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Published in:
The Journal of neuroscience : the official journal of the Society for Neuroscience

DOI:
10.1523/JNEUROSCI.6003-08.2009

2009

Link to publication

Citation for published version (APA):

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Neural Stem and Progenitor Cells Retain Their Potential for Proliferation and Differentiation into Functional Neurons Despite Lower Number in Aged Brain

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Neurogenesis in the subventricular zone (SVZ), which gives rise to new neurons in the olfactory bulb, continues throughout life but declines with increasing age. Little is known about how aging affects the intrinsic properties of the neural stem and progenitor cells (NSCs) in SVZ and the functional characteristics of their neuronal progeny. Here, we have compared the properties of NSCs isolated from embryonic lateral ganglionic eminence and adult and aged SVZ in mice using in vivo and in vitro systems, analyzed their gene expression profile, and studied their electrophysiological characteristics before and after differentiation into neurons. We show a loss of NSCs in SVZ from aged mice accompanied by reduced expression of genes for NSC markers, developmentally important transcription factors, and neurogenic factors. However, when isolated in vitro, the NSCs from SVZ of aged animals have capacity for proliferation and multilineage differentiation, including production of functional neurons, similar to that of NSCs in adult mice, albeit with lower efficacy. These properties are of major importance when considering therapeutic applications of neuronal replacement from endogenous NSCs in the injured, aged brain.

Introduction

Neural stem and progenitor cells (NSCs) in the lateral ganglionic eminence (LGE) and adult and aged subventricular zone (SVZ) ensure neurogenesis during embryonic development and throughout adult life. However, NSC proliferation and neuroblast formation in SVZ are decreased in old animals (Tropepe et al., 1997; Enwere et al., 2004). Aged mice show deficits in olfactory discrimination, most likely attributable to reduced neurogenesis (Enwere et al., 2004). The age-dependent decline of neurogenesis has been attributed to decreased proliferation and growth factor signaling (Tropepe et al., 1997; Enwere et al., 2004; Shetty et al., 2005), increased levels of corticosteroids (Cameron and McKay, 1999; Montaron et al., 1999, 2006), and senescence of neural progenitors (Molofsky et al., 2006). Whether there is a loss of NSCs in aged brain is not clear. Reports are conflicting regarding NSCs in the dentate gyrus (Shetty et al., 2005; Olariu et al., 2007; Hattiangady and Shetty, 2008). In SVZ, the number of NSCs was described to be unaltered in old animals (Tropepe et al., 1997), but studies of proliferation, neurosphere formation, and ultrastructure (Enwere et al., 2004; Maslov et al., 2004; Luo et al., 2006) have indicated decreased number of NSCs with age.

Although neurogenesis is reduced in old animals, it can be increased or even restored to adult levels by enriched environment (Kempermann et al., 2002), infusion of growth factors (Jin et al., 2003), or stroke (Jin et al., 2004; Darsalia et al., 2005). Seizure-induced enhancement of neurogenesis is not altered between 1 and 3 months of age (Gray et al., 2002), but a recent study (Rao et al., 2008) reported lack of increased neurogenesis in response to seizures in aged (24 months old) rats. Together, available data indicate that aged NSCs retain the ability to respond to extrinsic cues similar to NSCs in adult animals, which is important in the perspective of a potential future therapeutic use of neuronal replacement from endogenous NSCs in human neurodegenerative disorders (Lindvall and Kokaia, 2008), which mainly affect older people. Recently, Morgenstern et al. (2008) reported that despite a sharp decline in hippocampal neurogenesis in aged mice, the density of dendritic spines in the new neurons was similar to that observed in young adult animals. These findings suggest that the age-related mechanisms leading to reduced neurogenesis do not affect the morphological properties of the new neurons. However, very little is known about the influence of aging on the intrinsic properties of NSCs and the functional characteristics of their neuronal progeny.

The aims of the present study were twofold: First, to explore whether the age-related decline of neurogenesis is associated with changes in numbers and gene expression of NSCs in SVZ; second, to determine the intrinsic properties of NSCs and their progeny in aged compared with embryonic and adult mice. Using NSCs from embryonic LGE and adult and aged SVZ, we show...
that the age-dependent reduction in neurogenesis is accompanied by loss of NSCs and decreased expression of NSC markers and developmentally important transcription factors. However, the aged NSCs can proliferate, differentiate along all neural lineages, and produce functional neurons in vitro, similar to their adult counterparts, although with lower efficacy.

**Materials and Methods**

**Animals.** Embryonic (E13.5, day of vaginal plug detection designated as E0.5), adult (3–5 months), and aged (22–26 months) C57BL/6 (B&K Universal) and nestin-green fluorescent protein (GFP; gift from Dr. G. Enikolopov, Cold Spring Harbor Laboratory, Cold Spring Harbor, NY) mice were bred at Lund University Biomedical Center. All experimental procedures were approved by the Malmö–Lund ethical committee.

**Bromodeoxyuridine administration.** Mice were given bromodeoxyuridine (BrdU; 50 mg/kg, i.p.), dissolved in potassium PBS (KPBS), four times with 2 h interval, and were killed 2 h thereafter.

**Immunohistochemistry.** Animals received an overdose of sodium pentobarbital and were transcardially perfused with saline followed by ice-cold 4% paraformaldehyde (PFA) in 0.1 M KPBS. Brains were removed, postfixed overnight, and placed in 20% sucrose in 0.1 M phosphate buffer for 24 h. Coronal sections (30 μm) were cut on a freezing microtome and stored in cryoprotective solution. Free-floating sections were preincubated in 0.25% Triton X-100 in KPBS containing 5% appropriate secondary serum for 1 h. Sections were then incubated with primary antibodies overnight at 4°C. Primary antibodies included the following: rat anti-BrdU (1:100; Sigma), rabbit anti-phosphorylated histone 3 (p-H3; 1:400; Millipore), goat anti-doublecortin (Dcx; 1:400; Santa Cruz Biotechnology), rabbit anti-Sox2 (1:500; Millipore Bioscience Research Reagents), and rabbit anti-Gsh2 (1:2500; gift from Dr. Kenneth Campbell, University of Cincinnati, Cincinnati, OH). For Brdu staining, DNA was denatured in HCl for 30 min at 65°C. For Sox2 staining, heat-induced (microwave) epitope retrieval in citrate buffer, pH 6.0, was used. After overnight incubation, sections were rinsed and incubated in darkness for 2 h with Cy3-conjugated donkey anti-rat, donkey anti-rabbit, and donkey anti-goat antibodies (1:200; Jackson ImmunoResearch). Nestin–GFP + cells were visualized by endogenous GFP fluorescence and counterstained using TO-PRO-3 nuclear marker (Invitrogen). For double-label immunohistochemistry with Brdu and Sox2, free-floating sections were denatured in HCl and incubated with both primary antibodies overnight as described above. Sections were then rinsed and incubated in darkness for 2 h with Cy3-conjugated donkey anti-rat and biotinylated donkey anti-rabbit antibodies. Biotinylated antibodies were visualized by incubating sections with streptavidin–Alexa 488 (Invitrogen) for 2 h at room temperature. For Brdu and Dcx double-label immunohistochemistry, sections were incubated first with goat anti-Dcx antibody overnight. Then sections were incubated with biotinylated donkey anti-goat, streptavidin–Alexa 488, fixed in PFA for 20 min at room temperature, and BrdU immunostaining was performed. After final washing, sections were mounted on glass slides and coverslipped with glycerol-based mounting medium. When needed, autofluorescence elimiter reagent (Millipore Bioscience Research Reagents) was used to reduce lipofuscin autofluorescence in aged tissue.

**Microscopical analysis.** All assessments were performed by a blinded observer. Immunostaining was examined using an Olympus BX61 fluorescence light microscope. Briefly, four coronal sections throughout the SVZ, located 0.74, 0.5, 0.14, and 0.02 mm anterior to bregma (Paxinos and Watson, 1997), were analyzed bilaterally. Stereological estimations of the optical fractionator method (Gundersen and Jensen, 1987) using the optical dissector constituting a 5-μm-thick fraction of the total section thickness. Because of their very low numbers in the aged brain, all p-H3 + and Gsh2 + cells were counted in the same sections as for the stereological estimations, and the total number of cells was calculated by multiplying with the total number of sections. Percentage of Sox2 + and Dcx + cells colabeled with Brdu was estimated in ~100 cells in high magnification fields in the SVZ in four sections per brain.

**Tissue dissection, cell isolation, and flow cytometry.** Animals were deeply anesthetized using halothane and killed with cerebral dislocation. Embryos or brains were removed and placed in ice-cold L-15 (Invitrogen). The brains of adult and aged animals were cut into 1-mm-thick coronal sections using a dissecting microscope. The LGEs of each hemisphere were dissociated in basic medium in 37°C for 15 min, and mechanically dissociated into a single-cell suspension. The SVZ lining the lateral ventricle of adult and aged animals was dissected from the sections and enzymatically dissociated in HBSS with 0.15 μM Heps, 5.4 mg/ml d-glucose, 1.33 mg/ml Trypsin (Invitrogen), 80 U/ml DNase, 0.7 mg/ml hyaluronidase, and 0.2 mg/ml kynurenic acid (Sigma) at 37°C for 30 min. The cell suspension was purified on a sucrose gradient (30% sucrose in 0.5× HBSS) followed by a BSA gradient (4% BSA in Earle’s balanced salt solution with 1× Heps). The GFP + cells from LGE, SVZ, or cultures were sorted on a fluorescence-activated cell sorter, FACS DiVa (Becton Dickinson). An initial gate based on forward and side scatter was set to exclude debris and cell aggregates, 7AAD was used to exclude dead cells. A sorting gate was set around the main GFP + population, at least 1 log higher than GFP − controls. Sorted cells were processed for cell culture or RNA isolation as described below.

**Cell culture.** Neurospheres were grown in DMEM/F12 supplemented with B27, 0.6% glucose (Sigma), 2 mM Glutamun, 1.125% sodium bicarbonate, 15 mM Heps, and 0.05 mg/ml Gentamycin (Basic Medium). For proliferative conditions, 20 ng/ml epidermal growth factor, 10 ng/ml basic fibroblast growth factor (BD Systems), and 2 μg/ml Heparin (Sigma) was added (proliferative medium). For neurosphere assays, cells were grown at 10 cells/μl in uncoated cell culture flasks for 7 d. For counting, spheres were transferred to 48-well plates with grids. Bulk cultures were grown at 20–50 cells/μl. Neurospheres were fed every other day and passed every week using accutase (PAA Laboratories).

For differentiation and electrophysiological measurements, neurospheres were plated in proliferative medium overnight on poly-γ-lysine (PDL) laminin (Sigma)-coated coverslips or chamberslides (Nunc). After overnight incubation (proliferative state), medium was changed to basic medium supplemented with N2 and 1% FBS and cells were left to differentiate for 7–14 d (differentiated state). All reagents were from Invitrogen, if not otherwise stated.

**For in vitro proliferation, cells were grown in 96-well plates, and total cell number was assessed at 0, 2, 4, and 6 d in vitro (DIV) using 3-(4,5-dimethylthiazyl)2,5-diphenyltetrazolium bromid (MTT) test (Roche) with absorbance at 450 nm.**

**For survival assays, cells were grown under proliferative or differentiation conditions on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips. Levels of apoptosis were determined using TUNEL staining on PDL/laminin-coated coverslips.
RNA isolation, reverse transcriptase-PCR, and quantitative PCR. Ten thousand GFP + cells were sorted directly into RT-Lysis buffer (Qiagen) and frozen at −80°C. RNA was isolated using the RNeasy micro kit (Qiagen) with DNase treatment according to the manufacturer’s guidelines. RNA was reverse transcribed using oligoDT primers and superscript-II (Invitrogen). Quantitative PCR (Q-PCR) was performed with TaqMan universal master mix and TaqMan Gene expression assays (supplemental Table 1, available at www.jneurosci.org as supplemental material). cDNA from 150 cells was used for each Q-PCR, and all experiments were run in triplicate. cDNA input was normalized to glyceraldehyde-3-phosphate dehydrogenase (GAPDH). Relative gene expression was calculated using the ∆∆CT method. Results are means from two to four independent experiments.

Electrophysiology. All measurements were performed in aCSF solution (pH = 7.4 and 290 mOsm) containing (in mM) 119 NaCl, 2.5 KCl, 1.3 MgSO4, 1 NaH2PO4, 26 NaHCO3, 2.5 CaCl2, and 11 glucose. The internal solution used to record intrinsic membrane properties and voltage-gated potassium channel kinetics contained (in mM) 122.5 K-gluconate, 17.5 KCl, 10 KOH–HEPES, 0.2 KOH–EGTA, 2 MgATP, 0.3 Na3GTP, and 8 NaCl (pH = 7.2 and 300 mOsm). Potassium currents were measured in the presence of 1 μM tetrodotoxin (TTX) to block sodium channels. When sodium channel kinetics was investigated, potassium was replaced by cesium in the internal solution, and 20 mM tetraethylammonium (TEA) and 2 mM 4-aminopyridine (4AP) were added to the bath solution. To measure EPCs, we used an internal solution, pH = 7.2, containing (in mM) 117.5 Cs-glucuronate, 17.5 CsCl, 10 CsOH–HEPES, 0.2 CsOH–EGTA, 2 MgATP, 0.3 Na3GTP, 8 NaCl, and 5 QX314, and 100 μM glutamate was added to the external solution. To measure IPSCs, Cs-glucuronate was replaced by CsCl, and 100 μM GABA was added to the external solution. In some experiments, the internal solution contained 0.05% biocytin for later identification of recorded cells.

All recordings were performed at room temperature using the whole-cell patch-clamp technique (Hamill et al., 1981). Pipettes were pulled from borosilicate glass and had a resistance of 3–6 MΩ when filled with internal solution. Data were filtered at 2.9 kHz and sampled at 10 kHz with EPC9 amplifier (HEKA Electronik) and stored on a G4 Macintosh computer. Off-line analysis was performed using IgorPro (version 5.02, WaveMetrics) and FitMaster (version 2.05, HEKA Electronik). Leak currents were subtracted online. Capacitive transients were compensated using the circuitry of the patch-clamp amplifier. Cell capacitance was measured shortly after establishing the whole-cell configuration. Subsequently, the current-clamp configuration was used to measure resting membrane potential. The input resistance was calculated as the slope of the linear fit to the current-voltage curve resulting from 500 ms current injections of −15 to 0 pA in 5 pA increments. Current densities were expressed as the ratio between maximal current amplitude and whole-cell membrane capacitance (pA/pF) at given voltage depolarizations. Action potentials (APs) were elicited by 500 ms current injections in 10 or 20 pA increments, and threshold, amplitude, duration, and afterhyperpolarization of the first AP were analyzed. Duration of AP was measured between the fast upstroke and downstroke at the 50% level of maximal amplitude.

Total potassium currents (IK(DR+A)) were elicited by a voltage protocol comprising 200 ms pre-pulses of −110 mV followed by 200 ms voltage steps from −70 to +60 mV in 10 mV increments. Delayed rectified potassium currents (IK(DR)) were isolated using a −40 mV pre-pulse to inactivate A-type potassium channels (IK(A)). IK(DR) was obtained subtracting IK(DR+A) for IK(DR+A) + IK(A) was obtained subtracting IK(DR) for IK(DR+A). Measured currents (I) were divided by current maxima (Imax) to obtain normalized values. Blockers affinities were estimated using 20 mM TEA or 2 mM 4AP and a pre-pulse of −110 mV followed by a depolarizing step at +40 mV. Sodium currents were elicited by 10 ms voltage steps from −60 to +60 mV in 10 mV increments. To characterize the voltage dependence of inactivation, the conductance (G) at each command voltage was put in relation to the maximal conductance Gmax. The conductance was calculated as G = I/(E − Erev), where G is the conductance, I the measured current, E the applied voltage, and Erev the reversal potential for Na+. Using Nernst equation, we calculated the reversal potential for Na+ to be +74 mV. The inactivation kinetics were studied using a prepulse protocol with conditioning steps to membrane potentials between −110 and −40 mV for 200 ms followed by a 10 ms voltage step at 0 mV.

Occurrence of functional glutamate and GABA receptors was assessed...
by adding 100 μM glutamate or 100 μM GABA to the bath solution for 15–20 s by switching the perfusion medium. Excitatory and IPSCs were measured in the presence of 100 μM picrotoxin (PTX) and 10 μM 2,3-dihydroxy-6-nitro-7-sulfonyl-benzo[f]quinoxaline (NBQX) together with 50 μM 3-AP, respectively. In all voltage-clamp experiments, the holding potential was −70 mV. Liquid junction potentials were not compensated.

**Statistical analysis.** Comparisons were performed using one-way ANOVA followed by Fisher’s post hoc test or repeated-measures ANOVA. Data are presented as means ± SEM, and differences are considered significant at p < 0.05.

**Results**

**Reduction in cell proliferation and neuroblast formation and loss of stem/progenitor cells in aged subventricular zone**

We initially investigated in vivo cell proliferation in the SVZ of adult and aged mice. To label all proliferating cells over an 8 h period, BrdU was injected four times, 2 h apart, and mice were perfused 2 h thereafter. Sections were stained against BrdU and the proliferation marker p-H3, which visualizes cells in active mitosis at the time of perfusion. We found 65 and 79% decrease in the number of p-H3+ and BrdU+ cells, respectively, in aged compared with adult SVZ (Fig. 1A, B). These findings with two different markers show that SVZ cell proliferation is markedly decreased in aged animals.

Next, we analyzed the formation of neuroblasts in the adult and aged SVZ. Sections were stained with an antibody against Dcx, a microtubule-associated protein transiently expressed in neuroblasts up to 4 weeks after their formation (Brown et al., 2003). We found 49% reduction in the number of Dcx+ cells in SVZ of aged animals compared with adult ones (Fig. 1C). Thus, also the formation of neuroblasts has declined in the SVZ of aged animals.

To explore whether the reduced cell proliferation and neuroblast formation in SVZ of old animals were attributable to loss of NSCs, we analyzed the expression of markers that have been widely used to identify NSCs. The intermediate filament nestin (Lendahl et al., 1990) and the sry homeobox transcription factor Sox2 (Ellis et al., 2004) are expressed in NSCs both in the embryo and the adult SVZ. The homeodomain transcription factor Gsh2 is expressed in NSCs in LGE and adult SVZ, which give rise to striatal projection neurons and/or olfactory bulb interneurons, respectively (Stenman et al., 2003).

The number of nestin–GFP+, Sox2+, and Gsh2+ cells were determined in the adult and aged mouse SVZ. We detected reduced number of GFP+ cells in the SVZ of aged nestin–GFP mice (Fig. 1D). The number of nestin–GFP+ cells was 50% lower in aged compared with adult nestin–GFP mice (Fig. 1D). Strikingly, the number of Sox2+ cells was decreased by 49% (Fig. 1E) and the number of Gsh2+ cells by 96% in the SVZ of aged mice (Fig. 1F). Together, these data indicate that aging leads to a loss of NSCs in the SVZ.

To identify which populations of cells in the SVZ decrease their proliferative activity during aging, we analyzed sections double stained for BrdU and markers for NSCs and neuroblasts, Sox2 and Dcx, respectively. The percentage BrdU+/Sox2+ and BrdU+/Dcx+ cells of the total number of Sox2+ and Dcx+ cells in SVZ was reduced, by 38 and 58%, respectively, in aged compared with adult animals (Fig. 1G). These findings together with the decreased numbers of Sox2+ and Dcx+ cells indicate that NSCs and neuroblasts contribute to the age-related reduction of cell proliferation in SVZ.

**Figure 2.** Aging causes reduction of NSC and neurogenic factor gene expression in SVZ stem/progenitor cells. A, B, Gene expression levels for stem/progenitor cell markers (nestin, Sox2, Musashi1), developmentally important transcription factors (Gsh2, Er81), and neurogenic factors (Dlx2, Mash1) measured using quantitative PCR in primary (A) or expanded (B) nestin–GFP+ cells sorted from embryonic LGE, and adult and aged SVZ of nestin–GFP mice. Data are presented as relative expression normalized to GAPDH levels and compared with embryonic LGE levels (set at 1). Means from two to four individual experiments. ND, Not detected.

**Reduction in neural stem cell and neurogenic factor gene expression in aged subventricular zone**

To identify differences in intrinsic properties between NSCs from animals of various ages, we sorted nestin–GFP+ cells from embryonic LGE and from adult and aged SVZ tissue. We isolated mRNA and by quantitative PCR compared the gene expression levels of known stem cell markers and neurogenic factors (Fig. 2A). Nestin gene expression decreased to 40 and 20% of embryonic levels in adult and aged animals, respectively, whereas Sox2 expression was 50 and 41% higher (Fig. 2A). The RNA-binding protein Musashi 1 (Msi1) is expressed in NSCs, both during embryonic development and in adulthood (Sakakibara et al., 1996; Sakakibara and Okano, 1997). Expression of Msi1 was 60 and 50% lower and of Gsh2 30 and 70% lower in adult and aged animals, respectively, compared with embryonic levels (Fig. 2A). In the embryonic brain, the transcription factor Er81 is expressed in progenitors giving rise to olfactory interneurons, separating them from progenitors expressing Isl1, which generate striatal projection neurons. Er81 continues to be expressed in the postnatal SVZ, whereas Isl1 expression is lost. We found the expression of Er81 to be almost eightfold higher in adult SVZ NSCs compared with embryonic LGE cells. Interestingly, the expression of Er81 decreased by 32% in aged compared with adult cells (Fig. 2A). In accordance to previous findings (Parmar et al., 2003), where no Isl1 protein expression could be found in the SVZ, we detected gene expression of Isl1 only in embryonic cells.
(data not shown). Thus, there is a switch in the fate of nestin+ cells, from production of striatal projection neurons in the embryonic brain to generation of olfactory bulb interneurons in the adult brain, the latter process declining with age.

The proneural transcription factor Dlx2 is expressed in the embryonic LGE and in transit amplifying cells and neuroblasts in the adult mouse SVZ (Doetsch et al., 2002). Gene expression of Dlx2 was similar in adult and embryonic NSCs but was 42% lower in aged compared with adult cells (Fig. 2 A). Also, the expression of another proneural factor, Mash1, was reduced but in both adult and aged cells by 40 and 50%, respectively (Fig. 2 A). To summarize, the most consistent pattern in our gene expression data is a decline in the levels of NSC markers, developmentally important transcription factors, and proneural genes in the aged SVZ.

To explore to what extent the changes in gene expression in SVZ NSCs during aging are attributable to environmental influences, we performed the same gene expression analysis on in vitro-expanded cells (Fig. 2 B). Interestingly, the lower gene expression for nestin, Sox2, Gsh2, and Er81 in aged compared with adult primary SVZ cells was no longer observed after expansion. Expression of Gsh2 was lost in expanded cells from all ages (Fig. 2 B). Moreover, Sox2 and Er81 gene expression, which was higher in primary adult and aged SVZ cells compared with embryonic LGE cells, was similar or lower after expansion. Conversely, the lower Msi1 gene levels in primary adult and aged SVZ cells were not detectable in expanded cells. These data show that several of the differences in gene expression between embryonic, adult, and aged NSCs observed in primary cells disappear during expansion and, therefore, most likely are attributable to environmental factors rather than intrinsic NSC properties.

Aged neural stem cells proliferate slower but generate neurospheres and differentiate similar to adult ones after in vitro expansion

We assessed neurosphere formation from NSCs in embryonic LGE and SVZ of adult and aged brain. The yield of primary neurospheres was markedly lower from both adult and aged SVZ than from LGE (61 and 69% reduction, respectively) (Fig. 3 A). Aged SVZ gave rise to 22% fewer primary neurospheres than adult SVZ. Moreover, although secondary neurosphere formation did not differ between embryonic LGE and adult SVZ, it was 28% less in aged compared with adult tissue. Interestingly, after expansion of neurospheres from embryonic LGE and adult or aged SVZ for 10 passages, we could not detect any differences between the groups in the number of generated neurospheres (Fig. 3 A). We then wanted to investigate the neurosphere-forming capacity of enriched NSCs isolated from embryonic LGE and adult and aged SVZ, and, therefore, sorted GFP+ cells from nestin–GFP mice. In accordance with the findings in nonsorted, wild-type cells (Fig. 3 A), we found that the yield of primary neurospheres was 77 and 80% lower from sorted adult and aged cells, respectively, compared with embryonic NSCs (Fig. 3 B). Surprisingly, we found no differences between the groups in secondary neurosphere formation or in number of neurospheres after 10 passages (Fig. 3 B). When we plated GFP− cells under the same conditions as GFP+ cells, they failed to form even single neurospheres (data not shown). Together, our findings indicate that SVZ in adult and, in particular, aged mice contains fewer NSCs with neurosphere-forming capacity compared with embryonic LGE. However, after expansion and sorting, NSCs from all age groups have the same potential for neurosphere formation.

Next, we compared the in vitro proliferation capacity of NSCs from the different groups. Cells from embryonic LGE and adult and aged SVZ proliferated and were expanded for at least 15 passages without any obvious signs of senescence (data not shown). The rate of proliferation was measured at 0, 2, 4, and 6 DIV using the MTT assay (Fig. 4 A). Repeated-measures ANOVA revealed progressive increase of cell numbers with time in all three groups. However, the magnitude of the increase differed between the groups with embryonic showing the highest and aged NSCs the lowest rate of proliferation.

We also assessed the apoptotic death of NSCs of different origins. Under proliferating conditions, we found no differences in the number of apoptotic, TUNEL+ cells between neurospheres from embryonic LGE and adult and aged SVZ (Fig. 4 B). Under differentiating conditions, there was a trend toward increased apoptosis with age, and the number of TUNEL+ cells was significantly higher with aged compared with embryonic NSCs (Fig. 4 B).

One possible explanation to the age-related decline in neurogenesis could be that aged NSCs go into senescence. To test this hypothesis, we studied the expression of several cyclin-dependent kinase inhibitors implicated in senescence and cell cycle arrest in sorted primary nestin–GFP+ cells from the LGE, and adult and aged SVZ, as well as in neurospheres expanded from the corresponding cells. Furthermore, we measured SA-β-gal activity, an indicator of replicative senescence, in NSCs from all ages expanded as neurospheres. In contrast to previous reports (Molofsky et al., 2006), we could not detect p16INK4A in primary cells of any age group (Fig. 4 C). The expression of p19ARF was higher in adult and aged SVZ cells, compared with embryonic SVZ cells. Expression of p21Cip1 was undetectable in aged SVZ cells but sevenfold higher in adult compared with embryonic SVZ cells. Interestingly, levels of p27Kip1 increased progressively with age (Fig. 4 C). After in vitro expansion of NSCs as neurospheres, expression of p16INK4A was detected in all age groups (Fig. 4 D). The levels in adult and aged cells were 3.7- and 3.6-fold higher, respectively, than in embryonic cells. In contrast, levels of p19ARF were lower in adult and aged cells. Expression of p21Cip1 was 20 and 80% higher, whereas levels of p27Kip1 was 21 and 25% lower
cultures were used for quantification. SA-GFP hoc test. Scale bar, 30 μm. ND, Not detected.

**Figure 4.** Aging causes reduction of NSC proliferation and survival during differentiation in vitro. Neurosphere cells were grown under proliferative and differentiating conditions. A, Relative number of cells assessed by MTT test. Proliferation was significantly higher in embryonic compared with adult NSCs and in adult compared with aged NSCs at 0, 2, 4, and 6 DIV. Means ± SEM; *p < 0.05, repeated-measures ANOVA. Four independent cultures were used for quantification. B, Survival assessed by numbers of TUNEL + apoptotic cells under proliferative (at 4 DIV) or differentiating conditions (at 7 DIV). Means ± SEM. Four independent cultures were used for quantification. C, D, Expression of genes involved in senescence in primary (C) and expanded (D) Nestin–GFP + sorted cells. E, F, Photomicrograph (E) and percentage (F) of neurosphere cells, from the different age groups, expressing SA-β-gal. Mean ± SEM. Four independent cultures were used for quantification. *p < 0.05, one-way ANOVA with Fisher’s post hoc test. Scale bar, 30 μm. ND, Not detected.

in adult and aged cells, respectively, compared with embryonic levels (Fig. 4D). Analysis of SA-β-gal activity revealed a progressive increase with age (Fig. 4E, F). The percentage of neurosphere cells expressing SA-β-gal increased from 5% in embryonic cells to 15% in adult and 30% in aged cells (Fig. 4F). Together, our data indicate that NSCs in SVZ exhibit an age-dependent increase of gene expression for senescence markers in vivo and in vitro and of age-dependent replicative senescence in vitro.

To evaluate the differentiation potential of NSCs from various ages, we withdrew growth factors and added serum and N2 supplements to the culture medium. Cells were let to mature for 1 week and then stained with markers for neurons (β-III-tubulin), astrocytes (GFAP), and oligodendrocytes (CNPase). In all age groups, the majority of cells differentiated into GFAP + cells with mature astrocytic morphology (Fig. 5A). A small fraction of the cells became CNPase + with typical oligodendrocytic morphology. All age groups also generated β-III-tubulin + cells with multipolar processes and morphological characteristics of neurons (Fig. 5A). Thus, the multipotency and capacity to generate cells of all three neural lineages is maintained in aged brain. We assessed the neurogenic potential of these neurosphere cultures in more detail by quantifying the number of β-III-tubulin + cells formed from the different age groups. We found that embryonic LGE cells gave rise to 125 and 101% more β-III-tubulin + cells than adult and aged SVZ cells, respectively. However, there were no differences between adult and aged NSCs in the formation of neurons (Fig. 5B). To explore if NSCs of all age groups generated different subtypes of neurons, we differentiated for 2 weeks and cultures then stained for GABA and glutamate. We found that embryonic, adult, and aged neurospheres generated cells with mature neuronal morphology and that a fraction of these cells were GABA + or glutamate + (Fig. 5C).

**Figure 5.** Differentiation of neurosphere cells into the three neural lineages. A, Relative number of cells assessed by MTT test. Proliferation was significantly higher in the proliferating adult compared with embryonic and aged cells. After differentiating for 2 weeks, mean % cell proliferation was 69 ± 1% in embryonic (n = 13), 62 ± 2% in adult (n = 12), and 54 ± 3% in aged (n = 13). B, Survival assessed by numbers of TUNEL + apoptotic cells under proliferative (at 4 DIV) or differentiating conditions (at 7 DIV). Means ± SEM. C, D, Expression of genes involved in senescence in primary (C) and expanded (D) Nestin–GFP + sorted cells. E, F, Photomicrograph (E) and percentage (F) of neurosphere cells, from the different age groups, expressing SA-β-gal. Mean ± SEM. Four independent cultures were used for quantification. *p < 0.05, one-way ANOVA with Fisher’s post hoc test. Scale bar, 30 μm. ND, Not detected.

Adult and aged neural stem cells generate functional neurons in vitro, albeit at reduced frequency compared with embryonic cells. Whole-cell patch-clamp recordings showed that proliferating and differentiating cells from all ages exhibited similar input resistance and resting membrane potential with no significant differences related to age or degree of cell differentiation (Table 1).

The membrane capacitance was significantly higher in the proliferating adult compared with embryonic cells. After morphological differentiation, the membrane capacitance almost doubled in all groups. As determined in the current-clamp mode of the patch-clamp technique, none of the proliferating embryonic (n = 16) or aged cells (n = 13) fired AP in response to current injections, whereas 2 of 13 adult cells fired immature AP (Fig. 6A). In 7 of 20 adult proliferating NSC cells, we identified a small inward, fast-inactivating current, of 200–500 pA, which seemed to have the characteristics of sodium currents. Moreover, 2 of 21 aged, proliferating SVZ cells had large sodium currents of 1 and 6 nA, respectively (data not shown). After 2 weeks of differentiation, 88% of the neurons with mature morphology from embryonic NSCs fired mature and 12% fired
immediate APs during membrane depolarization (Fig. 6B). Aging lead to decreased percentage of differentiated cells that fired APs and increased the percentage that did not, from 58 and 26%, respectively, in adult group to 26 and 60%, respectively, in aged group (Fig. 6B). The percentage of differentiated cells firing immature APs was similar in all groups (12–16%) (Fig. 6B).

The properties of APs did not differ between cells from various groups (Table 1). They were completely blocked by 1 μM TTX, and were elicited with current injections of 10–20 pA. However, the relatively long duration and low amplitude of action potentials indicate that the cells most likely are still not functionally fully mature. A distinctive feature of differentiated neurons was the presence of a fast-inactivating inward current with the characteristics of Na+ current (Fig. 7C). The current showed a peak between −20 and 10 mV and an average maximum density of 97 ± 19 pA/pF (n = 6), 105 ± 16 pA/pF (n = 7), and 181 ± 37 pA/pF (n = 7) in neurons differentiated from embryonic, adult, and aged NSCs, respectively. We did not find any differences in activation and inactivation kinetics of Na+ channels between neurons derived from the different groups (data not shown).

To identify other ion channels underlying active membrane properties, we performed voltage-clamp experiments. In proliferating cells, we identified the outward current (Fig. 7A) composed of two K+ channel populations: a non-inactivating delayed-rectified current (I_{K(DR)}) that constitutes most of the total K+ current, and a fast-inactivating A-type K+ current (I_{K(A)}). There was higher percentage in adult and aged NSCs compared with embryonic ones for A-type current (Fig. 7B) but not for delayed-rectified current (Fig. 7C). Interestingly, total K+ current density increased with age of NSC source (Fig. 7D). Thus, the total K+ current density calculated at +60 mV increased with age, being 239 ± 18 pA/pF (n = 19), 296 ± 15 pA/pF (n = 20), and 386 ± 36 pA/pF (n = 21) in LGE, adult, and aged NSCs, respectively. Differentiating cells of all age groups exhibited robust outward K+ currents consisting of I_{K(DR)} and I_{K(A)}. In contrast to proliferating conditions, we observed that the distribution of I_{K(DR)} and I_{K(A)} and total K+ current density were similar in the three groups (data not shown). However, we found decreased K+ current density of 191 ± 23 pA/pF (n = 12), 207 ± 27 pA/pF (n = 9), and 155 ± 22 pA/pF (n = 9) in neurons differentiated from embryonic, adult, and aged NSC cells, respectively, compared with the proliferating conditions.

To further characterize K+ channels, we used two specific blockers, TEA and 4AP. Addition of TEA to proliferating NSCs of all age groups blocked 60% of total potassium current (I_{K(DR+)}). Of remaining K+ current, 60% (25% of total) was blocked by
Differentiated cells of all age groups, channels are sensitive to both TEA and 4AP, and a minor portion together, our data indicate that more than 50% of total potassium blocked in embryonic, adult, and aged NSCs (Fig. 8). If AP was added first, 4AP. If AP was added first, ~70% of the K\(^+\) current was blocked. The rest constituted a very slow-inactivating current of which 50% (15% of total) was blocked by TEA. With both TEA and 4AP in bath solution, more than 80% of the total K\(^+\) current was blocked in embryonic, adult, and aged NSCs (Fig. 8A–C). Together, our data indicate that more than 50% of total potassium channels are sensitive to both TEA and 4AP, and a minor portion to either TEA or 4AP.

In contrast to proliferating cells, after addition of TEA to differentiated cells of all age groups, ~60–70% of fast-inactivating K\(^+\) current remained (Fig. 8D), of which 70% (45% of total) were blocked by 4AP. Application of 4AP to the total current blocked ~20% of the K\(^+\) current. The remaining 80% constituted a very slow-inactivating current of which 75% (60% of total K\(^+\) current) was blocked by TEA. Using both TEA and 4AP, ~80% of total K\(^+\) current was blocked in neurons differentiated from embryonic, adult, and aged NSCs (Fig. 8D). These results indicate that during differentiation of NSCs from all ages, the composition of membrane K\(^+\) channels changed, being blocked by either TEA or 4AP.

We finally investigated whether the neurons generated from NSCs obtained from various ages had functional glutamate and GABA receptors. Application of 100 µM glutamate (Fig. 8G) or GABA (Fig. 8H) induced inward currents in neurons differentiated from NSCs of all ages. The glutamate-induced current was only partially blocked by 10 µM NBQX and 50 µM D-AP5.

**Discussion**

Here, we show age-dependent loss of NSCs and reduction in SVZ neurogenesis, accompanied by decreased gene expression of NSC markers, developmentally important transcription factors, and neurogenic factors. However, when the NSCs are expanded *in vitro*, they behave similarly, although with lower efficacy, as their counterparts from adult animals with respect to sphere-forming

### Table 1. Passive electrical properties of cells from embryonic, adult, and aged neural stem cell cultures under proliferation and differentiation conditions

<table>
<thead>
<tr>
<th></th>
<th>Embryonic Proliferation</th>
<th>Embryonic Differentiation</th>
<th>Adult Proliferation</th>
<th>Adult Differentiation</th>
<th>Aged Proliferation</th>
<th>Aged Differentiation</th>
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<tbody>
<tr>
<td>Capacitance (pF)</td>
<td>11.9 ± 0.7 (n = 30)</td>
<td>23.3 ± 2.3 (n = 27)</td>
<td>15.1 ± 0.8 (n = 27)</td>
<td>19.1 ± 2.1 (n = 17)</td>
<td>13 ± 0.1 (n = 24)</td>
<td>23.2 ± 1.8 (n = 15)</td>
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<tr>
<td>Input resistance (GΩ)</td>
<td>1.7 ± 0.3 (n = 16)</td>
<td>1.8 ± 0.3 (n = 19)</td>
<td>1.2 ± 0.4 (n = 13)</td>
<td>2 ± 0.3 (n = 13)</td>
<td>2.9 ± 0.8 (n = 8)</td>
<td>2 ± 0.4 (n = 12)</td>
</tr>
<tr>
<td>Resting membrane potential (mV)</td>
<td>−69.7 ± 2.8 (n = 20)</td>
<td>−62.5 ± 1.8 (n = 25)</td>
<td>−65.1 ± 4.5 (n = 15)</td>
<td>−65.6 ± 2.8 (n = 16)</td>
<td>−56.8 ± 5.4 (n = 14)</td>
<td>−66.7 ± 2.3 (n = 12)</td>
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<tr>
<td>Action potential threshold (mV)</td>
<td>−31.3 ± 1.2 (n = 23)</td>
<td>−31.3 ± 1.2 (n = 23)</td>
<td>−29.3 ± 1.5 (n = 16)</td>
<td>−29.3 ± 1.5 (n = 16)</td>
<td>−29.4 ± 1.0 (n = 11)</td>
<td>−29.4 ± 1.0 (n = 11)</td>
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<tr>
<td>Action potential amplitude (mV)</td>
<td>49.9 ± 2.2 (n = 23)</td>
<td>49.9 ± 2.2 (n = 23)</td>
<td>45.2 ± 3.9 (n = 16)</td>
<td>45.2 ± 3.9 (n = 16)</td>
<td>39.8 ± 2.3 (n = 11)</td>
<td>39.8 ± 2.3 (n = 11)</td>
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<tr>
<td>Action potential duration (ms)</td>
<td>4.5 ± 0.3 (n = 23)</td>
<td>4.5 ± 0.3 (n = 23)</td>
<td>5.0 ± 0.5 (n = 16)</td>
<td>5.0 ± 0.5 (n = 16)</td>
<td>3.9 ± 0.4 (n = 11)</td>
<td>3.9 ± 0.4 (n = 11)</td>
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<tr>
<td>Action potential after-hyperpolarization (mV)</td>
<td>14.6 ± 1.2 (n = 23)</td>
<td>14.6 ± 1.2 (n = 23)</td>
<td>13.2 ± 1.8 (n = 16)</td>
<td>13.2 ± 1.8 (n = 16)</td>
<td>14.3 ± 2.1 (n = 11)</td>
<td>14.3 ± 2.1 (n = 11)</td>
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capacity, proliferation, multilineage differentiation, and production of electrophysiologically functional neurons.

Reduction of SVZ cell proliferation has been reported previously by several groups (Tropepe et al., 1997; Enwere et al., 2004; Maslov et al., 2004; Luo et al., 2006). However, the changes in the number of cells expressing the cell cycle markers used in these studies could also reflect age-related alterations in the duration of different phases of the cell cycle as observed in adult SVZ cells after stroke (Zhang et al., 2006, 2008). We used both BrdU labeling and the mitotic marker, p-H3, which is specifically expressed by cells undergoing mitotic division. Importantly, our finding of reduced number of p-H3 cells for the first time directly shows that the number of NSCs undergoing actual mitotic division at a given time is decreased severalfold in the aged SVZ. Furthermore, by using double staining with BrdU and phenotypic markers, we demonstrated that the mitotic activity of Sox2+ NSCs and Dcx+ neuroblasts is decreased in the aged compared with the adult brain.

Although various mechanisms underlying the reduced neurogenesis in the aged brain have been suggested (Tropepe et al., 1997; Cameron and McKay, 1999; Montaron et al., 1999, 2006; Enwere et al., 2004; Shetty et al., 2005; Molofsky et al., 2006), it has been unclear whether the SVZ NSC pool diminishes with age. Together, our experimental evidence strongly indicates that aging leads to a loss of NSCs in the SVZ. First, we showed for the first time that the numbers of nestin-, Sox2-, and Gsh2-immunoreactive cells were decreased in the SVZ of aged animals. Second, we found overall reduction in gene expression of several NSC markers, i.e., nestin, Sox2, Gsh2, and Er81 in GFP cells sorted from the SVZ of nestin–GFP mice, indicating a loss of NSCs within the nestin–GFP population or reduced gene expression in the NSCs themselves. Third, we observed a decrease in primary neurosphere formation in aged animals, partially reflecting the number of NSCs in the SVZ, and neurosphere formation was rescued in these animals by enriching for NSCs from SVZ using cell sorting in nestin–GFP mice. These findings are at variance with two previous studies in which no change in number of NSCs was found in the aged rat hippocampus or mouse SVZ (Tropepe et al., 1997; Hattiangady and Shetty, 2008). However, our findings are supported by the work of Luo et al. (2006), who detected a loss of transient amplifying cells (so-called C-cells) in the aged mouse SVZ using electron microscopy.

The NSCs from embryonic LGE proliferated faster and gave rise to higher numbers of more mature and functional neurons than adult and aged cells. These findings identify embryonic LGE as the most suitable cell source for transplantation purposes. However, also NSCs from adult and aged mice proliferated extensively in vitro, despite the higher levels of SA-β-gal, increased gene expression of several factors involved in cell cycle arrest, and decreased proliferation with age, differentiated into cells of all lineages, including functional neurons. Thus, when taken out of their normal environment and challenged with growth factors,

Figure 7. Expression of K+ currents in proliferating cells. A, Total K+ current (I\( _{\text{K(DR+A)}} \)) (left), non-inactivating delayed-rectified current (I\( _{\text{K(DR)}} \)) (middle), and a fast-inactivating A-type K+ current (I\( _{\text{K(A)}} \)) (right) obtained by voltage steps from \(-70\) to \(+60\) mV after a hyperpolarizing prepulse to \(-110\) mV in proliferating embryonic, adult, and aged NSCs. I\( _{\text{K(DR)}} \) was separated from total I\( _{\text{K(A)}} \) using a depolarizing prepulse to \(-40\) mV. Calibration: 1 nA, 50 ms. B, K+ current density at \(+60\) mV. C, Mean ± SEM. Mean proportion I\( _{\text{K(A)}} \) (C) and I\( _{\text{K(DR)}} \) (D) out of total current plotted against the applied voltage steps for embryonic, adult, and aged NSCs, respectively. In C, comparison with repeated-measures ANOVA. n = 19, 20, 21 in embryonic, adult, and aged, respectively.
NSCs from adult and aged SVZ respond similarly. These findings are consistent with in vivo data showing that NSCs in aged SVZ increased their proliferation and neuroblast formation in response to brain damage or infusion of growth factors (Jin et al., 2003, 2004; Darsalia et al., 2005).

Although our findings indicate that the environment in the aged brain impairs NSC survival, proliferation, and differentiation, we also obtained evidence suggesting that age-dependent, intrinsic changes in the NSCs themselves play a role. Thus, we detected reduced expression of NSC markers, developmentally important transcription factors, and neurogenic factors. Also, aged NSCs did not generate as many functional neurons as adult cells despite the same in vitro conditions. This could be partly explained by the increased mutational load described in aged

Figure 8.  K^+ channel sensitivity to TEA and 4AP in embryonic, adult, and aged NSCs. A. Percentage of the current left unblocked by 20 mM TEA, 2 mM 4AP, or 20 mM TEA and 2 mM 4AP in proliferating embryonic, adult, and aged NSCs. Means ± SEM; n = 11, 10, 7 and 9, 7, 17, 13 in embryonic, adult, and aged, respectively. B, C, Representative traces of outward K^+ currents activated by a depolarizing voltage step from the resting membrane potential to +40 mV in proliferating NSCs. The currents are blocked by 20 mM TEA (B) or 2 mM 4AP (C). Calibration: 500 pA, 25 ms. D, Percentage of the current left unblocked by 20 mM TEA, 2 mM 4AP, or 20 mM TEA and 2 mM 4AP in neurons differentiated from embryonic, adult, and aged NSCs. Means ± SEM; n = 5, 5, 5 and 6, 7, 8, 10, 11 in embryonic, adult, and aged, respectively. E, F, Representative traces of outward K^+ currents activated by a depolarizing voltage step from the resting membrane potential to +40 mV in neurons differentiated from aged NSCs. The currents are blocked by 20 mM TEA (E) or 2 mM 4AP (F). Calibration: 250 pA, 25 ms. G, Effect of 100 μM glutamate (left) or glutamate together with 10 μM NBQX and 50 μM D-AP5 (right) on neurons differentiated from embryonic, adult, and aged NSCs. Calibration: for embryonic and aged, 100 pA, 50 ms; for adult cells, 20 pA, 50 ms. n = 6, 8, and 6 in embryonic, adult, and aged, respectively. H, The effect of 100 μM GABA (left) or 100 μM GABA together with 10 μM PTX (right) on neurons differentiated from embryonic, adult, and aged NSCs. Calibration: for embryonic and aged, 500 pA, 50 ms; for adult cells, 40 pA and 50 ms. n = 8, 7, and 5 in embryonic, adult, and aged, respectively.
neurospheres, which most likely affects normal function (Bailey et al., 2004).

Previous studies have shown similar dendritic spine density (Morgenstern et al., 2008) and input resistance (Couillard-Despres et al., 2006) of newly formed neurons in adult and aged hippocampus. Before our study, no electrophysiological recordings or functional assays had been performed on neurons derived from NSCs of aged SVZ either in vivo or in vitro. Here, we report an extensive electrophysiological characterization of both proliferating NSCs and neurons differentiated from NSCs in SVZ of embryonic, adult, and aged mice. We find that the NSCs derived from embryonic LGE, and adult and aged SVZ, generate functional neurons with similar electrophysiological properties. Under proliferating conditions, the cells of all age groups have similar passive membrane properties, characterized by hyperpolarized resting membrane potential and high input resistance. We found slightly higher membrane capacitance in adult and aged NSCs and also in the cellular membrane, high density of voltage-activated potassium channels, mainly delayed rectified channels, which are indistinguishable by their passive and active electrophysiological properties. Differences regarding the expression of K+ channels between the age groups. One characteristic feature of the differentiated neuron populations be- tween the age groups. Although K+ channel populations were similar to the ones observed in the proliferating stage, TEA and 4AP sensitivity changed during differentiation. Thus, the channels were less sensitive to TEA and even to a lesser extent to 4AP, indicating that the K+ channel population of differentiated neurons is constituted from both Kv2 and Kv3 families (Grissmer et al., 1994; Coetzee et al., 1999). However, TEA and 4AP sensitivity was similar in neurons differentiated from cells of the three age groups. One characteristic feature of the differentiated neuron populations was the presence of a high density of Na+ channels. Mature neurons with complex morphologies were generated from NSCs of all age groups. After 2 weeks of differentiation, these neurons were firing repetitive APs that could be abolished by TTX and were able to respond to applications of glutamate or GABA. Importantly, despite the decrease in the number of differentiated neurons with age, we could not detect any changes in the functional electrophysiological properties of the individual neurons. Thus, our findings indicate that neurons differentiated from NSCs of embryonic LGE and adult and aged SVZ have similar potential for functional integration.

Together, our findings show that aging is associated with a loss of NSCs in the SVZ but that the remaining cells retain their ability for proliferation and differentiation if removed from the aged environment. Moreover, aged NSCs generate functional neurons, which are indistinguishable by their passive and active electrophysiological properties. This knowledge is potentially useful as a basis for future attempts to use endogenous neurogenesis for brain repair in elderly patients with acute or chronic neurodegenerative disease.

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