



Diversity of Neural Precursors in the Adult Mammalian Brain

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Aided by advances in technology, recent studies of neural precursor identity and regulation have revealed various cell types as contributors to ongoing cell genesis in the adult mammalian brain. Here, we use stem-cell biology as a framework to highlight the diversity of adult neural precursor populations and emphasize their hierarchy, organization, and plasticity under physiological and pathological conditions.

The adult mammalian brain displays remarkable structural plasticity by generating and incorporating new neural cell types into an already formed brain (Kempermann and Gage 1999). Largely restricted within the subventricular zone (SVZ) along the lateral ventricle and the subgranular zone (SGZ) in the dentate gyrus (DG), neural genesis is thought to arise from neural stem cells (NSCs) (Ming and Song 2011). Stem cells are defined by hallmark functions: capacity to self-renew, maintenance of an immature state over a long duration, and ability to generate specialized cell types (Fig. 1). These

features distinguish stem cells from committed progenitor cells that more readily differentiate into specialized cell types (Fig. 1). Stem and progenitor cells (collectively called precursors) are additionally characterized by their lineage capacity. For example, multipotential neural precursors generate neurons and glia, whereas unipotential cells produce only one cell type, such as neurons (Gage 2000; Ma et al. 2009). The classical NSC definition is based on cell culture experiments in which a single cell can self-renew and generate neurons, astrocytes, and oligodendrocytes (Gage 2000; Ma et al. 2009).

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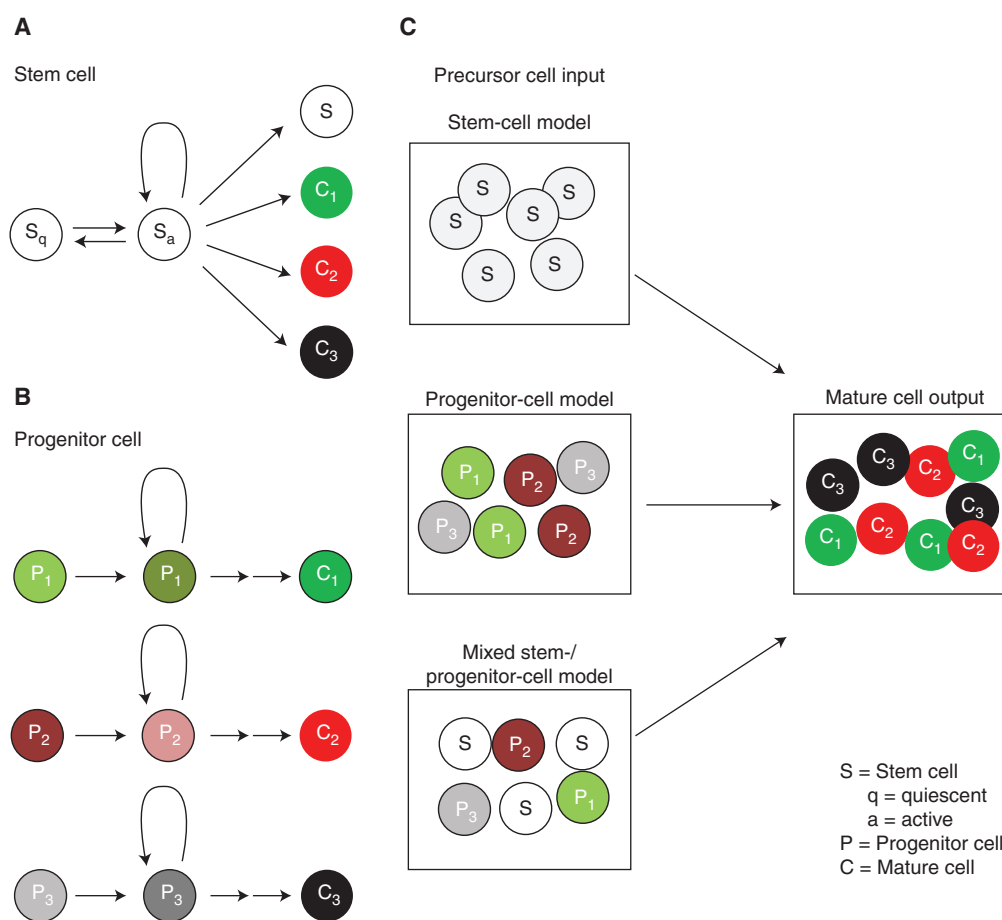


Figure 1. Models of generating cell diversity in the adult tissues. (A,B) Definitions of stem and progenitor cells. In A, quiescent stem cells (S_q) become active stem cells (S_a) that proliferate to generate different types of specialized cells (C_1 , C_2 , C_3) and new stem cells (S). The active stem cell can return to quiescence and remain quiescent over long periods of time. In B, lineage-restricted progenitor cells lacking self-renewal capacity (P_1 , P_2 , P_3) each give rise to distinct populations of specialized cells (C_1 , C_2 , C_3). (C) Generation of specialized cells in a tissue could be explained by three models. (1) The stem-cell model, in which multipotent stem cells give rise to all the specialized cells in the tissue. (2) The progenitor cell model, in which diverse, lineage-restricted progenitor cells give rise to different cell types in the tissue. (3) A hybrid model, in which a mixture of stem cells and lineage-restricted progenitor cells generate specialized cells of the adult tissue.

Yet, reprogramming studies have raised the question of whether cultured lineage-restricted neural progenitors acquire additional potential not evident in vivo (Palmer et al. 1999; Kondo and Raff 2000; Gabay et al. 2003). As a result, various lineage models have been proposed to explain cell generation in the adult brain (Fig. 1) (Ming and Song 2011). In one model, bona fide adult stem cells generate multiple lineages at the individual cell level. In another, cell genesis rep-

resents a collective property from a mixed population of unipotent progenitors. Importantly, these models are not mutually exclusive as evidence for the coexistence of multiple precursors has been observed in several adult somatic tissues, in which one population preferentially maintains homeostasis and another serves as a cellular reserve (Li and Clevers 2010; Mascré et al. 2012). Recent technical advances, including single-cell lineage tracing (Kretschmar and



Watt 2012), have made it possible to dissect basic cellular and behavioral processes of neural precursors in vivo (Table 1) and have revealed diverse neural precursor populations coexisting within classical neurogenic zones (see Fig. 4) (Bonaguidi et al. 2012). In this work, we review our current knowledge of precursor cell identity, hierarchical organization, and regulation to examine the diverse origins of cell genesis in the adult mammalian brain.

HIERARCHICAL PRECURSOR CELLS IN THE ESTABLISHED ADULT NEUROGENIC NICHE

Classically, cell lineages are arranged in a hierarchy with cells of the greatest plasticity at the top and fully differentiated cells at the bottom (Waddington 1957). In the adult brain, NSCs represent the top layer followed by restricted neural progenitors and finally specialized cell types: astrocytes, oligodendrocytes, and neurons (Fig. 2). In the prevalent model of adult neurogenesis, radial cells (SGZ: radial glia-like

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cell [RGL], or type 1 cells; SVZ: type B cells) act as NSCs to generate committed proliferative precursors known as intermediate progenitors (SGZ: intermediate progenitor cells [IPCs], or type 2 cells; SVZ: type C cells), which give rise to neuroblasts (SGZ: type 3 cells; SVZ: type A cells) and then immature neurons (Fig. 2). Emerging in vivo evidence indicates that astroglial and oligodendroglial lineages could also arise from lineage-restricted progenitor cells (Zhu et al. 2011). Although these hierarchical relationships exist under physiological conditions, neural precursors' capacity and therefore hierarchical position may alter on injury or exposure to ectopic molecular cues (Palmer et al. 1999; Kondo and Raff 2000; Gabay et al. 2003). Therefore, each experimental approach used to examine stem-cell properties and establish hierarchical relationships requires careful consideration of the cellular and molecular context as well as an understanding of the precursor cells initially targeted. Precursor diversity may occur at any of the three hierarchical levels owing to (1) different cell-type identities, and the heterogeneity within a cell type based on its (2) behavior/lineage capacity, and/or (3) function under environmental alterations.

Putative NSCs

Radial glia-like precursor cells are characterized by the expression of nestin, glial fibrillary acidic protein (GFAP), Sox2, and possess a defining radial branch that extends through the granule cell layer in the dentate or contacts blood vessels in the SVZ (Fuentealba et al. 2012). In both niches, radial cells display a relatively quiescent nature that distinguishes them from nonradial and intermediate progenitors (Morshead et al. 1994; Lugert et al. 2010). Evidence from both physiological and nonphysiological approaches, including population level transgenic fate mapping (Ahn and Joyner 2005; Lagace et al. 2007; Dranovsky et al. 2011), genetic ablation (Garcia et al. 2004), and antimitotic treatment recovery (Seri et al. 2001), suggest radial cells exist atop the cellular hierarchy. In the adult SGZ, the neuronal and astroglial lineages, but not the oligodendrocyte lineage, are generated under physi-

Table 1. Comparison of different methods used to study the generation of new cells in the adult mammalian nervous system

- (1) In vivo imaging allows real-time visualization of cells in their natural environment.
- (2) Lineage tracing is the utilization of transgenic animals to label single precursor cells and retrospectively analyze the fate choices made by these cells.
- (3) Fate mapping entails the study of lineage decision made by populations of cells, utilizing either using transgenic animals or administration of thymidine analogues.
- (4) Adenovirus, lentivirus, and retrovirus, when injected into the brain, can be used to trace single cells or population of cells depending on the virus used and the amount of virus injected into the animals.
- (5) Transplantation of precursor cells is a useful tool to examine the intrinsic and extrinsic regulation of precursor cells in the brain.
- (6–7) Ex vivo methods involve sections in the brain being maintained in culture media, whereas in in vitro studies, the dissociated cells are cultured either as neurospheres or in a monolayer culture system.

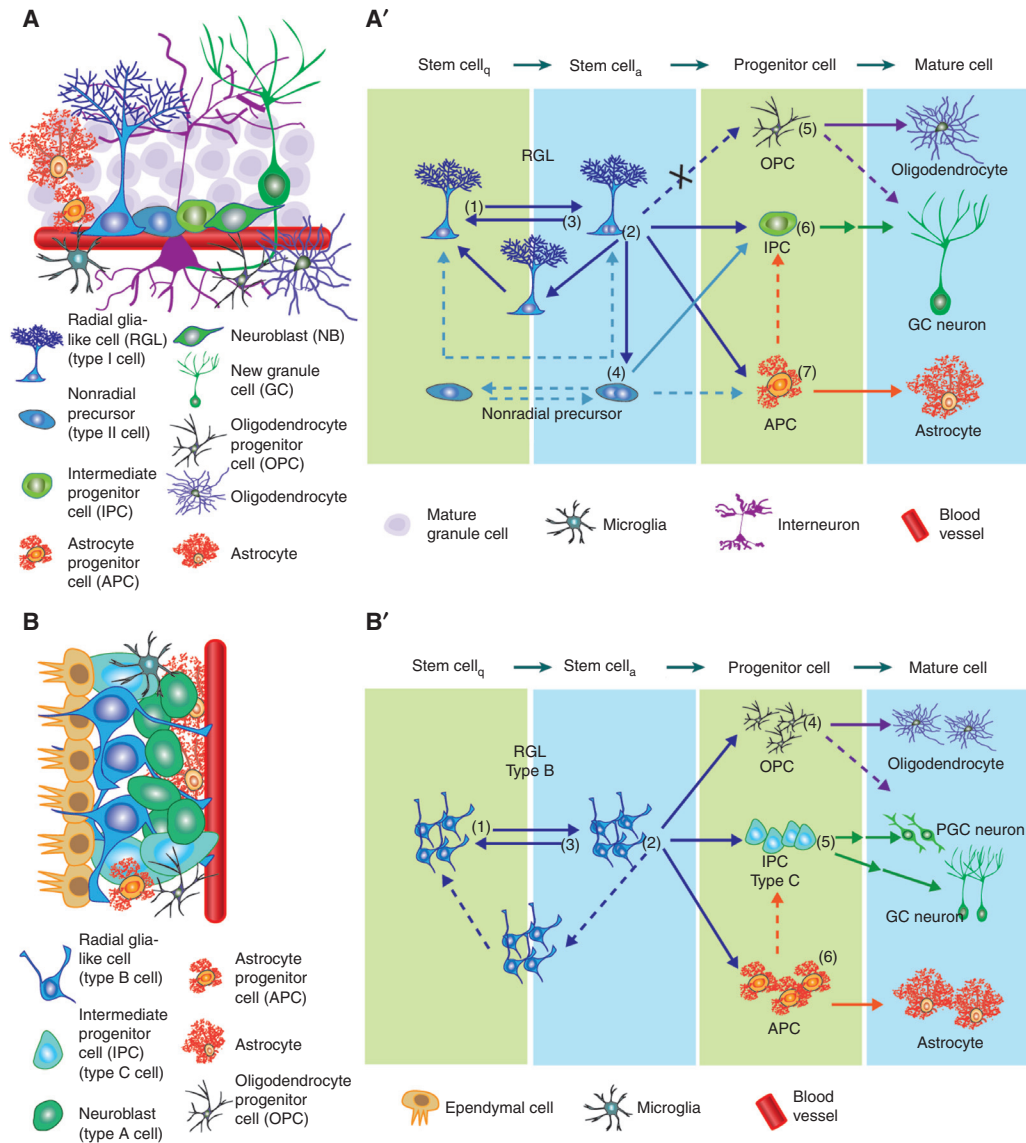


Figure 2. The stem-cell niche and lineage hierarchy among stem cells and their progeny. (A) Schematic illustration of the cell types present in the neurogenic niche of subgranular zone (SGZ) in the dentate gyrus (DG). Shown in A' is a model of the neural genesis in the adult SGZ under basal conditions. On the single-cell level, quiescent radial glia-like cells (RGLs) can become active and enter cell cycle (1). The active RGLs can generate an intermediate progenitor cell (IPC), an astrocyte progenitor cell (APC), or generate an additional RGL (2), which can become either quiescent or active. The active RGLs can remain in a proliferative state or return to quiescence (3). Nonradial, nestin⁺ cells produce IPCs and may cycle between quiescent and active states (4). Oligodendrocyte progenitor cells (OPCs) can enter cell cycle and generate mature oligodendrocytes (5). IPCs generate excitatory granule cell (GC) neurons (6). Newborn APCs can generate mature astrocytes (7). (B) A schematic illustration of the cell types present in the neurogenic niche of the subventricular zone (SVZ). Shown in B' is a model of the neural genesis in the adult SVZ under basal conditions. On the population level (indicated by more than one cell), quiescent RGLs (type B cells) can become active and enter cell cycle (1). Progeny of active type B cells can give rise to IPCs (type C cells), APCs, and OPCs. Whether the RGLs directly make these fate choices remains unknown. RGLs may also generate additional type B cells, which could be either active or quiescent. Active type B cells can return to quiescence (3). OPCs can enter cell cycle and generate mature oligodendrocytes (4). IPCs generate classes of inhibitory neurons including periglomerular cell (PGC) and GC neurons, although it remains unclear whether the fate decisions occur within RGLs or IPCs (5). Newborn APCs can generate mature astrocytes (6). Arrows indicate direct cell generation. Dotted arrows represent potential choices requiring further experimental evidence. Double arrows represent multistep cell generation. Arrows with an X represent choices not experimentally observed.



ological conditions (Suh et al. 2007; Jessberger et al. 2008). This has been confirmed by in vivo clonal analysis within the SGZ where it has been revealed that individual RGLs have the potential to be multipotent stem cells and divide not just asymmetrically but also symmetrically, self-renewing over a long duration (Bonaguidi et al. 2011; Song et al. 2012). Similarly, RGLs have been suggested to expand symmetrically over time at the population level (Dranovsky et al. 2011). However, alternative RGL properties have been proposed, including a lack of long-term maintenance. In one model, once activated, RGLs repeatedly enter cell cycle and generate only neurons before terminally differentiating into astrocytes (Encinas et al. 2011). Together, these findings suggest the existence of several RGL subpopulations that display varying levels of self-renewal and differentiation capacity (Bonaguidi et al. 2012). For example, RGLs labeled using different Cre drivers display discrete proliferation responses to running and antimetabolic treatment, suggesting that RGLs also display functional heterogeneity (DeCarolis et al. 2013).

Evidence also implicates diversity among the radial cell pool within the radial B cells in the adult SVZ (Codega et al. 2014; Mich et al. 2014). Unlike the SGZ, B cells have the capacity for generating oligodendrocytes in addition to neurons and astrocytes in vivo (Ahn and Joyner 2005; Menn et al. 2006). However, single-cell tracing of B cells ex vivo has recently indicated that discrete B cells may give rise to the neuronal and oligodendroglial lineages, whereas each can also generate the astroglial lineage (Ortega et al. 2013). These results suggest that a single B cell may not be able to clonally generate all three neural cell types. Interestingly, various putative classes of B cells can be marked by differential expression of brain lipid-binding protein (BLBP), GFAP, epidermal growth factor receptor (EGFR), Hes5, CD133, TLX, and ID1 (Nam and Benezra 2009; Pastrana et al. 2009; Li et al. 2012; Giachino et al. 2013). It is still unclear how these molecules define radial subtypes and can represent either discrete cell types or different molecular states of a single-cell type. Whether multiple NSC types with diver-

gent behaviors coexist within the adult radial population remains a fundamental question in stem-cell biology. One powerful approach to distinguish stem cells from more committed progenitors and establish cellular hierarchy is clonal analysis combined with computational modeling (Table 1) (Blanpain and Simons 2013).

Nonradial precursors have also been proposed as a stem-cell population in the adult SGZ, although their identity has not yet been described in the SVZ (Suh et al. 2007). Nonradial precursors lack any radial processes and may contain parallel extensions to the dentate granule cell layer. These precursors express Sox2, but not GFAP, and are labeled in the *Hes5:gfp* reporter mice (Lugert et al. 2010). They are more mitotic than RGLs, but most are not in cell cycle at any given time (Suh et al. 2007; Lugert et al. 2010). The identity and potential of nonradial precursors is not clearly delineated from early IPCs (type 2a cells), which share similar morphological and molecular characteristics and may represent an earlier state of IPCs (Kronenberg et al. 2003; Steiner et al. 2006; Lugert et al. 2010). In vivo clonal analysis indicates that RGLs can give rise to nonradial Sox2⁺ cells (Bonaguidi et al. 2011). Meanwhile, retrovirus-mediated lineage tracing of individual Sox2⁺ cells in the adult SGZ suggests limited capacity as most labeled clones exhibited limited self-renewal and unipotent differentiation, whereas no clones displayed both self-renewal and multipotentiality (Suh et al. 2007). Nonradial precursor identity, differentiation potential, and lineage relationship remain elusive and therefore require future studies using alternative lineage-tracing approaches (Table 1).

Found in the SVZ, but not in the SGZ, ependymal cells are postulated to be a third cell type displaying NSC properties. Defined by their proximity to the lateral ventricle and function in regulating cerebrospinal fluid (CSF) motility, ependymal cells are essential niche components for neurogenesis (Sawamoto et al. 2006; Paez-Gonzalez et al. 2011). Ependymal cells also possess characteristic cilia and express high levels of Prominin-1/CD133 along with S100B and tubulin- β -IV (Pfenninger et al. 2007). Originally proposed as an NSC population under physio-

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logical conditions, more recent studies suggest that ependymal cells contribute to neurogenesis and astrogenesis *in vivo* only under injury conditions, (Johansson et al. 1999; Coskun et al. 2008; Carlen et al. 2009). These results are also consistent with their role in the adult spinal cord where, on injury, ependymal cells proliferate and their progeny migrate toward the site of injury (Horner et al. 2000; Barnabe-Heider et al. 2010). Intriguingly, developmental studies suggest that ependymal cells and radial B cells derive from a common radial glia lineage and therefore may regain additional NSC behavior if given sufficient cues (Spassky et al. 2005). Ultimately, ependymal cells may represent a reserve neural stem-cell pool, although this requires further confirmation.

Lineage Committed Neural Progenitor Cells

Neural progenitor cells are generally considered to be restricted to a single lineage under physiological conditions. For example, IPCs, oligodendrocyte progenitor cells (OPCs), and astrocyte progenitor cells (APCs) generate neurons, oligodendrocytes, and astrocytes, respectively (Fig. 2). Common among them, with the possible exception of APCs, is a high level of proliferation—especially relative to NSCs (Bonaguidi et al. 2011; Costa et al. 2011; Encinas et al. 2011; Ponti et al. 2013). IPCs are the most numerous cell type in cell cycles within the adult SVZ and SGZ, and undergo ~3 divisions in both regions (Encinas et al. 2011; Ponti et al. 2013). These cells possess small tangential processes, predominately express Tbr2 in the SGZ, Mash 1 and/or *Dlx 2* (type C cells) in the SVZ, and preferentially incorporate BrdU (Doetsch et al. 2002; Hodge et al. 2008; Ponti et al. 2013). Derived from both radial and nonradial precursors (Fig. 2) (Kronenberg et al. 2003; Luger et al. 2010; Bonaguidi et al. 2011; Encinas et al. 2011), IPCs are considered transient amplifying precursors because they soon express doublecortin (DCX), a marker of committed immature neurons (Hodge et al. 2008). It remains unclear whether they can also produce astroglia or oligodendrocytes and self-renew over a long duration under basal conditions.

However, IPCs may acquire additional differentiation capacity *in vivo* following changes in the local environment, such as mediated by epidermal growth factor (EGF) infusion into the SVZ (Craig et al. 1996; Kuhn et al. 1997; Doetsch et al. 2002; Pastrana et al. 2009), suggesting that stem-cell competence may extend into normally lineage-restricted cells.

NG2⁺ oligodendrocyte progenitors constitute the major proliferative population in the adult nonneurogenic areas and are also found in substantial number in both the SVZ and SGZ (Dawson et al. 2003; Encinas et al. 2011). OPCs possess long wispy processes, express the characteristic proteoglycan NG2, as well as *Olig2*, platelet-derived growth factor receptor α (PDGFR- α), and *Sox10* (Zhu et al. 2011). Despite substantial debate over OPC cell potential (Belachew et al. 2003; Kang et al. 2010), the emerging consensus is that they produce astroglia and oligodendrocytes during development, but are restricted to oligodendrocyte generation in adult under physiological conditions (Fig. 2) (Zhu et al. 2011). OPCs and oligodendrocytes are generated from B cells in the SVZ, which migrate great distances into the corpus callosum, striatum, and fimbria fornix (Menn et al. 2006; Costa et al. 2011). So far, OPCs and oligodendrocytes have not been observed arising from RGLs in the adult SGZ (Fig. 2) (Bonaguidi et al. 2011), suggesting that they may represent a discrete precursor population with different embryonic origins. Nevertheless, OPCs are a dynamic cell population that can self-renew or differentiate into oligodendrocytes (Hughes et al. 2013). Remarkably, endogenous NG2 cells undergo extensive proliferation in response to various demyelinating conditions and act as a surveillance network on injury (Nait-Oumesmar et al. 1999; Menn et al. 2006; Hughes et al. 2013). Whether OPCs can also show expanded potential for cell generation under injury or other pathological conditions, as their potential role in the origin of glioma suggests, remains an interesting question (Liu et al. 2011).

Astroglia are a potential third precursor population. In the adult SGZ, these cells exhibit horizontal or bushy morphology and express GFAP, S100 β , and *Aldh1l1* (Seri et al. 2004).



Under basal conditions, astroglia are largely not considered to be a neuronal precursor cell type because they lack nestin expression and are remarkably quiescent (Steiner et al. 2004; Bonaguidi et al. 2011; Encinas et al. 2011). However, rare astroglia can be labeled with cell-cycle markers and share many immunohistological similarities with RGLs, including the neuronal determinate *Ascl1* (*Mash1*) (Seri et al. 2004; Kim et al. 2011). Interestingly, new astroglia are also generated from RGLs in both the SGZ and SVZ and may be considered as APCs (Levison and Goldman 1993; Bonaguidi et al. 2011; Tsai et al. 2012). Whether these cells can act as transient precursors before leaving the neurogenic niche or even remain as precursors for an extended period remains to be determined (Fig. 2). Interestingly, thrombospondin 4 (*Thbs4*) expression may distinguish APCs born in the adult SVZ from mature cortical astrocytes and is a necessary component of wound healing following ischemic conditions. In addition, *Thbs4* removal decreases astrogenesis with a concomitant increase in neurogenesis in the SVZ (Benner et al. 2013). Because APCs have only recently been characterized in the adult brain, it will be interesting to examine their lineage capacity under physiological and nonphysiological conditions (Table 1).

ORGANIZATION OF NSC DIVERSITY

A central question in NSC biology is how their behaviors arise and this has been researched at both system and cellular levels. Computational analysis indicates that stem cells can be generally categorized as deterministic or stochastic populations (Klein and Simons 2011). Stem cells with a determined fate, such as developing *Drosophila* neuroblasts (Isshiki et al. 2001), possess intrinsic programming and make the same resolute divisions over time. Tissue diversity therefore arises from the coexistence of intrinsically diverse stem cells. Alternatively, stem cells displaying stochastic behavior—more common in mammalian systems, such as the developing retina (He et al. 2012)—undergo several division choices in a seemingly random manner. Stochastic decision making allows for diversity

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among individual stem cells (Simons and Clevers 2011). Although both models permit stem-cell regulation at intrinsic and extrinsic levels, stochastic mechanisms confer flexibility within the stem-cell pool and endow compensatory behavior as has been observed in the postnatal SVZ (Kuo et al. 2006). Given the regionalization and patterning of the mammalian brain, examples of each model may persist from early development to serve discrete functions under physiological and pathological conditions in the adult (Chen et al. 2012).

Regional Organization of Neural Precursor Cells

A defining feature of the brain is its remarkable compartmentalization with different regions displaying specialized cell types, molecular identities, varied circuit connections, as well as contributions to different behaviors. The development of tissue specialization implies that adult NSC properties are, in part, established by neural patterning (Kriegstein and Alvarez-Buylla 2009). Evidence of this principle that developmental processes may be maintained into adult neurogenic zones arises from multiple observations. Adult radial precursors are derived from radial glia of similar embryonic region suggesting that adult precursors may be remnants from development (Merkle et al. 2004; Young et al. 2007). Moreover, similar transcript factor codes that specify neural precursors in the embryonic ventricular zone are conserved into the adult SVZ (Schuurmans and Guillemot 2002; Hack et al. 2005; Young et al. 2007), where the production of certain interneuron subtypes are enriched within SVZ domains (Fig. 3) (Merkle et al. 2007; Brill et al. 2009; Ihrle and Alvarez-Buylla 2011). Similarly, in the adult SVZ, astroglia arise from distinct postnatal spatial domains and undergo limited radial migration to maintain their positional fate (Tsai et al. 2012). In addition, NG2⁺ OPCs also display distinct spatial origins in the adult SVZ, preferentially arising from the dorsal SVZ compared with the lateral wall (Ortega et al. 2013). Collectively, these data suggest a model in which the SVZ displays mosaic organization not only among

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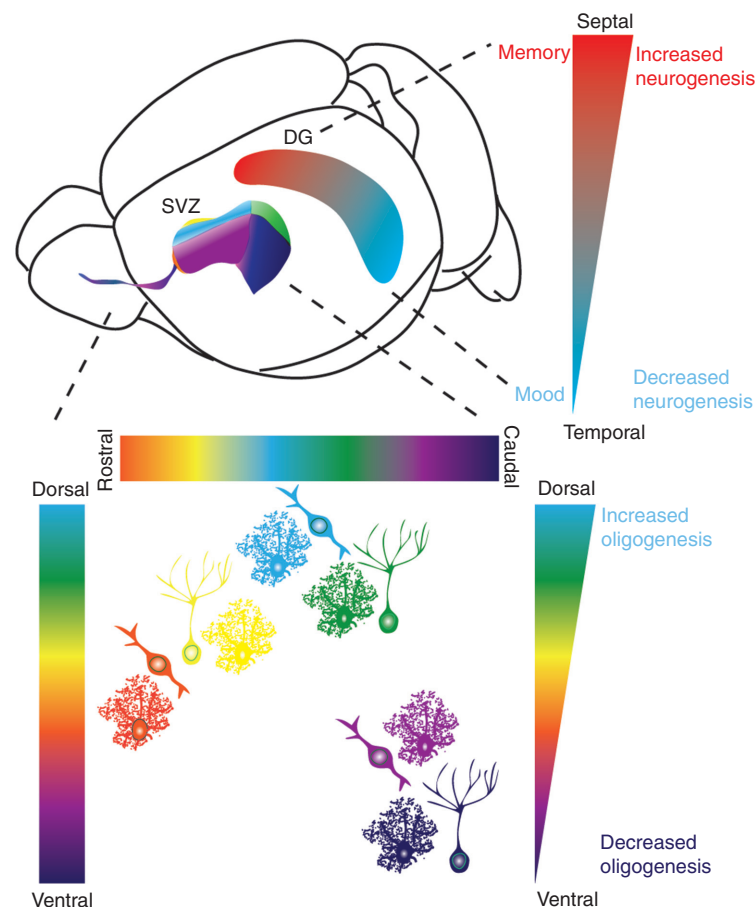


Figure 3. Regionalization of adult neural genesis in classical neurogenic zones. A schematic illustration of the mouse brain indicating the location of SVZ/RMS (subventricular zone/rostral migratory stream) and subgranular zone (SGZ) neurogenic niches. The different colors denote spatial heterogeneity of neural precursor cells. Precursor cells within the SVZ generate different interneuron subtypes, granule cells (GCs) or periglomerular (periglomerular cells [PGCs]; see Fig. 2). Studies have further divided these cell types in which precursor cells originating from the ventral SVZ generate deep granule cell (purple) and calbindin-expressing periglomerular cells (magenta). Precursor cells from the dorsal SVZ on the other hand, generate superficial granule cells (green) and TH⁺ periglomerular cells (light blue), whereas precursor cells in the medial parts of the ventricular wall generate calretinin-expressing superficial granule cells (yellow) and periglomerular cells (orange). Contrary to spatial heterogeneity and migration of long distances of neuron subtypes, astrocytes are generated throughout the SVZ and migrate more locally. Oligodendrogenesis is higher in the dorsal SVZ compared with the lateral and ventral SVZ. As schematically illustrated, the septal part of the dentate gyrus (DG) is associated with memory, has higher neurogenesis within the SGZ compared with the temporal part, which in turn has lower neurogenesis and is associated with mood regulation.

adult neurogenesis, but also astrogenesis and oligogenesis (Fig. 3).

Regionalization of NSCs in the adult SGZ is an emerging model. Differences in neural connectivity and behavior along the longitudinal (septotemporal) axis of the hippocampus are

well documented (Fanselow and Dong 2010; Wu et al. 2015). The septal (dorsal) portion is preferentially engaged in cognitive process of learning and memory, whereas the temporal (ventral) hippocampus is associated with motivational and emotional behavior. Moreover,

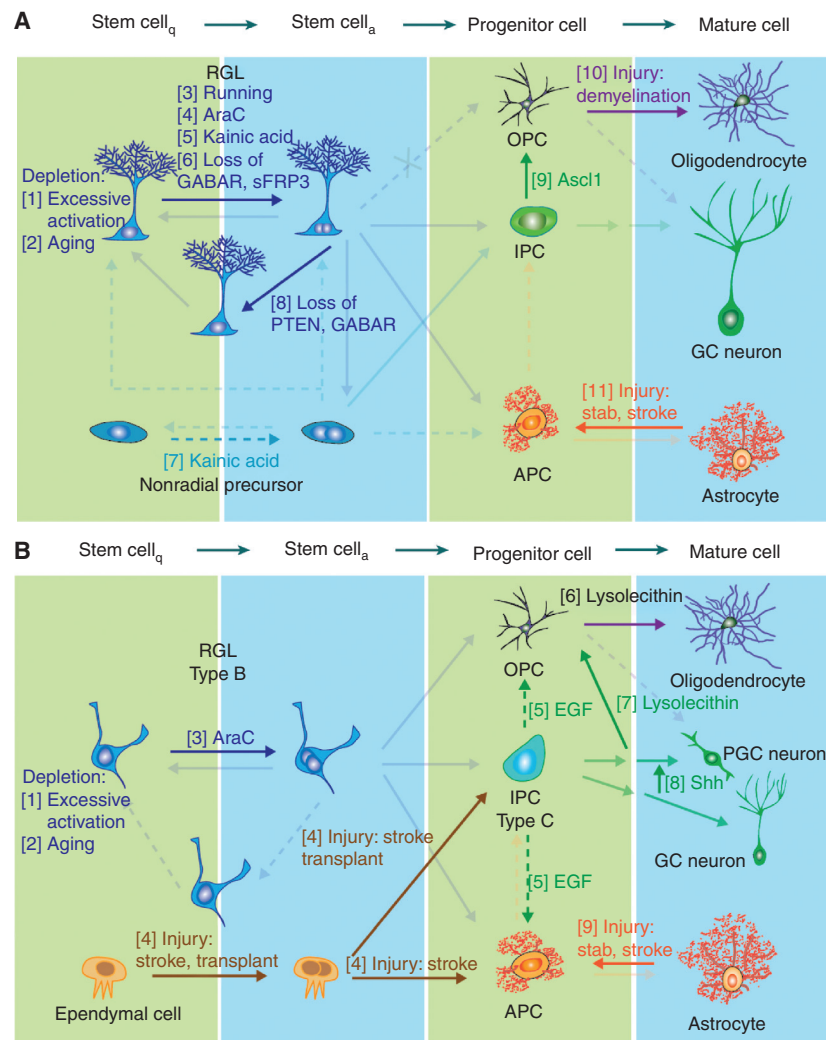


Figure 4. Regulation of neural precursor plasticity within the classical neurogenic zones. Schematic illustration of example factors and manipulations known to regulate cell genesis in the adult subgranular zone (SGZ) (A) and subventricular zone (SVZ) (B). Numbers denote examples known to affect lineage decisions at the stage indicated in the figure. (A) Stem-cell loss occurs when their proliferation is highly induced, such as through Notch and FoxO deletion (1) (Paik et al. 2009; Renault et al. 2009; Ehm et al. 2010; Imayoshi et al. 2010), or in aged mice (2) (Kuhn et al. 1996; Encinas et al. 2011; Villeda et al. 2011). Mobilization of quiescent radial glia-like cells (RGLs) occurs during voluntary running (3) (Kempermann et al. 1997; van Praag et al. 1999); brain injury, such as injection of the antimetabolic drug Ara-C (Seri et al. 2001) (4) or seizure-inducing Kainic acid (5) (Steiner et al. 2008; Jiruska et al. 2013). Molecular inhibitors of RGL activation include SFRP3 and GABA signaling (6) (Song et al. 2012; Jang et al. 2013). Kainic acid-induced seizures activate nonradial progenitor cells (7) (Lugert et al. 2010). Increasing Akt signaling or decreasing tonic GABA signaling alters the division mode of RGLs, fostering the symmetric fate (8) (Bonaguidi et al. 2011; Song et al. 2012). Ectopic expression of *Ascl1* changes the fate of intermediate progenitor cells (IPCs) to generate oligodendrocyte progenitor cells (OPCs) (9) (Jessberger et al. 2008) and demyelination injury induces OPC proliferation (10) (Nait-Oumesmar et al. 1999; Menn et al. 2006; Hughes et al. 2013). Stab wound, stroke and ischemic injuries activate astrocytes into reactive astroglia (11) (reviewed in Robel et al. 2011). (B) In the SVZ excessive activation (1) (Paik et al. 2009; Renault et al. 2009; Ehm et al. 2010; Imayoshi et al. 2010) and aging (2) (Kuhn et al. 1996; Molofsky et al. 2006; Villeda et al. 2011) leads to stem-cell loss. Ara-C promotes RGL cell-cycle entry (3) (Doetsch et al. 1999) and stroke injury activates the normally quiescent ependymal cells (4) (Johansson et al. 1999; Coskun et al. 2008; Carlen et al. 2009). Infusion of EGF increases production of astroglia and OPCs while reducing proliferation of IPCs (5) (Craig et al. 1996; Kuhn et al. 1997). Demyelination injury increases OPC proliferation (6) and doublecortin (DCX)⁺ neural progenitors to switch fate into OPCs (7) (Nait-Oumesmar et al. 1999; Menn et al. 2006; Jablonska et al. 2010; Hughes et al. 2013). Manipulation of the Sonic hedgehog (SHH) signaling pathway can change the fate of a subset of neural progenitors from granule cell (GC) neurons to periglomerular cell (PGC) neurons (8) (Ihrle et al. 2011). Stab wound, stroke, and ischemic injuries activate astrocytes into reactive astroglia (9) (reviewed in Robel et al. 2011).

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detailed gene expression supports a segregation of the hippocampus and the SGZ in particular into septal, temporal, and intermediate zones (Fanselow and Dong 2010). Interestingly, adult neurogenesis is not uniform throughout the SGZ, with fewer neural precursors in the temporal DG (Fig. 3) (Tashiro et al. 2007; Jinno 2011a). Meanwhile, the tempo of neurogenesis is slower in the temporal region compared with the septal region and behaves differently to neuronal activity and aging (Jinno 2011b; Piatti et al. 2011; Snyder et al. 2012). Dentate formation arises from distinct developmental locations within the septotemporal plane (Altman and Bayer 1990) and from the temporal pole of the ventral hippocampus (Li et al. 2013) implicating mosaic spatial patterning as a putative mechanism underlying adult SGZ neural precursor diversity.

The relative contribution of intrinsic and extrinsic cues in conferring divergent adult neural precursor behavior is starting to emerge. When neural precursors are put in cell culture or transplanted across SVZ regions, interneuron subtype generation remains the same as the region of origin (Merkle et al. 2007). These observations are consistent with a timed program encoded, at least in part, by intrinsic factors (Qian et al. 2000). Meanwhile, changes in extrinsic cues are also instructive of cell fate decisions. Ectopic activation of Sonic hedgehog signaling (Shh) is sufficient to respecify dorsally derived interneuron progeny to more ventrally defined fates (Ihrie et al. 2011). Additional examples of cell fate alteration by extrinsic regulation occur in the SGZ. RGLs that normally undergo asymmetric neuronal and astroglial cell division instead preferentially divide symmetrically on increasing Akt signaling or decreasing tonic GABA signaling (Song et al. 2012). Therefore, it appears that intrinsic cues prime cell fate, whereas extrinsic cues can modulate that program, a model consistent with stochastic asymmetry regulation of stem-cell behavior.

Local Organization within Neurogenic Regions

Further local organization of neural precursor cells may allow for diverse individual stem-cell

behavior and coexistence of neural precursor populations. First suggested within the hematopoietic system (Schofield 1978), the niche is a microenvironment that houses and supports stem-cell behavior and largely restricts high levels of adult mammalian neurogenesis to the SVZ and SGZ under basal conditions (Riquelme et al. 2008; Fuentealba et al. 2012). The niche constituents can also promote neural precursor diversity (Fig. 3). Vasculature, astroglia, microglia, ependymal cells, and neural precursor subtypes create niche microdomains. Dynamic access to these components can influence NSC decisions, such as radial cell contact with vasculature mediating a switch between quiescence and proliferation (Palmer et al. 2000; Mirzadeh et al. 2008; Shen et al. 2008; Tavazoie et al. 2008). These associations could also serve as an attractant for more proliferative precursors and potentially a hotspot for discrete mitotic stem-cell populations. In addition, cells present throughout the adult brain can have different functions within the niche. Astroglia within the adult SVZ and SGZ have the unique capability to promote neurogenic fate specification, neuroblast migration, and survival (Lim and Alvarez-Buylla 1999; Song et al. 2002; Barkho et al. 2006; Kaneko et al. 2010; Platel et al. 2010). As astroglia remain local to their region of origin (Fig. 3) (Tsai et al. 2012), it remains unclear whether subpopulations of astroglia within the niche exhibit specialized functions and whether such heterogeneity could further diversify neural precursor cells (Zhang and Barres 2010).

The role of cellular components within the niche can also change depending on the environment. For instance, microglia normally phagocytose and remove dying IPCs (Sierra et al. 2010) and can, within different contexts, have either a beneficial or detrimental effect on neural precursor development (Ekdahl et al. 2009). Likewise, the composition of the CSF secretome is altered on various physiological and pathological conditions (reviewed in Falcao et al. 2012; Zappaterra and Lehtinen 2012). Given that many extrinsic cues, including growth factors, neurotransmitters, cytokines, and hormones, change during injury (reviewed in Zhao et al. 2008), the state of the local environment itself



can determine neural precursor diversity. An important function of these niche factors under physiological conditions is to maintain NSC quiescence, as excessive NSC activation can lead to impaired NSC maintenance (Paik et al. 2009; Renault et al. 2009; Ehm et al. 2010; Imayoshi et al. 2010). The neuronal circuit itself may act in this manner where interneurons and mature dentate granule neurons in the SGZ repress RGL activation through tonic GABA signaling and the WNT antagonist secreted frizzled related protein 3 (sFRP3), respectively (Song et al. 2012; Jang et al. 2013). These mechanisms act far reaching (GABA) or locally (sFRP3) to fine-tune stem-cell response to environmental change. How neural network activity controls stem-cell behavior directly or indirectly through the niche is an emerging area of research.

Moreover, coordination among the cellular constituents within the niche likely also contributes to homeostasis. For example, elimination of IPCs/neuroblasts causes activation of the quiescent radial cells, which rapidly repopulate IPCs (Doetsch et al. 1999; Seri et al. 2001). The feedback mechanisms from IPCs/neuroblasts to radial precursor cells could include GABA or Notch signaling (Liu et al. 2005; Basak et al. 2012) and suggests that NSC descendants are active constituents of the adult niche. As IPCs and neuroblasts are transient in nature, their waxing and waning provides another layer of local dynamics within the niche. Ultimately, these principles provide a collective context for how local cellular cues can promote individual NSC diversity.

REGULATION OF ADULT NEURAL PRECURSOR CELL PLASTICITY

A hallmark of neurogenesis in the adult brain is its sensitivity to external changes. Environmental or pathological alterations can have stage-specific effects on precursor proliferation, cell migration, cell death, neuronal maturation, and circuit integration (Zhao et al. 2008). Less well studied are the roles of extrinsic cues on specific neural precursor types and how precursor plasticity adjusts to meet shifting tissue de-

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mands. An emerging general theme across multiple somatic-stem-cell compartments is the coexistence of precursor cell populations for differential responses to physiological and injury conditions (Leung et al. 2007; Wilson et al. 2008; Li and Clevers 2010; Lu et al. 2012; Mascré et al. 2012; Buczaczk et al. 2013). Technical advances in neural precursor cell-type identification, lineage tracing, and cell-type-specific manipulation provide an opportunity to reveal the cellular and molecular substrates of external changes beyond describing general cell proliferation (Table 1). Indeed, recent studies suggest that diverse neural precursor cells also exhibit differential responses to the environment under physiological or pathological conditions (Lugert et al. 2010; DeCarolis et al. 2013).

Physiological Regulation of Neural Precursor Plasticity

Adult neurogenesis is influenced by many physiological stimuli including exercise, environmental enrichment (EE), social isolation, aging, and learning paradigms (reviewed in Ming and Song 2011). The exact neural precursor targets of these experiences are largely unknown, but general principles are beginning to emerge. From studies of the SGZ, it appears that experience can dictate the properties of neural precursors. For example, running either alone or as a component of EE has been long known to promote neurogenesis (Kempermann et al. 1997; van Praag et al. 1999; Steiner et al. 2008; Kobilov et al. 2011; Mustroph et al. 2012). Although the hippocampal IPC cell populations increase after running, this may be a result of mobilizing quiescent RGLs (Kronenberg et al. 2003; Lugert et al. 2010). Curiously, the effect of running is observed in $Hes5^{+}$ and not $Nestin::GFP^{+}$ RGLs suggesting functional heterogeneity within the radial cell population (Kronenberg et al. 2003; Steiner et al. 2008; Lugert et al. 2010). It remains unclear whether running also facilitates a fate switch of RGLs from glial to neuronal production. Appealing candidates to mediate the effect include Wnt, bone morphogenetic protein (BMP), and brain-derived neurotrophic factor (BDNF) signaling (Rossi et al. 2006; Gobeske

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et al. 2009; Jang et al. 2013). Conversely, social stress or glucocorticoids have profound negative effects on neurogenesis and cell genesis has a role in efficient stress recovery (Dranovsky and Hen 2006; Stranahan et al. 2006; Snyder et al. 2011). In addition to decreasing neurogenesis, social isolation in juvenile mice promotes expansion of the adult RGL population (Dranovsky et al. 2011). This expansion occurs through increased RGL activation and a switch from asymmetric to symmetric divisions via tonic GABA signaling (Song et al. 2012). Together, results from exercise and social isolation experiments suggest another emerging principle: environmental. Environmental factors can have positive and negative effects on neurogenesis, partially through targeting the NSC populations.

Aging studies provide another platform to investigate mechanisms regulating neural precursor behavior because neurogenesis strongly declines as a rodent ages (Kuhn et al. 1996). This decline in both SGZ and SVZ neurogenesis is thought to result from the loss of stem cells, including the radial precursors (Molofsky et al. 2006; Encinas et al. 2011; Villeda et al. 2011). Interestingly, RGLs in the SGZ do not appear to change their proliferation activity with age (Encinas et al. 2011; but see Hattiangady and Shetty 2008). Instead, changes in the niche may underlie the decrease in neural precursor number. One experimental approach used to test niche contribution is parabiosis, in which the blood systems of two mice are surgically combined. Remarkably, old mice paired with young mice show an increase in the number of immature neurons in the DG, whereas the opposite is seen in the young mice in the same pair (Conboy et al. 2013). This result is associated with the age-dependent increase in chemokines, such as eotaxin (Villeda et al. 2011). In addition, physiological activity, such as EE and running, also increase proliferation in aged mice, although the mechanisms are unclear (Kempermann et al. 1998). These data suggest that a decline in neural precursor number with age may be partially ameliorated by altering the aged cellular niche, and it will be important to understand the cellular and molecular niche involved in aging.

Pathological Regulation of Precursor Plasticity

Neurogenesis in the adult brain has also been investigated in response to numerous pathological conditions including seizures, cytosine- β -D-arabinofuranoside (Ara-C) injury, stab wound injury, traumatic brain injury (TBI), stroke, psychological disorders, and neurodegenerative diseases. A common theme among injury studies is that, unlike lower vertebrates, such as fish and amphibians, mammals do not significantly repair damaged neuronal tissue (Berg et al. 2010; Zupanc and Sirbulescu 2011; Illis 2012). Repair in mammals appears limited by deficits in neuronal maturation and survival, (Kokaia et al. 2006; Kazanis et al. 2013; although see Magavi et al. 2000) and can be initially robust in both the SVZ and SGZ (Liu et al. 1998; Arvidsson et al. 2002; Parent et al. 2002). Importantly, production of astroglia and neuroblasts contributes to minimizing initial damage during injury, suggesting an essential role of cell genesis in buffering the brain from a more severe insult (Jin et al. 2010; Benner et al. 2013). Increased proliferation is observed on many injury models, even in areas that are not neurogenic under physiological conditions (Magavi et al. 2000; Chen et al. 2004). At present, the precise cellular and molecular events mediating diverse precursor behavior under pathological conditions have not been explored extensively.

A general premise to describe the neurogenic response to injury may be that the severity of an injury or acute pathological condition determines the magnitude of the initial neural precursor response. In less severe conditions, such as AraC and epilepsy, NSCs are recruited in attempts to promote homeostasis. For example, the antimitotic drug Ara-C has been used to study regeneration of neurogenic populations in the adult brain. Because Ara-C treatment primarily kills the proliferating IPCs, quiescent radial precursor cells were identified as cellular origins of neural precursor recovery in both the SVZ and SGZ (Doetsch et al. 1999; Seri et al. 2001). Intriguingly, RGLs labeled using different Cre drivers selectively proliferate on antimitotic treatment indicating that neural precursors with similar identity can display



functional heterogeneity (DeCarolis et al. 2013). In addition, epilepsy and seizures are known to affect many aspect of adult neurogenesis, including proliferation of neural precursor cells and the migration pattern of immature neurons (Parent et al. 1997). Injection of seizure-inducing substances, such as kainic acid and pilocarpine have a propoliferative effect on radial cells, IPCs, and neuroblasts (Steiner et al. 2008; Jiruska et al. 2013). Interestingly, kainic acid increases the size of the nonradial, but not radial, precursor pool, suggesting that neural precursors of different identities can display diverse response to a common injury (Lugert et al. 2010). Yet an unanswered question is whether radial and nonradial cells change their basal stem-cell properties on injury.

Greater cell plasticity appears to occur on more severe injury, such as TBI and stroke. For instance, ependymal cells have been observed to change their morphology radial glial-like shape after stroke (Zhang et al. 2007). Normally, quiescent ependymal cells activate and generate both astrocytes and neuroblasts suggesting that they might act as a reservoir stem-cell population (Zhang et al. 2007; Carlen et al. 2009). Astrocytes represent a second cell population displaying enhanced cell plasticity after injury. In response to a stab wound or cerebral ischemia, astrocytes can become reactive and display NSC markers including nestin, vimentin, and BLBP (Robel et al. 2011). Reactive astroglia remain restricted to gliogenesis in vivo, yet become primed to behave as NSCs on exposure to growth factors in vitro, whereas astrocytes under basal conditions do not (Buffo et al. 2008; Sirko et al. 2013). Astrocytes are a heterogeneous population of cells, and it is not known whether diverse populations of astrocytes react differently to the various types of injury (Zhang and Barres 2010). In vivo imaging has revealed that astrocytes found in close proximity to vasculature proliferate in response to a stab wound injury (Bardehle et al. 2013). Perhaps the greatest example of neural precursor injury-induced plasticity in the adult mammalian brain is the fate switch of DCX⁺ neuroblasts in the rostral migratory stream (RMS). Lysolecithin-induced demyelination of corpus callosum redirects

DCX⁺ progenitors from neuronal to glial fates to generate new oligodendrocytes in the corpus callosum (Jablonska et al. 2010). However, changes in neural precursor developmental potential do not appear to be sufficient to significantly repair damaged neuronal tissue and may require additional plasticity, such as in vivo reprogramming (Jessberger et al. 2008; Niu et al. 2013).

CONCLUSIONS

Diverse origins of cell genesis is an emerging principle in many adult somatic tissues, which can display differences among precursor identity, potential, and regulation (Alvarez-Buylla et al. 2008; Li and Clevers 2010). A common theme among different somatic systems is the coexistence of multiple precursor cells within a tissue to satisfy particular local demands. Neural stem and progenitor cells take multiple forms in the adult brain: from radial to nonradial precursor cells; lineage-restricted neuronal, astroglial, and oligodendroglial progenitors; and ependymal cells. Further, diversity may also exist within neural precursors of similar cellular identity. Recent data indicates that cellular hierarchy exists among cell types under physiological conditions. Meanwhile, neural precursor competence is partially conferred through cellular interactions, both locally within the niche and patterning across various spatial domains. Finally, dormant plasticity among diverse cell types can be unleashed during pathological response or by genetic alterations. These studies collectively suggest that a diverse range of neural precursors are organized to collectively meet the dynamic needs of specialized remodeling functions. Future investigations should take into account the diversity of cell types to understand how the adult brain undergoes dynamic remodeling to mediate an individual's interactions with the outside world.

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