hypothalamic ACM deriving from WT and p50KO astrocytes on adult hippocampal WT NPC fate specification.

A second line of investigation was aimed at assessing the differentiation of WT hypothalamic NPC in response to hippocampal ACM deriving from WT and p50KO astrocytes cultures.

Materials and Methods

Adult mouse hippocampal (Hp) and hypothalamic (Hyt) neural progenitor cell (NPC) cultures. Hippocampi and hypothalami from three adult (3-4 month-old) C57BL/6 male mice were dissected and cell suspension was prepared as previously described (Valente et al, 2012). Cells were maintained as floating neurospheres through subsequent passages (P1-P30).

NPC differentiation experiments. Differentiation experiments were carried out as previously described (Valente et al., 2012). Briefly, neurospheres (passages 5-30) were dissociated into single NPC and plated onto laminin-coated Lab-Tek 8-well permanox chamber slides (Nunc) at a density of 43,750 cells/cm² in NPC differentiation medium (NDM, Neurobasal-A medium supplemented with B27, 2 mM L-glutamine and 100 U/100 µg/ml Penicillin/Streptomycin). NPC were treated in presence of astrocyte conditioned media (ACM) derived from WT or NF-κB p50KO glial cultures (referred to as hippocampal, Hp – or hypothalamic, Hyt - ACM), or in presence of NDM (referred to as standard, STD medium) for 24 h.

Primary Astrocyte Cultures. Primary mixed glial cultures were prepared from hippocampus and hypothalamus of neonatal (P1-2) C57BL/6 mice and grown in DMEM high glucose, 10% FBS, 15 mM HEPES, 2 mM L-glutamine and 100 U/100 µg/ml Penicillin/Streptomycin at a density of 78,000 cells/cm². When cells reached confluence (around 10-12 D.I.V.), proliferation of non-astrocytic cells was blocked by 10 µM cytosine arabinoside (Sigma-Aldrich) for 96h. Then cultures were switched to NDM. Phenotypic characterization of cultures was performed by immunocytochemistry with antibodies against GFAP (mouse monoclonal, 1:600, Millipore) and CD11b (rat monoclonal; 1:150, Millipore). Contaminating microglial cells were below 3% of total cells, with the remaining cells being GFAP⁺.

Preparation of astrocyte-conditioned media (ACM). To prepare astrocyte-conditioned medium we used cells at confluence. ACM was collected after 48h of conditioning of enriched glial cultures in presence of Neurobasal-A medium with B27 supplement. Medium collected was centrifuged for 10 min x 16,100 x g at 4°C, to detach cells and large debris, and filtered using 0.20 µm filter before use in NPC differentiation studies. Collected conditioned medium was stored at -80°C. The ACM was used with 1:2 dilutions in all experiments.

Immunolocalization studies in adult hippocampal and hypothalamic NPC. Phenotypic characterization of NPC-derived cells was carried out by immunolocalization for MAP-2 (rabbit polyclonal, 1:600; Millipore), GFAP (mouse monoclonal, 1:600, Millipore), nestin (chicken monoclonal, 1:1500, Neuromics), NG-2 (rabbit polyclonal, 1:500; Millipore). Nuclei were counterstained with 0.8 ng/ml Hoechst (Thermo Fisher scientific) diluted in PBS. In each experiment, 5 fields/well (corresponding to about 150-200 cells/well) were counted with a 60X objective by a Leica DMIRB inverted fluorescence microscope. Cells positive for MAP-2, GFAP and NG-2 were counted and their percentage over total viable cells was calculated.

Statistical analysis. All experiments were run in triplicate using different cell preparations and repeated at least three times. Data were analyzed using Student's *t*-test when only two independent groups were compared, or by one-way analysis of variance (ANOVA) followed by Tukey's post-hoc when comparing more than three groups. Results were expressed as mean ± S.D. Statistical significance level was set for *p* values < 0.05.

Results

Hippocampal but not hypothalamic ACM can promote neuronal differentiation of hippocampal NPC.

Soluble factors released by astrocytes have been shown to modulate differentiation of adult neural progenitor cells (NPC) (Song et al., 2002). Based on these observations we evaluated the effect of astrocyte-generated soluble factors on the differentiation rate of adult hippocampal NPC toward neuronal and non neuronal lineages. Multipotent Sox2⁺/nestin⁺/GFAP⁻ NPC isolated from adult mouse hippocampi can be maintained for several passages in an undifferentiated proliferative state (Cuccurazzu et al., 2013). Upon removal of growth factors from medium, NPC stop dividing and differentiate onto laminin-coated dishes. By double immunolabeling for markers of neurons (MAP-2) and undifferentiated progenitors (nestin), the appearance of newly generated MAP-2⁺/nestin⁻ neurons and MAP-2⁺/nestin⁺ neuroblasts, can be evaluated and quantified. Under these experimental conditions we tested the effect of hippocampal (Hp) and hypothalamic (Hyt) astrocyte-conditioned medium (ACM) in comparison with standard differentiation medium (STD). After 24 h, a significant increase in the percentage of mature neurons and neuroblasts was observed in cells exposed to Hp ACM, but not to Hyt ACM, compared to standard medium (mean % \pm S.D. over viable cells of MAP-2⁺/nestin⁻ cells: 3.49 \pm 0.40, 5.69 \pm 0.67, 3.46 \pm 0.33 % in presence of STD medium, Hp and Hyt ACM, respectively; ANOVA, $p < 0.001$ vs STD; Fig. 1 A; MAP-2⁺/nestin⁺ cells: 32.10 \pm 4.08, 40.46 \pm 2.66, 33.49 \pm 2.69 %, in presence of STD medium, Hp and Hyt ACM, respectively; ANOVA, $p < 0.001$ vs STD; Fig. 1 B). Under the same experimental *in vitro* conditions adult hippocampal NPC can also differentiate toward the astrocytic lineage. Interestingly, both Hp and Hyt ACM resulted in an increased percentage of GFAP⁺ cells when compared to standard medium (GFAP⁺ cells: 14.53 \pm 1.48, 29.18 \pm 6.74, 36.42 \pm 5.10 in presence of STD medium, Hp and Hyt ACM, respectively; ANOVA, $p < 0.001$ vs STD;

Fig. 1 C). Conversely, the NG-2⁺ population of oligodendroglial precursor cells was not affected by the treatment with Hp and Hyt ACM compared to STD medium (NG-2⁺ cells: 30.35 ± 6.00; 29.15 ± 6.04, 34.91 ± 4.59 %, in presence of STD medium, Hp and Hyt ACM; Fig. 1 D). To investigate whether ACM could promote cell survival in addition to neuronal differentiation, we also analyzed NPC apoptotic rate. No difference in the percentage of apoptotic cells was reported in presence of the different media (mean % ± S.D. over total cells: 9.08 ± 1.23, 7.13 ± 1.43, 6.86 ± 1.73 % in presence of STD medium, Hp and Hyt ACM, respectively; Fig. 1 E). Altogether these data suggest that a certain degree of region-specificity in the proneurogenic but not in the astroglial signals provided by astrocytes on adult hippocampal NPC. Apparently, only hippocampal astrocytes but not hypothalamic astrocytes can secrete proneurogenic signals for ahNPC. Conversely, both hippocampal and hypothalamic astrocytes can trigger astroglialogenesis from ahNPC.

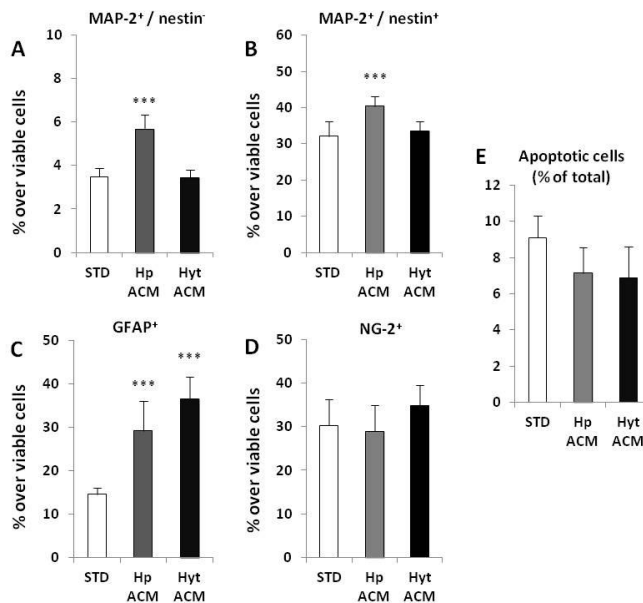


Figure 1. Hypothalamic ACM, unlike Hippocampal ACM, does not promotes hippocampal NPC neuronal differentiation. Both ACM promote astroglialogenesis. A-B Under differentiating conditions, 24h treatment of WT

hippocampal NPC with Hp ACM significantly increased the percentage of the subpopulations of MAP-2⁺/nestin⁻ neurons (**A**) and MAP-2⁺/nestin⁺ neuroblasts (**B**) compared with STD. Conversely, Hyt ACM was devoid of proneurogenic effect. **C**, Astroglial population of GFAP⁺ cells increased upon both Hp and Hyt ACM treatment. **D**, No significant differences was observed in the percentage of NG-2⁺ cells upon different media conditions. **E**, No significant difference was observed in the apoptotic rate of cultures exposed to STD medium, Hp and Hyt ACM. Data are expressed as mean values \pm S.D. of $n = 3$ experiments, run in triplicates and analyzed by one-way ANOVA followed by Tukey's post-hoc test. *** $p < 0.001$ versus STD medium.

Hypothalamic NPC do not respond to proneurogenic soluble signals coming from the neurogenic astrocytes of the hippocampal niche.

Recently hypothalamus was identified as a novel adult proneurogenic niche where newly formed neurons might contribute to the homeostatic control of many physiological behaviors related to hypothalamic functions, including energy balance regulation (Pierce and Xu 2010; Whalley 2012). In our laboratory we have set up and characterized an *in vitro* model of adult hypothalamic (Hyt) NPC (Salem et al, unpublished observations).

We then asked how adult Hyt NPC respond to astrocyte-secreted signals. In order to do so, we tested hypothalamic WT NPC differentiation rate toward neuronal and glial lineages upon treatment with astrocyte conditioned media derived from the hippocampal niche. When we tested the effect of Hp ACM on hypothalamic NPC, we did not observe any effect on neuronal differentiation (mean % \pm SD over viable cells of MAP-2⁺/nestin⁻ cells: 4.61 ± 0.54 , 4.48 ± 0.46 %, in presence of STD medium and Hp ACM, respectively; Fig. 2 *A*; MAP-2⁺/nestin⁺ cells: 35.27 ± 4.15 , 35.88 ± 3.73 %, in presence of STD medium and Hp ACM, respectively; Fig. 2 *B*). Under the same experimental conditions we observed a significant increase in newly generated astroglial cells upon ACM treatment (GFAP⁺ cells: 29.11 ± 6.03 , 44.09 ± 2.35 %, in presence of STD medium and Hp ACM, respectively; Fig. 2 *C*). Interestingly, a small, but significant increase in the percentage of NG-2⁺ cells in Hyt NPC cultures

was observed upon Hp ACM treatment (NG-2⁺ cells: 33.12 ± 3.49 , 37.19 ± 2.47 %, in presence of STD medium and Hp ACM, respectively; Student's *t*-test vs STD, Fig. 2 D). No difference in the percentage of apoptotic cells was reported between STD medium and Hp ACM treatment (mean % \pm S.D. over total cells: 6.91 ± 1.01 , 6.03 ± 1.32 %, STD medium and WT ACM, respectively; Fig. 2 E). Altogether, these preliminary data suggest that soluble molecules from hippocampal astrocytes are not able to promote neuronal differentiation of Hyt NPC. On the contrary, similarly to what we observed on Hp NPC, hippocampal ACM can promote astrogliogenesis from Hyt NPC. Additionally, Hp ACM increase the NG-2⁺ cell population only from Hyt NPC and not from Hp NPC. NG-2⁺ cells are commonly considered oligodendrocyte precursors (Nishiyama et al., 2009). Interestingly, it has been proposed that, although controversial, hypothalamic NG-2⁺ cells, distributed throughout the mediobasal hypothalamus, may represent a population able to give rise to new neurons (Robins et al., 2013). Future studies will be needed for understanding the nature of this population in our cell culture model, for example by assessing, under long-term culture conditions, the fate of these cells toward oligodendrocytic and neuronal lineages.

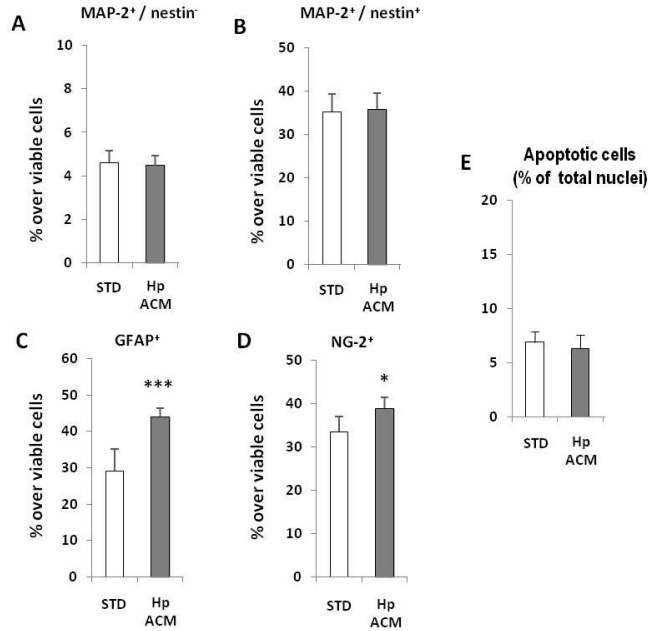


Figure 2. Hippocampal ACM promotes an increase in GFAP⁺ and NG-2⁺ cells from Hypothalamic NPC. *A-B* Under differentiating conditions, 24 h treatment of hypothalamic NPC with hippocampal ACM did not affect the percentage of the subpopulations of mature MAP-2⁺/nestin⁻ neurons (*A*) and MAP-2⁺/nestin⁺ neuroblasts (*B*) compared with standard differentiation medium (STD). *C*, Conversely, hippocampal ACM was able to promote astroglial differentiation. *D*, Hypothalamic NPC showed a significant increase in the percentage of NG-2⁺ cells upon treatment with hippocampal ACM. *E*, No significant difference was observed in the apoptotic rate of cultures exposed to STD medium or hippocampal ACM. Data are expressed as mean values \pm S.D. of $n = 3$ experiments, run in triplicates and analyzed by Student's *t*-test, * $p < 0.05$ and *** $p < 0.001$ versus STD medium.

Similar effects of p50KO Hp and Hyt ACM on WT hippocampal NPC.

We then evaluated if NF- κ B p50 absence has different consequences on the soluble signals released by hippocampal and hypothalamic astrocytes. In order to do so, primary astrocyte cultures were set up also from both regions of p50KO animals, ACM were collected and then tested in parallel on Hippocampal NPC. As previously shown (see Cvijetic et al. under submission) Hp p50KO ACM promotes astrogliogenesis (Fig. 3 B) but not neurogenesis (Fig. 3 A). Very similar effects were observed with p50KO Hyt ACM. In detail: MAP-2⁺/nestin⁻ cells: 3.31 ± 0.43 , 3.51 ± 0.82 , 3.34 ± 0.67 % in presence of STD medium, Hp and Hyt p50KO ACM, respectively (Fig. 3 A); GFAP⁺ cells: 13.16 ± 1.40 , 36.00 ± 3.24 , 30.38 ± 5.15 % in presence of STD medium, Hp and Hyt p50KO ACM, respectively (ANOVA, $p < 0.001$ vs STD; Fig. 3 B). When we checked for NG-2⁺ oligodendroglial precursor cells, we did not observe any significant effect on NG-2⁺ cells differentiation upon any treatment (NG-2⁺ cells: 30.36 ± 0.98 , 27.21 ± 5.21 , 35.58 ± 4.28 % in presence of STD medium, Hp and Hyt p50KO ACM, respectively; Fig. 3 C). No difference in the percentage of apoptotic cells was reported in presence of the different media (mean % \pm SD over total cells: 8.33 ± 0.54 , 6.92 ± 1.18 , 7.18 ± 1.52 % in presence of STD medium, Hp and Hyt p50KO ACM, respectively; Fig. 3 D). This subset of data suggests that absence of p50 subunit may not affect the astrogliogenic potential of both hippocampal and hypothalamic astrocytes on Hippocampal NPC.

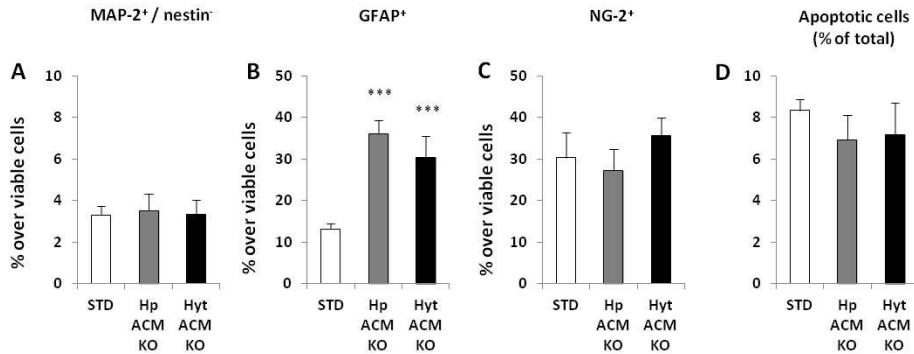


Figure 3. No significant differences in the effects of p50KO Hippocampal and Hypothalamic ACM on WT hippocampal NPC. *A*, Under differentiating conditions, 24h treatment of WT hippocampal NPC with Hp or Hyt ACM deriving from p50KO astrocytes does not induce neuronal differentiation compared with standard differentiation medium (STD). *B*, Conversely, both media are able to increase astroglial population of GFAP⁺ cells. *C*, No significant difference was observed in the percentage of NG-2⁺ cells upon different media conditions. *D*, No significant difference was observed in the apoptotic rate of cultures exposed to STD medium, Hp and Hyt ACM p50KO. Data are expressed as mean values \pm S.D. of $n = 3$ experiments, run in triplicates and analyzed by one-way ANOVA followed by Tukey's post-hoc test. *** $p < 0.001$ versus STD medium.

We extended our analysis by testing the effect of Hp p50KO ACM on WT hypothalamic NPC. Also in this experiment, we confirmed that hippocampal p50KO ACM (like WT hippocampal ACM on hypothalamic NPC) could not produce proneurogenic effects compared to STD condition. In detail the percentages of newly generated populations were as follows: MAP-2⁺/nestin⁻ cells, 4.61 ± 0.54 , 3.99 ± 0.57 % in presence of STD medium and Hp p50KO ACM, respectively (Fig. 4 A); MAP-2⁺/nestin⁺ cells: 35.27 ± 4.15 , 33.57 ± 3.83 % in presence of STD medium and Hp p50KO ACM, respectively (Fig. 4 B); GFAP⁺ cells: 29.12 ± 6.03 , 47.35 ± 1.95 % in presence of STD medium and Hp p50KO ACM, respectively (Student's *t*-test, *** $p < 0.001$ vs STD; Fig. 4 C). Interestingly, Hp p50KO, unlike Hp WT ACM, was devoid of effect on the NG-2⁺ population (NG-2⁺ cells: 33.12 ± 3.50 , 37.11 ± 5.13 % in presence of STD

medium and Hp p50KO ACM, respectively; Fig. 4 D). No difference was also present in the apoptotic rate in presence of the different media conditions (mean \pm SD over total cells: 7.54 ± 1.20 , 6.32 ± 1.53 % in presence of STD medium and Hp p50KO ACM, respectively; Fig. 4 E). These very preliminary data suggest that the absence of p50 may affect astrocyte secretion of soluble molecules which may promote an increase in the NG-2⁺ population originating from Hyt NPC.

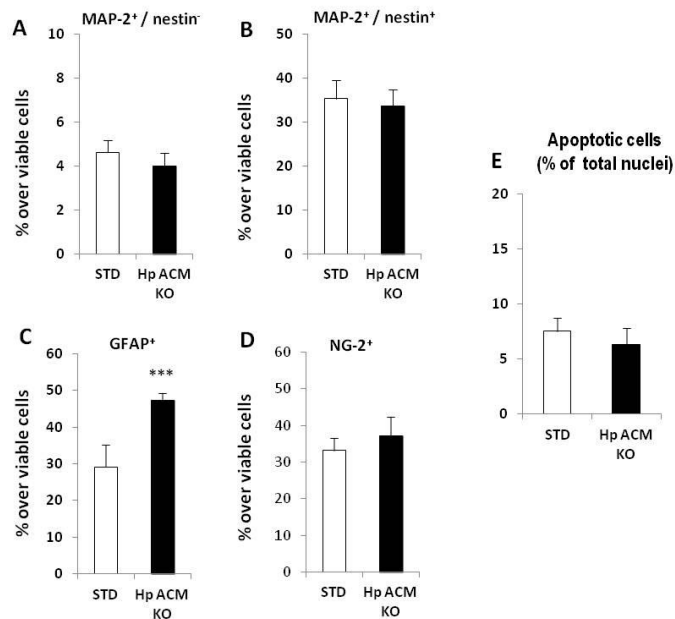


Figure 4. Hypothalamic WT NPC, in presence of Hippocampal ACM originating from p50KO astrocytes, differentiate only toward the astroglial lineage. *A-B* Under differentiating conditions, 24 h treatment of WT hypothalamic NPC with Hp ACM deriving from p50KO astrocytes does not induce neuronal differentiation compared with standard differentiation medium (STD). *C*, Conversely, Hp p50KO ACM is still able to increase astroglial population of GFAP⁺ cells. *D*, No significant differences was observed in the percentage of NG-2⁺ cells upon Hp p50KO ACM treatment. *E*, No significant difference was observed in the apoptotic rate of Hyt NPC cultures exposed to STD medium versus Hp p50KO ACM. Data are expressed as mean values \pm S.D. of $n = 3$ experiments, run in triplicates and analyzed by Student's *t*-test *** $p < 0.001$ versus STD medium.

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Chapter 5.

Discussion

Scientific background

Adult neurogenesis is a plastic and dynamic process that persists throughout post-natal life in mammalian species, including humans. Active sites of adult neurogenesis are the subventricular zone (SVZ) of the lateral ventricles, and the subgranular zone (SGZ) of the dentate gyrus in the hippocampus (Ming and Song, 2005; Bond and Song, 2015). Growing evidence suggests the presence of another neurogenic area in the hypothalamus, along the walls of the third ventricle (Kokoeva et al., 2005). The process of adult neurogenesis is supported by a pool of neural stem/progenitor cells (NSPC), which reside in a permissive microenvironment called “neurogenic niche”, where the different cellular types and molecular components provide a supportive milieu that balance NSPC quiescence along with proliferation and differentiation and survival (Lin & Iacovitti, 2015).

Astrocytes represent one major contributor within the hippocampal neurogenic niche. Astrocytes isolated from neurogenic and non-neurogenic brain areas have been reported to exhibit different effects on the neuronal differentiation of NSPC suggesting that functional heterogeneity of astrocytes may reflect the different brain region environment (Song et al. 2002; Barkho et al. 2006).

In previous publications our lab proved that members of the NF- κ B family of transcription factors are important contributors of signaling pathways in the SGZ neurogenic niche (Denis-Donini et al., 2005). Within the NF- κ B family, the p50 subunit appears to act as a crucial regulator of adult hippocampal neural progenitor cells (ahNPC). Specifically focusing on NF- κ B p50 subunit, our group has demonstrated that NF- κ B is involved in mediating the proneurogenic and antidepressive-like effect of some clinically relevant drugs (i.e. α 2 δ ligands; acetyl-L-carnitine) (Cuccurazzu et al., 2013; Valente et al., 2012), in the proneurogenic effects of damage-associated signals (i.e. HMGB-1 and A β ₁₋₂) (Meneghini et al., 2013), and that the absence of p50 affects the *in vitro*

response to those drugs/signals. *In vivo* our group also proved that p50KO mice display a dramatically reduced adult hippocampal neurogenesis which is in association with hippocampal-dependent short-term memory defects (Denis-Donini et al., 2008).

Working hypothesis

The major project for the PhD thesis developed from the observation that unlike *in vivo*, hippocampal NPC derived from WT and p50KO adult mice have similar rate of differentiation, proliferation and apoptosis *in vitro*. We hypothesized that deficient signaling within the niche and/or intrinsic defects of NPC, might contribute to the reduced neurogenesis observed *in vivo* in p50KO mice. In particular, we focused on astrocytes as one of the relevant components in niche signaling and well-studied modulators in neurogenesis. We decided to explore ahNPC-astrocyte cross-talk *in vitro* in order to further dissect the contribution of non-cell autonomous signaling and cell autonomous defects, played by NF- κ B p50, in fate specification.

Principal findings of the thesis work

Based on these observations, we exploited the advantages of using primary cell culture systems to investigate the contribution made by astrocyte conditioned media (ACM), under basal conditions, on adult NPC fate specification. The most relevant finding concerning WT NPC cultures, was that neuronal fate determination is mediated only by WT ACM treatment, whereas p50KO ACM has no proneurogenic activity. On the contrary, both WT and p50KO ACM display astroglial activity. The effect we observe in our experimental conditions are merely differentiative since ACM had no influence on the survival of NPC and their progeny. Interestingly, when we exposed p50KO NPC to WT and p50KO ACM, we did not observe any difference in lineage specification compared to STD condition, suggesting that also cell autonomous defects may affect the responsiveness to proneurogenic and astroglial signals secreted from the astrocytes. Previously our group demonstrated that $\alpha 2\delta 1$, a

thrombospondin-1 (TSP-1) receptor (Eroglu et al., 2009), is functionally expressed by ahNPC (Valente et al., 2012). During this PhD thesis we proved that $\alpha 2\delta 1$ /TSP-1 receptor expression is downregulated in p50KO NPC. When we tested thrombospondin-1, a well-known proneurogenic glycoprotein of the extracellular matrix released by astrocytes (Lu & Kipnis, 2010) we observed a defective response to the proneurogenic molecule only in p50KO NPC and not in WT NPC. Overall these data suggest that absence of p50 may also reduce ahNPC responsiveness to TSP-1, an astrocyte-derived signal.

A step forward in this study was done by analyzing the secretory profile of WT and p50KO astroglial cultures. By analyzing WT and p50KO ACM, we wanted (i) to explore if there were proteins differentially modulated under the control of NF- κ B p50, (ii) to identify, among the detected proteins, potential proneurogenic or antineurogenic molecules, to be further validated in our *in vitro* system. Through LC-MS/MS proteomic analysis of ACM, we identified four proteins that were upregulated in p50KO compared to WT ACM: neutrophil gelatinase-associated lipocalin-2 (NGAL/LCN-2), C-C motif chemokine-2 (CCL-2), H-2 class I Histocompatibility antigen, K-K alpha chain (HA1K) and retinoic acid receptor responder 2 (RARRES2, chemerin).

As a starting point, we focused our attention on Lipocalin-2 (LCN-2). No information is currently available on its role on adult neurogenesis. LCN-2 is mainly known as an acute phase protein secreted from activated microglia, reactive astrocytes and neurons. In the CNS, LCN-2 has been suggested to play a modulatory action in cognitive functions (Ferreira et al., 2013), neuronal excitability and anxiety (Mucha et al., 2011) and depression in human beings (Naudé et al., 2013). However, its precise role in the pathophysiology of the CNS remains to be outlined. We discovered for the first time that LCN-2 is able to elicit neuronal differentiation of WT ahNPC in a concentration-dependent fashion. Conversely, LCN-2 protein was found less active on p50KO ahNPC. These results, suggest for the first time that LCN-2 is an astroglial-derived signal that may promote neuronal differentiation of ahNPC and the lack of p50 results in a reduced response to its proneurogenic activity.

As described in the unpublished section of this thesis, we also collected some preliminary data on the cross-talk between ahNPC and ACM originating from the hypothalamic area. The hypothalamic region has been discovered recently as a new neurogenic site, where neurogenesis might be regulated by growth factors, diet and hormones, given the involvement of this region in homeostatic and physiological processes (Kokoeva et al., 2005; Lee et al., 2014). As a follow up of the experiments performed before, we compared the proneurogenic and progliogenic properties of hippocampal and hypothalamic ACM deriving from WT astrocytes on adult hippocampal WT NPC fate specification. Additionally we assessed the contribution in NPC cell fate determination of hippocampal and hypothalamic astrocytes cultures deriving from NF- κ B p50KO mice.

Our finding reports that hippocampal but not hypothalamic ACM can promote neuronal differentiation of hippocampal NPC, confirming that astrocytes may provide region-specific signals for the neurogenic process. This subset of data suggested us also that the astroglial signals are still retained in absence of p50 subunit and independently of the origin of the astrocytes. We further explored the lineage fate determination of adult hypothalamic WT NPC in basal conditions and in response to hippocampal WT and p50KO ACM. We found out that hypothalamic WT NPC do not respond to proneurogenic soluble signals coming from WT hippocampal astrocytes. Conversely, hypothalamic NPC in presence of both WT and p50KO hippocampal ACM only differentiate toward astroglial lineage. Furthermore, we observed for the first time a pro-differentiative effect of WT hippocampal ACM on oligodendroglial NG-2 cell population of WT hypothalamic NPC, and this effect appeared p50-dependent.

Conclusions and future perspectives

This work supports the concept that non-cell autonomous defects in astrocyte signaling might be relevant for the neurogenic process *in vitro* and, presumably, *in vivo*. We suggest that the absence of NF- κ B p50 subunit in astrocytes is a critical aspect that negatively influence the proneurogenic activity of cortico-hippocampal and hippocampal ACM.

On the other hand we confirm that NPC intrinsic defects might be at the basis of the deregulated neurogenesis observed in p50KO mice, since hippocampal NPC have downregulated α 2 δ 1 expression levels. Additionally we can confirm the importance of NF- κ B p50 in the regulation of proteins secreted by astrocytes, including new proneurogenic molecules such as LCN-2.

We also confirm the concept of region-specific regulation of ahNPC by molecular cues arising from the same neurogenic region. The field of region-specific proneurogenic potential deserves further investigation. Moreover, it would be interesting to assess the responsiveness of hypothalamic NPC to proneurogenic molecules relevant in hippocampal NPC neuronal differentiation (i.e. TSP-1, LCN-2).

Since LCN-2 was identified as an astrocyte-derived factor that exerts proneurogenic activity in WT NPC and to a lesser extent in p50KO NPC, we would like to further explore lipocalin-2 receptor expression, BOCT1/24p3R and megalin, in hippocampal WT and p50KO NPC.

Potential future work could also be aimed at understanding the involvement in neuronal fate determination of ahNPC for other proteins identified in astrocyte secretome, for example CCL-2. There are many reports claiming both negative and positive modulatory effects of the chemokine CCL-2 in neuronal differentiation of ahNPC. On one hand increased systemic levels of CCL-2 observed during aging have been associated with decreased neurogenesis and age-related cognitive impairment (Villeda et al., 2011), on the other hand other results indicate that CCL-2 can influence both the migration, survival and fate choice of SVZ-derived neural precursor cells (Gordon et al., 2012). Therefore,

it could be interesting to evaluate WT and p50KO NPC responsiveness to this chemokine.

Another potential field of future investigation is that of intercellular communication via extracellular vesicles (EV), such as exosomes and shedding microvesicles, which are involved in neuronal protection, neuronal development, as well as neurodegeneration (Kalani et al., 2014; Rajendran et al. 2014). It could be of interest exploring the involvement of EV as a possible novel mechanism of glia-to-NPC transfer of molecules which may affect NPC fate determination. Interestingly, LCN-2 and TSP-1 have both been reported to be secreted in exosomes (Burke et al., 2014).

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Chapter 6.

Manuscript under submission to Journal of Neuroscience

Astrocyte-mediated neuronal fate specification of adult hippocampal neural progenitors is regulated by NF- κ B p50.

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Poster sessions

NF- κ B-mediated regulation of astrocyte-secreted signals affects the neurogenic potential of adult hippocampal neural progenitors. V. Bortolotto, **S. Cvijetic**, M. Manfredi, E. Ranzato, E. Marengo, P.L. Canonico, M. Grilli. (45th annual meeting of the Society for Neuroscience, Chicago, USA, October 17-21, 2015)

NF- κ B-mediated regulation of astrocyte-secreted signals modulating the differentiation potential of adult hippocampal neural progenitor cells. **S. Cvijetic**, V. Bortolotto, E. Marengo, M. Manfredi, E. Ranzato, P.L. Canonico, M. Grilli. (37° Congresso Nazionale della Società Italiana di Farmacologia “I Nuovi Orizzonti Della Ricerca Farmacologica: Tra Etica E Scienza” Napoli, October 27-30, 2015)

Involvement of the NF- κ B p50 subunit in adult Neural Progenitor Cell-astroglia cross-talk. **S. Cvijetic**, V. Bortolotto, S. Lovecchio, R. Salem, P.L. Canonico, M. Grilli. (Abcam event: Adult Neurogenesis: Evolution, Regulation and Function, Dresden, Germany, May 6-8, 2015)

Potential role of NF- κ B p50 in astrocyte-neural progenitor cell communication within the neurogenic niche. V. Bortolotto, **S. Cvijetic**, S. Lovecchio, P.L. Canonico, M. Grilli. (44th Annual meeting of the Society for Neuroscience, Washington DC, USA, November 2014)

Neural progenitor cell-astroglia cross-talk: involvement of NF- κ B p50 subunit and extracellular vesicles. **S. Cvijetic**, V. Bortolotto, S. Lovecchio, P.L. Canonico, M. Grilli. (17° Seminario SIF Dottorandi, Assegnisti di Ricerca, Postdottorandi e Specializzandi, Rimini, September 16-18, 2014)

Role of NF- κ B p50 in microglia-neural progenitor cell cross-talk. **S. Cvijetic**, V. Bortolotto, S. Lovecchio, P.L. Canonico, M. Grilli. (9th FENS Forum of Neuroscience, Milan, July 5-9, 2014)

NF- κ B p50 subunit contribution to microglia-neural progenitor cells cross-talk: potential relevance in adult neurogenesis deregulation. **S. Cvijetic**, V. Bortolotto, S. Lovecchio, P.L. Canonico, M. Grilli (Monotematico SIF: Mood disorders: from neurobiology to novel therapeutic strategies, Modena, March 20-21, 2014)

Awards

Premio miglior comunicazione orale al 17° Seminario SIF - Dottorandi, Assegnisti di Ricerca, Postdottorandi e Specializzandi (Rimini, September 16-18, 2014). Titolo del lavoro premiato: Neural progenitor cell-astroglia cross-talk: involvement of NF- κ B p50 subunit and extracellular vesicles. **S. Cvijetic**, V. Bortolotto, S. Lovecchio, P.L. Canonico, M. Grilli

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