



Engineering of Adult Neurogenesis and Gliogenesis

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Neural stem/progenitor cells (NSPCs) retain their ability to generate newborn neurons throughout life in the mammalian brain. Here, we describe how recently developed virus- and transgenesis-based techniques will help us (1) to understand the functional effects of neurogenesis in health and disease, (2) to design novel approaches to harness the potential for NSPC-associated endogenous repair, and (3) to induce the generation of neurons outside the main neurogenic niches in the adult brain.

The fact that neural stem/progenitor cells (NSPCs) persist in the adult brain, challenged previously held concepts of brain function in health and disease and opened novel possibilities to treat neurological disease by harnessing the NSPC-associated potential for brain repair (Gage 2000; Alvarez-Buylla and Lim 2004; Gage and Temple 2013). However, over the last decade, it has become clear that targeting NSPCs for repair requires the detailed understanding of the biology of NSPCs and the underlying cellular and molecular mechanisms that govern the distinct steps of neural development from dividing NSPCs to integrating neurons (Ming and Song 2011). Based on findings obtained by aiming to understand the basic mechanisms of adult neurogenesis, a number of technical advancements have substantially increased our toolbox to manipulate NSPC

activity in vivo and/or subsequent neuronal integration in the adult brain. Furthermore, our understanding of the mechanisms regulating neural differentiation has substantially improved over the last years making it now possible to redirect the fate of NSPC-derived, newborn cells in the adult brain (Hack et al. 2005; Jessberger et al. 2008; Brill et al. 2009). Besides these novel tools that target NSPCs in their endogenous niches, basic understanding of the underlying neuronal differentiation of NSPCs in the adult brain enabled novel approaches to induce the generation of neurons also in nonneurogenic regions throughout the adult brain (Jessberger and Gage 2009; Amamoto and Arlotta 2014; Heinrich et al. 2015). In addition, novel cellular sources such as NSPCs derived from induced pluripotent stem cells (iPSCs), will enable us to study mechanisms of NSPC activity and neuro-

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nal differentiation in healthy and diseased human tissue and may also represent a novel source for exogenous cell-replacement strategies (Yu et al. 2013, 2014).

MANIPULATING LEVELS OF ADULT NEUROGENESIS

Over the last decade, a plethora of approaches have been developed to reduce or to enhance levels of adult neurogenesis with the aim to test experimentally the functional contribution of newborn neurons to adult behavior (Deng et al. 2010). Initially, cytostatic drugs or ionizing irradiation impairing cell proliferation were used (and are still in use, e.g., Shors et al. 2001;

Snyder et al. 2005; Saxe et al. 2006). However, given that the whole organism depends on proper levels of cell proliferation and the fact that irradiation also injures postmitotic cells, these approaches are likely to be associated with unwanted side effects that are bound to complicate the interpretation of behavioral tests (Fig. 1). Thus, the focus over the years has been to develop genetic or virus-based strategies to manipulate levels of neurogenesis. The most common principle used is to either kill proliferating NSPCs or to impair (or in the case of enhancing neurogenesis to increase) the survival of newborn neurons (Deng et al. 2010). For example, a number of different promoters were transgenically modified to drive thymidine kinase expres-

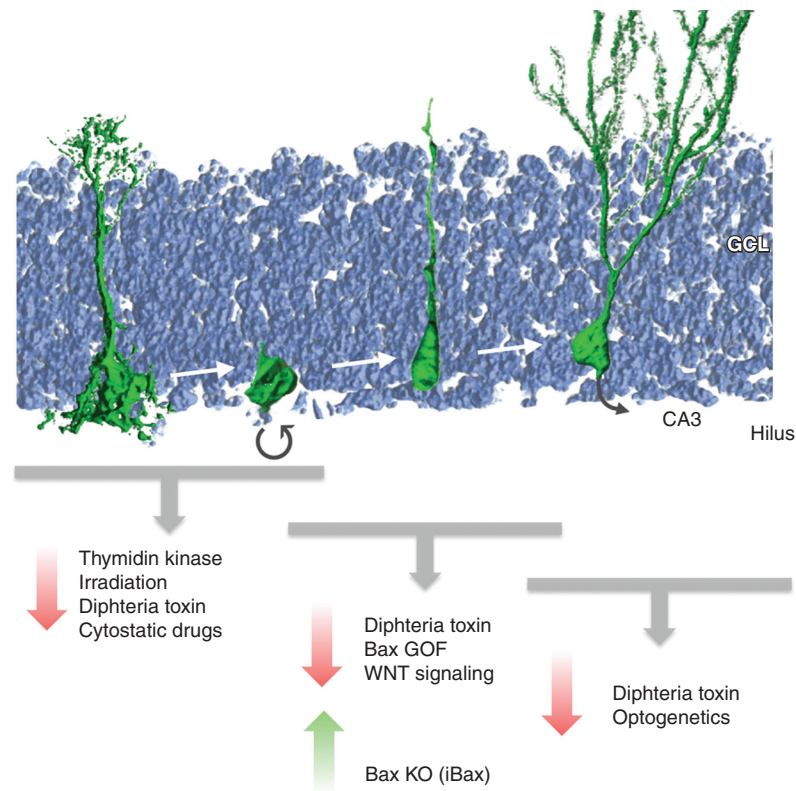


Figure 1. Approaches to engineer levels of adult neurogenesis. Depicted are the developmental steps from largely quiescent radial glia-like neural stem cells through proliferative nonradial neural stem/progenitor cells (NSPCs) generating newborn neurons that mature and functionally integrate over the course of several weeks in the rodent hippocampus. Experimental strategies to reduce or to enhance levels act on distinct stages from the dividing NSPC to the newborn neuron. GCL, granule cell layer; CA3, cornu ammonis area 3. (Figure courtesy of Dr. S.M.G. Braun, Brain Research Institute, Zurich, Switzerland.)



sion in adult NSPCs that on gancyclovir (GCV) administration results in the generation of a toxic metabolite that selectively kills dividing cells (Fig. 1) (Garcia et al. 2004; Saxe et al. 2006, 2007; Deng et al. 2009). Most laboratories used the regulatory elements of the *nestin* gene (coding for an intermediate filament that is highly enriched in NSPCs) or the promoter sequences driving glial fibrillary acidic protein (GFAP) (another intermediate filament that is expressed in NSPCs but also in a large set of classical astrocytes throughout the brain).

Using these approaches, a substantial (and even complete) loss of neurogenesis can be achieved that is either chronic or transient (Garcia et al. 2004; Deng et al. 2009). However, one potential disadvantage of using the *nestin* or *gfap* promoter is that these regulatory sequences may mediate also active sites of transgene transcription in other somatic stem-cell compartments, such as in the intestines or skin stem-cell niches (Bush et al. 1998; Mignone et al. 2007). The same concern holds true for *nestin*- or *gfap*-promoter-mediated deletion of survival genes resulting in the loss of newborn cells (Dupret et al. 2008). An elegant approach to restrict cell death to the neuronal progeny of *nestin*-expressing cells has been achieved by selectively expressing diphtheria toxin A (DTA) (killing cells on expression) in *nestin* expression-fate-mapped cells that have an active neuron-specific enolase (NSE) promoter (Imayoshi et al. 2006, 2008).

In addition to transgenesis-based approaches, viral vectors have been successfully used to reduce levels of neurogenesis, for example, by impairing proper WNT signaling (Fig. 1) (Lie et al. 2005; Jessberger et al. 2009). Despite the advantages of high temporal and spatial control (and the usability of rats which is advantageous for a number of behavioral experiments), there is the apparent risk for off-target effects if the targeted signaling pathways are not completely specific for NSPC/neuronal differentiation (e.g., WNT signaling appears to be also involved in synaptic function/remodeling of mature neurons and not only for the generation of new neurons) (Cerpa et al. 2011). Very recently, optogenetic tools have been developed allowing

the selective expression of hyperpolarizing (or depolarizing) opsins in newborn neurons in the adult brain (Gu et al. 2012). This represents a novel and exciting tool to characterize the functional contribution of newborn neurons (Fig. 1). However, it is also clear that a circuitry certainly responds “differently” to the acute silencing of neuronal populations compared with the situation in which this population is “missing,” which may be the more relevant situation when it comes to experimentally testing the role of new neurons in the disease context. An interesting alternative to optogenetics is the use of designer receptors exclusively activated by designer drugs (DREADDs) (Sternson and Roth 2014). For instance, virus-mediated expression of a designer G protein-coupled receptor (hM3Dq) selectively activated by the pharmacologically inert, orally bioavailable drug clozapine-*N*-oxide (CNO) allowed for assessing the ability of newborn dentate granule neurons to activate parvalbumin-positive interneurons (Temprana et al. 2015).

In contrast, to the progress made over the last decades to reduce neurogenesis, there are very few tools available to enhance neurogenesis. One elegant approach has been to delete proapoptotic genes selectively in adult-born cells, leading to a doubling of newborn neurons (Fig. 1) (Sahay et al. 2011). Excitingly, these transgenic mice (called iBaxNes) allowed for testing of potential gain-of-function effects of increased neurogenesis that could be indeed experimentally observed in the behavioral context of dentate gyrus-mediated pattern separation (Sahay et al. 2011). However, a potential concern with this approach may be that deletion of *bax* increases survival of cells that would be otherwise sorted out—by poorly understood, activity-dependent selection criteria (Tashiro et al. 2007). Thus, it is possible that cells are rescued that do not meet certain criteria for proper integration. Thus, future approaches may target rather NSPC proliferation, allowing for subsequent proper functional selection with regard to survival and functional integration.

Be that as it may, over the last decade, immense progress has been made to develop novel tools allowing for manipulating levels of neuro-

genesis and to test the functional contribution of the resulting new neurons for adult brain behavior. However, the specificity and temporal control of loss- and gain-of-function will need to be further optimized to improve our understanding how new neurons shape the functional connectivity of the adult dentate circuitry in health and disease.

ENABLING NEUROGENESIS OUTSIDE ITS NATURAL NICHES

Neurogenesis is restricted to few areas within the adult brain (Zhao et al. 2006), but an unfulfilled clinical need for new neurons arises in many more brain regions in the context of human disease or trauma. Although transplantation is potentially a viable approach to address this need (Lindvall 2013), here, we discuss recent work focusing on the possibility of converting endogenous nonneuronal cells into induced neurons (reviewed in Heinrich et al. 2015). Likewise, a similar approach may serve to regenerate oligodendroglia for those disease scenarios in which this cell type degenerates.

Reprogramming of Central Nervous System (CNS) Glia into Induced Neurons

One of the earliest studies in this direction was work by Kondo and Raff showing that sequential exposure to specific extracellular signals could reprogram oligodendrocyte precursor cells (OPCs) to multipotential CNS (central nervous system) stem cells (Kondo and Raff 2000). It was recognized that this reprogramming is accompanied by chromatin remodeling resulting in the reactivation of the *sox2* gene (Kondo and Raff 2004), a key transcription factor in neural stem cells, but also of pivotal importance during reprogramming toward pluripotency (Sarkar and Hochedlinger 2013).

Based on the recognition that neural stem cells both during development and in the adult have a radial glia identity sharing many molecular and cellular features with parenchymal astrocytes (Doetsch et al. 1999; Malatesta et al. 2000; reviewed in Kriegstein and Alvarez-Buylla 2009), Magdalena Götz and colleagues

hypothesized that astroglia may perhaps retain some potential of generating neurons with appropriate transcriptional cues provided that elicit a neuronal program (Costa et al. 2010). Having shown that *pax6* is required for neurogenesis from radial glia of the cerebral cortex, *pax6* was also the first transcription factor gene tested for its capacity to reprogram early postnatal mouse astroglia into induced neurons in vitro. Indeed, forced expression of *pax6* elicited a neurogenic response in these cells (Heins et al. 2002). Using this experimental paradigm, several studies then examined the effect of other neurogenic transcription factors, such as *neurog2*, *ascl1*, *dlx2*, and *neurod1* (Berninger et al. 2007; Heinrich et al. 2010; Guo et al. 2014), well known for their important roles during embryonic and adult neurogenesis and thus qualifying them as potential reprogramming factors (Amamoto and Arlotta 2014). It turned out that retrovirus-mediated *neurog2* expression could induce in early mouse astroglia the expression of a cascade of transcription factors (*Tbr2* and *Tbr1*) that hallmark glutamatergic neurogenesis (Berninger et al. 2007) and convert these into fully functional synapse-forming neurons of glutamatergic neuron identity (Heinrich et al. 2010). This very same astroglia could be reprogrammed into neurons of the GABAergic lineage by forced expression of *ascl1* or *dlx2*. This data thus show that a single transcription factor suffices to reprogram early postnatal astrocytes into functional neurons. But recent work indicates that even in culture, astrocytes quickly lose their permissiveness to single transcription factor reprogramming (Maserdotti et al. 2015). Five days after exposure to differentiating conditions, astrocytes derived from mouse postnatal cortex no longer reprogram following induced *Neurog2* expression. This loss of reprogramming permissiveness appears to be mediated by REST (RE-1 silencing transcription factor), which prevents *Neurog2* of binding to its target sites. However, failure of reprogramming can be bypassed by directly activating some of the *Neurog2* targets, such as *NeuroD4*. This data then support the notion that reprogramming permissiveness is strongly dependent on the epigenetic state of the cells to

be lineage-converted. Likewise, the highly different permissiveness of fibroblasts and keratinocytes to *Ascl1*-induced reprogramming appear to be consequence of their specific and distinct chromatin signature at *Ascl1* targets (Wapinski et al. 2013). After all, this leads to the, probably not very surprising, conclusion that cells that have undergone considerable epigenetic maturation will prove hardest to reprogram (Fig. 2).

Thus, it is encouraging that the adult human brain does contain cells that appear to be amenable to reprogramming into induced neurons. Aiming at converting cells isolated from human brain biopsies, we identified cells expressing a battery of pericytic markers in vitro that can be converted following combined expression of *sox2* and *ascl1* (Karow et al. 2012). Here, as in maturing astrocytes, the proneural gene (*ascl1*) alone fails to reprogram, suggesting that the effect of *sox2* coexpression may consist in rendering *Ascl1* targets more accessible (see discussion below). Interestingly, pericytes emerge as new players in CNS scar formation, a

response that requires deattachment from microvessels to which these cells are associated under healthy conditions (Goritz et al. 2011). In addition to glial cells, such “reactive” pericytes may represent an interesting target for in vivo reprogramming (Karow et al. 2012).

In Vivo Reprogramming in the Adult Brain

Attempts of direct lineage reprogramming of glial cells into neurons have now been carried on to the in vivo level in the cerebral cortex, striatum, and spinal cord (Buffo et al. 2005; Grande et al. 2013; Guo et al. 2014; Heinrich et al. 2014). Already, an early study from the Götz laboratory addressed the possibility of converting proliferative cells in the injured cerebral cortex into induced neurons and found that expression of *Pax6* or repression of *Olig2* transcriptional targets resulted in the transient emergence of doublecortin (DCX)-positive cells (Buffo et al. 2005). More recently, it was found that in the injured cerebral cortex in vivo fate conversion of nonneuronal cells by

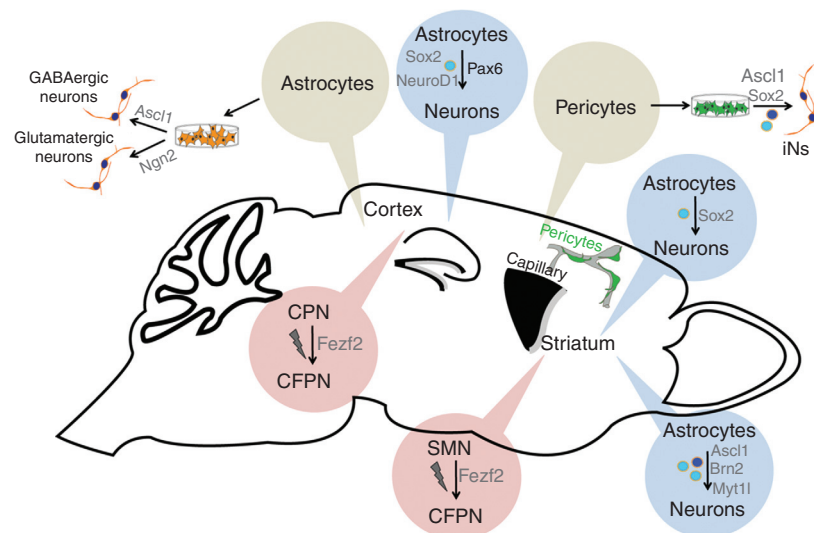


Figure 2. Engineering neurogenesis outside the classical neurogenic niches. Depicted are the different in vitro and in vivo approaches scrutinized so far for reprogramming nonneuronal cells into induced neurons. Likewise, highlighted are in vivo experiments assessing the possibility of converting distinct types of neuronal progenitors or neurons into another via in utero electroporation. MSN, medium spiny neuron; CPN, callosal projection neuron; CFPN, corticofugal projection neuron. (Figure courtesy of Dr. M. Karow, Institute of Physiology, LMU Munich, Germany.)



neurogenic fate determinants can be influenced by simultaneous delivery of growth factors (Grande et al. 2013). However, in both studies, the precise cellular source of the induced neuronal cells remained unknown.

Previously, pioneering work from the Wernig laboratory had shown that three factors, *ascl1*, *brn2*, and *myt1l*, can reprogram fibroblasts into induced neurons (Vierbuchen et al. 2010). Torper et al. (2013) now delivered this factor triad into the adult striatum via cre-inducible lentiviral vectors. Indeed, they found NeuN-positive cells that were also reporter-positive (an indicator of recombination) in a transgenic mouse line expressing *cre* under control of regulatory elements of the *gfap* gene (*gfap-cre*), providing evidence for direct astrocyte-to-neuron conversion in situ (Fig. 2). With a caveat for some leakiness of *gfap* promoter activity, this study provided compelling evidence for an astroglial origin of the reprogrammed cells. However, it remains unknown whether the virus-targeted glial cells were reactive (as a response to the virus injection) or nonreactive parenchymal astrocytes. In fact, recent work from the Frisén laboratory has suggested that striatal astrocytes possess a latent neurogenic program, under healthy conditions restrained by Notch signaling, but unleashed following injury (Magnusson et al. 2014).

In contrast, a recent study from the Zhang laboratory has provided evidence that forced expression of *sox2* alone can drive striatal astrocytes (as evidenced by several lines of genetic fate-mapping experiments) toward early stages of neurogenesis (Fig. 2) (Niu et al. 2013). Surprisingly, these data suggest that the reprogrammed cells do not acquire a neural-stem-cell identity (as evidenced by absence of nestin expression at any stage), as one might have expected at first glance, given the known role of *sox2* in neural stem cells. Nonetheless, these cells incorporated bromodeoxyuridine (BrdU) and apparently proliferated as DCX-positive neuroblasts. Although these cells fail to acquire a mature neuronal state when left alone, they develop into postmitotic neurons with remarkable electrophysiological properties when exposed to the neurotrophin brain-derived neuro-

trophic factor (BDNF) and the natural bone morphogenetic protein (BMP) inhibitor noggin or treated with a histone deacetylase inhibitor. Of note, we found that forced *sox2* expression can also convert reactive glia in the cerebral cortex of mice following stab wound injury (Heinrich et al. 2014). Remarkably, this effect was strictly dependent on prior injury suggesting that the lesioning rendered the cells more permissive to glia-to-neuron conversion. Moreover, to our surprise, genetic fate mapping revealed that the cell population undergoing fate conversion was not the astrocytic compartment but NG2 glia. Despite the fact that the physiological role of these cells in grey matter is still largely enigmatic, they can function as oligodendrocyte progenitor cells (Dimou and Gotz 2014). Hence, it might be possible that it is their status as progenitors that renders these cells more permissive to lineage conversion. Finally, this work confirmed the neurogenic effect of Sox2 first reported by Niu et al. (2013). Although this is somewhat puzzling, given the classical role of Sox2 as a stem-cell maintenance factor, recent work also shows that it primes the epigenetic landscape in adult hippocampal progenitors for expression of neurogenic genes (Amador-Arjona et al. 2015). Perhaps Sox2 does the same work also in the above-described reprogramming scenarios.

Finally, the Chen laboratory made the very surprising observation that retrovirus-mediated NeuroD1 alone was sufficient to drive reactive astroglia and NG2 glia (using *gfap* and *NG2* promoter-controlled retroviral vectors) of the adult mouse cerebral cortex with unprecedented efficiency toward neurogenesis in the absence of additional injury, besides that caused by virus injection, and the induced neurons fairly rapidly developed remarkable electrophysiological properties (Guo et al. 2014). Given the emerging concept that glial cells acquire stem-cell-like properties following injury (Dimou and Gotz 2014), it will be of obvious interest to assess the contribution of the virus injection-induced injury, if any, to the reprogramming process.

An altogether different approach developed by the Arlotta laboratory aims at reprogram-

ming one neuron type into another. Rouaux and Arlotta (2010) showed that striatal progenitors that normally give rise to medium spiny neurons can be reprogrammed into pyramidal-like neurons on forced expression of the transcription factor *Fezf2* (Fig. 2). Remarkably, these neurons can project to subcortical targets, including the spinal cord, consistent with the role of *Fezf2* in imposing a corticofugal projection neuron identity. This *Fezf2* effect extends into early postmitotic life, as illustrated by the fact that callosal projection neurons can still be converted into corticofugal neurons during the first postnatal days, but lose such potential on closure of a critical, molecularly not-yet-defined window of nuclear plasticity (Rouaux and Arlotta 2013). A key question will be how to reopen this window of plasticity, evidently imposed by epigenetic modification in the course of cell differentiation. One exciting possibility may be to render these cells capable of undergoing cell proliferation. Although this sounds a priori violating the fact that neurons are per definition postmitotic entities, studies in the adult heart have shown that bona fide postmitotic cells can be enticed to enter cell cycle without losing their cellular specification even in vivo on expression of specific microRNAs (Eulalio et al. 2012). It will be of great intellectual and technical value to learn whether postmitotic neurons are susceptible to similar manipulations.

Future Perspectives of Engineering Neurogenesis Outside Physiological Neurogenic Niches

Although the above-discussed findings provide strong evidence for the principal possibility of engineering neurogenesis in regions of the adult brain, which lack any physiologically relevant degree of neuron production, the question remains to what extent this will enable us to utilize reprogramming strategies for brain repair. There are several issues to be dealt with before we can realistically hope to translate the experimental studies into a clinical context.

First of all, although reprogramming transcription factors are powerful tools to change

the identity of cells, it remains to be explored whether the new phenotype corresponds to a “real” identity or rather reflects a chimeric state. Transcriptome studies of direct reprogrammed fibroblasts typically emphasize the similarity between the induced and naturally born neurons, but, nonetheless, cannot but acknowledge differences that appear to be the more pronounced the more terminally differentiated the cell of origin has been (Marro et al. 2011). This notion is of great significance when potentially considering reprogramming strategies within the adult human brain in which most cells are postmitotic, possibly with exception of slowly cycling OPCs (Yeung et al. 2014). The precise mechanisms underlying reprogramming are still poorly understood. A study in fibroblasts revealed that *ascl1*-induced conversion into neurons relies on this transcription factor’s capacity of accessing closed chromatin, but only when these loci are in a defined trivalent state (as defined by specific active and repressive histone modifications), reprogramming can occur (Wapinski et al. 2013). However, it remains unclear whether any of the non-neuronal cells retain such trivalent state into adulthood, especially in aged human beings, in which the need for new neurons may become most pressing. If the molecular underpinnings of reprogramming are still ill understood, even more mysterious is how morphologically complex cells will metamorphose into neurons. Considering the tremendous complexity of adult human astrocytes (Oberheim et al. 2009), it will be fascinating to see how these cells remodel their astrocytic process into neuronal ones, an issue perhaps to be addressed by two-photon live imaging in models of cortical reprogramming, which allow for the access required for in vivo imaging. Yet, given that rather complex radial glial cells manage to generate neurons (Noctor et al. 2001), it may not be entirely hopeless that other cells of equal complexity could achieve this. Live imaging could also reveal crucial information as to whether reprogramming takes place directly, as suggested by in vitro studies (Heinrich et al. 2010), or requires proliferation. After all, a proliferative metamorphosis may ultimately allow for asym-

metric cell division, by which not only a new neuron may be generated but ideally a cell of the original identity regenerated, thus avoiding depletion of functionally important cells.

A further challenge concerns the issue of functional integration into the neuronal circuitry. Functional integration may indeed be a prerequisite for long-term survival (Tashiro et al. 2006). Although early data suggest that induced neurons derived from *in vitro* as well as *in vivo* reprogrammed cells exhibit electrical properties of neurons, receive synaptic input, and, at least *in vitro*, are capable of generating synaptic output, it remains to be shown whether these cells incorporate in a meaningful manner. Failure to do so may not only compromise their chances for survival, but compromise circuit function. Many neurological diseases are recognized to be caused by defects in connectivity that may be as severe as in epilepsy or more subtle like in autism or schizophrenia. In this context, physiological adult neurogenesis may serve as a model of how new neurons are incorporated to smoothly modify circuit function. The studies of the Song laboratory have shown the important “chaperone” role of parvalbumin-positive interneurons for regulating genesis and differentiation of adult-born dentate granule neurons (Ge et al. 2006; Song et al. 2012, 2013). More generally, integration takes place in a very orderly fashion (Deshpande et al. 2013), suggesting that integration is a process of reciprocity between new neurons and pre-existing circuitries. Incorporation of induced neurons derived from reprogrammed cells is likely to require similar interactions with chaperone cells.

Finally, for most neurodegenerative diseases, such as Alzheimer’s disease, stroke and multiple sclerosis (MS), do not affect single cell types in clean isolation, but involve the entire nervous tissue comprising neurons, glia, microglia, and blood-borne immune cells, reprogramming may not be a viable strategy for brain repair after all, if this does not take into account the micro-environmental changes, such as, for example, the inflammatory response that these diseases induce. Thus, reprogramming strategies must go hand in hand with interventions that create

a conducive microenvironment (Heinrich et al. 2015).

DIRECTED DIFFERENTIATION OF ADULT NEURAL STEM CELLS

The focus on targeted engineering of endogenous NSPCs (or cells that do not generate neuronal cells under physiological conditions) has been to enhance the generation of neurons or to induce neuronal differentiation. However, diseases of the CNS obviously also affect the functional integrity or lead to a loss of glial cells. Given this, especially the NSPC-derived generation of oligodendrocytes, that are, for example, disease targets in chronic demyelinating diseases such as MS, has gained significant attention (Jadasz et al. 2012). Notably, NSPCs in the SVZ generate oligodendrocytes under physiological conditions, and the formation of oligodendrocytes is enhanced after demyelinating lesions (Menn et al. 2006; Xing et al. 2014). Thus, the identification of signaling pathways that mediate oligodendrocytic differentiation is important and may lead to the development of novel strategies to enhance oligodendrogenesis in the context of demyelinating disease (Fig. 3). This is true for the recently identified role for WNT signaling to expand the NSPC-derived oligodendrocytic lineage, the up-regulation of chordin that directs cells toward an oligodendrocytic fate on demyelination in the corpus callosum, and for the finding that loss of SIRT1 enhances oligodendrogenesis (even though it remains unclear whether the enhanced generation of oligodendrocytes on loss of SIRT1 depends on NSPCs) (Fig. 3) (Jablonska et al. 2010; Ortega et al. 2013; Rafalski et al. 2013).

The situation in the hippocampal stem-cell niche is distinct compared with the SVZ as only very few (if any) oligodendrocytes are generated under normal conditions (Zhao et al. 2006). However, it has been recently shown that the fate of newborn cells can be redirected from a neuronal toward an oligodendrocytic fate by the overexpression of the basic helix–loop–helix (bHLH) transcription factor *ascl1* (Fig. 3) (Jessberger et al. 2008). Even though induced oligo-

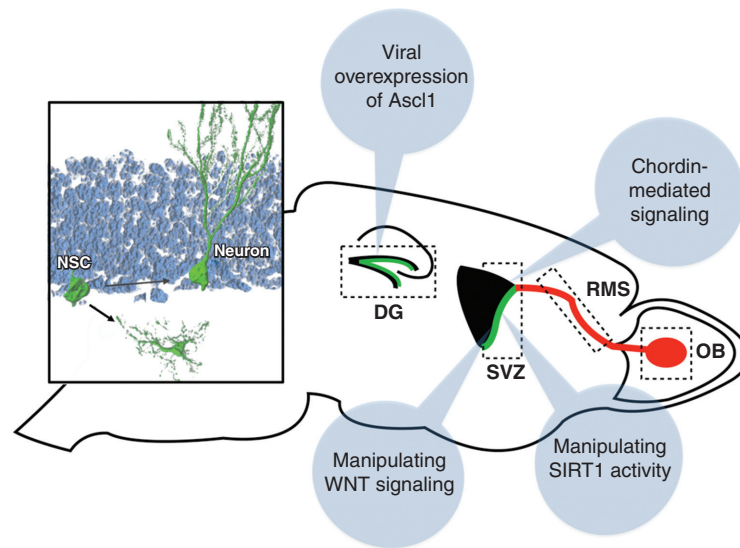


Figure 3. Mechanisms to direct the differentiation of endogenous neural stem/progenitor cells (NSPCs). Shown are several experimental strategies to enhance neural subtype specific differentiation (e.g., through modulating WNT signaling) or to direct the fate of newborn cells from a neuronal toward the glial lineage (e.g., through *ascl1* overexpression in hippocampal NSPCs; shown in the *left inset*). NSC, Neural stem cell; DG, dentate gyrus; SVZ, subventricular zone; RMS, rostral migratory stream; OB, olfactory bulb. (Figure courtesy of Dr. S.M.G. Braun, Brain Research Institute, Zurich, Switzerland.)

endrogenesis appears to be capable to generate myelin, it remains unclear whether newborn oligodendrocytes generated through directed differentiation turn out to be sufficient to enhance remyelination in the context of injury (Goldman and Natesan 2008; Jessberger and Gage 2009).

However, focusing the attempts to generate functional glial cells—in addition to enhanced or induced generation of neurons—will certainly be of growing interest over the next year but will require the development of novel tools to faithfully control the generation of glial cells within the adult brain.

FUTURISTIC BRAIN REPAIR

In some of the most devastating brain diseases like Alzheimer or stroke, it is not just a single type of neuron that degenerates but an entire circuit that needs to be replaced. How to rebuild such highly structured assemblies of neurons (and glia) by reprogramming? Will it be required to provide point-by-point instructions

in the form of specific transcription factors that instruct the genesis of all the plethora of neuronal and glial subtypes to recreate for instance a piece of infarcted cortex? Here, we can open only a highly speculative window into the solution by referring to the remarkable degree of self-organization that neural stem cells derived from embryonic stem cells exhibit when exposed to an appropriately permissive environment eventually resulting in the formation of cerebral organoids (Kadoshima et al. 2013; Lancaster et al. 2013). Too little is known of the underlying logic of this self-organizing process, but the fact that somatic cells such as fibroblasts can be reprogrammed into multipotent neural-stem-cell-like cells (Lujan and Wernig 2012) gives some cause for hope that this may ultimately go along with gaining not only the capacity for neuro- but even “cerebrogenesis.”

SUMMARY

Injury to the adult CNS, through acute or chronic disease, often results in the loss or func-

tional impairment of neuronal and/or glial cells. Recent advances indicated the possibility to engineer the fate and potency of NSPCs or other resident cells to serve as a source for cell replacement of lost structures, opening novel possibilities in the context of regenerative medicine. However, most approaches currently pursued require invasive strategies to genetically modify adult brain cells for fate direction or reprogramming, indicating that these highly experimental approaches are still far from clinical application. Additional proof-of-concept experiments are required that evaluate the potential of NSPC engineering to generate new cells with the aim to replace lost tissue in the context of acute or chronic diseases such as stroke or MS.

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