Tangential migration of neuronal precursors of glutamatergic neurons in the adult mammalian brain

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In a classic model of mammalian brain formation, precursors of principal glutamatergic neurons migrate radially along radial glia fibers whereas GABAergic interneuron precursors migrate tangentially. These migration modes have significant implications for brain function. Here we used clonal lineage tracing of active radial glia-like neural stem cells in the adult mouse dentate gyrus and made the surprising discovery that proliferating neuronal precursors of glutamatergic granule neurons exhibit significant tangential migration along blood vessels, followed by limited radial migration. Genetic birthdating and morphological and molecular analyses pinpointed the neuroblast stage as the main developmental window when tangential migration occurs. We also developed a partial "whole-mount" dentate gyrus preparation and observed a dense plexus of capillaries, with which only neuroblasts, among the entire population of progenitors, are directly associated. Together, these results provide insight into neuronal migration in the adult mammalian nervous system.

The nervous system is formed by migration of neuronal precursors and immature neurons to specific locations during development. The classic radial unit hypothesis of mammalian brain development postulates that in the developing neocortex, glutamatergic, excitatory, principal neurons migrate radially to form discrete information-processing columns of ontogenetic origin (1), whereas GABAergic, inhibitory, modulatory interneurons migrate tangentially across columns (2). Neurogenesis persists in the adult mammalian brain in two primary regions and is thought to follow the classic migration model (3, 4). In the subventricular zone (SVZ) of the lateral ventricles, new neurons generated from neural precursors migrate tangentially to the olfactory bulb to become GABAergic interneurons (5, 6). In contrast, in the subgranular zone (SGZ) of the dentate gyrus, new neurons generated from radial glia-like neural stem cells (RGLs) migrate radially into the granule cell layer to become principal glutamatergic granule neurons (7). Due to technical challenges, migratory patterns have only been examined at the cell-population level, and thus we still lack detailed information about the spatial relationship between individual precursors and their progeny in vivo. Both adult neurogenic niches are highly vascularized, and this property is hypothesized to play a critical role in adult neurogenesis (3). In both adult SVZ (8–10) and SGZ (11), proliferating progenitor cells are in close association with the vasculature, yet the functional role of the vasculature in the niche remains to be fully explored.

Contrary to the classic model, our recent clonal lineage tracing of individual quiescent RGLs showed tangential distribution of glutamatergic granule neurons with respect to their parental RGL in the adult dentate gyrus (12). We therefore systematically investigated the migration pattern and trajectory of these newborn cells. Using a clonal lineage-tracing approach that preferentially targets active RGLs in the adult mouse dentate gyrus, thereby birthdating their newborn progeny in vivo, we found significant tangential distribution of newborn neuroblasts from their parental RGL. Furthermore, neuroblasts directly contact the vascular network, suggesting an important function of blood vessels as a substrate for migration. Together, our results reveal a previously unidentified mode of glutamatergic neuronal migration under physiological conditions in the adult mammalian brain.

Results

Spatial Distribution of Clonally Related Newborn Neuronal Precursors with a Defined Birthdate. We used a tamoxifen-inducible Ascl1CreERT2 knockin mouse line (13) to sparsely label (~8–16 cells per dentate gyrus) and lineage-trace clones of neural precursors in the adult dentate gyrus (Fig. L4), following our previous strategy of clonally labeling RGLs using the NestinCreERT2 mouse line (12). At 1 d post-tamoxifen injection (1 dpi), labeled cells in the adult SGZ consisted of over 90% Nestin+GFAP+ RGLs and a few early intermediate progenitor cells (IPCs; n = 8 dentate gyrus), as assessed by morphology and clonal lineage tracing.

Significance

Mammalian brain development is a complex, ordered process whereby newborn neurons follow stereotyped migration modes to organize into specific patterns required for complicated neural circuit formation. Classically, principal excitatory neurons are thought to organize into radial columns that underlie the basic brain circuits, whereas inhibitory neurons disperse tangentially across these columns to modulate the principal circuits. These principles are thought to be fundamental to the genesis of the complex mammalian brain. Surprisingly, we found that precursors for excitatory principal neurons exhibit tangential migration in the adult mammalian brain. Our findings enrich our understanding of neurodevelopment and lay important conceptual groundwork for studies of brain plasticity, disease, and repair.


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and molecular marker expression (Table S1). Computational simulations (12) estimated that given eight initially labeled cells, any two labeled cells within 200 μm of each other had a 99% probability of belonging to the same clone (Fig. 1B). Thus, although few isolated IPCs were labeled at 1 dpi, the probability of their misassignment to an RGL clone was 1% or lower. Crucially, given our simulation results, we conservatively conducted all our analyses by excluding clones in which progeny were >150 μm away from their parent RGL, thus only focusing on clones with a maximum diameter of 300 μm.

In contrast to NestinCreERT2-based clonal lineage tracing of quiescent RGLs that exhibited stochastic activation over time (12), at 1 dpi the majority of Ascl1CreERT2-labeled RGLs (96 ± 2%, n = 10 dentate gyri) had just divided (with one adjacent progeny) or were in the process of cell division (Fig. 1A). Therefore, our current approach selectively targets activating RGLs, allowing, to our knowledge for the first time, birthdating of RGL progeny to track their development with high temporal precision. At 1 month posttamoxifen injection (mpi), we observed clones in which Prox1+NeuN+ mature dentate granule cells were distributed a long distance (more than 100 μm) away from their mother RGL along the SGZ plane (Fig. 1C and Movie S1). These results suggest, in contrast to the classic model, tangential migration of a glutamatergic neuronal subtype in the adult mammalian brain under physiological conditions.

Developmental Stage-Specific Tangential Distribution. Newborn cells may exhibit tangential distribution due to passive mechanisms such as displacement upon cell division. To provide further evidence for active tangential migration, we explored the dynamics of cell distribution by time-course analysis. At 3 dpi, labeled cells exhibited little displacement from each other, despite all being identified as highly proliferative IPCs (14) (Fig. 1D). In contrast, at 7 dpi there was significant distance along the tangential plane of the SGZ between labeled newborn neuronal progenitors and their parental RGLs, the distribution of which was very similar to that at 1–2 mpi (Fig. 1D). Only ~15% of labeled new neurons exhibited measurable radial displacement away from the SGZ into the granule cell layer even at 1 or 2 mpi (all <25 μm), yet among these cells ~47% spread more than 25 μm tangentially away from their parental RGL (Fig. 1E). The average tangential distance exceeded that of radial distance by almost 18-fold. Notably, radial migration was absent at 7 dpi, suggesting that tangential movement preceded any radial migration (Fig. 1E); no correlation between tangential and radial migration was observed among individual cells that exhibited both (Fig. 1E; R = 0.08, P = 0.85). Taken together, these results establish that tangential distribution of newborn progeny from their parental RGL could not have arisen by chance and instead represents the dominant mode of migration in the adult dentate gyrus.

We next asked whether tangential migration occurred during a specific developmental stage (or stages) during adult neurogenesis. Adult hippocampal neurogenesis proceeds in a stereotypic sequence—dividing RGLs give rise to IPCs, which in turn develop through neuroblast and immature neuron stages before becoming mature glutamatergic granule neurons (Fig. 2A) (4). Using morphological features and molecular markers to distinguish labeled cells at different developmental stages (Fig. 2B and Table S1), we found that neuroblasts, corresponding to 3- to 7-d-old DCX+ cells with a bipolar, elongated morphology, were the earliest cells in the developmental lineage that showed tangential distribution away from their parental RGL (Fig. 2B). On average, neuroblasts, immature neurons, and mature neurons exhibited similar tangential distances away from their parental RGLs (Fig. 2B). These results suggest that the majority of tangential migration occurs within the neuroblast stage during adult hippocampal neurogenesis.

Pattern and Magnitude of Tangential Distribution. A consequence of tangential migration of neuronal progeny is that individual clones disperse over areas beyond that of a radial unit, potentially allowing individual RGLs to modify a large domain of the hippocampal circuitry via neurogenesis. We visualized 2D SGZ projections of clones over time in 300-μm-square windows (Fig. S1; excluding clones with single RGLs that had not completed division). We also included non-RGL-containing clones only for 1–2 mpi, because in some clones RGLs became differentiated (12). Qualitatively, clones dispersed over large areas, strongly suggestive of widespread tangential migration (Fig. S1).
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tedentate gyrus, BrdU-incorporating cells are known to be in close

searched for potential substrates that could support tangential

explored heterotypic cellular interactions, such as migration in chains and

investigated a potential cellular mechanism underlying tangential

migrated in chains and 3D rendering of GFP

A

B

C

D

Fig. 2. Developmental stage-specific tangential distribution of newborn neuronal progeny. (A) Summary of molecular markers used for identification of each cell type during adult hippocampal neurogenesis. Newborn cells are generated from GFAP+Nestin+ RGLs that undergo asymmetrical neuronal di-

visions, which develop into Tbr2+ intermediate progenitor cells with short, multipolar processes within 3 d of birth. Within 3–7 d, newborn cells possess long, bipolar processes and elongated somas in a Tbr2+/DCX+ neuroblast stage before penetrating the granule cell layer and becoming a polarized Prox1+/DCX+ immature neuron with axon and dendrite. Over the next month, newborn cells mature into Prox1+/NeuN+ neurons with spiny dendrites and long axons that project to CA3. See also Table S1. (B) Summary of the tangential distance of each neural progeny from its parental RGL at each development stage. IN, immature neuron; N, mature granule neuron; NB, neuroblast. Values represent mean ± SEM (*P < 0.01; †P < 1; Wilcoxon rank-

sum test with Bonferroni correction for multiple comparisons; test statistic, z = −5.58 and −0.72, respectively). (C) Histogram and cumulative distribution plot of the maximum distance between clonally related neural progeny for all clones, including non–RGL-containing clones. See Fig. S1 for a 2D SGZ plane projection of representative clones (*P < 0.1; two-sample Kolmogorov–Smirnov test statistic: 0.38).

We next quantitatively determined the extent of tangential
distribution of neuronal progeny in each clone under 2D SGZ
projection. As opposed to measuring the distance of each
newborn progeny to its mother RGL as in Fig. 1, we instead mea-
sured the maximum distance among all newborn progeny within
each clone. The mean values were 43 ± 16 μm and 69 ± 19 μm,
whereas the maximum values were 149 μm and 210 μm at 7 dpi
and 1–2 mpi, respectively. The distribution of maximum dist-
tances was modestly shifted from 7 dpi to 1–2 mpi (Fig. 2C). The

considerable spacing among neural progeny supports our model
of neuronal migration as opposed to primarily motility of RGLs.

Vascular Substrate for Putative Tangential Migration. We then in-
vestigated a potential cellular mechanism underlying tangential
migration of neuronal precursors in the adult hippocampus in vivo. Tangential migration of GABAergic interneurons has been shown to be facilitated by a combination of homotypic and heterotypic cellular interactions, such as migration in chains and along existing axonal processes, respectively (15). We therefore searched for potential substrates that could support tangential migration during adult hippocampal neurogenesis. In the adult dentate gyrus, BrdU-incorporating cells are known to be in close

Fig. 3. Close association between tangentially migrating neuroblasts and blood vessels in the adult dentate gyrus. (A) Sample confocal images (Left) and 3D rendering (Far Right) of a GFP+ clone at 7 dpi with a neuroblast whose soma and tangential process closely associated with a CD31+ blood vessel. (B) Sample confocal image and 3D rendering of a GFP+ clone at 7 dpi containing a parental RGL and dispersed Tbr2+DCX+ neuroblasts (open arrowheads) in close association with CD31+ blood vessels (Movie S2). Note that CD31 and Tbr2 shared the same channel due to limited availability of channels but could be distinguished by different morphology (Tbr2, nuclear staining; CD31, tubular staining). (C) Sample confocal image and 3D rendering of a GFP+ clone at 7 dpi with a neuroblast whose tangential process extended along a blood vessel and contained polarized GM130+ Golgi apparatus (closed arrowheads) at its base, proximal to the cell soma. (D) Sample confocal image and 3D rendering of GFP+ neuroblasts at 7 dpi with polarized γ-tubulin+ centrosomes (closed arrowheads) and GM130+ Golgi apparatus (open arrowheads) and in close association with CD31+ blood vessels. (Scale bars, 10 μm.)
immature neurons (Fig. 6). Tangentially migrating neuroblasts may use the vasculature as a migration substrate, revealing an important role of this neurogenic niche component (Fig. 6).

Our finding of dominant tangential migration of neural precursors of principal glutamatergic dentate granule cells in the adult brain is a departure from the classic radial unit hypothesis of embryonic cortical development (1). Although early studies described tangential migration of presumed excitatory neurons in the embryonic cortex, they may have actually been describing tangentially migrating inhibitory neurons, which were discovered later (17). Interestingly, at least some rare (18), although usually transient (19, 20), populations of excitatory cortical neurons migrate tangentially in the developing brain. Tangential migration of neural precursors in the developing hippocampus has previously only been observed in early postnatal developmental stages, during which the hippocampus is still immature and marked by ongoing growth and development (21, 22). It may be that radial and tangential migration modes are not specific to neurons releasing a specific neurotransmitter. Indeed, in the cerebellum, inhibitory principal Purkinje neurons exhibit radial migration, whereas excitatory modulatory granule neurons migrate tangentially as well as radially (23). A very small number of glutamatergic modulatory interneurons are also generated in the adult SVZ and migrate tangentially to the olfactory bulb (24). Motor neurons in the spinal cord possess both radial and tangential migration (25). Additionally, neurons originating from the diencephalon can migrate tangentially to populate the amygdala, a telencephalic nucleus (26).

**Discussion**

In contrast to the prevailing model that glutamatergic neurons migrate radially whereas interneurons migrate tangentially during development, we demonstrated to our knowledge for the first time in the adult mammalian brain that neuroblast precursors of principal glutamatergic neurons exhibit significant tangential distribution away from their parental stem cells under physiological conditions. We revealed a two-step migration process during adult hippocampal neurogenesis in which significant tangential migration of neuroblasts is followed by limited radial migration of
Here we have described an example of tangential migration of neuronal precursors of excitatory principal neurons in the adult mammalian brain. The magnitude of migration observed, although significant and not due to cell displacement upon division, is much less than that of migrating SVZ neuroblasts or even radially migrating neocortical neurons. This may be due to differences in tissue architecture. Neuroblasts in the SVZ are born far from their target brain region and the neocortex is a multilayered structure, both requiring cells to travel longer distances to populate all target regions. By comparison, the dentate gyrus is a thin single-layer sheet whereby newborn neurons were presumed, before the present study, to be born, develop, and integrate all in the same location. Due to the limitation of single-color reporters in resolving different clones and the probabilities of clonality calculated from our simulations, we took an inherently conservative approach by imposing a 300-μm maximum for clone diameter. Our reported magnitude of tangential migration distance may therefore be an underestimation.

The vasculature has been proposed to be a crucial niche component for adult neurogenesis in both the SGZ and SVZ (8–11); however, its functional role is not well-understood. We showed that neuroblasts exhibit several features of migrating cells, such as polarized organelle distribution, and that they display a tight, direct association with the vasculature at cellular and ultrastructural levels. Although our data leave open the possibility of vasculature-independent tangential migration, our findings suggest that one function of the vascularized niche is to support migration of neuronal progeny away from their parental origin. Interestingly, GABAergic neurons generated in the adult SVZ have also been shown to migrate along blood vessels in the rostral migratory stream (27, 28) and toward injury sites following stroke (29). Thus, in the adult brain, the vasculature may serve as a common substrate for migration of different neuronal subtypes. This mechanism to support cell migration may also be important during early development, because vascular outgrowth often precedes neural outgrowth, and migrating cells can be found in regions lacking classic substrates such as glial fibers (30, 31).

We and others have previously only documented radial migration during adult hippocampal neurogenesis (32, 33), having been limited to population-level analyses that lacked lineage relationship information of individual stem cells and their progeny. Our current approach permits clonal analysis of activating adult dentate RGLs using the Ascl1creERT2 knockin mouse line (13) and direct examination of cell dispersion of individual newborn progeny with respect to their parental RGL. With the high temporal and spatial resolution our approach allows, we were able to directly compare tangential and radial migration of individual cells for the first time, to our knowledge. Interestingly, the two forms of migration appeared to be temporally segregated, with tangential migration preceding and exceeding radial migration. Together, these data suggest that independent mechanisms may regulate the two modes of migration. Future studies combining our clonal analysis with conditional gene inactivation could elucidate shared or independent molecular mechanisms governing different forms of migration.

We also developed a partial whole-mount preparation to optimize visualization of the SGZ niche and showed that the general population of neuroblasts is in close contact with the blood vessel network and exhibits polarized Golgi apparatus, characteristic of migrating cells. This preparation will allow for systematic characterization of the complete global architecture of the hippocampal niche in the future. Such a characterization could potentially reveal important motifs for understanding the underlying structure of the vascular network and migrating cells that we have observed. Ultimately, our technique can help broaden our knowledge of general principles of neurogenic niches, especially in combination with knowledge gained from similar approaches in the adult SVZ (8–10).

Our clonal analyses of activating radial glia-like neural stem cells reveal a previously unrecognized mode of neuronal migration during adult hippocampal neurogenesis. Given that adult neural stem cells are dynamically regulated by their local environment, including neuronal activity modulation (34, 35), our findings have significant implications for understanding how individual neural stem cells could impact the function of the existing microcircuitry via delivery of newborn neurons that are primed to undergo a critical period of enhanced plasticity (36). Our findings also provide a framework in which to investigate the aberrant migration of newborn granule cells under pathological conditions, such as epilepsy (37, 38). In addition, the identification of a novel mode of glutamatergic cell migration tangentially along the vasculature suggests a strategy for overcoming current challenges in targeting glutamatergic neurons to injured or degenerating regions for successful cell-transplantation therapy in the adult mammalian brain (39). Together, our results suggest that diverse modes of migration may be ubiquitous among different neuronal classes in the adult mammalian brain, providing important conceptual groundwork for future studies of neural development, plasticity, regeneration, repair, and brain disorders.

Materials and Methods

Animals and Tamoxifen Administration. Ascl1creERT2;Rosa-YFP; and Ascl1creERT2;Confetti reporter; mice and Nestin-GFP transgenic mice were used. All mice in the study were backcrossed to the C57BL/6 background for at least six generations. Animals were housed in a 14-h light/10-h dark cycle with free access to food and water. Tamoxifen (66.67 mg/mL; Sigma; T5648) was prepared in a 5:1 corn oil (Sigma)-to-ethanol mixture. To achieve sparse labeling for clonal analysis (8–16 clones per hemisphere), a single dose of tamoxifen (77.5 mg/kg for the Rosa-YFP reporter; 250 mg/kg for the Confetti reporter) was intraperitoneally injected into adult 8- to 10-wk-old male and female mice. Animal procedures were performed in accordance with institutional guidelines of the Johns Hopkins University School of Medicine.

Tissue Processing, Immunostaining, and Confocal Imaging. Serial 40-μm-thick coronal sections were prepared for immunohistochemistry as described (12). For partial whole-mount dentate gyrus SGZ preparations the whole hippocampus was dissected from the cortex and mounted dentate side-up on a frozen sliding microtome, and serial 50-μm horizontal sections were collected. All antibodies used are listed in SI Materials and Methods. Confocal images...
were taken to confirm cell identity according to immunohistological and morphological properties (Fig. 2A and Table S1).

**Image Processing and Data Analysis.** Clonal analysis and simulations to estimate the probability of two cells belonging to a clone from 3D reconstructed adult dentate gyrus were performed as previously described (12). Clones that spanned multiple serial sections were reconstructed using Reconstruct software (John C. Fiala, NIH, Bethesda). Aligned images were exported at full resolution for 3D visualization into Imaris (Bitplane). Cells and vasculature were identified, localized, and digitized into 3D space in Imaris according to distinct morphological and molecular markers (Fig. 2A and Table S1). Data were exported to Matlab (The MathWorks) for analysis. Distance measures at the clonal level were calculated from 7 hemispheres (five animals) at 3 dpi, 9 hemispheres (six animals) at 7 dpi, and 12 hemispheres (seven animals) at 1–2 mpi. For vasculature association analysis at the clonal level, all labeled clones and percentages of vasculature association were quantified and averaged across 10 hemispheres (five animals). For vasculature association analysis at the population level, all cells in a horizontal section from three Nestin-GFP animals were quantified. Simulated vasculature associations (Fig. 5D) were generated by randomly placing cells in the same 2D SGZ space, as estimated by a third-degree polynomial plane fit. One hundred iterations of the simulation were performed for each cell type.

**Electron Microscopy Analysis.** Adult C57BL/6 female mice were stereotactically injected with retrovirus expressing GFP (40) and processed 2 d after injection. Fifty-micrometer-thick coronal sections containing clearly labeled newborn progeny in close association with the CD31+ vasculature were sequentially incubated in biotinylated anti-GFP, avidin–biotin–peroxidase complex, 3,3-diaminobenzidine (DAB) peroxidase, and 1% osmium tetroxide before being dehydrated and lifted into resin. Seventy-nanometer-thick serial sections were collected onto copper grids and contrasted with a lead stain, and serial images were taken using a Philips CM10 transmission electron microscope to create 3D reconstructions with Fiji imagej software (41).

**Statistical Analysis.** Statistical analysis methods can be found in *SI Materials and Methods.*

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Supporting Information

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SI Materials and Methods

Animals and Tamoxifen Administration. Ascl1CreERT2; Rosa-YFP<sup>+</sup> and Ascl1CreERT2; Confetti<sup>+</sup> mice were generated by crossing Ascl1CreERT2 [Ascl1<sup>tm1.1(Cre/ERT2)jejo</sup>]; knockin line; Jackson Laboratory] (13) with Rosa-YFP<sup>+</sup> [B6.129X1-Gt(Rosa)26So<sup>tm1(CAG-tdTomato)Egfp</sup>]; Jackson Laboratory] and Confetti<sup>+</sup> [B6.129P2-Gt(Rosa)26So<sup>tm1(CAG-BruIN2.EGFP)Egjo</sup>]; Jackson Laboratory] mice, respectively. We used two different reporters in the present study to validate reporter independence of cell-migration phenomena. Nestin-GFP transgenic mice were from Grigori Enikolopov, Cold Spring Harbor Laboratory, Cold Spring Harbor, NY (16). All mice in the study were backcrossed to the C57BL/6 background for at least six generations. Animals were housed under a 14-h light/10-h dark cycle and had free access to food and water. Tamoxifen (66.67 mg/mL; Sigma; T5648) was prepared in a 5:1 corn oil (Sigma)-to-ethanol mixture. To achieve sparse labeling for clonal analysis (∼8–16 clones per hemisphere), a single dose of tamoxifen (77.5 mg/kg for the Rosa-YFP<sup>+</sup> reporter; 250 mg/kg for the Confetti<sup>+</sup> reporter) was intraperitoneally injected into adult 8- to 10-wk-old male and female mice. Animals were analyzed at 1, 3, or 7 d posttamoxifen injection (dpi) or 1–2 mo posttamoxifen injection (mpi). All animal procedures were performed in accordance with institutional guidelines of the Johns Hopkins University School of Medicine.

Tissue Processing, Immunostaining, and Confocal Imaging. Animals were perfused with cold 4% paraformaldehyde [wt/vol; in 0.1 M phosphate buffer (PB), pH 7.4], and cryoprotected with 30% sucrose (wt/vol). Serial 40-μm-thick coronal brain sections were cut on a frozen sliding microtome (Leica; SM2010R) for immunohistology as previously described (12). For partial “whole-mount” dentate gyrus SGZ preparations, the whole hippocampus was dissected from the cortex and mounted dentate side-up on a frozen sliding microtome, and serial 50-μm horizontal sections were collected. Antibodies diluted in TBS (Tris buffered saline) with 0.05% Triton X-100 and 3% (vol/vol) donkey serum were used against GFP (Aves Labs, chicken, 1:500; Rockland, goat, 1:500), Tbr2 (Abcam, rabbit, 1:250), CD31 (BD Biosciences, rat, 1:500), DCX (Santa Cruz, goat, 1:100), GFAP (Millipore, mouse, rabbit, 1:1,000), GM130 (BD Biosciences, mouse, 1:250), Nestin (Aves Labs, chicken, 1:500), γ-tubulin (BioLegend, rabbit, 1:500), Proxl (Abcam, rabbit, 1:500), and NeuN (Millipore, guinea pig, 1:500). Nestin and NeuN antigens were retrieved by incubating brain sections in 1× target retrieval solution (Dako) at 68 °C for 20 min, followed by a 10-min cooling to room temperature. Serial sections from the entire dentate gyrus were first immunostained for GFP to allow identification of prospective clone-containing sections on an epifluorescence microscope (Zeiss; Axiovert 200M). Labeled clone-containing sections were taken for further processing and confocal imaging at 40× on a Zeiss LSM 710 confocal microscope using multitrack or spectral linear unmixing “online fingerprinting” configurations (Carl Zeiss).

Image Processing and Data Analysis. Clonal analysis of adult hippocampal neural stem cells and their progeny was performed as previously described (12). Simulations to estimate the probability of two cells belonging to a clone, based upon numbers of initially labeled cells per hemisphere, were performed from 3D reconstructed dentate gyri also as previously described (12). For data acquisition, clones that spanned multiple serial sections were reconstructed using Reconstruct software (John C. Fiala, NIH, Bethesda). All aligned images were exported at full resolution for 3D visualization to Imaris (Bitplane) and analyzed. For illustrative purposes, some cells and features were volume-rendered using the Surface module in Imaris. Distance measurements were collected from 7 hemispheres (five animals) at 3 dpi, 9 hemispheres (six animals) at 7 dpi, and 12 hemispheres (seven animals) at 1–2 mpi. The Spots module in Imaris was used to digitize cell-nucleus locations in the 3D space and to code cell-type classifications according to distinct morphological and molecular markers (Fig. 2A and Table S1). The Clipping Plane module in Imaris was used to estimate the local SGZ plane for 2D clone projections. Spots and clipping planes were exported to Matlab (The MathWorks) for further distance measurements, calculations, and analyses. The distance between an RGL and a given newborn progeny in the local SGZ plane was calculated for tangential distance measurements (Figs. 1 and 2). The shortest distance between a given labeled newborn cell and the SGZ was calculated for radial distance measurements (Fig. 1 C and E). Cells at the dentate gyrus genu were excluded from analysis due to the potential ambiguity of tangential versus radial migration. Clones with multiple RGLs were also excluded due to ambiguity in distinguishing the cell of origin.

For vasculature association analysis at the clonal level, cells were classified as one of three groups: (i) those with no CD31<sup>+</sup> vasculature contact, (ii) those with their soma directly contacting a CD31<sup>+</sup> blood vessel, and (iii) those that contacted the CD31<sup>+</sup> vasculature only through one or more processes. All labeled clones and the percentage of their vasculature association were quantified and averaged across 10 hemispheres (five animals). For vasculature association analysis at the population level, all cells in a horizontal section from three Nestin-GFP animals were quantified and the results were averaged across animals. Locations of cell nuclei were digitized according to GFP, Tbr2, and DCX expression using Spots. Blood vessels were digitized using the Filament Tracer module in Imaris. Three-dimensional coordinates were exported to Matlab and the distance between cells and vasculature was measured. Cells with nuclei within 7 μm of a blood vessel center line were considered to be in contact with the blood vessel. Simulated vasculature associations (Fig. 5D) were performed by randomly placing cells in the same 2D SGZ space, as estimated by a third-degree polynomial plane fit. One hundred iterations of the simulation were performed for each cell type.

Electron Microscopy Analysis. Adult C57BL/6 female mice were stereotactically injected with an engineered murine oncoretrovirus expressing GFP as previously described (40). Mice were processed 2 d after injection and transcardially perfused with 4% paraformaldehyde (wt/vol) with 0.1% glutaraldehyde (vol/vol; Sigma) in 0.1 M PB (pH 7.4). Coronal sections (50-μm-thick) were sectioned on a vibrating microtome (Leica Biosystems; VT1200S). Sections were immunostained in PBS for CD31 and examined using an epifluorescence microscope to identify GFP<sup>+</sup> retrovirally labeled newborn cells. Sections containing clearly labeled newborn progeny in close association with the CD31<sup>+</sup> vasculature were placed in cryoprotectant [25% sucrose (wt/vol), 10% glycerol (vol/vol), in 0.05 M PB] for 2 h and freeze–thawed three times in liquid nitrogen. After three washes in PBS and then 0.5% BSA in 0.1 M PB (vol/vol; BSA-C; Aurion) to block the tissue, sections were incubated overnight in the primary antibody (biontinated rabbit anti-GFP; Vector Laboratories; BA-0702; 1:1,000) in 0.1% BSA-C in 0.1 M PB, shaking at room temperature. Sections were washed out of the primary antibody...
with one wash in 0.1% BSA-C in 0.1 M PB and three washes in PBS before incubation for 90 min in avidin–biotin–peroxidase complex (Vector Laboratories; ABC Elite). After three washes in PBS and then Tris buffer, a 3,3-diaminobenzidine (DAB) peroxidase reaction was performed (Vector Laboratories Kit; 4 min). The sections were then washed three times in Tris buffer and three times in PBS and left overnight at 4 °C. The next day, sections were washed in 0.1 M PB in glass vials and postfixed in 1% osmium tetroxide in 0.1 M PB for 30 min. They were then dehydrated in ascending concentrations of ethanol (two 50% ethanol washes, one 70% wash with 1% uranyl acetate, one 95% wash, and two 100% washes) and three washes in acetone before being lifted into resin (Fluka; Durcupan ACM) and left overnight at room temperature. The resin was then warmed to decrease its viscosity, and sections were placed onto microscope slides with coverslips before the resin was cured at 65 °C over 3 d. Locations of clearly labeled newborn progeny in close association with CD31+ blood vessels were identified at the light microscopic level and followed through to the electron microscopic level. Serial sections (70-nm-thick) were collected onto Formvar-coated, single-slot copper grids and contrasted with a lead stain (15 min). Sections were then examined using a Philips CM10 transmission electron microscope, and serial images of the labeled structures were captured using a digital camera (Olympus; Morada SIS). These images were used to create 3D reconstructions with Fiji ImageJ software (41), and individual images were processed using Adobe Creative Suite software.

**Statistical Analysis.** Statistical analysis was performed with one-way ANOVA (with Tukey post hoc test), one-tailed unpaired Student’s t test, two-sample Kolmogorov–Smirnov test, or Wilcoxon rank-sum test (with Bonferroni correction for multiple comparisons, where applicable), as indicated in the text and figures. When applicable, validation of normality and homogeneity of variance assumptions was performed for all compared data groups using a Shapiro–Wilk test and Monte-Carlo simulation, respectively. All statistical analyses were performed in Origin software (OriginLab) or Matlab.

**Fig. S1.** Tangential distribution among newborn neural progeny at 3 dpi, 7 dpi, and 1–2 mpi in the adult dentate gyrus. Two-dimensional SGZ plane projection of all RGL-containing clones containing at least IPCs at (A) 3 dpi, (B) 7 dpi, and (C) 1–2 mpi, plotted in a 300-μm-square window. RGLs are represented as open circles; neural progeny are represented as closed circles. Different time points or developmental stages are encoded by color. Clones that lacked RGLs at 1–2 mpi are also plotted in gray boxes and included in the histogram in Fig. 2C.
Table S1. Summary of the morphological markers used in the current study for identification of each cell type during adult hippocampal neurogenesis

<table>
<thead>
<tr>
<th>Cell type</th>
<th>Molecular markers</th>
<th>Morphology</th>
</tr>
</thead>
<tbody>
<tr>
<td>Radial glia-like cell (RGL)</td>
<td>Nestin, GFAP</td>
<td>RGL has its soma in the SGZ, radial process in the GCL and ML, nuclear cleavage with only one GFAP+ nucleus</td>
</tr>
<tr>
<td>with neurogenic division</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Intermediate progenitor cell (IPC)</td>
<td>Tbr2 (GFAP-)</td>
<td>IPC is spherical with short, multipolar processes in the SGZ</td>
</tr>
<tr>
<td>Neuroblast (NB)</td>
<td>DCX, Prox1</td>
<td>NB has elongated soma, bipolar processes</td>
</tr>
<tr>
<td>Immature neuron (IN)</td>
<td>DCX, Prox1, NeuN</td>
<td>IN has soma in the apical GCL, thin dendrite through the GCL</td>
</tr>
<tr>
<td>Mature neuron (N)</td>
<td>Prox1, NeuN (DCX-)</td>
<td>N has multibranched spiny dendrites in the ML and axon to CA3</td>
</tr>
</tbody>
</table>

Movie S1. Widespread distribution of clonally related cells in the adult dentate gyrus. Shown is a sample 3D reconstruction of a GFP-labeled clone showing the distribution of adult-born dentate granule neurons away from their parental RGL at 1 mo posttamoxifen induction in an Ascl1CreERT2;Rosa-YFPf/+ mouse. The projection view of the same clone is shown in Fig. 1C.

Movie S2. Close association of neuroblasts with blood vessels in the adult dentate gyrus. Shown is a confocal image of immunostaining for GFP, Tbr2, CD31, and DCX in an Ascl1CreERT2;Rosa-YFPf/+ mouse at 7 dpi. Note the close association of neuroblasts on blood vessels in the SGZ, away from their parental RGL. The projection confocal image is shown in Fig. 3B. Also shown is a surface rendering of all cells and their association with the vasculature. GFP, green; Tbr2 and CD31, red; DCX, blue.

Movie S1

Movie S2
**Movie S3.** Partial whole-mount dentate gyrus SGZ preparation. Shown is a confocal image of a whole-mount preparation section from an adult Nestin-GFP mouse hippocampus immunostained with GFP (green) and Nestin (red). A schematic illustration of this preparation is shown in Fig. 5A.

**Movie S4.** Association of neuroblasts with the vasculature in a whole-mount dentate gyrus SGZ preparation. Shown is a confocal image of a whole-mount preparation section from an adult Nestin-GFP mouse hippocampus immunostained with GFP (green), Tbr2 and CD31 (red), and DCX (blue). Also shown are surface renderings of representative neuroblasts on blood vessels in the adult SGZ.