

1 Subventricular zone cytogenesis provides trophic support for neural repair
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16 **Abstract**

17 Stroke enhances proliferation of neural precursor cells within the subventricular zone (SVZ) and
18 induces ectopic migration of newborn cells towards the site of injury. Here we characterize the
19 identity of cells arising from the SVZ after stroke and provide insight into their function by
20 uncovering a mechanism through which they facilitate neural repair and functional recovery.
21 Using genetic lineage tracing, we show that SVZ-derived cells that migrate towards stroke-
22 induced cortical lesions in mice are predominantly undifferentiated precursors, suggesting that
23 the main function of post-injury cytogenesis is not cell replacement. We find that SVZ-derived
24 cells are a unique cellular source of trophic factors that instruct neural repair. Chemogenetic
25 ablation of neural precursor cells or conditional knockout of VEGF in the adult neural stem cell
26 lineage impairs neuronal and vascular reparative responses and worsens functional recovery after
27 stroke. In addition, normal aging markedly diminishes the cytogenic response to stroke, resulting
28 in worse functional recovery. Therapeutic replacement of VEGF in peri-infarct cortex is
29 sufficient to induce neural repair and functional recovery in mice with arrested cytogenesis.
30 These findings indicate that the SVZ cytogenic response following brain injury is a source of
31 trophic support that drives neural repair and recovery.

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36 **Introduction**

37 Limited recovery of function occurs after damage to the central nervous system.
38 Consequently, stroke and other forms of brain injury often cause long-lasting disabilities.
39 Following stroke, remodeling of residual tissue surrounding the site of injury is thought to
40 underlie recovery. For example, extensive plasticity of neural circuits and blood vessels occurs in
41 peri-infarct regions and these processes are associated with functional improvement (Brown et
42 al., 2007; Clark et al., 2019; Tennant et al., 2017; Williamson et al., 2020). Repair processes are
43 mediated by interactions across disparate cell types (Brown et al., 2007; Clark et al., 2019; Joy et
44 al., 2019; Kim et al., 2018; Williamson et al., 2021). However, the intercellular interactions that
45 orchestrate repair and recovery remain to be completely defined. A more complete understanding
46 of the mechanisms that govern neural repair could inform development of new treatment
47 strategies.

48 Cytogenesis, the formation of new cells, is limited in the adult mammalian brain. The
49 subventricular zone (SVZ) is one of a small number of regions that contains multipotent neural
50 stem and progenitor cells (collectively referred to here as precursors) that generate new neurons
51 and glia in adulthood (Doetsch et al., 1999; Garcia et al., 2004). Normally, the predominant
52 progeny arising from the SVZ are new neurons that migrate towards the olfactory bulb and
53 integrate into existing olfactory circuitry. However, injuries such as stroke markedly increase
54 SVZ precursor proliferation and induce ectopic migration of SVZ-derived cells towards the site
55 of injury (Arvidsson et al., 2002; Lagace, 2012; Li et al., 2010; Ohab et al., 2006; Parent et al.,
56 2002; Williamson et al., 2019). Past studies of this process after brain injury have largely focused
57 on neurogenesis — the formation of new neurons and their localization in peri-infarct regions
58 (Arvidsson et al., 2002; Parent et al., 2002). In general, experimental manipulations that increase

59 post-stroke neurogenesis are associated with enhanced functional recovery (Lagace, 2012). The
60 prevailing view has been that cell replacement, especially neuron replacement, by SVZ
61 precursors allows for partial brain regeneration and consequently improved function. However,
62 recent findings indicate that new neurons poorly integrate into existing circuits in peri-infarct
63 regions and receive little synaptic input (Kannangara et al., 2018), making their functional
64 importance unclear (cf. Liang et al., 2019). Moreover, other studies have found that glia
65 outnumber neurons among migrating SVZ-derived cells, but the entire population of SVZ-
66 derived cells has yet to be comprehensively characterized (Benner et al., 2013; Faiz et al., 2015;
67 Li et al., 2010). Overall, the identity and functional importance of new cells that arise from the
68 SVZ after injury are not well understood.

69 Here we investigate the SVZ cytogenic response following cortical photothrombotic
70 strokes in mice. Our goals were to characterize the types of cells produced by the SVZ after
71 stroke and mechanistically understand the role of SVZ cytogenesis in stroke recovery. We use
72 indelible lineage tracing to phenotype SVZ-derived cells that migrate towards the site of injury.
73 Unexpectedly, we find that the majority of these cells are undifferentiated precursors, suggesting
74 that cell replacement is limited. Reducing cytogenesis impairs motor recovery after stroke, at
75 least in part due to deficits in neuronal and vascular plasticity. With gain- and loss-of-function
76 manipulations, we show that VEGF produced by SVZ-derived cells drives repair and functional
77 recovery. These findings identify SVZ cytogenesis as a source of trophic support that facilitates
78 neural repair. Thus, our study demonstrates a mechanism other than cell replacement by which
79 endogenous neural precursor cells contribute to repair and recovery in the injured central nervous
80 system.

81

82 **Results**

83 *Cells arising from the subventricular zone after stroke are predominantly quiescent precursors*
84 *and astrocytes*

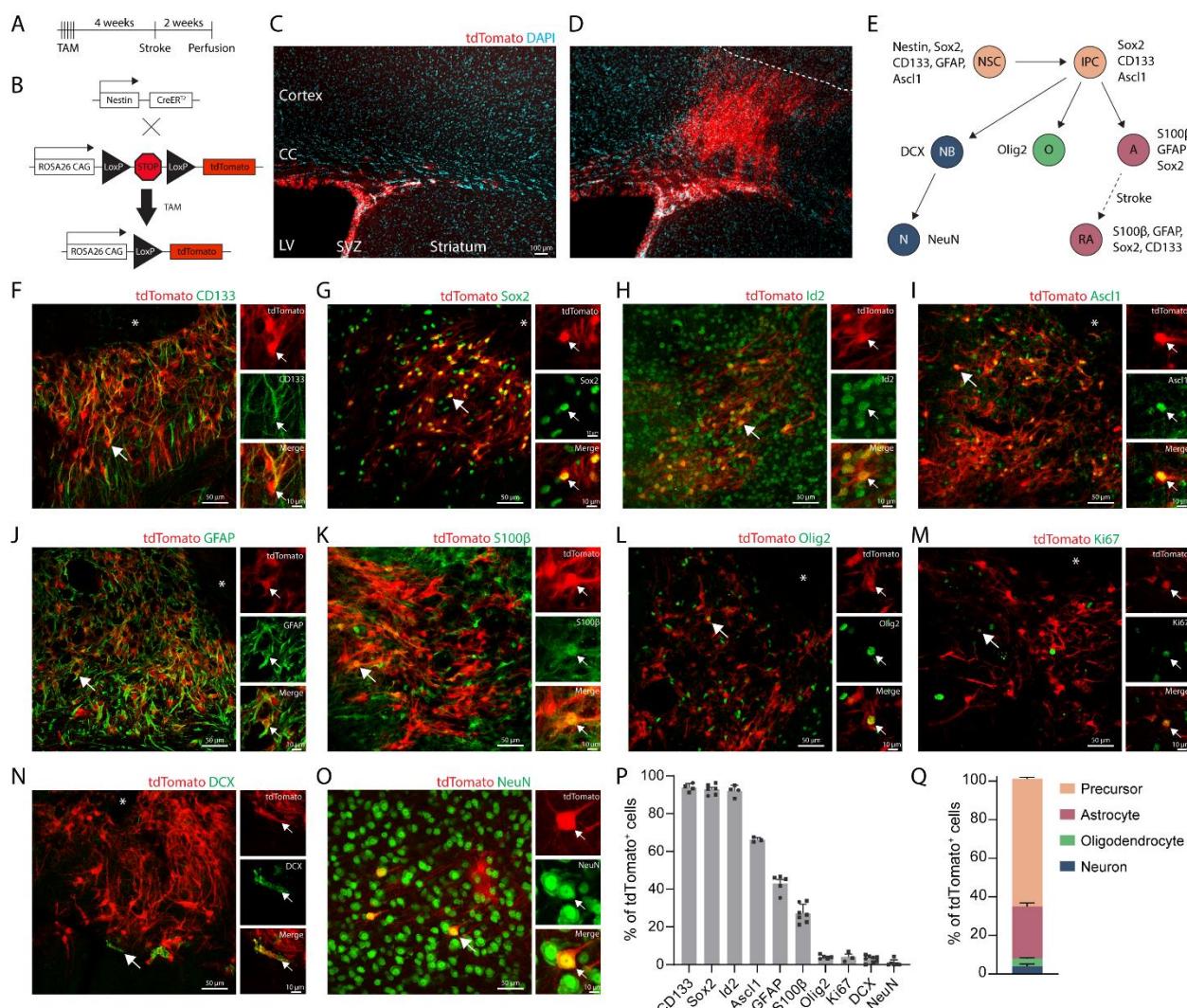
85 We used genetic lineage tracing to characterize the SVZ cytogenic response to stroke.

86 Young adult (3-6 months old) *Nestin-CreER*; Ai14 mice were injected with tamoxifen to induce
87 indelible tdTomato expression in neural stem cells and their progeny (Figure 1A, B) (Benner et
88 al., 2013; Li et al., 2010). Four weeks later, photothrombotic cortical infarcts were induced, and
89 tissue was collected two weeks post-stroke, which is a time when large numbers of SVZ-derived
90 cell are localized to peri-infarct cortex, substantial neural repair is ongoing, and functional
91 improvement is incomplete (Kannangara et al., 2018; Kim et al., 2018; Williamson et al., 2021).

92 While no cortical migration was seen in the absence of injury, unilateral strokes induced a
93 profound migration of tdTomato⁺ cells from the SVZ into peri-infarct cortex (Figure 1C, D). We
94 immunostained tissue to examine expression of an array of differentiation stage-specific and
95 proliferation-associated proteins in lineage-traced cells (Figure 1E-Q, Supplemental Figure 1,
96 Supplemental Figure 2).

97 Unexpectedly, the majority of tdTomato⁺ cells in peri-infarct cortex expressed precursor
98 cell-associated markers (93.8±1.1% were CD133⁺, 92.9±1.1% were Sox2⁺, and 66.2±0.7% were
99 Ascl1⁺). 27.1±1.9% of tdTomato⁺ cells were differentiated astrocytes based on expression of
100 S100 β , which defines astrocyte maturation and loss of multipotency (Lattke et al., 2021; Raponi
101 et al., 2007) and is not expressed by SVZ precursors (Codega et al., 2014). Astrocyte reactivity is
102 associated with re-expression of some precursor cell-associated proteins, including CD133 and
103 Sox2 (Götz et al., 2015; Robel et al., 2011), but reactive astrocytes do not express Ascl1

104 (Magnusson et al., 2014; Zamboni et al., 2020). Thus, the *Ascl1*⁺ subpopulation defined
105 undifferentiated precursors, the *S100β*⁺ subpopulation defined astrocytes, and *CD133/Sox2*
106 labeled both subpopulations (Figure 1E). Lineage-traced cells were largely quiescent as defined
107 by expression of the quiescence marker *Id2* in 92.1±1.5% of cells (Llorens-Bobadilla et al.,
108 2015), and rare expression of the proliferation marker *Ki67* (4.1±1.5%). Oligodendrocyte-lineage
109 (*Olig2*⁺, 4.1±0.5%) and neuron-lineage cells (*DCX*⁺, 3.0±0.6%, and *NeuN*⁺, 0.9±0.7%; (also see
110 Supplemental Figure 2)) made up the remainder of lineage-traced cells. We corroborated these
111 phenotyping results with parallel experiments in *Ascl1-CreER*; *Ai14* mice, in which a subset of
112 neural stem and progenitor cells are lineage-traced (Liang et al., 2019) (Supplemental Figure 1).
113 We also confirmed that few new neurons are formed up to 6 weeks post-stroke using *Nestin-*
114 *CreER*; *Sun1-sfGFP*^f mice in which neural stem cells and their progeny were labeled with a
115 nuclear membrane-targeted fluorophore (Supplemental Figure 2). These experiments identify
116 undifferentiated precursors as the predominant cell type produced by the SVZ in response to
117 stroke. Since the majority of new cells that localize to peri-infarct regions remain in an
118 undifferentiated state, cell replacement may not be the primary function of the cytogenic
119 response.



120

121 **Figure 1. SVZ-derived cells are predominantly undifferentiated precursors and astrocytes.**

122 A) Experimental timeline for lineage tracing neural stem cell progeny after stroke. Tissue was
 123 collected two weeks post-stroke. B) Schematic of inducible, neural stem cell-specific lineage
 124 tracing system in Nestin-CreER; Ai14 mice. Tamoxifen (TAM) administration induces tdTomato
 125 expression in neural stem cells and their progeny. C) Image of tdTomato-expressing cells in the
 126 subventricular zone in the absence of injury. Note the lack of cortical migration of tdTomato⁺
 127 cells. CC, corpus callosum; LV, lateral ventricle; SVZ, subventricular zone. D) Substantial
 128 migration of tdTomato⁺ cells towards the infarct after cortical photothrombotic stroke (dashed
 129 line indicates the approximate infarct border). E) Schematic of differentiation stages as defined
 130 by marker expression. Neural stem cells (NSC) produce intermediate progenitor cells (IPC),
 131 which give rise to cells of the three major neural lineages: neurons (NB, neuroblast; N, neuron),
 132 oligodendrocytes (O), and astrocytes (A). After stroke, reactive astrocytes (RA) re-express some
 133 neural stem cell markers. F-O) Representative confocal images of tdTomato⁺ cells and
 134 immunostaining for lineage and functional markers. Images are from peri-infarct cortex two
 135 weeks post-stroke. Asterisks indicate the lesion core. Most tdTomato⁺ cells (precursors and

136 astrocytes) expressed the stem cell-associated markers CD133 (F) and Sox2 (G), and the
137 quiescence marker Id2 (H). Id2⁺ tdTomato⁻ cells include mature resident cortical cells.
138 Undifferentiated precursors were uniquely identified by expression of Ascl1 (I). J) GFAP
139 marked astrocytes and a subset of precursors. K) Differentiated astrocytes were defined by
140 expression of S100 β . L) The oligodendrocyte lineage was defined by Olig2 expression. M) Few
141 cells expressed the proliferation marker Ki67. Neuron lineage cells, DCX⁺ (N) and NeuN⁺ (O),
142 were the least common. P) Quantification of marker expression by tdTomato⁺ cells. Data are
143 from n = 3-8 mice per marker, and > 100 tdTomato⁺ cells counted per mouse per marker. Q)
144 Estimate of phenotype distribution of lineage traced cells. Precursors were defined by Ascl1
145 expression. Astrocytes were defined by S100 β expression. Oligodendrocyte lineage was defined
146 by Olig2 expression. Neuron lineage was defined by DCX and NeuN expression. See also
147 Supplemental Figure 1 and Supplemental Figure 2. Data are presented as mean \pm SEM.
148 Datapoints representing males are shown as circles; datapoints representing females are shown as
149 squares.

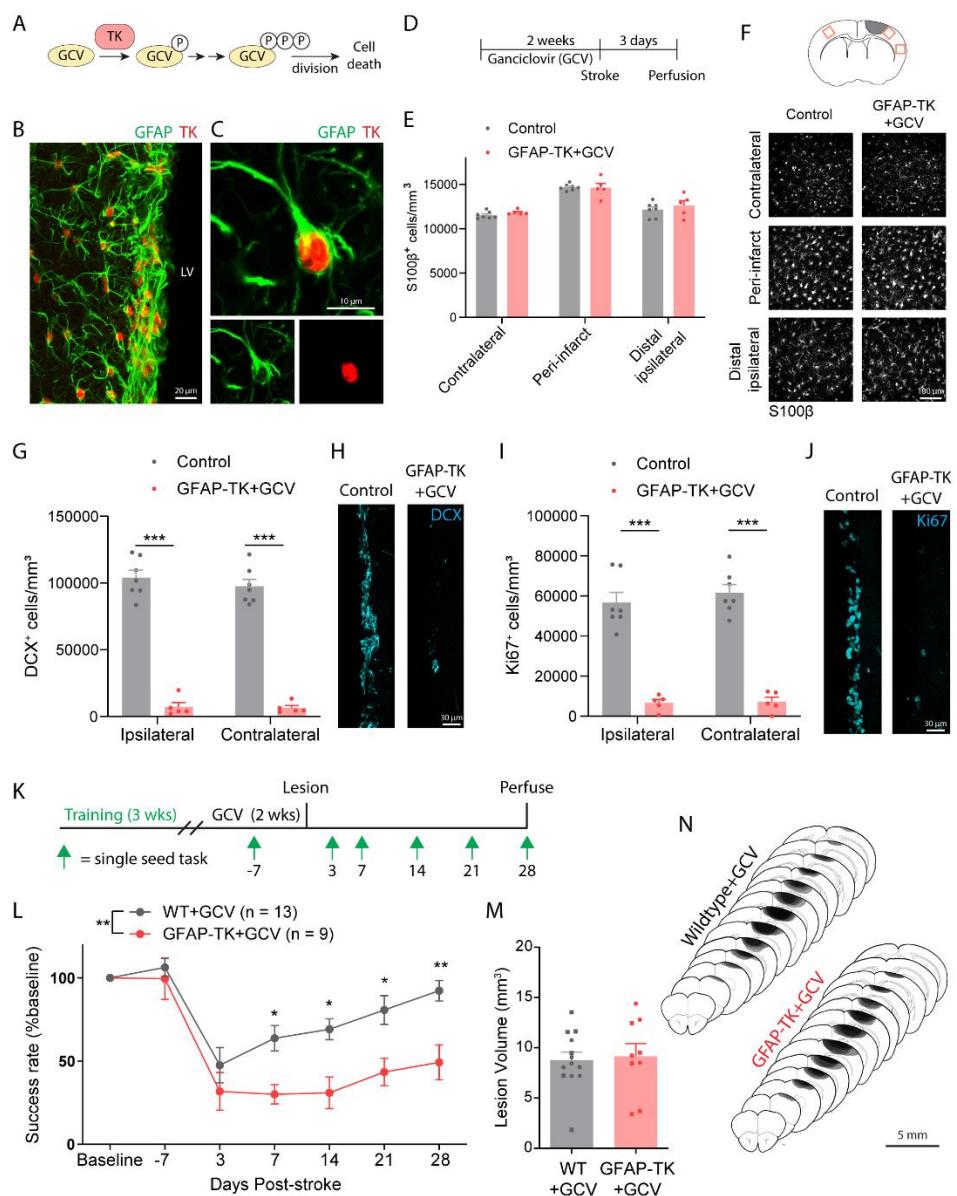
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151 *Chemogenetic ablation of neural stem cells impairs recovery after stroke*

152 Having characterized the phenotype of SVZ-derived cells, we next investigated whether
153 SVZ cytogenesis provides functional benefits for recovery after stroke. We used GFAP-TK mice
154 to selectively ablate neural stem cells prior to stroke following an established ganciclovir (GCV)
155 administration paradigm (Garcia et al., 2004; Swan et al., 2014) (Figure 2A-J). Delivery of GCV
156 via subcutaneous osmotic pumps for two weeks ablates mitotic TK-expressing cells (i.e. neural
157 stem cells), but spares non-mitotic GFAP-expressing cells, including cortical astrocytes (Garcia
158 et al., 2004; Swan et al., 2014) (Figure 2D-F; Supplemental Figure 3A-C). SVZ cytogenesis, as
159 measured by the density of SVZ DCX⁺ and Ki67⁺ cells, was substantially reduced in GFAP-
160 TK+GCV mice relative to littermate controls, which included wildtype mice given saline or
161 GCV and GFAP-TK mice given saline (Figure 2G-J).

162 Following GCV administration, we compared motor recovery between GFAP-TK mice
163 and wildtype littermates after photothrombotic cortical infarcts targeting the forelimb area of
164 motor cortex (Tennant et al., 2011; Williamson et al., 2021) (Figure 2K). Motor function was

165 assessed with the single seed reaching task, a highly sensitive and translationally relevant
166 measure of skilled reaching (Klein et al., 2012; van Lieshout et al., 2021). There was no
167 difference between groups in reaching performance during pre-stroke GCV delivery (Bonferroni-
168 corrected $p > 0.999$). Mice lacking cytogenesis showed significantly worse recovery out to four
169 weeks following stroke (Figure 2L). Lesion size and location did not differ between groups
170 (Figure 2M, N). These results demonstrate that SVZ cytogenesis promotes functional recovery
171 after stroke.



172

173 **Figure 2. Ablation of neural stem cells worsens recovery after stroke.**

174 A) Schematic illustrating the thymidine kinase (TK)/ganciclovir (GCV) system for conditional
 175 ablation of proliferating cells. B) Confocal image showing TK expression in GFAP⁺ cells of the
 176 subventricular zone in a GFAP-TK mouse. LV, lateral ventricle. C) High-resolution image of a
 177 TK-expressing SVZ stem cell. D) Experimental timeline for assessing the specificity and
 178 effectiveness of arresting SVZ cytogenesis with GFAP-TK mice (E-J). GCV or saline was
 179 delivered for 14 days via osmotic pump. Tissue was collected 3 days after stroke. n = 7 control
 180 (wildtype mice given GCV or GFAP-TK mice given saline), n = 5 GFAP-TK+GCV. E)
 181 Parenchymal astrocytes were not depleted in this paradigm. $t(10) < 1.53$, $p \geq 0.157$, t tests
 182 comparing groups for each region. F) Representative images of S100^β⁺ cells from three cortical

183 regions (see diagram at top) show lack of astrocyte ablation. G-J) GFAP-TK mice permit
184 conditional arrest of cytogenesis. The number of SVZ DCX⁺ (G, H) and Ki67⁺ cells (I, J) was
185 significantly reduced in GFAP-TK mice given GCV relative to controls. ***t(10) ≥ 7.99, p <
186 0.001, t tests. K) Experimental design for assessing motor recovery after photothrombotic stroke
187 with the single seed reaching task (n = 13 wildtype (WT), n = 9 GFAP-TK). L) Arrest of
188 cytogenesis significantly worsened recovery of motor function. Time x group interaction F(6,
189 120) = 3.61, p = 0.0025. *p < 0.05, **p < 0.01, Bonferroni tests. M) Lesion volume was not
190 different between groups. t(20) = 0.27, p = 0.787. N) Lesion reconstruction. Darker shades
191 represent more overlap between animals. Data are presented as mean ± SEM. Where individual
192 datapoints are shown, datapoints representing males are shown as circles; datapoints representing
193 females are shown as squares.

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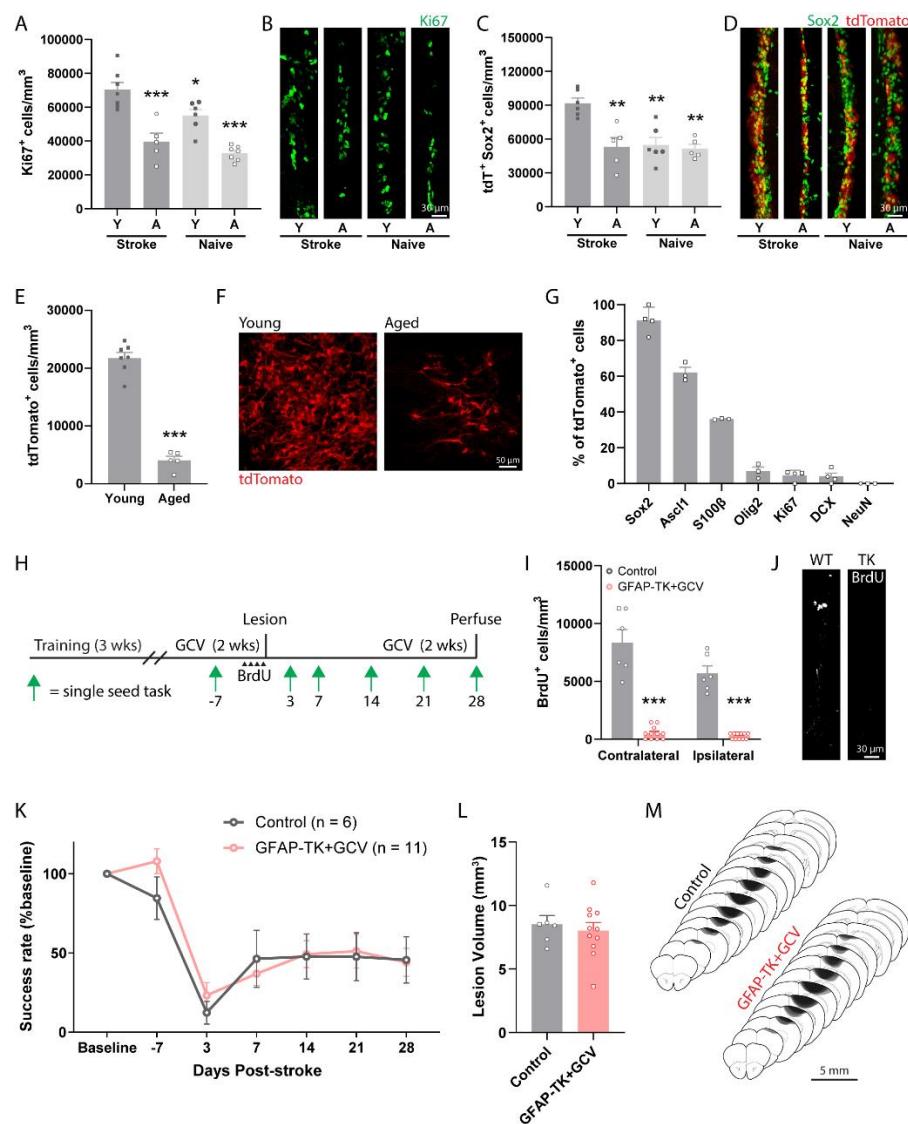
195 *Aging diminishes the SVZ cytogenic response to stroke and its functional benefits*

196 Aging is associated with a substantially higher incidence of stroke (Kissela et al., 2012)
197 and reduced recovery of function (Paolucci et al., 2003). In animals, aging is associated with
198 diminished SVZ cytogenesis (Bouab et al., 2011; Jin et al., 2004; Kalamakis et al., 2019; Luo et
199 al., 2006). We examined the impact of aging on normal and post-stroke SVZ cytogenesis by
200 comparing young adult (aged 3-6 months) and aged (12-16 months) *Nestin-CreER*; Ai14 mice.
201 Mice were either uninjured (naïve) or subjected to cortical stroke two weeks prior to tissue
202 collection. In young mice, stroke increased the number of Ki67⁺ proliferative cells and
203 tdTomato⁺Sox2⁺ precursors in the SVZ relative to young naïve, aged naïve, and aged stroke mice
204 (Figure 3A-D). By contrast, in aged mice, there was no significant difference in the number of
205 SVZ Ki67⁺ or tdTomato⁺Sox2⁺ cells after stroke relative to naïve mice (p = 0.604, p = 0.998,
206 respectively, Tukey tests). Thus, stroke increases SVZ proliferation and expands the precursor
207 cell pool in young, but not aged, mice.

208 We next examined the effects of aging on the migratory response of lineage-traced SVZ
209 cells after stroke. The density of tdTomato⁺ cells in peri-infarct cortex two weeks post-stroke
210 was about five-fold lower in aged mice relative to young mice (Figure 3E, F). Despite the

211 diminished cytogenic response, the phenotype distribution of tdTomato⁺ cells in peri-infarct
212 cortex of aged mice (Figure 3G) was similar to what we observed in young adult mice (Figure 1),
213 with the majority of cells identified as precursors and astrocytes. Thus, far fewer SVZ-derived
214 cells localize to peri-infarct regions in aged mice, but their phenotype distribution is similar to
215 that of young mice.

216 We next investigated whether the blunted cytogenic response observed in aged mice was
217 still functionally beneficial (Figure 3H). We used GFAP-TK mice to selectively ablate neural
218 stem cells prior to stroke. BrdU pulse labeling of proliferating cells revealed a near complete loss
219 of BrdU⁺ cells in the SVZ of GFAP-TK+GCV mice relative to control aged mice (Figure 3I, J;
220 all mice aged 12-16 months). We assessed motor recovery with the single-seed task and found no
221 differences between groups (Figure 3K). Notably, both aged groups performed significantly
222 worse on day 28 than young mice within intact cytogenesis (data from Figure 2; $F(3, 35) = 7.46$,
223 $p < 0.001$ one-way ANOVA; $p < 0.012$, Tukey tests), but similarly to young GFAP-TK+GCV
224 mice ($p > 0.978$, Tukey tests). Lesion size was not different between aged groups (Figure 3L,
225 M), and was similar to that of young adults (Figure 2M, N). Altogether, our results indicate that
226 aging diminishes the cytogenic response to stroke and its functional benefits. Together with our
227 findings in young mice, our results show that reduced SVZ cytogenesis, by neural stem cell
228 ablation or aging, is associated with worse functional recovery. Reduced cytogenesis may
229 contribute to worse outcome after stroke with aging.



230

231 **Figure 3. Aging diminishes the cytogenic response to stroke and its contribution to**
 232 **recovery.**

233 A-D) Assessment of SVZ cytogenesis in young (Y, 3-6 months) and aged (A, 12-16 months)
 234 Nestin-CreER; Ai14 mice (n = 5-7 mice/group). Tissue was collected two weeks post-stroke.
 235 The number of proliferating cells (Ki67⁺, A and B) and neural precursor cells (tdTomato⁺ Sox2⁺,
 236 C and D) in the SVZ was significantly higher in young mice after stroke compared to all other
 237 groups. *p < 0.05, ** p < 0.01, ***p < 0.001 relative to young stroke mice, Tukey tests. E)
 238 Fewer tdTomato⁺ cells were observed in peri-infarct cortex of aged mice two weeks post-stroke.
 239 ***t(10) = 13.4, p < 0.001. F) Representative images of tdTomato⁺ cells in peri-infarct cortex. G)
 240 Quantification of lineage marker expression by tdTomato⁺ cells in peri-infarct cortex of aged
 241 mice at two weeks post-stroke (n = 3-4 mice per marker). Similar to what was observed in young
 242 mice (Figure 1), most SVZ-derived cells were undifferentiated precursors or astrocytes. H)
 243 Experimental timeline for examining the effects of neural stem cell ablation in aged mice (n = 6

244 controls, n = 11 GFAP-TK given GCV). I) GFAP-TK+GCV mice had fewer BrdU⁺ cells in the
245 SVZ, validating stem cell ablation. *** $t(15) \geq 9.36$, p < 0.001. BrdU was given twice per day for
246 two days prior to stroke. J) Representative images of BrdU immunostaining in the SVZ. K) Both
247 aged controls and GFAP-TK+GCV mice showed poor recovery following stroke. There was no
248 difference between groups. Group main effect $F(1,15) = 0.17$, p = 0.690. L) Lesion volume was
249 not different between groups ($t(15) = 0.50$, p = 0.628). M) Lesion reconstruction. Darker shades
250 represent more overlap between animals. Data are presented as mean \pm SEM. Where individual
251 datapoints are shown, datapoints representing males are shown as circles; datapoints representing
252 females are shown as squares.

253

254 *Arrest of cyogenesis disrupts neuronal and vascular repair*

255 Our results indicate that most SVZ-derived cells are undifferentiated precursors and
256 reactive astrocytes. We hypothesized that these cell types may influence repair processes to
257 promote behavioral improvement. In particular, synaptic plasticity and vascular remodeling are
258 two functionally important aspects of neural repair that could potentially be augmented by
259 factors produced by SVZ-derived cells. Indeed, past studies have demonstrated beneficial effects
260 of transplanted stem cells and resident cortical astrocytes on these processes (Andres et al., 2011;
261 Bacigaluppi et al., 2016; Horie et al., 2011; Lin et al., 2017; Llorente et al., 2021; Pluchino et al.,
262 2005; Roitbak et al., 2011; Sabelström et al., 2013; Williamson et al., 2021).

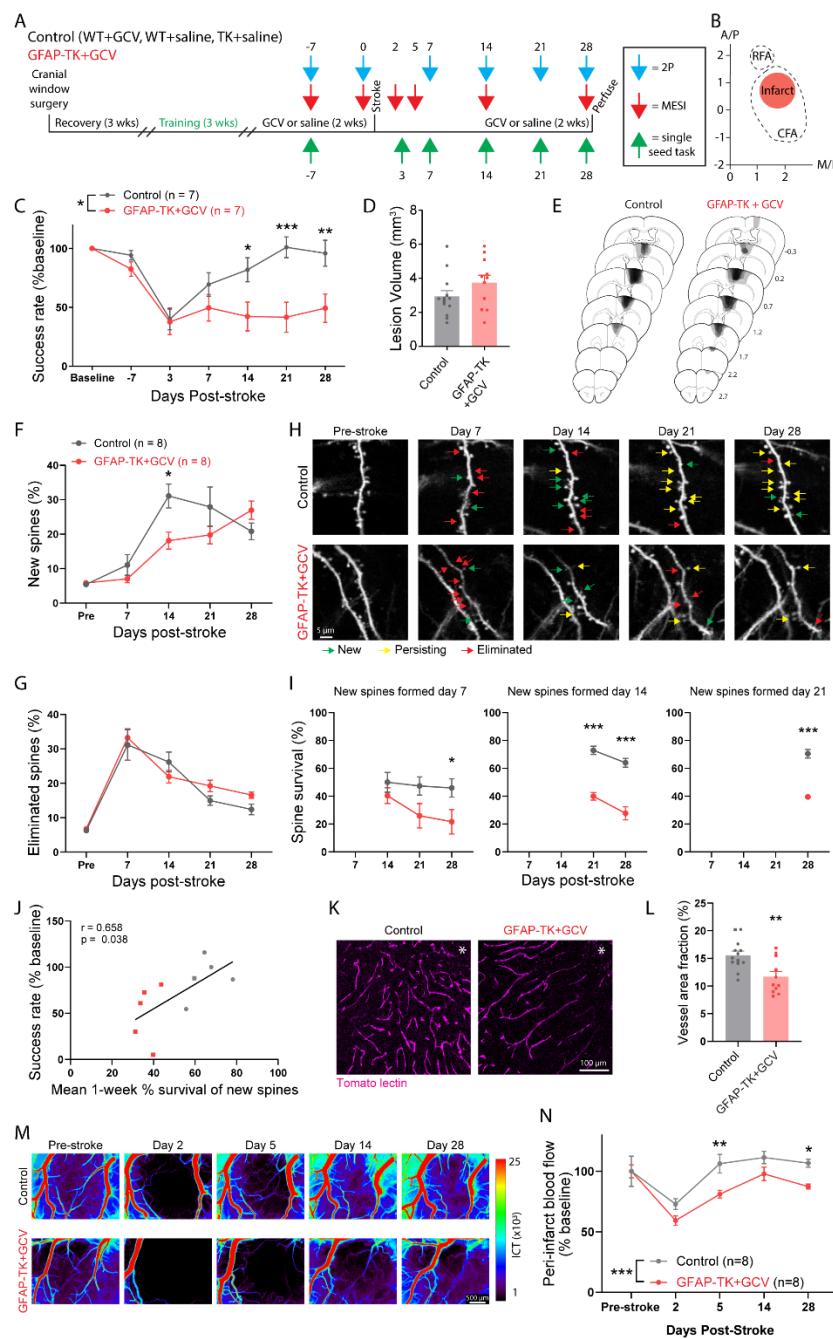
263 We investigated the consequences of neural stem cell ablation on synaptic and vascular
264 plasticity in residual cortex surrounding photothrombotic infarcts with a longitudinal imaging
265 approach (Figure 4A, B). We bred GFAP-TK mice with Thy1-GFP mice, which have sparse,
266 GFP-labeled pyramidal neurons. Resulting double transgenic mice allowed us to monitor
267 synaptic remodeling at single-synapse resolution with repeated 2-photon imaging of dendritic
268 spines on apical dendrites before and after stroke (Brown et al., 2007; Clark et al., 2019;
269 Mostany et al., 2010) with the ability to conditionally arrest cyogenesis. Blood flow was tracked
270 with multi-exposure speckle imaging (MESI), a quantitative, optical, contrast-free technique that

271 yields high-resolution blood flow maps (He et al., 2020; Williamson et al., 2020). Prior to stroke,
272 control and TK^{+/−} mice were unilaterally implanted with cranial windows, trained on the single-
273 seed reaching task, administered GCV or saline, and subjected to baseline imaging. We then
274 induced strokes in forelimb motor cortex and periodically imaged dendritic spines and blood
275 flow and assessed behavioral performance during recovery.

276 Motor recovery was again significantly impaired in GFAP-TK+GCV mice, with no
277 differences in lesion size observed between groups (Figure 4C-E). 2-photon imaging of dendritic
278 spines revealed increased spine turnover in peri-infarct cortex (Figure 4F-H), consistent with past
279 work (Brown et al., 2007; Clark et al., 2019; Joy et al., 2019; Mostany et al., 2010). There were
280 no group differences in spine dynamics before stroke (Supplemental Figure 4). New spine
281 formation peaked during the second week after stroke, and was significantly higher in mice with
282 intact cytogenesis. Spine elimination was greatest during the first week post-stroke, and
283 subsequently declined with time, without significant differences between groups. A
284 subpopulation of new spines formed after stroke persists long-term, and the persistence of newly
285 formed spines is associated with greater functional recovery (Clark et al., 2019). The survival of
286 new spines was significantly reduced in GFAP-TK+GCV mice regardless of the day of spine
287 formation (Figure 4I). Moreover, spine survival rate was positively correlated with behavioral
288 performance on the final assessment day (Figure 4J). Thus, SVZ cytogenesis supports synaptic
289 remodeling after stroke, particularly by promoting the long-term stabilization of new synapses.

290 Broad regions of reduced blood flow persist for days to weeks surrounding focal strokes
291 (He et al., 2020; Williamson et al., 2020). Remodeling of peri-infarct vasculature helps to restore
292 blood flow and is associated with behavioral improvement (Williamson et al., 2020). To examine
293 vascular remodeling, we injected mice with fluorophore-conjugated tomato lectin immediately

294 before euthanasia to label perfused vasculature. Vessel density in peri-infarct cortex was
295 significantly reduced in GFAP-TK+GCV mice relative to controls (Figure 4K, L). Furthermore,
296 longitudinal blood flow imaging demonstrated impaired recovery of blood flow at days 5 and 28
297 post-stroke (Figure 4M, N). Early vascular permeability was not affected by arresting
298 cytoplasma (Supplemental Figure 3). Overall, these findings demonstrate that SVZ-derived cells
299 beneficially shape neuronal and vascular remodeling processes after stroke in order to promote
300 recovery.



301

302 **Figure 4. SVZ cytogenesis supports neuronal and vascular remodeling.**

303 A) Experimental design for longitudinal imaging and behavioral testing. Subgroups of mice were
304 subjected to longitudinal imaging and behavioral testing (n = 5/group), imaging only (n =
305 3/group), behavioral testing only (n = 2/group), and neither (histology only; n = 3 control, n = 1
306 GFAP-TK+GCV). B) Schematic illustrating infarct placement relative to caudal (CFA) and
307 rostral (RFA) forelimb areas in motor cortex. Axes indicate mm relative to Bregma. C)
308 Performance on the single-seed reaching task. GFAP-TK+GCV mice showed significantly worse

309 recovery relative to control mice (significant group x time interaction, $F(6, 72) = 4.7$, $p < 0.001$).
310 * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$, Bonferroni tests. D) Lesion volume was not different
311 between groups ($t(22) = 1.5$, $p = 0.159$). E) Lesion reconstructions. Darker shades represent more
312 overlap between animals. F-J) Longitudinal imaging of dendritic spines revealed altered spine
313 dynamics in mice with ablated neural stem cells. F) New spine formation was significantly
314 higher in control mice on day 14 (significant time x group interaction, $F(4,52) = 4.1$, $p = 0.006$).
315 * $p < 0.05$, Sidak's multiple comparison tests between groups for each day. G) Spine elimination
316 was not significantly different between groups (group effect, $F(1, 66) = 1.2$, $p = 0.285$). H)
317 Longitudinal 2-photon images of GFP-expressing dendritic spines illustrating spine formation
318 (green arrows), persistence of new spines (yellow arrows), and spine elimination (red arrows). I)
319 Survival of new spines was significantly reduced in GFAP-TK+GCV mice. Plots show survival
320 of new spines formed on days 7, 14, and 21 after stroke. * $p < 0.05$, *** $p < 0.001$, Sidak's
321 multiple comparison tests. J) Spine survival was significantly positively correlated with
322 behavioral performance on the final day of testing. K) Representative images of peri-infarct
323 vasculature. L) Peri-infarct vessel density was significantly reduced in GFAP-TK+GCV mice
324 (** $t(22) = 3.3$, $p = 0.004$). M) Representative MESI images of blood flow. N) Peri-infarct blood
325 flow was significantly reduced on days 5 and 28 in GFAP-TK+GCV mice (group effect, $F(1, 14)$
326 = 32.98, $p < 0.001$). * $p < 0.05$, ** $p < 0.01$, Sidak's multiple comparison tests. Data are presented
327 as mean \pm SEM. Where individual datapoints are shown, datapoints representing males are
328 shown as circles; datapoints representing females are shown as squares.

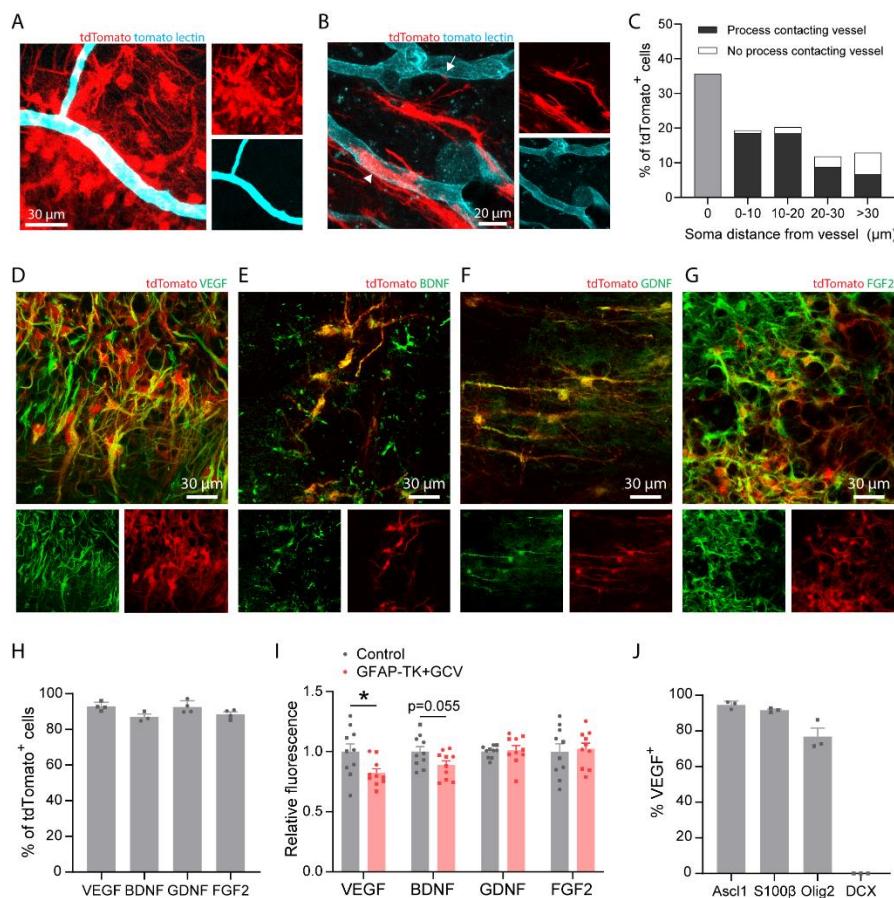
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330 *SVZ-derived cells interact with vasculature and produce trophic factors*

331 Neuroblasts originating in the SVZ migrate along vascular scaffolds towards the
332 olfactory bulb in the healthy brain (Bovetti et al., 2007) and towards peri-infarct regions after
333 stroke (Ohab et al., 2006; Thored et al., 2007). We observed frequent contact between lineage-
334 traced SVZ progeny and blood vessels in peri-infarct cortex, including contact of nearby vessels
335 by the processes or cell bodies of 88.2% of SVZ-derived cells (Figure 5A-C). Thus, contact with
336 blood vessels may underlie some of the reparative effects of SVZ-derived cells. In addition,
337 stroke may induce expression of migratory cues in endothelial cells to drive migration of
338 reparative cells from the SVZ (Ohab et al., 2006).

339 Transplanted neural precursors of various sources and in diverse disease settings have
340 been reported to express trophic factors, which may be implicated in their therapeutic effects

341 (Andres et al., 2011; Bacigaluppi et al., 2009; Drago et al., 2013; Horie et al., 2011; Llorente et
342 al., 2021; Martino and Pluchino, 2006; Roitbak et al., 2011). To prospectively investigate
343 molecular mechanisms underlying the facilitation of post-stroke repair by SVZ cytogenesis, we
344 examined the expression of four trophic factors known to drive neuronal and vascular growth.
345 ~90% of all SVZ-derived lineage traced cells in peri-infarct cortex expressed VEGF, BDNF,
346 GDNF, and FGF2 (Figure 5D-H). We compared relative abundance of each of these proteins in
347 peri-infarct cortex between control and GFAP-TK+GCV mice at 28 days post-stroke to evaluate
348 production of these factors by SVZ-derived cells relative to other cell types. VEGF protein was
349 significantly reduced in GFAP-TK+GCV mice, indicating that SVZ-derived cells are a major
350 source of VEGF (Figure 5I). Notably, among SVZ-derived cells, all lineages except for new
351 neurons produced VEGF (Figure 5J). These findings suggest that SVZ cytogenesis may facilitate
352 neural repair at least in part through production of trophic factors, particularly VEGF.



353

354 **Figure 5. SVZ-derived cells interact with vasculature and produce trophic factors.**

355 A) Confocal image showing clustering of lineage traced cells around blood vessels in peri-infarct
 356 cortex. B) tdTomato-expressing cells were frequently observed with cell bodies abutting vessels
 357 (arrowhead) or extending processes that terminated on nearby vessels (arrow). C) Quantification
 358 of SVZ-derived cell interaction with vasculature in peri-infarct cortex two weeks post-stroke.
 359 Data are from 661 cells across 3 Nestin-CreER; Ai14 mice. D-G) Confocal images from peri-
 360 infarct cortex showing expression of the trophic factors VEGF (D), BDNF (E), GDNF (F), and
 361 FGF2 (G) in tdTomato⁺ cells. H) Quantification of trophic factor expression in tdTomato⁺ cells.
 362 I) Quantification of trophic factor expression in peri-infarct cortex four weeks post-stroke
 363 between control mice and GFAP-TK+GCV mice (n = 10/group). Fluorescence intensity is
 364 reported relative to controls. VEGF protein fluorescence was significantly reduced in GFAP-
 365 TK+GCV mice. * $t(18) = 2.4$, p = 0.025. J) Quantification of VEGF expression by phenotype in
 366 lineage traced cells. VEGF was expressed by Ascl1⁺ precursors, S100 β ⁺ astrocytes, and Olig2⁺
 367 oligodendrocyte-lineage cells, but not by neuronal lineage cells (DCX⁺). Data are presented as
 368 mean \pm SEM. Where individual datapoints are shown, datapoints representing males are shown
 369 as circles; datapoints representing females are shown as squares.

370

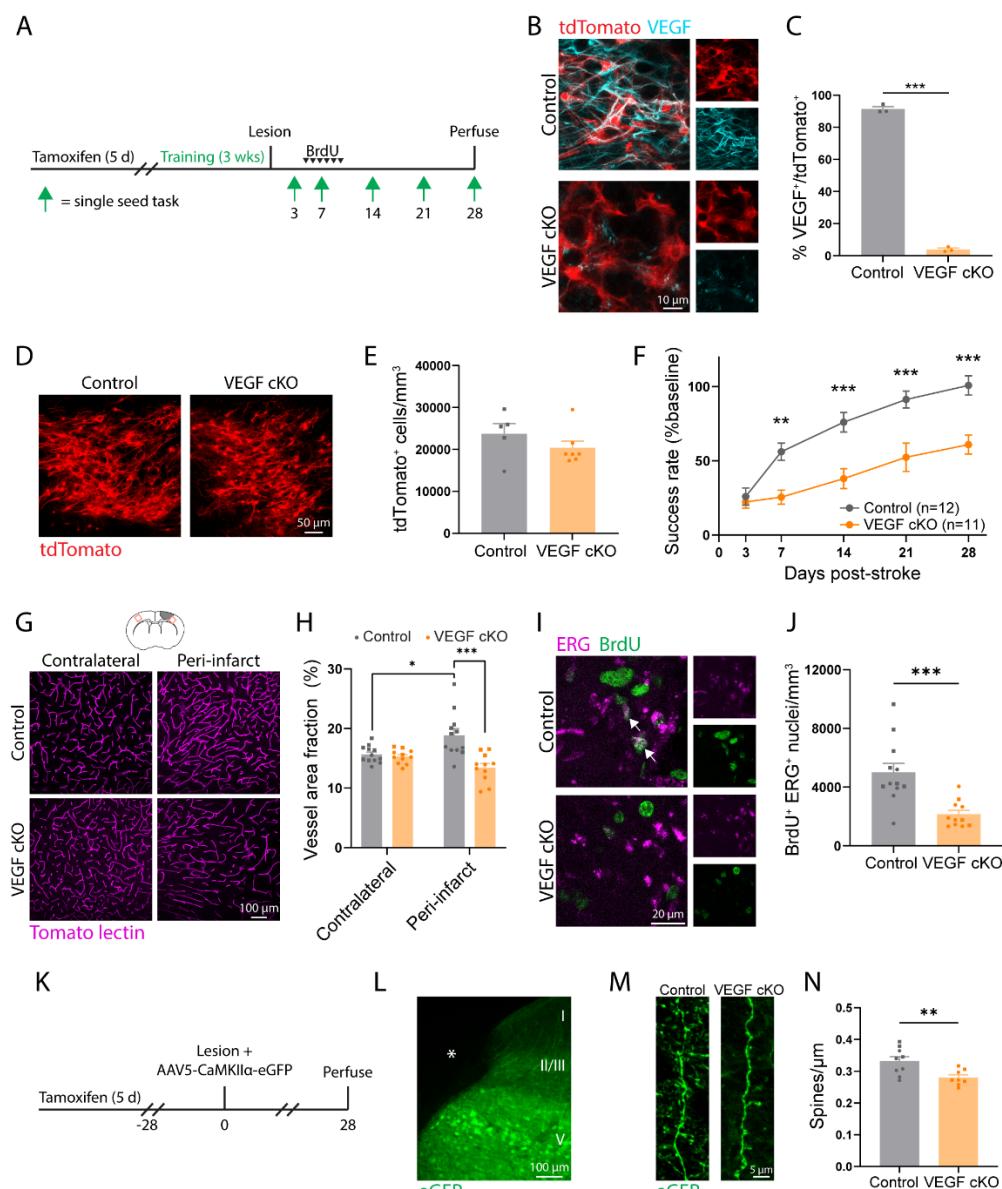
371 *Conditional deletion of Vegf in adult neural stem cells impairs recovery and repair*

372 VEGF promotes the growth of blood vessels and neurons (Gerber et al., 1999; Raab et
373 al., 2004; Rosenstein et al., 2003; Sun et al., 2003). We next investigated whether VEGF
374 produced by SVZ-derived cells was involved in post-stroke recovery and repair. We generated
375 *Nestin-CreER; Vegf^{f/f}* mice to permit inducible deletion of *Vegf* in adult neural stem cells and
376 their progeny (VEGF cKO; Figure 6A). Lineage tracing in *Nestin-CreER; Vegf^{f/f}*; Ai14 mice
377 showed a near complete loss of VEGF in tdTomato⁺ cells in peri-infarct cortex, but no change in
378 the number of migratory cells relative to controls at two weeks post-stroke (Figure 6B-E).
379 Recovery of forelimb motor function was significantly worse in VEGF cKO mice relative to
380 controls as measured with the single-seed reaching task (Figure 6F). Lesion size and location
381 were not different between groups (Supplemental Figure 5). Vascular remodeling following
382 stroke is seen by increases in peri-infarct vascular density (Williamson et al., 2021, 2020). While
383 vessel density in peri-infarct cortex was increased in control mice relative to the contralateral
384 hemisphere, this was not seen in VEGF cKO mice, and peri-infarct vessel density in VEGF cKO
385 mice was reduced relative to controls (Figure 6G, H). We injected mice with BrdU daily during
386 the peak in angiogenesis from days 5-10 after stroke to quantify new vessel formation. The
387 number of new endothelial cells (BrdU⁺ERG⁺) was significantly reduced in VEGF cKO mice,
388 confirming impaired angiogenesis in mice lacking VEGF in SVZ-derived cells.

389 In a separate experiment, we examined the effects of conditional *Vegf* deletion on
390 dendritic spine density after stroke (Figure 6K). Peri-infarct layer V cortical pyramidal neurons
391 were labeled by injections of AAV5-CaMKIIa-eGFP (Figure 6L). Four weeks post-stroke, we
392 quantified spine density on apical dendrites in layer II/III. We previously found that changes in
393 spine density in this region parallel functional recovery (Clark et al., 2019). Spine density was
394 significantly reduced in VEGF cKO mice (Figure 6M, N). Altogether, our results identify VEGF

395 produced by SVZ-derived cells as a key driver of repair and recovery after stroke. More broadly,
 396 our findings position newborn cells arising from the SVZ in response to injury as a unique
 397 cellular source of trophic support that instructs neural repair.

398



399

400 **Figure 6. Adult neural stem cell-specific deletion of Vegf impedes recovery and repair.**

401 A) Timeline of experiments to examine recovery in control (Nestin-CreER^{+/−}; VEGF^{fl/+}, Nestin-
 402 CreER^{−/−}; VEGF^{fl/+}, or Nestin-CreER^{+/−}; VEGF^{+/+}) and VEGF cKO (Nestin-CreER^{+/−}; VEGF^{fl/fl})

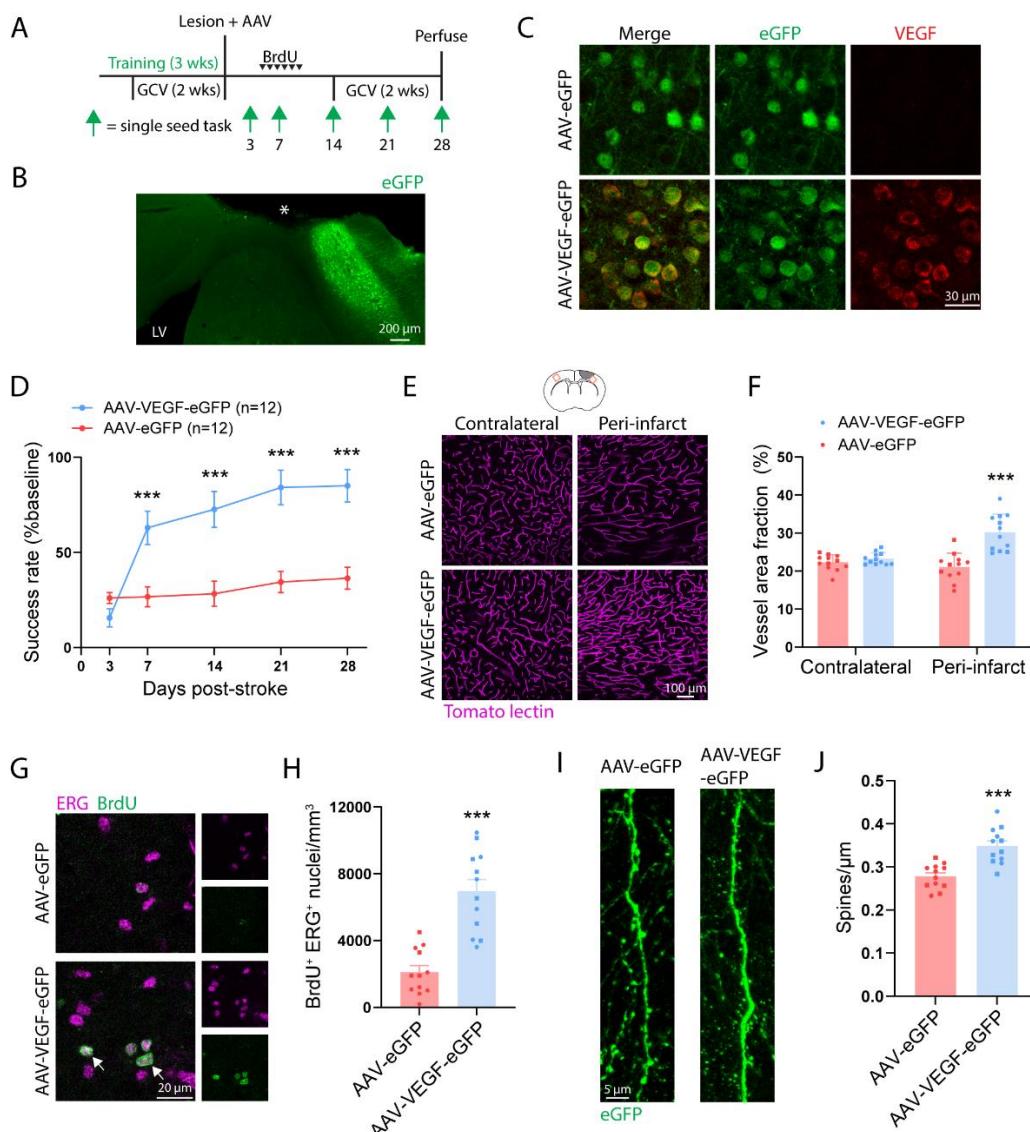
403 mice. B) Confocal images demonstrating VEGF loss in tdTomato⁺ cells in Nestin-CreER^{+/−};
404 VEGF^{fl/fl}; Ai14 mice. Images are from peri-infarct cortex two weeks post-stroke. C)
405 Substantially fewer tdTomato⁺ cells express VEGF in VEGF cKO mice. n = 3 mice/group.
406 ***t(4) = 51.23, p < 0.001. D) Representative images of tdTomato⁺ cells in peri-infarct cortex
407 two weeks post-stroke. (E) There was no difference in peri-infarct tdTomato⁺ cell density
408 between controls and VEGF cKO mice, indicating that the cytogenic response was unaffected. n
409 = 5 controls, n = 7 VEGF cKO. t(10) = 1.18, p = 0.266. F) VEGF cKO significantly impaired
410 motor recovery measured with the single-seed reaching task (n= 12 control, n = 11 VEGF cKO).
411 Time x group interaction F(4, 84) = 9.2, p < 0.001. **p < 0.01, ***p < 0.001, Sidak's tests
412 between group for each timepoint. G) Representative confocal images and quantification (H) of
413 vasculature show that VEGF cKO markedly reduced peri-infarct vessel density relative to
414 control mice. ***t(21) = 4.1, p < 0.001. Within control mice, peri-infarct vessel density was
415 significantly greater than in contralateral cortex, consistent with stroke-induced
416 neovascularization. *t(22) = 2.8, p = 0.011. By contrast, VEGF cKO mice had diminished vessel
417 density in peri-infarct cortex relative to the intact contralateral cortex, indicating a failure of
418 neovascularization (t(20) = 2.3, p = 0.030). I) Representative confocal images of new endothelial
419 cells (BrdU⁺ ERG⁺) in peri-infarct cortex. J) The number of BrdU⁺ ERG⁺ nuclei was
420 significantly less in VEGF cKO mice. ***t(15.0) = 4.2, p < 0.001, Welch's corrected t-test. K)
421 Timeline of experiments for evaluating peri-infarct spine density in control (n = 9) and VEGF
422 cKO (n = 8) mice. L) Layer V pyramidal neurons were labeled by intracortical injections of
423 AAV5-CaMKiiα-eGFP. Image shows eGFP labeling in peri-infarct cortex. Asterisk indicates
424 lesion core. M) Representative images and quantification (N) of dendritic spine density. Spine
425 density was significantly lower in VEGF cKO mice. **t(15) = 3.0, p = 0.009. Apical dendrites
426 were sampled from layer II/III between 100-700 μm from the infarct border. 2631 spines were
427 counted along 7.9 mm total length of dendrite in controls. 1934 spines were counted along 6.9
428 mm total length of dendrite in cKO mice. Data are presented as mean ± SEM. Where individual
429 datapoints are shown, datapoints representing males are shown as circles; datapoints representing
430 females are shown as squares.

431

432 *AAV-mediated expression of VEGF in peri-infarct cortex enhances recovery in mice with*
433 *arrested cytogenesis*

434 We next tested whether replacement of VEGF would be sufficient to enhance recovery in
435 mice in which cytogenesis was arrested. GFAP-TK mice were trained on the single-seed
436 reaching task and administered GCV to ablate neural stem cells. Immediately after stroke, mice
437 were injected with either AAV5-EF1α-VEGF-P2A-eGFP (AAV-VEGF-eGFP), to induce VEGF
438 and eGFP expression, or AAV5-EF1α-eGFP (AAV-eGFP), to induce only eGFP expression, into

439 layer V of peri-infarct cortex (Figure 7A-C). AAV-VEGF-eGFP induced rapid and sustained
440 motor recovery as measured by the single-seed reaching task, whereas AAV-eGFP injected mice
441 showed little improvement up to four weeks post-stroke (Figure 7D). Lesion size and location
442 were not different between groups (Supplemental Figure 6). We examined vascular density in
443 homotopic contralateral and peri-infarct cortex 28 days post-stroke. Vascular density was
444 significantly greater in peri-infarct cortex of AAV-VEGF-eGFP injected mice relative to AAV-
445 eGFP mice (Figure 7E, F). Moreover, there was no difference in vascular density between
446 contralateral and peri-infarct cortex in the AAV-eGFP group, indicating a failure of
447 neovascularization ($t(22) = 0.97$, $p = 0.344$). Mice were given daily injections of BrdU during
448 days 5-10 post-stroke to label new blood vessels. AAV-VEGF-eGFP mice had substantially
449 more angiogenesis in peri-infarct cortex than AAV-eGFP mice as measured by the number of
450 BrdU⁺ERG⁺ nuclei (Figure 7G, H). Finally, we examined spine density of eGFP-expressing
451 pyramidal neurons on apical dendrites in layer II/III. Spine density was significantly higher in
452 mice given AAV-VEGF-eGFP (Figure 7I, J). In wildtype mice subjected to a sham stroke
453 procedure, AAV-VEGF-eGFP increased vessel density but did not affect motor function
454 (Supplemental Figure 7). Overall, our findings indicate that replacement of VEGF is sufficient to
455 enhance repair and recovery in mice lacking cytogenesis. More broadly, these findings suggest
456 that replacement of factors produced by SVZ-derived cells may constitute an effective therapy.



457

458 **Figure 7. VEGF rescues poor recovery due to neural stem cell ablation.**

459 A) Experimental timeline. Both groups used GFAP-TK mice given GCV to ablate neural stem
 460 cells (n = 12 mice per group). Animals were injected with either AAV5-Ef1 α -eGFP (AAV-
 461 eGFP) or AAV5-Ef1 α -VEGF-P2A-eGFP (AAV-VEGF-eGFP) in layer V of peri-infarct cortex.
 462 B) Image illustrating AAV targeting of peri-infarct cortex. Asterisk indicates lesion core. LV,
 463 lateral ventricle. C) Confocal images validating that AAV-VEGF-eGFP induces VEGF
 464 expression. D) AAV-VEGF-eGFP improved motor recovery on the single seed reaching task.
 465 Significant time x group interaction $F(4, 88) = 15.9$, $p < 0.001$. ** $p < 0.01$, *** $p < 0.001$,
 466 Sidak's multiple comparison tests. E) Representative confocal images and quantification (F) of
 467 vasculature show that AAV-VEGF-eGFP increased peri-infarct vessel density relative to AAV-
 468 eGFP mice. *** $t(22) = 5.3$, $p < 0.001$, t test. G) Representative confocal images of new
 469 endothelial cells (BrdU⁺ ERG⁺, arrows) in peri-infarct cortex. H) The number of BrdU⁺ ERG⁺

470 nuclei was significantly greater in AAV-VEGF-eGFP mice. *** $t(22) = 6.0$, $p < 0.001$. I)
471 Representative images and quantification (J) of dendritic spine density. Spine density was
472 significantly higher in AAV-VEGF-eGFP mice. *** $t(22) = 4.8$, $p < 0.001$. Apical dendrites were
473 sampled from layer II/III between 100-800 μm from the infarct border. 2825 spines were counted
474 along 10.2 mm total length of dendrite in AAV-eGFP mice. 3231 spines were counted along 9.3
475 mm total length of dendrite in AAV-VEGF-eGFP mice. Data are presented as mean \pm SEM.
476 Where individual datapoints are shown, datapoints representing males are shown as circles;
477 datapoints representing females are shown as squares.

478 **Discussion**

479 Our study revealed that a previously underappreciated class of cells, undifferentiated
480 precursors, constitutes the majority of cells that arise from the SVZ and migrate towards the site
481 of injury following stroke. The migration of primarily undifferentiated cells towards the site of
482 injury suggests that the main function of post-injury cytogenesis is likely not cell replacement.
483 We found that reducing SVZ cytogenesis, by neural stem cell ablation or aging, leads to poor
484 functional recovery. Moreover, synaptic and vascular repair were disrupted in mice with
485 deficient cytogenesis. SVZ-derived cells produced trophic factors, most notably VEGF. Loss-of-
486 function experiments demonstrated that VEGF produced by SVZ-derived cells is crucial for
487 effective repair and recovery. Finally, gain-of-function experiments showed that replacement of
488 VEGF was sufficient to enhance recovery in mice lacking cytogenesis. We conclude that trophic
489 support from SVZ-derived cells drives neural repair and functional recovery after stroke. Thus,
490 newborn cells formed in response to injury enable recovery by acting as a unique source of
491 trophic cues that instruct neural repair.

492 With lineage tracing of adult neural stem cells and extensive phenotyping, we identified
493 undifferentiated precursors as the largest subpopulation of SVZ-derived cells after stroke.
494 Undifferentiated neural precursor cells may be better suited than differentiated neural cell types
495 to facilitate neural repair. In culture, neural precursors secrete factors that facilitate vessel
496 formation and neuronal outgrowth (Kirby et al., 2015; Roitbak et al., 2011; Rosenstein et al.,
497 2003). In addition, transplantation of neural stem cells enhances repair and recovery in models of
498 stroke without differentiation of transplanted cells (Andres et al., 2011; Bacigaluppi et al., 2016;
499 Horie et al., 2011; Llorente et al., 2021; Roitbak et al., 2011). Collectively, these studies

500 illustrate the reparative abilities of undifferentiated precursors and suggest that post-stroke
501 cytogenesis facilitates neural repair without substantial cell replacement.

502 Further work is needed to understand how the fate of individual cells is decided among
503 the total population of newborn cells arising from the SVZ. Principles of neural development
504 may apply to cell fate decisions during post-injury cytogenesis in adulthood. For example, the
505 transcription factor NFIA controls gliogenesis during development (Deneen et al., 2006), and is
506 also necessary for SVZ astrogenesis after stroke (Laug et al., 2019). In addition, NFIA inhibits
507 neurogenesis via Notch effectors (David-Bercholz et al., 2021; Deneen et al., 2006).
508 Accordingly, interfering with Notch signalling in neural stem cells biases SVZ progeny towards
509 a neuronal fate and away from an astrocytic fate after stroke (Benner et al., 2013). Additional
510 work will be needed to clarify the mechanisms that dictate phenotypes of SVZ cells responding
511 to stroke, and whether different cell types have distinct reparative functions.

512 Our finding that aging reduces SVZ cytogenesis could be translationally relevant given
513 that stroke incidence increases with age (Kissela et al., 2012). We observed the loss of stroke-
514 induced SVZ proliferation and precursor cell pool expansion in aged mice, which suggests
515 deficient activation of neural precursor cells with aging. Inflammatory signals have been
516 implicated in controlling quiescence/activation of precursors. Chronic inflammatory signals,
517 including interferons, promote stem cell quiescence in aging (Kalamakis et al., 2019). By
518 contrast, acute interferon signaling after stroke stimulates precursor activation (Belenguer et al.,
519 2021; Llorens-Bobadilla et al., 2015). Therefore, inflammatory signaling pathways may be a
520 target to restore cytogenesis in aged animals. We also observed a reduction in the number of
521 cells localized in peri-infarct regions in aged mice that was disproportionate relative to the

522 diminishment of SVZ cytogenesis. Thus, there may also be an age-dependent reduction in either
523 the migratory ability of SVZ cells or the expression of migratory cues at the site of injury.

524 SVZ-derived cells interacted closely with peri-infarct blood vessels and produced trophic
525 factors to facilitate their growth after stroke. The ectopic migration of cells from the SVZ is
526 caused in part by expression of migratory cues in peri-infarct vasculature after stroke (Ohab et
527 al., 2006; Thored et al., 2007). This may be an adaptive response to attract precursors that
528 facilitate tissue growth and repair. Similar processes are mirrored in at least two cases during
529 development. First, neuroepithelial cells produce VEGF to stimulate initial embryonic brain
530 angiogenesis (Breier et al., 1992; Raab et al., 2004). Second, endothelial cues drive
531 oligodendrocyte precursor cell attachment and migration along vessels in the embryonic nervous
532 system (Tsai et al., 2016), and oligodendrocyte precursor-derived HIF-dependent factors,
533 including VEGF, subsequently drive early postnatal angiogenesis (Yuen et al., 2014; Zhang et
534 al., 2020). The ectopic migration of reparative SVZ cells towards injury may represent a reuse of
535 tissue growth mechanisms from development.

536 We have demonstrated that the SVZ cytogenic response to stroke primarily produces
537 undifferentiated precursors that localize to peri-infarct regions – the site of neural repair. SVZ-
538 derived cells produce VEGF that is critical for effective vascular and synaptic plasticity, and
539 ultimately behavioral recovery. Thus, our findings position SVZ cytogenesis as a mechanism that
540 promotes recovery via trophic support rather than cell replacement. These findings provide
541 insight into a fundamental brain repair process and may be relevant for informing treatment
542 strategies.

543 **Materials and Methods**

544 *Subjects and experimental design*

545 Young adult (3-6 months) and aged (12-16 months) mice of both sexes were used. All
546 mice were on a predominantly C57BL/6 background. Transgenic strains were Rosa-CAG-LSL-
547 tdTomato (Ai14, JAX #007914), Nestin-CreER (JAX #016261), ASCL1-CreER (JAX #012882),
548 Rosa-CAG-LSL-Sun1-sfGFP (JAX #021039), GFAP-TK (JAX #005698), Thy1-GFP M-line
549 (JAX #007788), floxed Vegfa (Genentech) (Gerber et al., 1999) (Supplemental Table 1). Mice
550 were bred locally. Animals were housed 2-5 per cage with free access to food and water, except
551 during periods of restricted feeding for behavioral training and assessment. Animals were
552 randomized to groups except when group assignment was dependent on genotype.
553 Experimentation and analysis were done blinded to group allocation. Experiments consisted of 1-
554 5 cohorts of animals. Sample sizes were based on past work using similar methods (Benner et al.,
555 2013; Brown et al., 2007; Clark et al., 2019; Williamson et al., 2021, 2020).

556

557 *Drug administration*

558 100 mg/kg of 20 mg/mL tamoxifen dissolved in corn oil was given (i.p.) daily for 5 consecutive
559 days. 100 mg/kg of 10 mg/mL BrdU dissolved in saline was given (i.p.) once or twice per day
560 for 2 or 6 consecutive days. Ganciclovir dissolved in saline was delivered continuously for 14
561 days via subcutaneous osmotic pumps (Azlet) at a rate of 6.25 µg/hr. In some experiments, a
562 second course of ganciclovir was given beginning two weeks after stroke to maintain stem cell
563 ablation.

564

565 *Cranial window implantation*

566 Chronic glass cranial windows were placed over forelimb motor cortex (Clark et al.,
567 2019; Williamson et al., 2021, 2020). Isoflurane (3% induction, 1-2% maintenance) in oxygen
568 was used for anesthesia. Circular craniotomies (~4.5 mm diameter) were made 1.5 mm lateral
569 from Bregma. 4 mm glass windows (Warner Instruments) were secured in place with
570 cyanoacrylate, and exposed skull was covered with dental cement. Carprofen (5 mg/kg, i.p.) was
571 given daily for 7 days to minimize inflammation.

572

573 *Ischemic stroke*

574 To model stroke, unilateral photothrombotic lesions were induced in the forelimb region
575 of motor cortex (Tennant et al., 2011; Williamson et al., 2021). Isoflurane (3% induction, 1-2%
576 maintenance) in oxygen was used for anesthesia. Body temperature was maintained with a heated
577 pad for the duration of anesthesia. Stroke was induced through the intact skull by making a scalp
578 incision, administering rose bengal (0.15 mL, 15 mg/mL, i.p.), and illuminating the skull 2 mm
579 lateral from Bregma with a surgical lamp (Schott KL 200) for 15 minutes through a 3 mm
580 aperture. For animals with cranial windows, penetrating arterioles supplying motor cortex were
581 identified by live speckle contrast imaging, and subsequently targeted with a 20 mW 532 nm
582 laser for 15 minutes after administering rose Bengal (0.2 mL, 15 mg/mL, i.p.) (Williamson et al.,
583 2020). Sham stroke procedures involved omitting either illumination or rose bengal.

584

585 *Virus injections*

586 Cortical layer V was targeted for virus injections with a Drummond Nanoject II
587 microinjector through a pulled pipette. Injections were made 0.7 mm below the pial surface at
588 three locations relative to Bregma: 2 mm anterior, 2 mm lateral; 0.5 mm anterior, 3.2 mm lateral;
589 and 1 mm posterior, 2.8 mm lateral. 230 nL was injected per site at a rate of 46 nL/min. The
590 pipette was left in place for 2 minutes after the final injection at each site before it was slowly
591 removed. See Supplemental Table 1 for details on viruses.

592

593 *2-photon imaging of dendritic spines*

594 Mice were anesthetized with isoflurane (3% induction, ~1.5% maintenance) in oxygen
595 and head-fixed to minimize breathing artifacts. Imaging was done with a Prairie Ultima 2-photon
596 microscope with a Ti:Sapphire laser (MaiTai, Spectra Physics) tuned to 870 nm. Image stacks
597 were acquired with 512 x 512 pixel resolution and 0.7 μm z step size using a water-immersion
598 20 \times /1.0 (Olympus) objective. 4x magnification yielded a 117.2 μm x 117.2 μm field of view.

599 Imaging was done weekly, including two pre-stroke and four post-stroke imaging
600 sessions. During pre-stroke imaging, typically 6-10 regions of unobstructed dendrites were
601 imaged to a depth of ~150 μm . Regions were selected based on a predicted proximity of <1 mm
602 from the infarct. After stroke, the infarct border was identified from blood flow maps and loss of
603 GFP fluorescence (Supplemental Figure 4). Regions within 700 μm of the infarct border were re-
604 imaged at subsequent time points in order to track spine dynamics during recovery. In some
605 animals, additional imaging regions were added after stroke. Time lapse images of dendrites >30
606 μm in length with clearly visible spines were analyzed to quantify spine formation and

607 elimination, and persistence of newly formed spines (Clark et al., 2019; Joy et al., 2019; Tennant
608 et al., 2017).

609

610 *Blood flow imaging*

611 Blood flow was imaged through cranial windows with multi-exposure speckle imaging, a
612 label-free, quantitative, optical method (Clark et al., 2019; Williamson et al., 2021, 2020).
613 Anesthetic level was consistent across all imaging sessions (1.25% isoflurane in oxygen). Two
614 pre-stroke images were collected to establish baseline blood flow. Post-stroke images were
615 collected on days 2, 5, 14, and 28. Each imaging session lasted <10 mins. Blood flow was
616 measured in parenchymal regions and tracked over time as before (Williamson et al., 2021).

617

618 *Behavioral testing*

619 Skilled forelimb use was assessed with the single seed reaching task, which is highly
620 sensitive to deficits caused by motor cortical damage and is translationally relevant (Clark et al.,
621 2019; Klein et al., 2012; van Lieshout et al., 2021; Williamson et al., 2020). Animals were food
622 restricted to ~90% free feeding weight to encourage reaching. First, animals were shaped on the
623 task and the preferred paw for reaching was determined over 2-5 days. Training was done over
624 15 sessions, once per day, five days per week. Each session consisted of 30 trials. For each trial,
625 animals were allowed up to two reach attempts. A successful reach was defined as the animal
626 grasping the seed and bringing it to its mouth. Failure was defined as missing the seed, knocking
627 it out of the well, or releasing it before it was brought to the animal's mouth. Baseline

628 performance was defined as the mean success rate per trial over the last three training sessions.
629 Inclusion criteria was a minimum baseline success rate of 30%. 1 mouse (GFAP-TK+GCV) from
630 the experiment in Figure 2,8 mice (n = 7 control, n = 1 GFAP-TK+GCV) from the experiment in
631 Figure 3, and 2 mice (n = 1 AAV-eGFP, n = 1 AAV-VEGF-eGFP) from the experiment in
632 Supplemental Figure 7 failed to meet this threshold and were excluded. Test sessions were done
633 on days 3, 7, 14, 21, and 28 post-stroke.

634

635 *Histology and image analysis*

636 Mice were euthanized by overdosed with a pentobarbital/phenytoin solution followed by
637 perfusion with 0.1M phosphate buffer and 4% paraformaldehyde in phosphate buffer. Brains
638 were postfixed overnight at 4°C. 35 µm coronal sections were collected with a vibratome
639 (VT1000S, Leica). To label vasculature, mice were retro-orbitally injected with 0.1 mL of
640 Dylight 594- or 649-conjugated tomato lectin 5 minutes prior to perfusion (Williamson et al.,
641 2021, 2020). To examine vascular permeability, mice were retro-orbitally injected with 0.1 mL
642 of 50 mg/mL FITC-conjugated albumin 2 hours prior to perfusion.

643 To create lesion reconstructions and quantify lesion volume, one set of every fifth section
644 was Nissl stained. Lesions were reconstructed as previously described (Kim et al., 2018). Lesion
645 volume was calculated using Cavalieri's method as the difference in volume between uninjured
646 and injured cortex.

647 Immunohistochemical staining was done by washing tissue in phosphate buffered saline
648 (PBS), blocking with 10% donkey serum in PBS with 0.25% Triton for 60 minutes, incubating
649 with primary antibodies overnight (antibodies and dilutions are reported in Supplemental Table

650 1), washing in PBS, incubating with species appropriate 405-, 488-, 594, or 647-conjugated
651 secondary antibodies, and washing a final time in PBS. Tissue was pretreated with 2 N HCl (30
652 mins) followed by 0.1 M boric acid (10 mins) when staining for BrdU.

653 Confocal images with 1-2 μ m Z-step size were collected with a Leica TCS SP5
654 microscope. 20 \times /0.7 NA and 40 \times /1.0NA objectives were used. Acquisition settings were
655 consistent between samples. Typically, 3 sections were imaged per region of interest per mouse.

656 FIJI was used for image analysis. Area fraction and fluorescence intensity were
657 quantified as before (Williamson et al., 2021). The optical disector method was used to count cell
658 density. Cell density was calculated by number of cells / (frame area \times section thickness).

659

660 *Statistics*

661 Data are expressed as mean \pm S.E.M. Measurements from individual animals are shown on plots
662 as datapoints where possible. Data were analyzed with GraphPad Prism version 9.3. Independent
663 samples were compared with two-tailed t tests. Variance was assessed with F tests, and Welch's
664 corrected t tests were used in cases where variance was significantly different. One- and two-way
665 ANOVAs, mixed-effects analyses, and linear regressions were used as noted in the text. Post hoc
666 tests were used following significant ANOVA as noted in the text. Details on the statistical tests
667 used for each experiment are located in the Results and figure legends. Alpha was set at P < 0.05.

668

669 *Study approval*

670 Animal use was in accordance with a protocol approved by the Institutional Animal Care and
671 Use Committee at the University of Texas at Austin.

672 **Author contributions**

673 Conceptualization: M.R.W.; Methodology: M.R.W., T.A.J., and M.R.D.; Investigation and
674 analysis: M.R.W., S.P.L., R.L.F., N.A.D., and J.L.R.; Writing – Original Draft: M.R.W.; Writing
675 – Review & Editing: all authors; Supervision: T.A.J. and M.R.D.; Funding Acquisition: A.K.D.,
676 M.R.D., and T.A.J.

677

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684 Austin.

685

686 **Competing Interests**

687 None.

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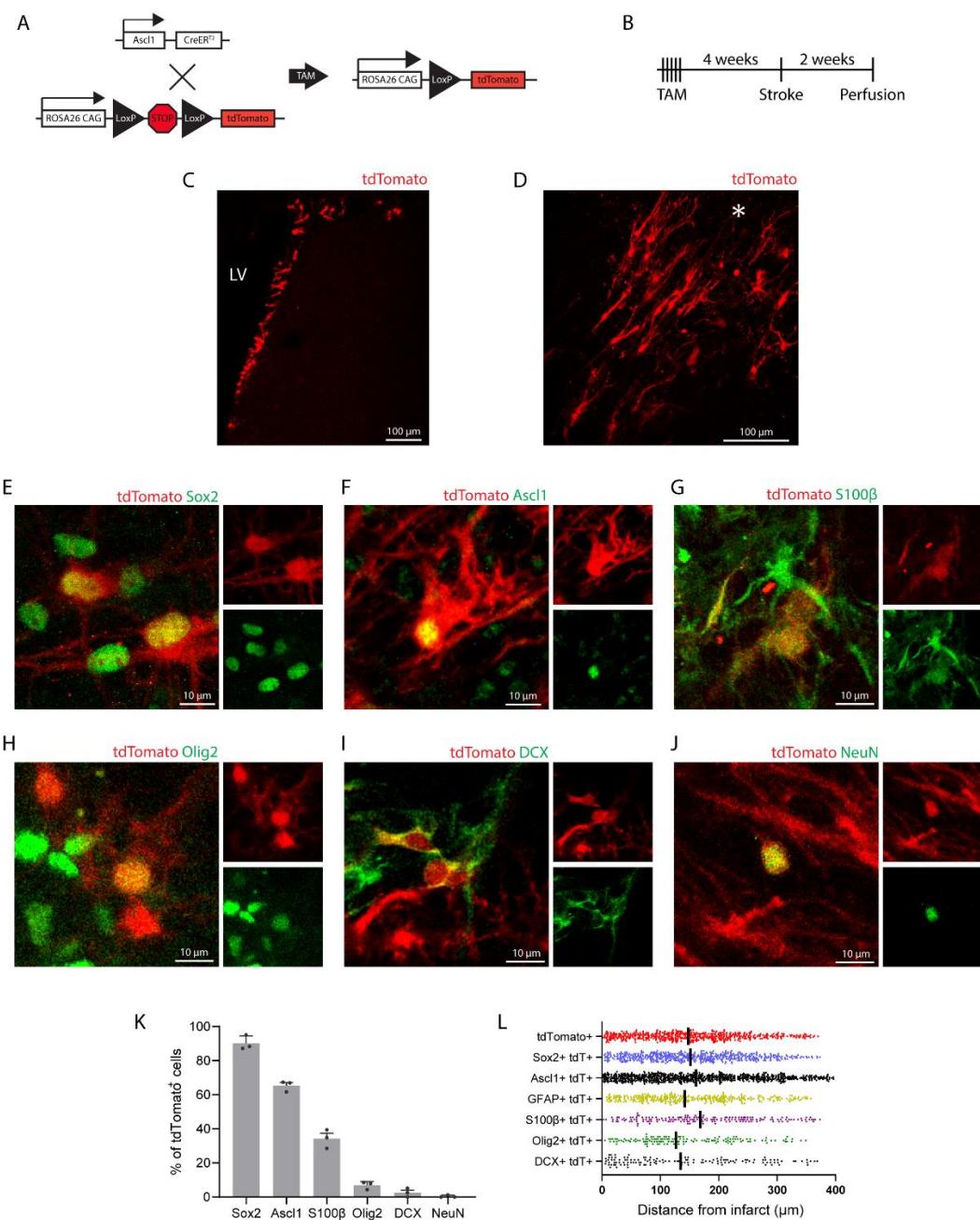
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903

904 **Supplemental figures and tables**

905



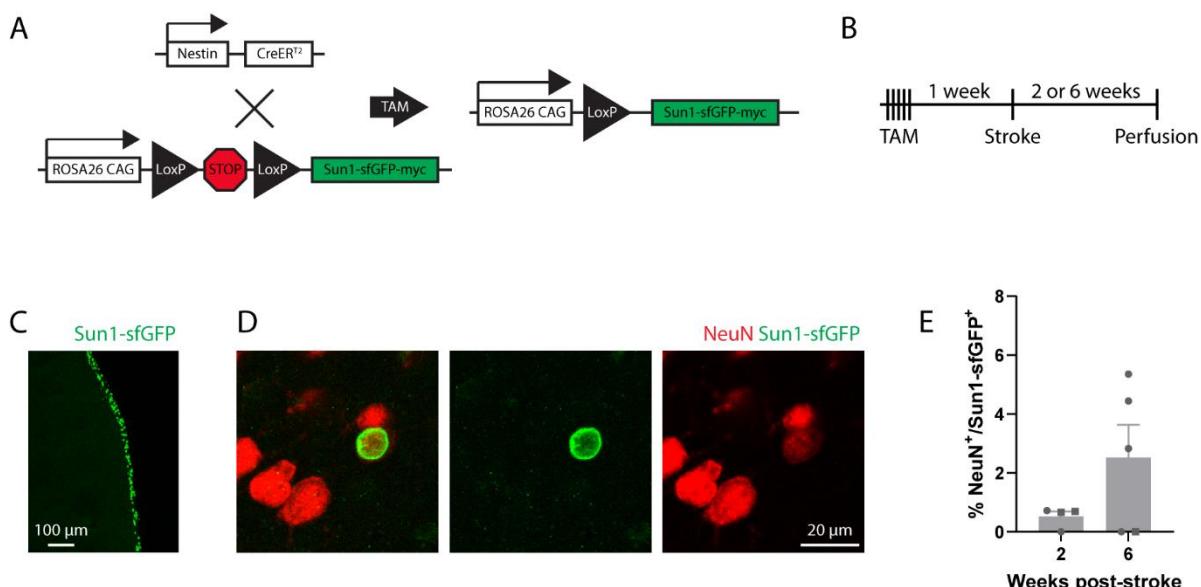
906

907 **Supplemental Figure 1. SVZ-derived cells are predominantly undifferentiated precursors**
908 **and astrocytes.**

909 A) Schematic of genetic lineage tracing system where tamoxifen (TAM) induces indelible
910 expression of tdTomato in *Ascl1*-expressing neural stem cells, intermediate progenitor cells, and
911 their progeny. B) Experimental design. C) Image showing tdTomato expression in the SVZ. LV,
912 lateral ventricle. D) Representative image of tdTomato⁺ cells localized to peri-infarct cortex 2
913 weeks after a cortical stroke (asterisk). E-J) Lineage tracing using *Ascl1*-CreER; Ai14 mice
914 corroborates the major finding of Figure 1, that SVZ-derived cells in peri-infarct cortex are
915 predominantly undifferentiated precursors and astrocytes. Representative images illustrating co-
916 labeling of lineage traced tdTomato⁺ cells with differentiation stage-specific markers at two
917 weeks post-stroke (E) *Sox2*, (F) *Ascl1*, (G) *S100 β* , (H) *Olig2*, (I) *DCX*, (J) *NeuN*). K) Quantification
918 of marker expression by % of tdTomato⁺ cells. Data are presented as mean \pm SEM. L) Spatial
919 distribution of cells relative to the infarct border by marker expression at 2 weeks post-stroke in
920 *Nestin*-CreER; Ai14 mice. Each point indicates a single cell. Vertical lines indicate medians.
921 There was no clear pattern of spatial organization by cell type. Distance was measured as the
922 distance to the nearest part of the infarct border within a given section for each cell. Only cells
923 within 400 μ m of the infarct border and within cortex were included in this analysis.

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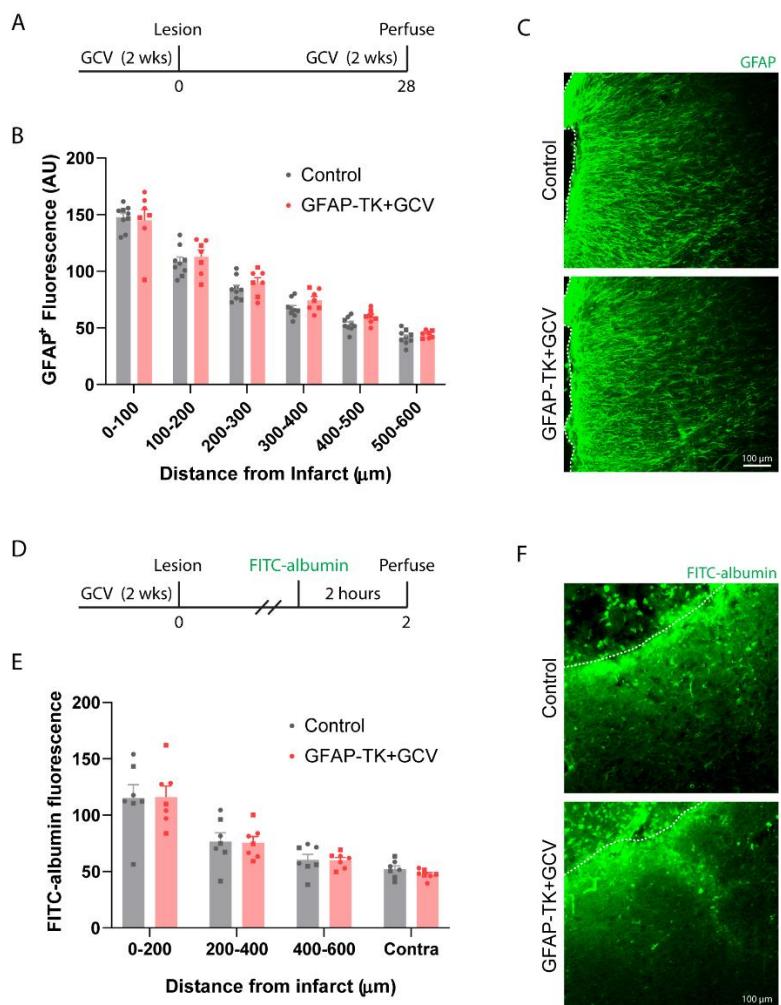


926

927 **Supplemental Figure 2. Few SVZ-derived cells become neurons.**

928 A) Schematic of genetic lineage tracing system where tamoxifen (TAM) induces indelible
929 expression of a nuclear membrane-bound Sun1-sfGFP in neural stem cells and their progeny. B)
930 Experimental design. C) Image showing Sun1-sfGFP expression in the SVZ. D) Representative
931 image of a NeuN⁺Sun1-sfGFP⁺ nucleus in peri-infarct cortex. E) Quantification of SVZ-derived
932 neurons. Data were derived from a combined 1014 nuclei counted across nine mice (n = 4 at two
933 weeks, n = 5 at six weeks). There was not a significant difference between time points ($t(4.2) =$
934 1.79, $p = 0.145$, Welch's corrected t test). Data are presented as mean \pm SEM. Datapoints
935 representing males are shown as circles; datapoints representing females are shown as squares.

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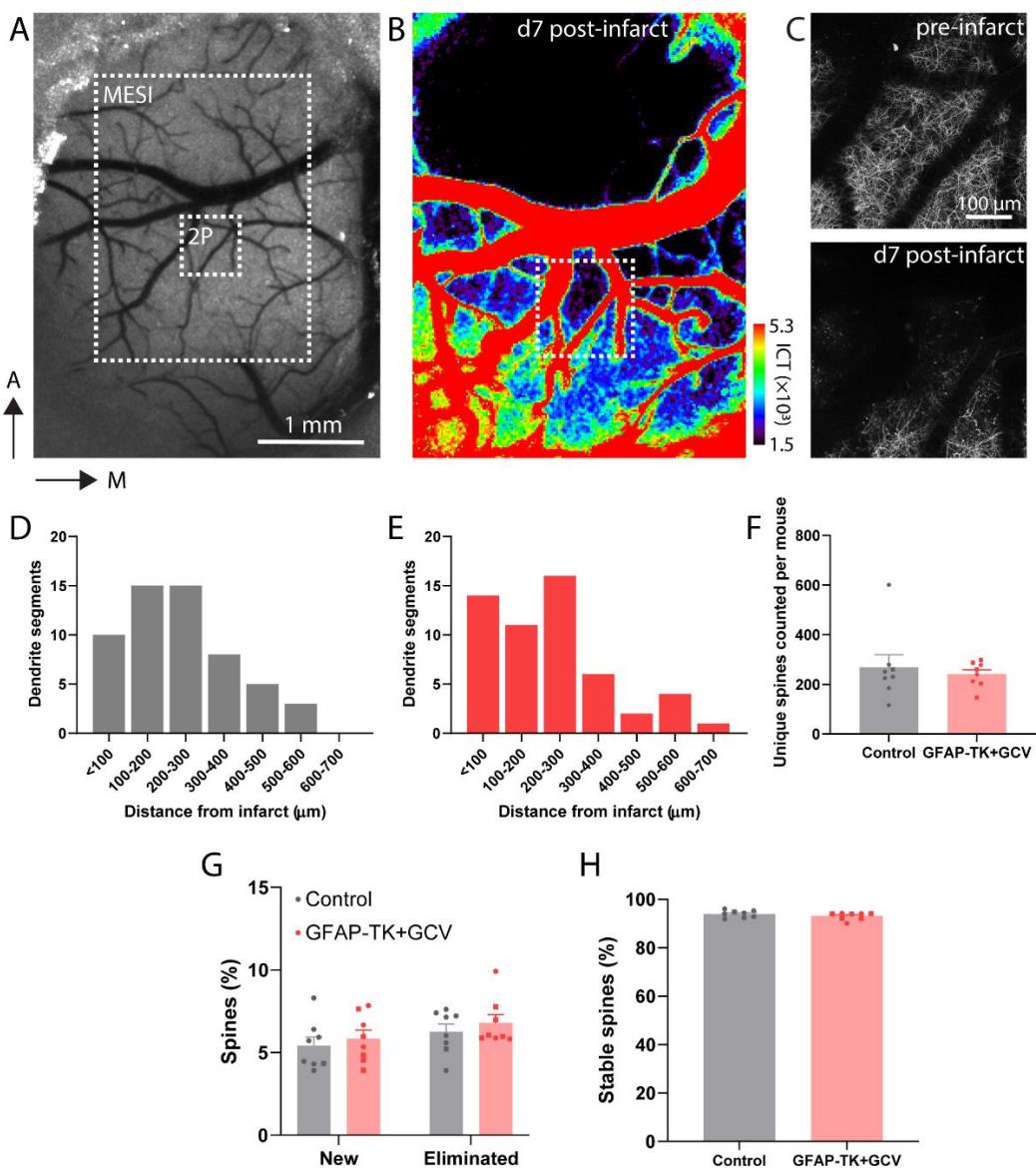


937

938 **Supplemental Figure 3. Neural stem cell ablation does not alter parenchymal astrocyte**
939 **reactivity or vascular permeability after stroke.**

940 A) Experimental timeline for examining GFAP fluorescence. Tissue was obtained 28 days post-
941 stroke (n = 9 controls (TK^{-/-}), n = 7 GFAP-TK^{+/+}). To assess parenchymal astrocyte reactivity,
942 images were taken in superficial peri-infarct cortex where few SVZ-derived cells localize. B)
943 GFAP⁺ fluorescence was increased near the lesion, but declined with distance away,
944 characteristic of astrocyte reactivity (distance effect $F(5, 84) = 172.9, p < 0.001$). There was no
945 significant group effect ($F(1, 84) = 2.8, p = 0.100$). C) Representative images of GFAP
946 immunostaining in peri-infarct cortex. Dashed lines indicate the lesion border. D) Experimental

947 design for examining vascular permeability. Two days post-stroke, animals were retro-orbitally
948 injected with FITC-conjugated albumin 2 hours before perfusion (n = 7 controls (TK^{-/-}), n = 7
949 GFAP-TK^{+/−}). E) FITC-albumin fluorescence was greatest near the infarct border and declined
950 with distance away, consistent with injury-induced vascular permeability (distance effect F(3,
951 48) = 36.9, p < 0.001). There was no significant effect of group (F(1, 48) = 0.07, p = 0.789). F)
952 Representative images of FITC-albumin fluorescence in peri-infarct cortex. Dashed lines
953 indicate the lesion border. Data are presented as mean ± SEM. Datapoints representing males are
954 shown as circles; datapoints representing females are shown as squares.



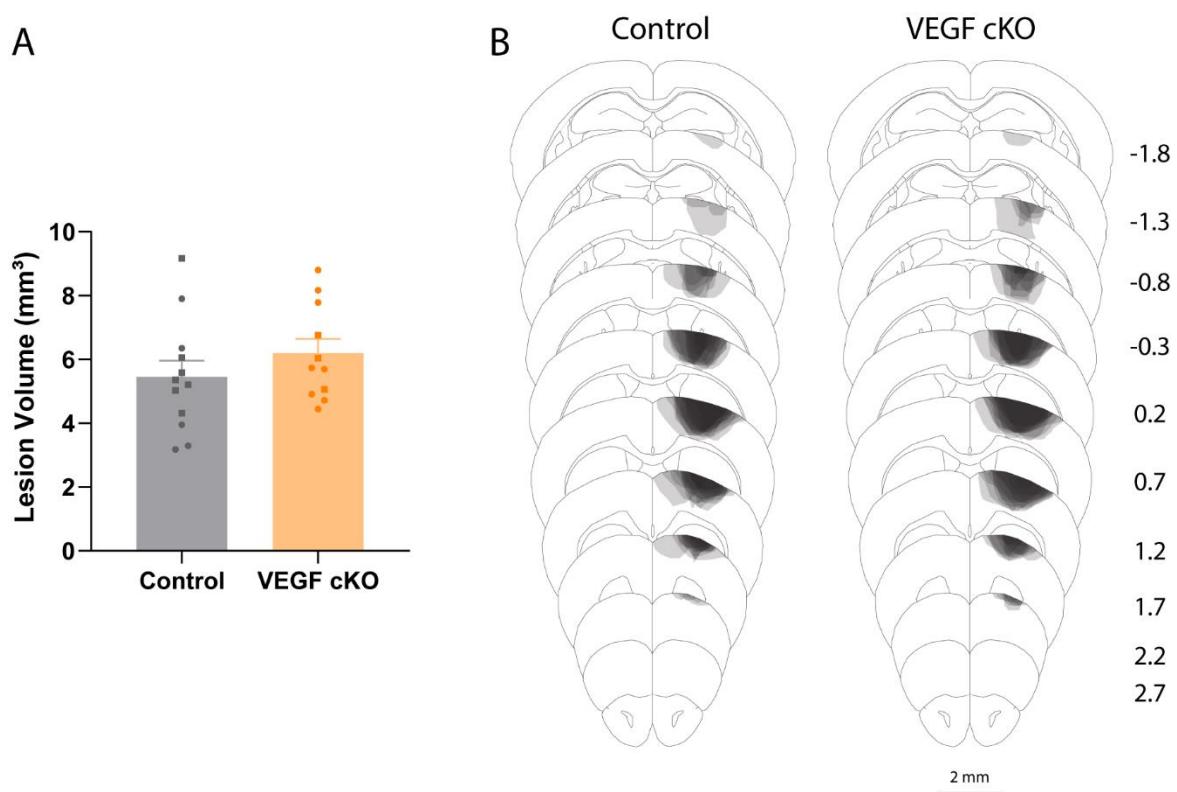
955

956 **Supplemental Figure 4. Additional data 2-photon imaging data.**

957 A) MESI accurately delineates the infarct border. Widefield laser speckle contrast image
958 showing the cortical surface through a cranial window. B) Multi-exposure speckle imaging
959 (MESI) image of blood flow corresponding to the region indicated in panel A. C) Two photon
960 image of GFP-labeled apical dendrites (Thy1-GFP) corresponding to the region indicated in
961 panels A and B. The infarct border revealed by MESI (black region in B) matches the region in

962 which GFP fluorescence is absent. D-E) Distribution of the locations of analyzed dendrite
963 segments relative to the infarct border for control (D) and GFAP-TK+GCV mice (E) (number of
964 segments is summed across mice). F) Numbers of unique longitudinally tracked dendritic spines
965 by group. Each point corresponds to an individual animal. Pre-stroke spine turnover (G) and
966 stability (H) were not different between groups ($t(14) \leq 1.1$, $p \geq 0.272$). Data are presented as
967 mean \pm SEM. Datapoints representing males are shown as circles; datapoints representing
968 females are shown as squares.

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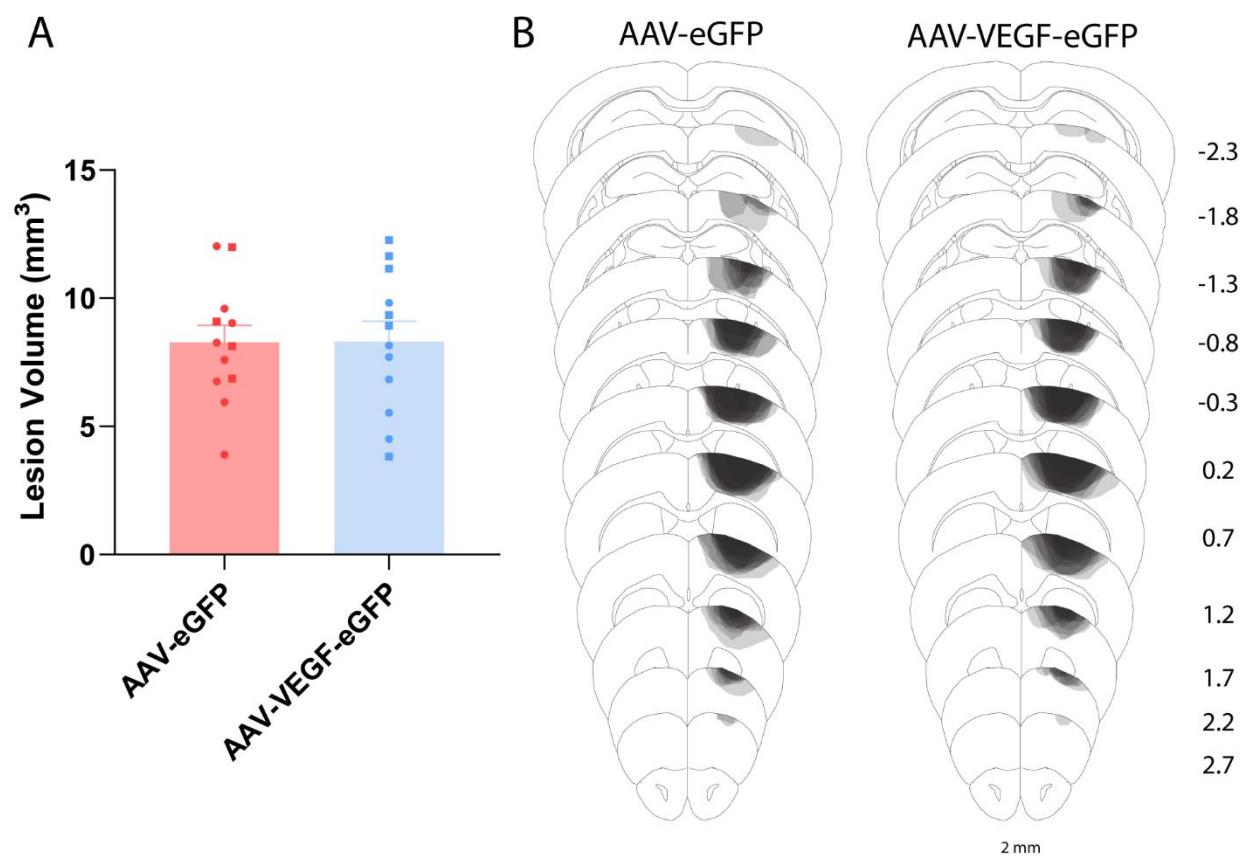


970

971 **Figure S5. VEGF cKO did not affect lesion size or location.**

972 A) Lesion size was not different between groups ($t(21) = 1.08$, $p = 0.292$). B) Lesion
973 reconstruction. Darker shades indicate greater overlap between animals.

974



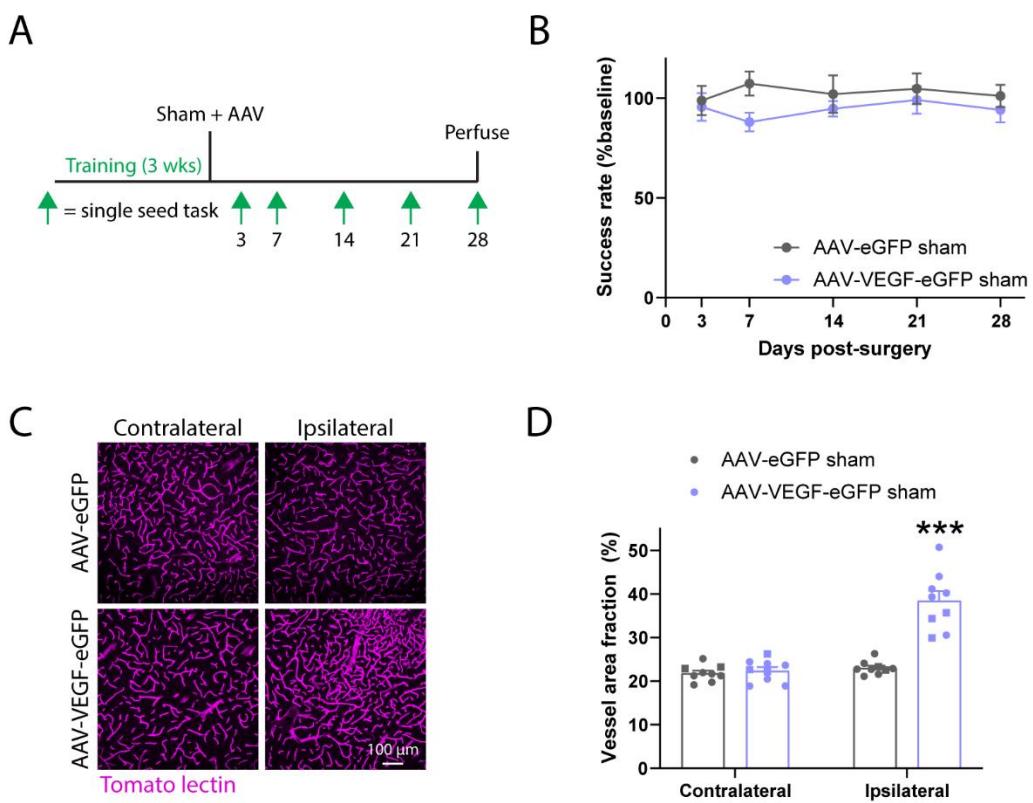
975

976 **Figure S6. AAV-VEGF-eGFP did not affect lesion size or location.**

977 A) Lesion size was not different between groups ($t(22) = 0.04$, $p = 0.967$). B) Lesion

978 reconstruction. Darker shades indicate greater overlap between animals.

979



980

981 **Figure S7. AAV-VEGF-eGFP does not affect motor function in sham-operated animals.**

982 A) Experimental timeline. Wildtype mice were trained on the single seed reaching task,
983 subjected to a sham stroke procedure, and injected with either AAV-eGFP or AAV-VEGF-eGFP
984 ($n = 8/\text{group}$). Performance on the single seed task was periodically tested up to 28 days post-
985 surgery. B) Performance on the single seed task was not different between groups ($F(1, 14) =$
986 3.8, $p = 0.071$). C, D) Representative images (C) and quantification (D) of vasculature in
987 contralateral and ipsilateral cortex (relative to AAV injection site). AAV-VEGF-eGFP increased
988 vessel density in ipsilateral cortex ($t(16) = 6.8$, $***p < 0.001$) ($n = 9/\text{group}$). Data are presented
989 as mean \pm SEM. Where individual datapoints are shown, datapoints representing males are
990 shown as circles; datapoints representing females are shown as squares.

991

992 **Supplemental Table 1. List of reagents.**

Antibodies	Source	Identifier
Rabbit polyclonal anti-ASCL1 (1:1000)	Abcam	ab74065
Rabbit polyclonal anti-ASCL1 (1:500)	Cosmo Bio	CAC-SK-T01-003
Rabbit polyclonal anti-CD133 (1:1000)	Abcam	ab19898
Rabbit monoclonal anti-BDNF (1:1000)	Abcam	ab108319
Rabbit polyclonal anti-BrdU (1:500)	Abcam	ab152095
Rat monoclonal anti-BrdU (1:500)	Abcam	ab6326
Goat polyclonal anti-DCX (1:500)	Santa Cruz Biotech.	Sc-8066
Rabbit monoclonal anti-ERG (1:500)	Abcam	ab92513
Rabbit polyclonal anti-FGF2 (1:500)	Sigma	F-3393
Rabbit polyclonal anti-GDNF (1:100)	Abcam	ab18956
Rabbit polyclonal anti-GFAP (1:1000)	Dako	Z0334
Chicken polyclonal anti-GFP (1:5000)	GeneTex	GTX13970
Goat polyclonal anti-HSV thymidine kinase (1:1000)	Santa Cruz Biotech.	Sc-28038
Rabbit monoclonal anti-Id2 (1:1000)	CalBioreagents	M213
Rabbit polyclonal anti-Ki67 (1:500)	Abcam	ab66155
Rabbit monoclonal anti-NeuN (1:2000)	Millipore	MABN140
Rabbit polyclonal anti-Olig2 (1:1000)	Millipore	AB9610
Rabbit monoclonal anti-S100 β (1:1000)	Abcam	ab52642
Rabbit polyclonal anti-Sox2 (1:1000)	Millipore	AB5603
Rabbit polyclonal anti-VEGF (1:1000)	Millipore	ABS82
Rabbit polyclonal anti-VEGF AF647 conjugate (1:1000)	Millipore	ABS82-AF647

Rabbit polyclonal anti-VEGF (1:250)	Sigma	07-1420
Alexa Fluor 488-conjugated donkey anti-chicken (1:500)	Jackson ImmunoResearch	703-545-155
Alexa Fluor 488-conjugated donkey anti-goat (1:500)	Jackson ImmunoResearch	705-545-147
Alexa Fluor 594-conjugated donkey anti-goat (1:500)	Jackson ImmunoResearch	705-585-147
Alexa Fluor 488-conjugated donkey anti-rabbit (1:500)	Jackson ImmunoResearch	711-545-152
Alexa Fluor 594-conjugated donkey anti-rabbit (1:500)	Jackson ImmunoResearch	711-585-152
Alexa Fluor 647-conjugated donkey anti-rabbit (1:500)	Jackson ImmunoResearch	711-605-152
Alexa Fluor 488-conjugated donkey anti-rat (1:500)	Jackson ImmunoResearch	712-545-153
Alexa Fluor 594-conjugated donkey anti-rat (1:500)	Jackson ImmunoResearch	712-585-153

Viruses	Source	Identifier
AAV5-CaMKIIa-eGFP	Addgene	50469-AAV5
AAV5-EF1 α -eGFP	Addgene	105547-AAV5
AAV5-EF1 α -VEGFA-P2A-eGFP	Vector Builder	N/A

Chemicals	Source	Identifier
Rose Bengal	Sigma	Cat# 330000
Bromodeoxyuridine	Sigma	Cat# B5002
FITC-conjugated albumin	Sigma	Cat# A9971
Dylight 594-conjugated tomato lectin	Vector Labs	DL-1177-1
Dylight 649-conjugated tomato lectin	Vector Labs	DL-1178-1

Ganciclovir	Roche	N/A
Mice	Source	Identifier
Ai14: B6.Cg-Gt(ROSA)26Sortm14(CAG-tdTomato)Hze/J	The Jackson Laboratory	JAX #007914
GFAP-TK: B6.Cg-Tg(Gfap-TK)7.1Mvs/J	The Jackson Laboratory	JAX #005698
Nestin-CreER: C57BL/6-Tg(Nes-cre/ERT2)KEisc/J	The Jackson Laboratory	JAX #016261
ASCL1-CreER: Ascl1tm1.1(Cre/ERT2)Jejo/J	The Jackson Laboratory	JAX #012882
CAG-LSL-Sun1-sfGFP: B6;129-Gt(ROSA)26Sortm5(CAG-Sun1/sfGFP)Nat/J	The Jackson Laboratory	JAX #021039
Thy1-GFP: Tg(Thy1-EGFP)MJrs/J	The Jackson Laboratory	JAX #007788
floxed Vegfa	Genentech, Gerber et al. 1999	N/A

993