



# A balanced evaluation of the evidence for adult neurogenesis in humans: implication for neuropsychiatric disorders

Alvaro Duque<sup>1</sup> · Reynold Spector<sup>2</sup>

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## Abstract

There is a widespread belief that neurogenesis exists in adult human brain, especially in the dentate gyrus, and it is to be maintained and, if possible, augmented with different stimuli including exercise and certain drugs. Here, we examine the evidence for adult human neurogenesis and note important limitations of the methodologies used to study it. A balanced review of the literature and evaluation of the data indicate that adult neurogenesis in human brain is improbable. In fact, in several high-quality recent studies in adult human brain, unlike in adult brains of other species, neurogenesis was not detectable. These findings suggest that the human brain requires a permanent set of neurons to maintain acquired knowledge for decades, which is essential for complex high cognitive functions unique to humans. Thus, stimulation and/or injection of neural stem cells into human brains may not only disrupt brain homeostatic systems, but also disturb normal neuronal circuits. We propose that the focus of research should be the preservation of brain neurons by prevention of damage, not replacement.

**Keywords** Adult neurogenesis · Neural stem cells · Memory · Bromodeoxyuridine · Homeostasis · Neuronal protection · DNA repair/methylation

## Abbreviations

|        |                                      |
|--------|--------------------------------------|
| AN     | Adult neurogenesis                   |
| BBB    | Blood–brain barrier                  |
| B-CSF  | Blood–cerebrospinal fluid barrier    |
| BrdU   | Bromodeoxyuridine                    |
| CP     | Choroid plexus                       |
| CSF    | Cerebrospinal fluid                  |
| DG     | Dentate gyrus                        |
| NSC    | Neural stem cell                     |
| SNc    | Substantia nigra pars compacta       |
| SGZ    | Subgranular zone                     |
| VZ-SVZ | Ventricular zone–subventricular zone |

## Introduction

In rodents, monkeys, and more recently humans, there have been hundreds of publications on neurogenesis [the birth of new neurons in the central nervous system (CNS) and spinal cord] during development through adulthood and into old age. Moreover, it is often assumed that newborn neurons in the adult CNS mature are integrated and function normally (Cope and Gould 2019). The purpose of this review is to analyze and synthesize the extant data on neurogenesis in normal and diseased adult humans. The surprising result is that there is no or minimal human adult neurogenesis (AN). To document this conclusion, we will first review the relevant human data, then the methodological issues for validly detecting neurogenesis, and finally the implications for treatment of neuropsychiatric diseases.

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- ✉ Alvaro Duque  
alvaro.duque@yale.edu
- ✉ Reynold Spector  
mspec007@gmail.com

<sup>1</sup> Department of Neuroscience, Yale University School of Medicine, SHM C317B, New Haven, CT 06520, USA

<sup>2</sup> Department of Medicine, Rutgers Robert Wood Johnson Medical School, Piscataway, NJ 08554, USA

## Adult human neurogenesis: what is the evidence?

Hundreds of studies in rodents have employed different interventions to stimulate AN (Chen et al. 2000; Arvidsson et al. 2002; Santarelli et al. 2003; Chen and Sun 2007; Young 2009; Chen and Wang 2016) with the implicit or

explicit assumption that new neurons will be beneficial and that positive results will be applicable to humans, especially adults with disease (e.g., Alzheimer's disease, Parkinson's disease, depression, and stroke) (Benraiss et al. 2001; Gutierrez-Fernandez et al. 2012; Boldrini et al. 2013; Chen and Wang 2016; Cope and Gould 2019). As discussed below, this assumption may be incorrect. In fact, the bulk of the adult human data collected over the last 20 years has failed to prove AN occurs and is of any functional significance in our species.

After considering it theoretically possible on the basis of studies in reptiles, birds, and small mammals, the possibility of human AN received again considerable attention in 1998 when Eriksson et al. took advantage of the fact that five

cancer patients infused with 250 mg of bromodeoxyuridine (BrdU) for diagnostic purposes gave informed consent for their brains to be studied after death. One cancer patient with no BrdU infusion was used for control. These patients died 16 days–2.1 years later (ages 57–72). In postmortem brain samples, occasional BrdU<sup>+</sup> cells were detected in the dentate gyrus (DG), some of which were GFAP<sup>+</sup> (astrocytes; see Table 1 for histological abbreviations and their meaning), while just over 20% of them were co-labeled with NSE or NeuN (putative neurons). The authors concluded that in humans over age 50 continued neurogenesis in the DG existed. They also suggested that there were neural stem cells (NSCs) in adult human brain ventricular–subventricular zone (VZ–SVZ), an area above the anterior horn of the

**Table 1** Common antigens, and their corresponding abbreviations, for the detection of cell proliferation and differentiation

|                       | Antigen          | Description   | Specificity  |
|-----------------------|------------------|---|--|
| Proliferative markers | Ki67             | Cell proliferation protein marker; marks cells in G1, S, G2, and M phases, i.e., marks dividing progenitors                                 | High   |
|                       | MCM-2            | Mini-chromosome maintenance Protein-2; marks cells in G1, S, G2, and M phases   | High   |
|                       | PCNA             | Proliferating cell nuclear antigen—cell proliferation protein marker; marks cells mostly in late G1-phase and S-phase                       | Low  |
|                       | PHH3             | Phosphohistone-H3 (PHH3); mitosis-specific marker   | High   |
|                       | RR               | Ribonucleotide reductase  | High   |
| Progenitor markers    | BLBP             | Brain lipid binding protein for neuronal precursors; highly specific marker of radial glia cells  | High   |
|                       | GFAP             | (See below); radial glia marker   | Medium   |
|                       | Nestin           | Intermediate filament protein; progenitor marker  | Medium   |
|                       | Vimentin         | Intermediate filament protein; progenitor marker; highly expressed in mesenchymal cells   | Medium   |
|                       | Tbr2             | Intermediate progenitor marker  | High   |
|                       | SOX2             | Marks immature, undifferentiated cells  | Medium   |
|                       | PAX6             | Early progenitor marker   | High   |
| Immature neurons      | DCX              | Microtubule-associated phosphoprotein; doublecortin, marker of early stage (immature) neurons, promotes dendritic growth and cell migration | Medium   |
|                       | MASH-1           | Transcription factor—mammalian achaete scute homolog-1; controlled by Hedgehog  | High   |
|                       | NeuroD1          | Transcription factor—specific for postmitotic cells; promotes neuronal differentiation; also marks radial glia cells                        | High   |
|                       | PSA-NCAM         | Polysialic acid—neural cell adhesion molecule; facilitates cell motility—essential for migration, cell growth and synaptogenesis            | Low  |
|                       | Reelin           | Glycoprotein secreted in developing cortex  | Medium   |
|                       | TBR1             | Member of T-box family of transcription factors; Involved in regional and lamina identity (deep cortical layers)                            | High   |
|                       | Neuronal markers | NeuN  | Mature neuron marker; neuronal nuclei marker (Aka, Fox3)—specific for postmitotic cells; Not expressed in Golgi, Purkinje, mitral cells, photoreceptors, cells of inferior olive, dentate nucleus, sympathetic ganglia cells, dopamine cells of substantia nigra |
| NSE                   |                  | Neuron specific enolase; expressed by cultured (and immature) oligodendrocytes and other glia under pathological conditions                 | Medium   |
| TUJ1                  |                  | Neuron marker for $\beta$ -tubulin 3 (immature neurons)   | High   |
| Non-neuronal markers  | GFAP             | Glial fibrillary acidic protein—for astrocytes; progenitor marker   | Low  |
|                       | GFAP $\delta$    | For stem cells not clearly dividing   | Medium   |
|                       | IBA-1            | Ionized calcium binding adaptor molecule 1—for microglia  | High   |
|                       | Olig-2           | For oligodendrocytes precursors and GABAergic progenitors   | Medium   |
|                       | S-100            | For astrocytes  | High   |

Antigen specificity is not absolute and may vary depending on whether use is in vitro or in vivo, developmental age, region, species, etc

lateral ventricle, the origin of the rostral migratory stream that continuously supplies neurons to the olfactory bulb in many animal species, whose olfactory ventricle remains open. In humans, the olfactory ventricle closes during development and the hypocellular region has characteristic gaps devoid of cell bodies. Eriksson and colleagues also pointed out substantial inter-individual variability in the number of BrdU<sup>+</sup> cells and a decline in the number of BrdU<sup>+</sup> cells detected in patients with the longest interval between BrdU injection and histological assessment. They interpreted this observation to indicate a progressive death of the newly generated cells over time. This interpretation seems in agreement with Gould et al. (2001) who reported that adult generated neurons had a transient existence and who also pointed out the difficulty in replicating AN results other than in already damaged brains. To us, a more plausible explanation for the observations in both studies is simply that damaged neurons, attempting to repair themselves, incorporate BrdU (see below) and continue to die, so that eventually, the longer the waiting period before histological evaluation, the less BrdU<sup>+</sup> cells there are. In view of the metabolic cost of producing new cells in a complex already mature neuronal circuit, it is questionable that the brain would invest resources in producing new neurons to have them die soon after production.

Starting in 1999, a few studies claimed that there was neurogenesis in the neocortex and hippocampus in adult primates and the olfactory system in humans (Gould et al. 1999, 2001; Gould 2007). However, these claims were not substantiated in non-human primates (Kornack and Rakic 2001; Rakic 2002a, b, c; Koketsu et al. 2003; Breunig et al. 2007) or in humans (Sanai et al. 2004, 2011; Bhardwaj et al. 2006; Kempermann 2006).

Sanai et al. (2004) confirmed (using cell cultures from a large sample of adult human SVZs [ $n=65$  neurosurgical resections and  $n=45$  autopsied brains]) that the SVZ contained NSCs that could differentiate into immature neurons, TUJ1<sup>+</sup>, PSA-NCAM<sup>+</sup>, and a few DCX<sup>+</sup> neuroblasts. However, the authors acknowledged a robust germinal capacity but no evidence of cells migrating in chains along the SVZ or olfactory peduncle to the olfactory bulb. In adult humans, the existence of NSC in the SVZ was later confirmed (van den Berge et al. 2011) as was the fact that these NSCs did not divide and migrate (Kempermann 2006; Spalding et al. 2013).

Knoth et al. (2010) studied neurogenesis in the DG employing DCX, as the screening marker in  $n=54$  deceased humans, ages 0–100. To confirm neurogenesis, they double labeled DCX<sup>+</sup> cells with other neuronal markers including PCNA and found within the neurogenic niche of the DG (the granule cells layer) on a log–log scale of DCX<sup>+</sup> cells/mm<sup>2</sup> versus age ( $n=45$ , 1 day–94 years), and there was a linear decline from hundreds in fetuses to  $\sim 1$  in the very

old. However, the authors acknowledged that both DCX and PCNA might not be specific for newborn neurons. In their conclusion, Knoth and colleagues were very circumspect. They stated: “Our data alone cannot prove or disprove the true presence or absence of neurogenesis (in humans) at any age...”

Employing atmospheric atomic bomb-test-derived <sup>14</sup>C in genomic DNA, Spalding et al. (2013) concluded that one-third of human DG neurons turnover at a rate of 1.75% per year after studying deceased adults aged 19–92 ( $n=55$  for hippocampal neurons and  $n=65$  for non-neuronal cells). However, their conclusion that “neurons (in the hippocampus) are generated through adulthood and that the rates are comparable in middle aged humans and (9 months) mice” is wrong. This conclusion is based on a calculation error that goes back to original data, in the mouse, reported by Ben Abdallah et al. (2010). As pointed out by Lipp and Bonfanti (2016) and further commented by Parolisi et al. (2018), about 700 new granule cells out of 20 million in a 40-year-old human corresponds to a mere 0.0035% of the total population. In a 9-month-old mouse, 416 new cells in a population of 0.5 million correspond to 0.083% of the population. Obtaining the ratio to compare between species ( $0.0832/0.0035=23.77$ ) makes it clear that the turnover rates are different by a very large factor, here, more than 20 fold in the mouse than in human. Hence, based on their own data, what Spalding et al. (2013) actually show is that the adult human neurogenesis turnover rate in the DG of the hippocampus is a tiny fraction of that in adult mice. Subsequently, using the same technique, they also detected minimal neurogenesis in the striatum (Ernst et al. 2014). The methodology involved in these studies is complex and requires separating neuronal from non-neuronal nuclei in a highspeed cell sorter, separation of rosettes, isolation of the DNA, and subsequent mass spectrometry to identify the <sup>14</sup>C in the DNA (Spalding et al. 2005). The problems here are that, aside of issues of contamination, there are multiple assumptions and corrections by as much as >20%. One important assumption that has been received with some skepticism is that <sup>14</sup>C in pine tree rings is an accurate reflection of <sup>14</sup>C in the air. However, the authors conceded that in industrial cities, this assumption may not be valid (Spalding et al. 2005). In addition, the authors underestimated the amount of DNA repair and removal of methyl groups in DNA which was unknown in 2005 when their methodology was worked out (Spector and Johanson 2014) and which could explain some of the variability in their data. Indeed, in a very active group of cells in the DG, the slow but continuous repair and removal/return of methyl groups explains best the incorporation of <sup>14</sup>C in these aging neurons. Importantly, if cells are <sup>14</sup>C negative, the results suggest that new neurons were not generated, since new neurons cannot be created without carbon incorporation, although it is formally

possible that new neurons were generated using carbon from dead cells as happens with the transfer of BrdU incorporated in the DNA of dead cells into living ones (Spector and Johanson 2007). However, if the cells are  $^{14}\text{C}$  positive, the positivity is not specific. The incorporation of the  $^{14}\text{C}$  isotope could be due to cell division or more likely to any one of several other possibilities including DNA repair/methylation and cell death, processes which are prominent in older specimens. This is why reports using this technique, suggesting that there are no new neurons in the adult human cerebral cortex (Bhardwaj et al. 2006) or in the human olfactory bulb (Bergmann et al. 2012) (negative results) are correct as noted in the experiments described below, while reports by the same group reporting human adult neurogenesis in the hippocampus (Spalding et al. 2013) and striatum (Ernst et al. 2014) are probably incorrect, or at least incomplete and controversial (also see below). Cellular DNA repair, methyl group removal/return, and apoptotic cell death differ in dissimilar structures. However, the result is that the  $^{14}\text{C}$  incorporation method cannot prove the birth of new neurons or the absence of new neurons in adult human brain.

Doorn et al. (2014) studied clinically diagnosed and pathologically verified Parkinson's disease patients ( $n = 14$ , ages 59–96), healthy controls that had some  $\alpha$ -synuclein pathology at autopsy ( $n = 6$ , ages 56–91), and healthy controls without  $\alpha$ -synuclein pathology ( $n = 9$ , ages 62–92). To assess cell proliferation in the DG, they employed MCM-2 staining to screen for newborn cells and used colocalization with IBA-1 to identify microglia. They found that, in the DG, over 90% of the very few MCM-2<sup>+</sup> cells also stained for IBA-1 and thus were microglia.

In an extremely careful study of 23 brains from deceased individuals ages 0.2–59 years, Dennis et al. (2016) studied the SVZ and the DG subgranular zone (SGZ). They employed immunohistochemistry and immunofluorescence in combination with unbiased stereology. To identify the phenotype of proliferating cells in the SVZ and SGZ, they double labeled cells with Ki67 and DCX, and then triple labeled them with TUJ1 or epidermal growth factor receptor, to identify neurons; or GFAP (astrocytes), IBA-1 (microglia) or Olig 2 (oligodendrocytes). Importantly, all studies were done blinded. The assumption was that triple staining would lessen the chance for false positives. By age 3, 60% of the Ki67<sup>+</sup> cells in the SGZ co-stained with IBA-1 (and thus were microglia). At older ages, > 99% of the Ki67<sup>+</sup> cells stained with IBA-1. Thus, this study is consistent with Doorn et al. (2014) that the principal proliferating cells in the adult human DG are microglia not neurons. In the SVZ, they replicated the results of Sanai et al. (2004) (up to 18 months) and the exponential decrease with aging for DCX immunostaining in the SGZ as shown by Knoth et al. (2010). The authors point out that the results are consistent with PCNA labeling in the infant

human SVZ and rostral migratory stream and that only a non-proliferative pool remains in the adult. They further indicate that PCNA-based immunohistochemistry studies overestimate proliferative events in human postmortem brain tissue.

A more recent paper by Mathews et al. (2017), looking at neurogenesis associated changes in gene expression ( $n = 26$ , ages 18–88), is consistent with the results of Doorn et al. (2014) and Dennis et al. (2016). Interestingly, they point out that strongly GFAP<sup>+</sup> profiles with large thick radiating processes found in many elderly adults are “consistent with the morphology of activated astrocytes”. Both, astrocyte activation and microglia activation, are common in neuroinflammation and especially prominent in neurodegenerative brain diseases (Jang et al. 2013; Liddel et al. 2017).

Sorrells et al. (2018) published a comprehensive study on hippocampal neurogenesis in humans. They used a total of  $n = 59$  postmortem and post-operative samples of the hippocampus and investigated the presence of progenitor cells and neurons from fetal age to adulthood. Their controls ( $n = 37$ ) spanned from 14 gestational weeks to 77 years and their samples from people with epilepsy ( $n = 22$ ) spanned 3 months–64 years. They also used  $n = 12$  Rhesus macaques. They employed multiple label detection with various combinations of Ki67, DCX, PSA-NCAM, BLBP, NeuN, GFAP, and other markers, as reported in supplementary Table 3. The methodology included light, confocal and electron microscopy, RNAscope in situ hybridization, and comparative gene transcription analysis. Their careful and exhaustive examination provided data in support of a rapid decrease of DG neurogenesis with age. They stated that in the monkey, hippocampus SGZ proliferation was found in early postnatal life and diminished during juvenile development, as was previously reported (Eckenhoff and Rakic 1988; Kornack and Rakic 1999); importantly, they concluded that “...neurogenesis in the dentate gyrus does not continue, or is extremely rare, in adult humans”. In a very thoughtful discussion, they suggested that earlier studies that found low-level DG neurogenesis in human adults were probably in error. They pointed out that DCX can be expressed by glial cells and that the BrdU<sup>+</sup> cells in the Eriksson et al. (1998) paper “...could possibly be explained by processes not associated with cell division...” as suggested previously (Kuan et al. 2004; Breunig, et al. 2007; Gould 2007; Spector and Johanson 2014).

As commented by Snyder (2018), hopes of neuronal plasticity may not necessarily be lost, since the plasticity, due to the addition of new neurons in rodents, may in humans be provided by the prolonged development of neurons, which in some cases may last decades. Andrae (2018) and La Rosa et al. (2019) proposed focusing research efforts on methodologies to extend the period of normal neurogenesis, so that this can be used to treat neurological disorders in adults. Whether this is possible will require future intense

investigation. Similarly, Ovsepian (2019) suggested repurposing underutilized neurons for plasticity.

A study by Cipriani et al. (2018) reports results in agreement with those of Sorrells and colleagues. This study, which included  $n = 39$  controls spanning 13 gestational weeks to 72 years and  $n = 5$  Alzheimer's disease cases ages 74–89, concluded that pools of “morphologically, antigenically, and topographically diverse neural progenitor cells are present in the human hippocampus from early developmental stages until adulthood, including in Alzheimer's disease patients, while their neurogenic potential seems negligible in the adult”. As pointed out in a commentary by Arellano et al. (2018), the common finding of these separate, yet almost simultaneous studies, is very significant, because the DG was considered the only structure in the adult human brain, where the possibility of neurogenesis was still being considered. In non-human primates, new neurons could also not be detected in this structure after puberty (Eckenhoff and Rakic 1988).

Recently, three studies suggested that there is neurogenesis in the DG. A study by Boldrini et al. (2018), in autopsied persons ages 14–79 ( $n = 28$ ) with no neuropsychiatric disease or treatment, states that there is continuous sustained neurogenesis throughout aging in the DG. However, this study is fraught with problems. One example can be seen in Fig. 1h in which they claim that a Nestin<sup>+</sup> cell is an intermediate neuronal progenitor. This cell looks like a reactive astrocyte. However, since staining with IBA-1 or Olig-2 is not presented, one cannot be sure how glial cells were ruled out. Microglia and astrocytes can be Nestin<sup>+</sup> (Takamori et al. 2009). It is possible that they consistently misidentified non-neuronal dividing cells for neurons. In agreement with this, Parolisi et al. (2018) also reviewed the evidence for AN and used humans and dolphins as examples of long living species with large brains to discuss its evolutionary implications. Parolisi et al. state that in the Boldrini paper, “...various molecular markers were found associated to different stages of immature neurons, which do not show the typical aspect of recently generated neuroblasts”. In addition, Boldrini et al. claimed that they could not compare their findings to those of Knoth or Sorrells, because these studies did not use stereology. However, they did not compare their results to studies that did use stereology such as, for instance, (Dennis et al. 2016). Commentaries by Lee and Thuret (2018) and Kuhn et al. (2018) stress the advantages of using stereology, the shorter postmortem delays, and the healthier controls in Boldrini's study. However, they do not mention advantages of any of the additional methodologies used in the Sorrells paper. Paredes et al. (2018) points out that “...stereology is only a useful technique if what is being counted is correctly identified”.

Another recent study claims that hippocampal neurogenesis is abundant in neurologically healthy subjects and that it

drops sharply in patients with Alzheimer's disease (Moreno-Jimenez et al. 2019). This study shows clear DCX staining in adult human brains. However, as established previously, DCX positivity is not sufficient to demonstrate new neurons (see, for instance, Gomez-Climent et al. 2008; Luzzati et al. 2009; Klempin et al. 2011). In addition, the supposedly new neurons are in the wrong place, i.e., mostly away from the hilus and they look mature. Unfortunately, there is no documentation of any dividing cells or precursors and the authors offer no proof of the presence of cycling/dividing cells with progenitor morphology and molecular profiling (Sox2<sup>+</sup>, Nestin<sup>+</sup>, and GFAP<sup>+</sup>). In addition, what the authors called normal controls were in fact ill individuals (many had cancer). If there was abundant adult neurogenesis in the hippocampus, as the authors claim, it is puzzling that the hippocampus is not constantly growing. It seems logical to expect that, for it to stay the same size, old neurons must die and be removed at least at the same rate as the new ones repopulating it. Yet, we are unaware of any study showing “abundant” adult apoptosis concomitant with correspondingly abundant neurogenesis.

A third paper claims that human hippocampal neurogenesis persists in aged adults and Alzheimer's disease patients (Tobin et al. 2019). These authors are aware of the results of Moreno-Jimenez et al. (2019), but do not critically reflect on the differences found in reference to the amount of putative adult neurogenesis in the Alzheimer's patients. They do acknowledge that they do not observe a correlation between level of neurogenesis and amyloid deposits or fibrillary tangles. In a careful examination of their figures, we have failed to identify any clear evidence of newborn neurons on their way to become adult neurons in process of being incorporated into the DG circuitry. Needless to say, any progenitor could become glia instead of neuron and there is no IBA-1 staining to, at least, rule out microglia. While extensively critical of the results in Sorrells et al. (2018), the authors do not comment on the results of Cipriani et al. (2018), which are in agreement with those of Sorrells et al. (2018).

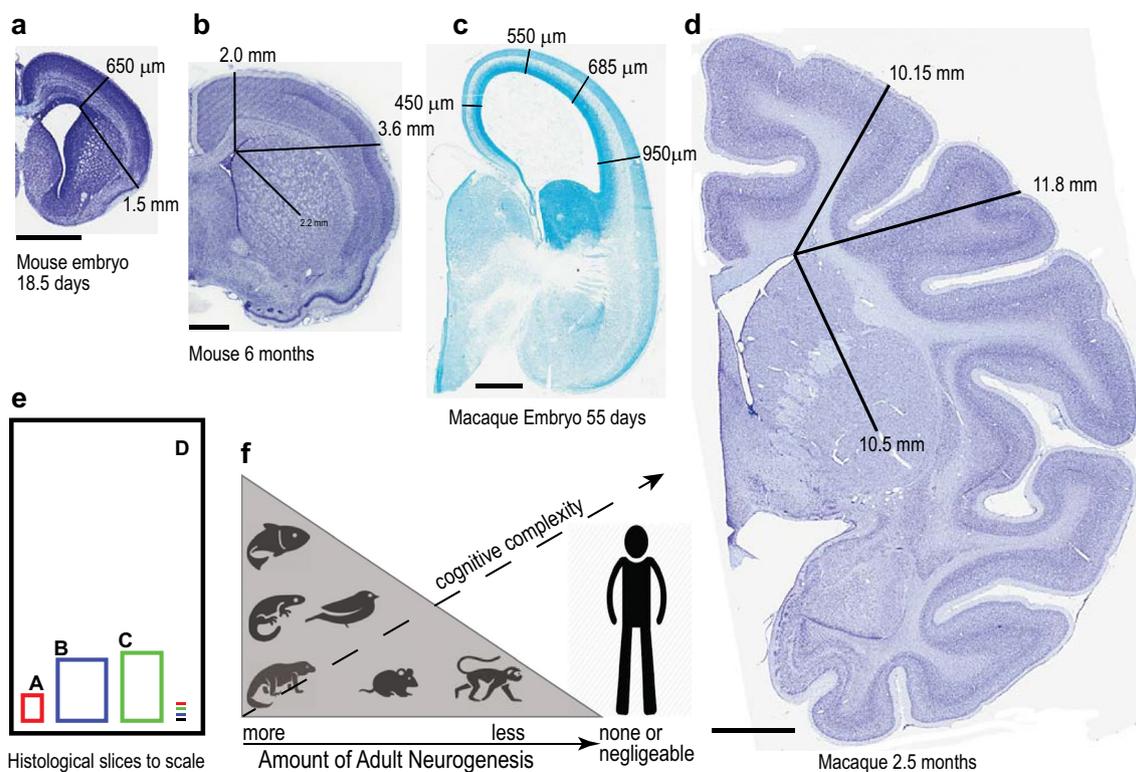
Thus, as it now stands, the studies by Boldrini et al. (2018), Tobin et al. (2019) and Moreno-Jimenez et al. (2019) are at odds with the works of Knoth et al. (2010), Doorn et al. (2014), Dennis et al. (2016), Mathews et al. (2017), Cipriani et al. (2018), and Sorrells et al. (2018). A commentary by Kempermann et al. (2018) points out the need to very seriously consider all technical issues “...for a full evaluation of the evidence”. We could not agree more. However, the assertion that the “...functional contribution that new neurons would make to human cognition is not negligible” is not consistent with the data summarized above. There is, in fact, no evidence that would clearly indicate human cognition would weaken or diminish if there was no AN in the hippocampus. In rats, it is worth noting that in a technical tour de force employing a pharmacogenetic model,

Groves et al. (2013) showed that ablating adult neurogenesis in the DG showed no difference from controls in spatial pattern separation, spatial learning, or contextual or cued fear conditioning. A meta-analysis of all published studies also showed no effect for ablation of adult neurogenesis on dentate function, but did find remarkable high levels of heterogeneity among studies of hippocampal function. Moreover, in agreement with the results of Groves et al. (2013), in a paper entitled “A transgenic rat for specifically inhibiting adult neurogenesis”, Snyder et al. (2016) reported that blockade of neurogenesis in the hippocampus did not affect anxiety levels or patterns of exploration. The Kempermann et al. (2018) commentary also fails to consider, for instance, Akers et al. (2014) who suggests that hippocampal AN may contribute to forgetting. Or as Mongiat and Schinder (2014) pointed out in their commentary on Akers’ paper, adding new neurons (in the DG) “...will still impose a cost to network stability”. Thus, adding new neurons might, in fact, be counterproductive. Hence, we agree that the standardization of methodologies is a crucial step towards a possible solution to the contradictions and controversies in the field and,

as Fred Gage (2019) suggests, “the creation of open-access brain banks with tissues more ideally suited to these types of studies” may be an essential first step.

Finally, it is worth noting that the studies of Knoth et al. (2010), Dennis et al. (2016), Sorrells et al. (2018), and Cipriani et al. (2018) had built in internal controls. Using the same methodology, they found abundant neurogenesis in unborn and young humans, which in all four studies declined to undetectable levels or minimal neurogenesis in adults and elderly humans, and thus, the methods they employed were robust in detecting neurogenesis when it occurred. In other words, the methodologies employed in these four studies have face validity and are responsive to change.

Human brain size has also been hypothesized to be an important limiting factor for AN (Paredes et al. 2016). These authors highlighted the tremendous differences in distances that separate cell origin and destination in the different species and discuss the fact that migration would be extremely difficult in developed large brains due to obstacles such as intervening white matter. We also show this in Fig. 1.



**Fig. 1** a–e Comparison of brain sizes and some migratory distances to illustrate the large differences among mouse and macaque monkey at different developmental points. f Amount of adult neurogenesis from “more” in fish, salamanders, and other reptiles to “substantial” in birds and rodents, to “less” in monkeys and finally to “none, negligible or not detectable” in humans. The capacity for adult neurogenesis seems negatively correlated with the cognitive capacity in the

different species, so that the less cognitive complexity the more likely adult neurogenesis is present. Rakic (1985) postulated that humans, throughout evolution, lost their capacity to regenerate neurons in exchange for stability in the neural networks, so that processes of memory, learning and higher cognitive functions were favored. Scale bars: 1.0 mm (A–C), 4.0 mm in D. Macaque tissue examples are from MacBrainResource.org

In summary, almost 100 years ago with their meticulous histological studies, anatomists such as Kolliker, His, and Ramon y Cajal correctly indicated that there is no human AN. They suggested that nerve cells are responsible for the most precious human mental functions and are irreplaceable under normal conditions. In other words, as Rakic (1985) suggested "...a prolonged period of interaction with the environment, as pronounced as it is in all primates, especially humans, requires a stable set of neurons to retain acquired experiences...". The question is why do some studies still provide controversial results interpreted as evidence that there is human AN? We believe that methodological issues are at the center of the controversy.

## Methodological issues

Among the technical issues that may contribute to, what we consider, the erroneous idea of human AN, there are problems unique to dealing with human samples (e.g., the difficulty in assuring homogenous postmortem delays before fixation and processing, while some markers are degrading), problems of translation, since the results in other species have not been translatable to humans and general technical problems of marker specificity, toxicity, detection, quantitative methodologies, proper use of controls, and even expression differences between brain regions or at different developmental stages. Because the renewed interest in AN was sparked by studies using BrdU, which is still the most commonly used of the thymidine analogues, and because much work on this subject has been done in mice and uses a host of different neuronal and non-neuronal markers, we focus our technical review on these issues.

### The use and misuse of BrdU

The use of BrdU as a specific and decisive marker of cell division has been particularly misleading in the large AN field. In part, because many authors continue to ignore the warnings expressed in many reviews (e.g., Taupin 2007; Spector and Johanson 2007; Levkoff et al. 2008; Ross et al. 2008; Duque and Rakic 2011, 2015; Lipp and Bonfanti 2016) about the possibility of false BrdU labeling or the erroneous interpretation of such labeling. Hence, many researchers take uncritically any immunolabeling of the cell nucleus, and even perinuclear cytoplasm, as a definitive sign of cell division and, therefore, as evidence of the production of new neurons. Of course, BrdU does label dividing cells. However, the possibility that BrdU positivity may be due to attempts of cells to repair themselves or a sign of cell death (apoptosis) are usually neglected or understated. Excitation of the brain by running and/or induced epilepsy, both which stimulate DNA synthesis and BrdU positivity, were

interpreted as production of new neurons (van Praag et al. 1999; Parent and Lowenstein 2002; Jakubs et al. 2006). The fact that, despite supposedly producing new useful neurons in the hippocampus in epileptic animals and human cases, the hippocampus may become smaller rather than bigger is neglected.

BrdU's toxicity has been well documented including, among others, its detrimental effects on chromosomes and DNA stability, the cell cycle, cell differentiation and survival (Hsu and Somers 1961; Webster et al. 1973; Bannigan and Langman 1979; Saffhill and Ockey 1985; Biggers et al. 1987; Kolb et al. 1999; Sekerkova et al. 2004; Breunig et al. 2007; Lehner et al. 2011; Morris 1991). Moreover, BrdU labeling may signify cell death rather than renewal. In an experiment in which unilateral ligation of the internal carotid artery caused a large number of BrdU<sup>+</sup> cells in the hippocampus on the side of the ligation with subsequent loss of labeled neurons (Kuan et al. 2004), many BrdU<sup>+</sup> cells could be double labeled with the tunnel method as a sign of cell death. Hence, to continue to ignore or downplay the toxicity of BrdU and caveats concerning its utility only serves to add to the confusion.

### The use and misuse of inbred animal models, particularly mice

One fundamental and very disturbing fact with most laboratory rat and mouse models is that these animals have been inbred for hundreds of generations. Hence, it is unlikely that the laboratory animal is even a good model of a true "wild" one. In fact, while many studies have indicated the exuberant existence of AN in the hippocampus of laboratory mice the concept of AN in the real "wild" mouse hippocampus is questionable. Take for instance the report by Hauser et al. (2009) in which the authors captured male and female wood mice in a park around the University of Zurich and found that wheel running had no impact on cell proliferation, neurogenesis, or cell death in their hippocampi.

There are some advantages to inbred species, especially when studying genetics, because the similarity among individuals may facilitate the isolation of genes involved in particular traits. However, it is clear that inbreeding has made our laboratory animals rather different than their wild counterparts. In humans, the same is true. Because of cultural, political, or geographic factors, human inbreeding has resulted in populations experiencing different sorts of genetic problems (e.g., early onset Alzheimer's disease) which can facilitate the isolation and study of the genes involved.

Mice and humans went on their own evolutionary paths tens of millions of years ago. Therefore, each species developed their own unique features. There are many analogous structures and systems, but they are not always homologous.

Rodents are not a small version of humans which may very well be why over, and over again, many treatments that work in rodents fail in humans.

### The limitations of antibodies and marker specificity

Specific markers for NSC and/or newborn cells may not be as specific as expected. Their expression could be transient, intermittent, species dependent or could be missed completely for technical reasons. Even if the labeling is not an artifact, the function of the marked cell may be different from the expectations of the experimentalist. For instance, microglia may be produced and detected as adult newborn cells, but they are not neurons and will not become neurons. They may have a transient and specific function such as in the clearing of dead cells (Streit 2000; Lu et al. 2011), which could also be cleared by immature neuroblasts (Sierra et al. 2010). An additional confound is the possibility of the retention of immature neurons which maintain their markers and morphology and which may mature much later in life. Unfortunately, events such as the birth of the organism are not good predictors of the brain's developmental state and it is difficult to detect and correctly interpret cellular events in the nonlinearly scaling lifespan of different species (Charvet and Finlay 2018).

Table 1 lists some of the commonly employed antigens for cell division and a few presumably specific neuronal markers. However, specificity is often not absolute and this also is either ignored or neglected by many authors. For instance, the neuronal marker NSE is expressed by cultured (and immature) oligodendrocytes and other glia under pathological conditions. DCX and PSA-NCAM, not only mark immature neurons in non-human species, they also mark mature neurons and glia cells in humans. To expand on the idea of the need to be careful when interpreting the expression and specificity of any sort of labeling, take as an example the case of the commonly used marker NeuN (for a review, see Guselnikova and Korzhevskiy 2015). Its expression is commonly believed to occur exclusively in neurons, but not all neurons. Notably, NeuN does not label cerebellar Golgi, Purkinje, or dentate nucleus cells; mitral cells of the olfactory bulb, photoreceptors and most cells of the inner nuclear layer of the retina, neurons of inferior olive, dorsal cochlear nucleus, gamma motor neurons, sympathetic chain ganglia cells, or cortical Cajal-Retzius cells, and was deemed unreliable to label dopaminergic cells of the (rat) substantia nigra. To complicate matters, *in vitro* experiments indicate that GFAP<sup>+</sup> (astrocytes) cells are also NeuN<sup>+</sup>. Therefore, are these cells *in vitro*, neurons or glia? Their morphology is consistent with that of astrocytes.

Because it is difficult to find a single method in neurobiology that stands alone and is sufficient to demonstrate any theory, further measures to minimize the possibility of error

include, for instance, the use of confocal microscopy with multiple planar views, when using immunofluorescence, to be certain an antigen is actually in the assigned cell. This avoids the common mistake of identifying a satellite glial cell for a neuron. Species-specific limitations are particularly important when employing genetic techniques, viral vectors, electroporation, recombinase-based systems, and transplants. For example, genetic models in animals include, among others, knock-in, knockout, transgenic, Cre/lox P, inducible Cre, and mosaic analysis (Kempermann 2006; Breunig et al. 2007). However, in general, these methods are not usable in humans, except when there are “experiments of nature” (e.g., human knockouts).

## Pharmacological implications for the treatment of neuropsychiatric diseases

### Brain protection, nourishment, and homeostasis

In view of the lack of, or minimal, neurogenesis in the adult human brain, only possibly in the DG, one might expect a really robust protective and nourishing environment for the brain. This would prevent damage and provide a stable environment, since the neurons must function smoothly and last intact for decades. Surrounding the adult human brain are a myriad of anatomical, physiological, and homeostatic systems that provide a near constant internal milieu (Table 2).

Anatomically, the skull encased brain is isolated from the blood by capillaries with tight junctions, i.e., the blood–brain barrier (BBB; Spector 2009, 2010; Spector and Johanson 2010, 2014). The cerebrospinal fluid (CSF) is also isolated from the blood by tight junctions between the epithelial cells in the choroid plexus (CP), which secretes the CSF, and the tight junctions in the layers of the arachnoid cell membrane which encompasses the CSF (Spector et al. 2015a, b). Thus, the CP and arachnoid membrane make up the blood–cerebrospinal fluid (B-CSF) barrier. There is, however, ready exchange of small molecules between CSF and brain; in other words, the brain and CSF, which also provides buoyancy for the brain, can be considered a single compartment isolated from the blood and protected by the skull (Spector et al. 2015a, b). Of course, within the central compartment, astrocytes and microglia constantly sculpt and protect neurons, and help provide a stable milieu.

Physiologically, these barriers provide anatomical protection against water-soluble, ionic, and large molecules entering the brain from the blood (Spector 2009; Spector and Johanson 2010). However, highly lipid soluble compounds and water can freely enter brain from blood. Moreover, at both the BBB and B-CSF barrier, there are multiple specialized active-transport “pumps” to extrude unwanted

**Table 2** Architecture and systems for brain protection, nourishment, and homeostasis

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|   |
|---|
| Anatomical protection   |
| (a) Skull   |
| (b) Cerebrospinal fluid (brain buoyancy with operational brain weight 40 g) |
| (c) Blood–brain barrier (BBB)—tight junctions in cerebral capillaries       |
| (d) Blood CSF (B-CSF) barriers (see text)                                   |
| Physiological protection  |
| (a) Multiple specific and non-specific efflux transport systems             |
| 1. At BBB   |
| 2. At choroid plexus (CP)   |
| 3. Via turnover of CSF  |
| (a) Metabolism of some compounds  |
| 1. At CP  |
| 2. At BBB (e.g., dopamine)  |
| Nourishment of brain  |
| (a) At BBB  |
| 1. Macronutrients (e.g., glucose, fatty, and amino acids)                   |
| 2. Micronutrients (e.g., riboflavin and thiamine)                           |
| 3. Ions (e.g., iron but not sodium or chloride)                             |
| (b) At B-CSF barrier via CP into CSF  |
| 1. Micronutrients (e.g., vitamin C and folate)                              |
| 2. Irons (e.g., sodium, chloride, and bicarbonate)                          |
| 3. Growth factors and peptide hormones (e.g., prolactin)                    |
| Remarkable mechanisms for brain homeostasis at or within                    |
| (a) CP and CSF  |
| (b) BBB   |
| (c) Brain cells   |

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endogenous and exogenous molecules from the central compartment back into the blood (Spector 2010; Spector and Johanson 2010); importantly for pharmacologists, these protective systems can also prevent the entry of therapeutic drugs. For instance, in the case of meningitis, penicillin and its congeners, which are potentially useful treatments, are vigorously transported by organic acid transporter-3 out of CSF (Spector 2010). This problem was overcome by the finding that the long half-life ceftriaxone was not transported by organic acid transporter-3 and thus became the drug of choice for most cases of meningitis. In other cases, e.g., the congenital vitamin transport disorders, exemplified by the lack of riboflavin, thiamine, or folate transporters into brain at the BBB, the barrier to vitamin entry (because of the missing transport system) was overcome by giving massive doses of the appropriate vitamin and thus preventing the severe neurological damage associated with these vitamin transport disorders (Spector 2014). There are some drugs, whose entry is facilitated by “riding” into brain on endogenous transport systems, e.g., diphenhydramine, or others that avoid the barriers, e.g., the lipid soluble pentothal. Therefore, the barriers that work effectively to protect the brain from toxic chemicals and drugs (e.g., ivermectin by P-glycoprotein at the BBB) (Spector 2010) are the same that

produce significant problems for the delivery of potentially useful drugs into the central nervous system.

In terms of brain nourishment, both the BBB and CP contribute via separate saturable systems (Spector 2009; Spector et al. 2015a, b). Many of these systems at the barriers also provide homeostasis and participate in maintaining a constant internal milieu for the brain. In fact, most nutrients including glucose and vitamins in the CSF and brain as well as ions are buffered from changes in concentrations in blood (Spector 2009; Spector et al. 2015a, b). For example, large increases or decreases in blood potassium minimally change the potassium concentration in CSF and the extracellular space of brain. Similarly, large drops in many vitamin blood levels are not reflected in brain because of the vitamin homeostatic systems. An excellent example is scurvy in which the rest of the body is scorbutic, but the brain until the very end retains enough vitamin C to function—a remarkably protective result (Spector 2009). This is not due to lack of turnover, but due to the efficacy of the sodium-dependent vitamin C transporter-2 in the CP to pump the vitamin into CSF and brain at very low blood concentrations (Spector 2009). With these multiple complex systems for protection, nourishment, and homeostasis, the adult human brain can function optimally. Without even one

of them (e.g., the thiamine transport-homeostatic system) at the BBB and B-CSF barriers due to congenital absence or severe deficiency states, irreversible neurological damage occurs (Spector 2014).

Thus, in the brain, unlike the gut, skin and liver which are exposed to both chemical and, in the case of skin, physical toxins (e.g., UV light), the non-replicating neurons in adult human brain are protected and nourished by a large number of discrete systems as noted above. These latter tissues especially the gut and skin, in fact, have programmed turnover mechanisms—with stem cells in the crypts in the gut and basal layers of the skin dividing, renewing, and replacing the dead superficial cells. Even bone slowly remodels. The neurons in the brain do not and thus require exquisite protection for longevity.

### Transplantation of neural stem cells: example of Parkinson's disease

Parkinson's disease is due to a decrease of dopamine input to the striatum from dopaminergic cells in the substantia nigra pars compacta (SNc). Transplantation of dopamine-producing NSC into the putamen or SN was tested in two careful double-blind trials (Freed et al. 2001; Olanow et al. 2003) in which cyclosporine was employed to prevent rejection. Both trials provided clear evidence of stem-cell survival with production of dopamine, but there was no clear benefit, although there was possibly a treatment effect in milder patients. However, in both trials, there was disabling “off-levodopa therapy” dyskinesia; in the later trial, 56% of the transplanted patients were affected and Olanow and colleagues concluded that “fetal (stem cell) transplantation cannot be recommended as a therapy for Parkinson's disease”. They hypothesized the dyskinesia was due to “incomplete or aberrant re-innervation of the striatum”. Since these trials were completed over 15 years ago, there has been substantial new information about the biology of the SN. For example, in rats, there is tonic somatodendritic dopamine release in the SN with the extracellular SN concentration of dopamine ~ 80 nM (Yee et al. 2019). This is enough to saturate DA D<sub>2</sub> receptors. The tonic release of DA is not due to action potentials. Whether this occurs in humans is unknown but probable. This finding greatly complicates the traditional understanding of SN biology. For example, how could transplantation of dopamine neurons into the putamen of Parkinson's patients influence dopamine levels in the SN?

In addition, since there is renewed interest in transplanting human iPS cell-derived dopaminergic neurons in Parkinson's disease, there are many issues that require resolution. In a primate model of Parkinson's disease employing human iPS cell-derived neurons from four patients with PD and four controls, Kikuchi et al. (2017) grafted these eight cell lines into the putamen of monkeys. The grafted monkeys

improved and there was no dyskinesia unlike humans as noted in above. The grafted cells from both normal and Parkinson's disease patients produced substantial dopamine. Kikuchi et al. used the immune-suppressive drug (FK506) to counter rejection. In this study, there was tremendous inter-individual variation in the number of DA cells in the putamen that were tyrosine hydroxylase positive (a putative measure of DA cells). One of the issues that became apparent is how an investigator should select “good donor cells” for transplantation. A further complication is the finding that how you grow the cells matters, e.g., the importance of timely vitamin C in the growth medium to establish crucial epigenetic modifications to ensure maximal utility of the donor cells (Wulansari et al. 2017). Finally, even if you can pick a “good donor” cell line for transplantation, the pharmaceutical issues of scaling up to potentially tens of thousands of transplants employing FDA-required good laboratory practice and keeping the cells stable are formidable challenges (Crow 2019). Whether transplantation of stem cells is hopeless in Parkinson's disease or if a different paradigm might work remains to be established.

### Is there neurogenesis after stroke?

Over 90% of strokes are ischemic due to interference with brain blood flow because of intra-arterial clotting or less commonly an embolus (Spector 2016). At present, there are two established (proven) treatments, although of modest efficacy: intravenous tissue plasminogen activator to dissolve the clot or endovascular therapy to remove or relieve the obstructing clot. Both require treatment within a few hours after the incident for a potential beneficial effect (Chen and Wang 2016). In animal models, with and without stimulation, many reports found meaningful neurogenesis in the penumbral region (the region surrounding the core necrotic region) after stroke (e.g., Pencea et al. 2001; Parent et al. 2002; Kernie and Parent 2010; George and Steinberg 2015). Although early reports suggested that there was neurogenesis in the penumbral region after stroke in humans (Jin et al. 2006; Lindvall and Kokaia 2015), these new neurons or old neurons unsuccessful attempts at dividing were not permanent. In the Spalding et al.'s (2013) study of <sup>14</sup>C dating of neurons in the penumbra of stroke for new neuron birth dating, the authors could not detect any (lasting) neurogenesis in the penumbra. Lindvall and Kokaia (2015) pointed out the difficult task for the production of useful replacement neurons in human stroke: the new neurons must survive, migrate to the appropriate place, differentiate into the phenotype of dead neurons that need to be replaced, and finally be integrated into functional circuits with correct synaptic connectivity. This apparently does not happen (Spalding et al. 2013). In summary, there is no evidence for human AN after a stroke. Moreover, in stroke, even if you could stimulate

neurons with growth factors to divide, the challenge of turning such newborn neurons into functioning integrated neurons in adult brain seems challenging and, so far, impossible. It is worth noting that Kondziolka et al. (2005) did a controlled transplantation trial in stable (for 2 months) post-stroke patients with 5–10 million human neuronal cells. This trial was negative and roundly criticized by Bakay (2005).

## Drug therapy and neurogenesis

As documented above, in adult monkeys, there is minimal and in humans undetectable AN, except possibly in the DG. In rodents, there are hundreds of reports describing interventions that increase neurogenesis, e.g., exercise, lithium, tricyclic anti-depressants and serotonin-selective receptor inhibitors like fluoxetine, and, in some studies, these interventions improved cognition (Gould et al. 2001; Santarelli et al. 2003; Foland et al. 2008; Young 2009). However, there are multiple technical problems with these studies. For example, focusing on lithium used to treat manic-depressive disease, Chen et al. (2000) employed high-dose BrdU to detect new neurons in the DG in control versus lithium-treated mice. After 4 weeks of lithium treatment, they found a 25% increase in BrdU<sup>+</sup> cells in the DG. Moreover, they found that 65% of the BrdU<sup>+</sup> cells in both control and lithium-treated mice were NeuN<sup>+</sup>, suggesting that they were neurons. However, they did not measure the blood levels of BrdU which could have been increased due to the lithium treatment (Kempermann 2006; Breunig et al. 2007; Spector and Johanson 2007). If the blood levels of BrdU were 25% higher in the lithium group, their results could be explained by the altered pharmacokinetics of BrdU which would also explain why there was no increase in the percentage of NeuN<sup>+</sup> cells in the control and lithium-treated mice. Similar types of issues can be raised in the putative increase in dentate neurogenesis in the anti-depressant studies (e.g., Santarelli et al. 2003). Moreover, even if there is increased neurogenesis in the DG of animals with lithium and/or anti-depressants, this does not prove neurogenesis in the DG gyrus is the cause of the improved animal cognition. However, many authors, as noted above, have either explicitly or implicitly suggested that these rodent study results can be applied to human adults (Santarelli et al. 2003; Chen et al. 2000; Cope and Gould 2019). Further complicating the rodent studies, Snyder et al. (2009) showed that adult born hippocampal neurons are more numerous and more involved in behaviors in rats than in mice. Hence, what to believe, rats or mice?

In an autopsy cross-sectional study of patients with major depression, Boldrini et al. (2013) reported a larger DG and more granule cell neurons in anti-depressant-treated patients than in untreated control patients. However, they concluded

that postmortem studies “are correlative” and do not prove causality, i.e., that the anti-depressant increased dentate size is due to neurogenesis. In fact, there are many serious problems associated with the hypothesis that anti-depressants work by rapidly increasing neurogenesis in the DG. In monkeys, if AN was to happen, then as reported by Kohler et al. (2011), it would take longer than 6 months for new neurons in the DG to connect and be integrated in a functionally meaningful way. Moreover, the well-known observation that ketamine can overcome severe depression in hours undercuts the neurogenic hypothesis (i.e., that anti-depressants work by increasing neurogenesis) and favors the chemical imbalance hypothesis.

In view of the finding that there is minimal or, in some studies, no detectable human AN, it is difficult to argue that several diseases (e.g., depression, manic-depressive disease, Alzheimer’s disease, and dementia) are due to insufficient neurogenesis in the DG. Even if one could increase neurogenesis with safe interventions, appropriate cellular integration into functioning systems is doubtful, since the distances axons must travel to correctly connect to their targets are much longer and more complex in human than in rodent brains (Fig. 1) and the developmental signals and chemicals to direct axonal growth are long gone. In addition, it seems unlikely that diseases of such diverse pathogenesis, affecting different cellular populations in dissimilar brain structures, would all be due to the lack of, or remediated by, AN. The literature favoring the existence of human AN seems to imply that it must exist, because AN exists in other species. However, this is intrinsically inconsistent with the uniqueness of species-specific capacities and even with evolution (La Rosa et al. 2019).

In our view, a beneficial result of stimulating neurogenesis or transplanting NSCs into adult human brain is, at present, just a hypothesis. The experimental finding in Parkinson’s disease noted above in which transplanted neurons functioned (and made dopamine), but made many patients worse is most instructive. Until there is clarification of this issue, i.e., whether transplantation or stimulation of neurogenesis is helpful or, as suggested by Rakic (1985) and Akers et al. (2014) potentially harmful, a better approach for neuroscientists and pharmacologists would be to promote optimal use of the brain by ensuring ideal care of the body (e.g., good diet, normal weight, and no smoking) and to develop treatments that prevent brain damage (e.g., from hypertension, stroke, atherosclerosis, arterial clotting, and nutritional deficiencies).

## What can neuroscientists and pharmacologists offer to preserve brain function?

Neuroscientists, pharmacologists, and clinical pharmacologists have contributed by, for example, employing their knowledge of the pathophysiology of stroke and its causes

to diminish risk factors. They have developed well-known preventive therapies for the control of hypertension and diabetes. Even arterial clots that lead to downstream tissue necrosis (e.g., stroke) can be prevented in some cases with statins and aspirin (Spector 2016). These therapies have reduced stroke in America from the second leading cause of death several decades ago to the fifth leading cause now (Benjamin et al. 2018). Put simply, the preventive approach focusing on well-known risk factors with effective, safe, and inexpensive generic drugs is a medical triumph. At present, a substantial part of the continuing incidence of stroke is due to medication non-compliance and lack of access of many patients who would benefit from preventive therapy. To improve medication compliance and access, pharmacologists need to work with additional health professionals. For instance, as in other areas of therapy, long acting anti-hypertensive drugs would almost certainly improve compliance and outcomes. Consider denosumab for the prevention of osteoporosis and fractures; this drug requires only one subcutaneous shot every 6 months. In fact, a highly durable RNAi inhibitor of PCSK9 for lowering serum cholesterol which potentially requires one subcutaneous injection every 6 months is being developed (Fitzgerald et al. 2017).

Of course, there are many brain diseases (e.g., Parkinson's disease, Alzheimer's disease, and manic depression), where the cause is not known, and thus, a good pharmacological target is not apparent. However, it is unlikely that transplantation or neurogenesis stimulation is the answer. Instead, the cause must be found and preventive therapy would be best, as in AIDS or the vitamin deficiency syndromes (e.g., pellagra and Wernicke's; niacin and thiamine deficiency, respectively).

## Conclusion

There is little convincing evidence to support the existence of functional AN in the human brain. There is a dramatic decrease in brain AN from salamanders (huge) to rodents (substantial) through monkeys (much less) to minimal or none in humans (Fig. 1). The notion that certain brain disorders are due to lack of human AN (e.g., depression and manic-depressive psychosis) is unlikely. The failure of stem-cell transplantation in Parkinson's disease suggests that even if new neurons were somehow introduced in adult human brain, they would have difficulty properly integrating into and functioning in extant brain circuitry. Since there are disorders (e.g., Alzheimer's disease, Parkinson's disease) in which particular neuronal populations are steadily lost, the challenge is to find methods to preserve them. Adult neuronal networks are composed of neurons that are born sequentially during development and slowly mature according to genetic and environmental cues. Once established,

those circuits, although plastic to some extent, cannot be dramatically altered. Hence, we suggest that more emphasis should be placed on the retention and preservation of the health of existing neurons, instead of the introduction of new ones.

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## Compliance with ethical standards

**Conflict of interest** The authors declare no conflict of interest.

**Ethical statement** The author declares that the manuscript is in complete compliance with the ethical standards of *Brain Structure and Function*.

**Ethical approval and inform consent** This is a review article and no procedures of any kind were performed on any animals or humans by the authors themselves. This review article is in accordance with general ethical standards of scientific conduct and scientific writing.

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