Frontal cortical regulation of neurogenesis and cellular proliferation in the ventral subventricular zone

Authors/Affiliations:

Moawiah M Naffaa[1,2,†], Rehan R Khan[2], Chay T. Kuo, Henry H. Yin[1,3,†]

1. Department of Psychology and Neuroscience; Duke University

2. Department of Cell Biology, Duke University, School of Medicine

3. Department of Neurobiology; Duke University School of Medicine

†**Correspondence**:

Moawiah Naffaa moawiah.naffaa@duke.edu

Henry Yin hy43@duke.edu

Key words: postnatal neurogenesis, subventricular zone, frontal cortex, anterior cingulate, subep-ChAT[+], neural stem cells proliferation, calretinin

Summary

Neurogenesis and differentiation of the neural stem cells (NSCs) in the subventricular zone (SVZ) are controlled by cell-intrinsic molecular pathways that interact with extrinsic signaling cues. Here we identified a novel circuit that regulates neurogenesis and cellular proliferation in the lateral ventricle SVZ (LV-SVZ). Our results demonstrate direct glutamatergic inputs from the frontal cortex as well as local inhibitory interneurons, control the activity of distinctive cholinergic neurons in the subependymal zone (subep-ChAT$^+$). *In vivo* optogenetic stimulation and inhibition in this circuit were sufficient to control local SVZ neurogenesis, LV NSCs proliferation, and SVZ cellular divisions in ventral SVZ. These findings shed light on local and distal neural circuit activity-dependent regulation of postnatal and adult SVZ neurogenesis and LV-SVZ cellular proliferation.

The rodent subventricular zone (SVZ) of the lateral ventricles (LV) is a major site of postnatal neurogenesis [1]. Neurogenesis in SVZ provides a useful model system for studying neuronal regeneration and tissue remodeling in the adult brain [2,3]. LV neural stem cells (NSCs) divide asymmetrically for self-renewal or differentiate to become transient amplifying intermediate progenitors (TAPs) [4-6]. These intermediate progenitors divide and differentiate into doublecortin-positive neuroblasts [7,8], which migrate to the olfactory bulb (OB) [7,9,10], where they become mature interneurons that are incorporated into the local circuitry [11,12]. The generation of adult-born neurons from the SVZ contributes to experience-dependent plasticity in the postnatal brain [13,14] and plays a critical role in social behavior in rodents [12,15,16].

The LV NSCs receive inputs from a variety of sources, including neighboring NSCs, TAPs, and neuroblasts[17,18]. Different neurotransmitters, including GABA, dopamine, and serotonin, released by neurons in different brain regions, are also known to regulate postnatal and adult SVZ neurogenesis [19-24]. However, little is known about the local and distal circuity which directly or indirectly influence the SVZ neurogenesis.

Recent work has identified a small population of cholinergic neurons in the subependymal space to the lateral ventricle. These neurons, which express Choline acetyltransferase (ChAT[+]), can modulate the proliferation of LV NSCs and neuroblasts in an activity-dependent manner [25]. They are unlike neighboring striatal cholinergic interneurons, and release acetylcholine (ACh) into the SVZ niche. We hypothesize that the population of subependymal ChAT[+] (subep-ChAT^{+}) neurons in the ventral domain of SVZ is a key node in the neural circuit regulation of SVZ neurogenesis and LV NSCs proliferation. Using neural circuit tracing, electrophysiology and *in vivo* optogenetic strategies, we identified a novel neural circuit involving anterior cingulate glutamatergic projections and local calretinin-positive interneurons

that can regulate the activity of subep-ChAT$^+$ neurons. Consequently, this circuit directly

modulate SVZ neurogenesis, activity of LV NSCs proliferation and SVZ cellular divisions in the

ventral SVZ. These findings revealed a novel circuitry by which frontal cortical inputs can

influence cholinergic signaling in ventral SVZ to modulate neurogenesis and cellular

proliferation.

Results

Excitatory and inhibitory presynaptic inputs to the subep-ChAT$^+$ neurons

Unlike striatal cholinergic neurons, subep-ChAT$^+$ neurons do not show tonic activity in

the absence of synaptic inputs, suggesting that excitatory inputs are needed to activate these

neurons[26]. However, it is currently unknown where the excitatory inputs are generated from and

how they influence the activity of subep-ChAT$^+$ neurons. We hypothesized that glutamatergic

inputs from cortical projection neurons may provide excitatory drive of the subep-ChAT$^+$

neurons. The cortical neurons mainly express vesicular glutamate transporter 1 protein

(VGlut1)[27]. To selectively stimulate these glutamatergic inputs, we crossed *VGlut1-Cre* mice

with *ChAT-eGFP* and ChR2-tdTomato transgenic mouse line (*Ai27*) that expresses

channelrhodopsin in a Cre-dependent manner[28]. The resulting *VGlut1-Cre; ChAT-eGFP; Ai27*

mice express channelrhodopsin in all VGlut1$^+$ neurons **(Fig. 1A)**.

Whole-cell recordings from subep-ChAT$^+$ neurons of *VGlut1-Cre; ChAT-eGFP; Ai27*

mice showed robust firing in response to 473-nm light activation of VGlut1$^+$ axon terminals

(firing frequency = 5.375 ± 0.324 Hz) **(Fig. 1B)**. In voltage-clamp mode (holding at -60 mV),

photo-stimulation evoked excitatory postsynaptic currents (EPSCs) in the subep-ChAT$^+$ neurons.

The induced currents can be blocked by glutamatergic receptors blockers AP5 (NMDA receptor

antagonist) and CNQX (AMPA receptor antagonist) (Current amplitude = 98.2 ± 10.7 pA, and

current latency = 8.6 ± 0.62 ms) **(Fig. 1C)**. These findings demonstrate the existence of cortical

glutamatergic inputs that drive and regulate the activity of subep-ChAT$^+$ neurons.

To examine the inhibitory drivers of subep-ChAT$^+$ neurons, we crossed VGat-ChR2-

eYFP (i.e.: channelrhodopsin expressed in GABAergic neurons) with *ChAT-Cre* and *Ai27*. The

result is VGat-ChR2-eYFP*; ChAT-Cre; Ai27* mice which allow recording from subep-ChAT$^+$

neurons in animals with channelrhodopsin expressed in all GABAergic neurons **(Fig. 1D)**.

Whole-cell recordings from subep-ChAT$^+$ neurons in voltage-clamp mode (holding at -60 mV)

and activating VGAT$^+$ axon terminals via 473-nm laser resulted in consistent inhibitory

postsynaptic currents (IPSCs). The evoked IPSCs in the subep-ChAT$^+$ neurons were blocked by

picrotoxin (GABA$_A$ receptors antagonist) (Current amplitude = 196.1 ± 24.13 pA, and current

latency = 4.5 ± 0.88 ms) **(Fig. 1E)**. These results suggest that subep-ChAT$^+$ neurons receive

direct GABAergic inputs that manage their release of ACh in the SVZ.

Developing rabies virus strategy for tracing neural circuit

Our initial electrophysiological recording results established the presence of excitatory

and inhibitory modulators to the activity of subep-ChAT$^+$ neurons **(Fig. 1F)**. However, it remains

unclear exactly where these inputs originated. To answer this question, we generated a Cre-

dependent *R26R-FLEX-TVA-2A-RabiesG-2A-tdTomato-FLEX (R26F-RTT)* mice to trace the

connectivity of subep-ChAT$^+$ neurons via a single Rabies viral injection **(Fig. 2A & B)**. This

new tracing tool allows for expressing TVA, RabiesG and tdTomato in all Cre-expressing

neurons. The *R26F-RTT* mice were first tested by injecting EnvA G-deleted Rabies-eGFP virus

into the striatum where no nonspecific rabies infections were detected at the injection site

(Extended Fig. 1A & B). For examining the specific expression of TVA, RabiesG and tdTomato

in Cre-dependent mice, *R26F-RTT* mice were crossed with *chat-Cre, VGat-Cre and D2-Cre* to

generate *R26F-RTT; chat-Cre, R26F-RTT; VGat-Cre* and *R26F-RTT; D2-Cre* mice, respectively.

In *R26F-RTT; chat-Cre mice,* all ChAT$^+$ neurons were only found to express tdTomato

(Extended Fig. 1C), which indicates that *R26F-RTT* mice are selectively expressed in Cre-

positive neurons**.** To assess the retrogradely trans-synaptic spread of rabies with *R26F-RTT*,

EnvA G-deleted Rabies-eGFP virus was injected into the striatum of *R26F-RTT; VGat-Cre* and

R26F-RTT; D2-Cre mice **(Extended Fig. 2A & C).** After 7-days, the cortical excitatory neurons

in cingulate and secondary motor cortices which are presynaptically connected to the striatal

VGat$^+$ and D2$^+$ neurons were found labeled with GFP **(Extended Fig. 2B & D)**. This indicates

that rabies can retrogradely spread from Cre-positive VGat$^+$ and D2$^+$ neurons when they are

crossed with *R26F-RTT* mice. Together, these results demonstrated that *R26F-RTT* mice can

express TVA, RabiesG and tdTomato selectively in Cre-expressing neurons and allows for

retrograde presynaptic neuronal tracing.

To allow for neural tracing of cholinergic neurons, we generated *ChAT-Cre; R26F-RTT*

mice to express TVA receptor, RabiesG protein, and tdTomato in all cholinergic neurons **(Fig.**

2B). We first injected EnvA G-deleted rabies-eGFP virus into the striatum of P30 *ChAT-Cre;*

R26F-RTT mice **(Extended Fig. 3A)**. As the striatal cholinergic neurons receive synaptic inputs

from each other[29], an infection of a striatal cholinergic neuron with EnvA G-deleted Rabies-

eGFP virus results in polysynaptic labeling of other neighbor striatal cholinergic neurons, which

subsequently pass rabies to other connected neurons **(Extended Fig. 3B & C)**. Then, EnvA G-

deleted Rabies-eGFP virus was injected into the LV of *ChAT-Cre; R26F-RTT* mice to target

subep-ChAT$^+$ neurons, immediately following mannitol injection to disrupt the ependymal layer

locally **(Fig. 2C & F)**. Using this strategy, we were able to selectively infect a subep-ChAT$^+$

neuron and trace their presynaptic inputs **(Fig. 2D & G)**. The local connectivity of subep-ChAT$^+$

neurons and their projections were checked at 7- and 14-days post rabies injection **(Fig. 2D, E, G**

& H). Unlike striatal cholinergic neurons, subep-ChAT$^+$ neurons in the SVZ niche are not

connected with neighboring striatal cholinergic neurons **(Fig. 2I)**. These results confirm that

subep-ChAT$^+$ neurons have distinct neuronal connectivity.

Local presynaptic inhibitory input to subep-ChAT$^+$ neurons

Injection of EnvA G-deleted Rabies-eGFP virus into the LV of *ChAT- Cre; R26F-RTT*

mice revealed local GFP$^+$ interneurons adjacent to the infected subep-ChAT$^+$ neurons **(Fig. 3A**

& B). While staining for markers of the known inhibitory neurons, colocalization between GFP$^+$

interneurons and calretinin (CR) antibody was noticed **(Fig. 3C)**.

To test if the CR$^+$ interneurons provide local inhibitory inputs to subep-ChAT$^+$ neurons,

we performed whole-cell patch clamp recording from subep-ChAT$^+$ neurons of *Cr-Cre; ChAT-*

eGFP; Ai29 mice **(Fig. 3D)**. A high-chloride internal solution was used to detect inhibitory

postsynaptic currents (IPSCs) while holding at -60 mV in voltage clamp. We stimulated local

CR$^+$ interneurons with blue light and revealed robust evoked IPSCs, which were completely

blocked by picrotoxin (current amplitude = 122.3 ± 9.0 pA, and current latency = 4.9 ± 0.33 ms)

(Fig. 3E). Then, we tested for other known inhibitory interneurons expressing calbindin (CB$^+$),

somatostatin (SST$^+$) or parvalbumin (PV$^+$) **(Extended Fig. 4A)**. Using whole-cell recording from

subep-ChAT$^+$ neurons, an optogenetic stimulation of GABA released from these inhibitory

interneurons (i.e.: CB, SST and PV) did not induce IPSCs **(Extended Fig. 4B)**.

To further confirm the local CR$^+$ interneuron inputs to the subep-ChAT$^+$ neurons, we

used a Patterned Light Stimulator LED controller (Mightex) to target CR$^+$ labelled neurons in

SVZ wholemount of *Cr-Cre; ChAT-eGFP; Ai29* mice. Focal optogenetic activation of a single

CR$^+$ interneuron generated IPSCs in subep-ChAT$^+$ neuron (current amplitude $= 40.14 \pm 3.78$ pA,

and current latency $= 4.7 \pm 0.4$ ms) **(Fig. 3F & G)**. Subsequently, we stained against CR and

ChAT using *C57BL/6J* mice, revealing the presence of 2-4 CR$^+$ interneurons surrounding each

subep-ChAT$^+$ neuron in the LV-SVZ **(Fig. 3H & I)**. Together these results confirm that local

CR$^+$ interneurons are the main source of inhibitory inputs to the subep-ChAT$^+$ neuron.

Presynaptic excitatory inputs from cingulate cortex area 1 (Cg1) to subep-ChAT$^+$ neurons

To illuminate long distance connections for subep-ChAT$^+$ neurons in the brain, EnvA

rVSV-eGFP (EnvA/RABVG-eGFP) virus[30] was injected into the LV of *ChAT- Cre; R26F-RTT*

mice **(Fig. 4A)**. Distal GFP$^+$ projection neurons from the anterior cingulate cortex area 1 (Cg1)

were observed in the ipsilateral hemisphere only **(Fig. 4B)**. To confirm the Cg1 input to subep-

ChAT$^+$ neurons, pAAV-CaMKIIa-hChR2(E123A)-mCherry virus was first injected into the Cg1

region of *C57BL/6J* mice to express mCherry in the GFP$^+$ neurons from Figure 4B **(Fig. 4C)**. By

tracing their projections in the SVZ region, we observed them adjacent to the subep-ChAT$^+$

neurons **(Fig. 4D)**.

To functionally examines a monosynaptic input from Cg1 to the subep-ChAT$^+$ neurons, a

general ChR2 viral vector (pAAV-CaMKIIa-hChR2(E123T/T159C) -EYFP) was injected into

the Cg1 region of *Chat-Cre; Ai9* mice to express ChR2 in the projections of Cg1 neurons **(Fig.**

4E). Optogenetic stimulation of the cingulate cortex projections reliably evoked action potentials

in subep-ChAT$^+$ neurons (firing frequency $= 2.5 \pm 0.327$ Hz) **(Fig. 4F)**. Using voltage clamp, the

optogenetic stimulation induced reliable EPSCs in subep-ChAT$^+$ neurons which were completely

blocked by the glutamate antagonists (AP5 and CNQX) (current amplitude = 64.86 ± 2.424 pA,

and current latency = 9.1 ± 0.7 ms) **(Fig. 4G)**. These results confirm the presence of

glutamatergic neurons in the ipsilateral Cg1 region that excite subep-ChAT$^+$ neurons.

To further confirm the (Cg1-subep-ChAT$^+$) circuit, pAAV-Ef1a-DIO

hChR2(E123T/T159C)-mCherry virus was injected into the Cg1 region of *VGlut1-Cre; ChAT-*

eGFP mice to express ChR2 in Cg1 VGlut1$^+$ neurons **(Fig. 4H)**. In the voltage clamp mode,

reliably EPSCs were generated in subep-ChAT$^+$ neurons upon optogenetic stimulation of Cg1

VGlut1$^+$ neurons projection **(Fig. 4I, Left)**. The induced EPS currents were completely blocked

by AP5 and CNQX, confirming the presence of Cg1 glutamatergic inputs (current amplitude =

53.29 ± 2.96 pA, and current latency = 8.7 ± 0.44 ms) **(Fig. 4I, Right)**. These findings show that

a specific population of cortical neurons in the Cg1 region provides excitatory drive to subep-

ChAT$^+$ neurons.

Calretinin-Cre (Cr-Cre) **mice label the distal Cg1 excitatory presynaptic input to the subep-**

ChAT$^+$ neurons

After identifying Cg1 glutamatergic input into subep-ChAT$^+$ neurons using rabies neural

tracing and electrophysiological recording, (Cg1-subep-ChAT$^+$) circuit was then further tested

by injecting AAVrg viruses (i.e.: retrograde tracing using AAV viruses) into LVs of *C57BL/6J,*

VGlut1-Cre and *CR-Cre* mice. Upon injection of pAAVrg-hSyn-mCherry and pAAVrg-hSyn-

DIO-mCherry viruses into LVs of *C57BL/6J* and *VGlut1-Cre* mice, respectively **(Extended Fig.**

5A & C), we observed mCherry$^+$ neurons in the ipsilateral Cg1 region **(Extended Fig. 5B & D)**.

Surprisingly, injecting pAAVrg-hSyn-DIO-mCherry virus into the LV of *CR-Cre* mice labels

neurons in the ipsilateral Cg1 region **(Fig. 5A &B)**. The detected mCherry$^+$ neurons in the Cg1

of *C57BL/6J, VGlut1-Cre* and *CR-Cre* mice post pAAVrg virus injections and the GFP$^+$ Cg1

neurons in **Fig. 4B** are located along the same rostro-caudal axis of the Cg1 region. These

findings strongly suggest that the Cg1 presynaptic glutamatergic drivers of subep-ChAT$^+$

neurons are calretinin$^+$ (CR$^+$) during the postnatal period.

To examine that these Cg1 glutamatergic inputs to the subep-ChAT$^+$ neurons are CR$^+$

neurons, whole-cell recordings from subep-ChAT$^+$ neurons were first performed in *Cr-Cre;*

ChAT-eGFP; Ai27 mice **(Fig. 5C)**. In voltage mode at -60 mV, optogenetic stimulation of CR$^+$

terminals reliably evoked EPSCs (Current amplitude = 106 ± 6.68 pA, and current latency = 5.13

± 0.44 ms), which were completely blocked by AP5 and CNQX **(Fig. 5D)**. This predicts that the

population of Cg1 glutamatergic that drives the activity of subep-ChAT$^+$ neurons are CR$^+$

postnatally. To further functionally testing whether these Cg1 excitatory CR$^+$ neurons regulate

the activity of subep-ChAT$^+$ neurons, we injected AAV-EF1a-DIO-hChR2(E123T/T159C)-

mCherry virus into the Cg1 region of *Cr-Cre; ChAT-eGFP* mice **(Fig. 5E)**. Whole-cell recording

from subep-ChAT$^+$ neurons in voltage clamp mode induced EPSCs upon optogenetic stimulation

(Current amplitude = 33 ± 2.46 pA, and latency = 5.13 ± 0.44 ms) **(Fig. 5F)**. All previous data

confirm that a specific population of cortical neurons in Cg1 region (VGlut1$^+$ and CR$^+$) provides

excitatory drive to subep-ChAT$^+$ neurons **(Fig. 5G)**.

The total number of labeled Cg1 cortical neurons with *Cr-Cre* mice during postnatal

period is must lower than the number of labeled neurons with *VGlut1-Cre* mice. Thereby, the *Cr-*

Cre mouse is a reliable tool to study the population of glutamatergic neurons from the Cg1

region that control the activity of subep-ChAT$^+$ neurons.

***in vivo* (Cg1-subep-ChAT$^+$) circuit stimulation regulate neurogenesis in the ventral domain**

of SVZ

Based on our earlier findings, we defined the Cg1 presynaptic excitatory drivers of

subep-ChAT$^+$ neurons. Thereafter, we studied the *in vivo* functional role of (Cg1-subep-ChAT$^+$)

circuit on SVZ neurogenesis and LV cellular proliferation. We hypothesized that this circuit

regulate SVZ neurogenesis surrounding subep-ChAT$^+$ neurons by modulating the cellular

proliferation in the ventral SVZ. To fulfill this aim, AAV-EF1a-DIO-hChR2 (E123T/T159C)-

mCherry virus was injected into ipsilateral Cg1, and optical fibers were implanted into the Cg1

regions of *Cr-Cre* mice **(Fig. 6A)**. The optogenetic stimulation was continuously conducted for

2- or 3-days and delivered by TTL control of 473-nm laser, 10 ms pulses at 10 Hz, lasting 10 s,

given once every 1 min **(Fig. 6B and Extended Fig. 6A)**.

In the 2-days circuit stimulation experiments, the mice were perfused immediately after

stimulation cessation and brain slices of the stimulated Cg1 sections were stained against ChAT,

phospho-S6 ribosomal (P-S6) and doublecortin (DCX) antibodies. To confirm (Cg1-subep-

ChAT$^+$) circuit activation in the studied coronal sections, the neuronal activity of subep-ChAT$^+$

neurons in these sections was determined by quantifying the intensity of P-S6 protein **(Extended**

Fig. 6B (B' & B''); upper). Among the included coronal sections in our studies, subep-ChAT$^+$

neurons on the ipsilateral side showed significantly higher P-6S intensity than their counterpart

on the contralateral side within the same brain (43.9 ± 7.3%) **(Extended Fig. 6C)**. In these

sections, the doublecortin (DCX$^+$) neuroblasts surrounding subep-ChAT$^+$ neurons on the

activation side were not clustered and were mainly disordered unlike their counterpart DCX$^+$

neuroblasts on the contralateral side **(Extended Fig. 6B (B' & B''); lower)**. This observed

disorganized pattern of the clustered local DCX$^+$ cells adjacent to the activated subep-ChAT$^+$

neurons may be related to the continuous stimulation of the circuit which result in a local

constant release of ACh. Although the circuit was stimulated for two days, the quantification of

the total number of DCX$^+$ neuroblasts showed no significant difference between the ipsilateral

(activated) and contralateral (control) sides of these coronal sections **(Extended Fig. 6D)**. The

total DCX$^+$ cells in the whole SVZ were also assessed by quantifying the intensity of DCX

staining on the ipsilateral SVZ (stimulated) vs. contralateral SVZ (control) **(Extended Fig. 6E)**.

The intensity of DCX was remarkably higher in the activated SVZ compared to the control SVZ

$(17.7 \pm 6.6\%)$ **(Extended Fig. 6F).**

In (Cg1-subep-ChAT$^+$) circuit stimulation experiment for 3-days, the neuronal activity of

subep-ChAT$^+$ neurons was first examined in the coronal sections which were used for functional

studies **(Fig. 6C (C' & C''); upper)**. These sections have shown remarkable higher P-S6

intensity in the ipsilateral (activated) subep-ChAT$^+$ neurons compared to the subep-ChAT$^+$

neuron in the contralateral (control) side of the same brain $(37.8 \pm 8.2\%)$ **(Fig. 6D)**. The DCX$^+$

neuroblasts surrounding the subep-ChAT$^+$ neurons on the activation side were highly

randomized and not clustered as their comparative DCX$^+$ cells on the contralateral side **(Fig. 6C**

(C' & C''); lower (red arrows)). Due to the persistent stimulation of subep-ChAT$^+$ neurons,

constant generation of DCX$^+$ neuroblasts was observed, thus many neuroblasts migrate before

they cluster together **(Fig. 6E; red arrows)**. Notably, the higher rate of newly produced DCX$^+$

neuroblasts in the activated side compared to the control side led to more accumulation of

neuroblasts on the margins of the dorsal and ventral SVZ domains **(Fig. 6E; blue arrows)**. The

total number of DCX$^+$ cells in the stimulated coronal section of the ipsilateral side was

significantly higher than on the contralateral side $(25.2 \pm 4.3\%)$ **(Fig. 6F)**. Although, the

intensity of DCX staining of the stimulated and control SVZ wholemounts does not appear to be

significantly different from each other **(Fig. 6G & H)**, it was observed to be less in the area

surrounding subep-ChAT⁺ neurons of the stimulated ventral SVZ compared to the control ventral

SVZ **(Fig. 6G; yellow dotted circle)**.

***in vivo* (Cg1-subep-ChAT⁺) circuit stimulation regulate cellular proliferation in the ventral**

domain of SVZ

So far, we demonstrated the role of (Cg1-subep-ChAT⁺) circuit in regulating the activity

of ventral SVZ neurogenesis. Subsequently, we studied the impact of this circuit on the regional

LV cellular proliferation around subep-ChAT⁺ neurons in the ventral SVZ. To achieve this goal,

AAV-EF1a-DIO-hChR2 (E123T/T159C)-mCherry virus was injected into ipsilateral Cg1, and

optical fibers were implanted into the Cg1 regions of *Cr-Cre* mice **(Fig. 7A; right)**. To test the

stimulation effect of (Cg1-subep-ChAT⁺) circuit on the LV-SVZ cellular proliferation, the circuit

was activated for one day and 5-Ethynyl-2′-deoxyuridine (EdU) was intraperitoneally (IP)

injected within the last 2-3 hours before mice were sacrificed. **(Fig. 7A; left)**. Due to the spatial

cellular arrangements of proliferative cells and subep-ChAT⁺ neurons in the LV-SVZ neurogenic

niche, we used SVZ wholemount to observe the cellular proliferation in the V-SVZ (i.e.: the

proliferative cells underneath ependymal cells and 4-6um above subep-ChAT⁺ neurons) and

SVZ (i.e.: the proliferative cells lining around the subep-ChAT⁺ neurons) **(Fig. 7B)**.

Prior to observing the cellular proliferation in the area around subep-ChAT⁺ neurons, we

first examined the neuronal activity of subep-ChAT⁺ neurons in the stimulated SVZ wholemount

compared to their counterpart in the control SVZ wholemount. A mouse in each group of the

stimulated mice was used to inspect and analyze the neuronal activity of subep-ChAT⁺ neurons

(Extended Fig. 7A). In these devoted mice, the activity of subep-ChAT⁺ neurons in the

stimulated SVZ was higher than in the control ones (31.6 ± 7.8%) **(Extended Fig. 7B)**. A

combination of EdU with either GFAP (glial fibrillary acidic protein) or pEGFR (epidermal

growth factor receptor (phospho Y1068)) markers were used to view the cellular proliferation

adjacent to subep-ChAT^{+} neurons in the ventral SVZ **(Fig. 7C & E)**. While GFAP^{+} cells (i.e.: A

marker of quiescent NSCs (qNSCs) and early activated NSCs (aNSCs))[6] were observed mainly

in the V-SVZ **(Fig. 7C)**, the number of spatially localized EdU[+]/GFAP^{+} cells above the subep-

ChAT^{+} neurons in the ipsilateral V-SVZ (activated) **(Fig. 7C')** were higher than in the

contralateral V-SVZ (control) **(Fig. 7C'')** ($3 \pm 0.56\%$) **(Fig. 7D)**. This shows that (Cg1-subep-

ChAT[+]) circuit regulates the proliferative activity of LV NSCs in the ventral SVZ.

pEGFR is a known marker for labeling both the aNSCs and transit amplifying cells

(TAC)[6] which was utilized here to study the cellular proliferation in the V-SVZ and SVZ post

(Cg1-subep-ChAT[+]) circuit stimulation **(Fig. E)**. In the ipsilateral V-SVZ, the number of

spatially localized EdU[+]/pEGFR^{+} cells above the subep-ChAT^{+} neurons were notably higher

than in the control V-SVZ **(Fig. 7E' & E''; upper)** ($2.23 \pm 0.5\%$) **(Fig. 7F; left)**. This strongly

suggests that (Cg1-subep-ChAT[+]) circuit has a role in the regulation of LV NSCs activity. In

addition, there are more observed EdU[+]/pEGFR^{+} cells around subep-ChAT^{+} neurons in the

stimulated SVZ than in control SVZ **(Fig. 7E' & E''; lower)** ($1.92 \pm 0.43\%$) **(Fig. 7F; right)**.

The presence of more proliferative cells in both the V-SVZ and SVZ following the (Cg1-subep-

ChAT[+]) circuit stimulation proposes that this circuit is involved in modulating the activity of

NSCs and cellular division in the LV-SVZ.

***in vivo* (Cg1-subep-ChAT[+]) circuit inhibition modulate SVZ neurogenesis and LV cellular**

proliferation in the ventral SVZ

Our previous results predicted that the (Cg1-subep-ChAT[+]) circuit controls neurogenesis

and cellular proliferative activity in the ventral domain of LV-SVZ. To further understand the

role of this circuit, we hypothesized that *in vivo* optogenetic inhibition of (Cg1-subep-ChAT[+])

circuit is sufficient to modulate the activity of SVZ neurogenesis around subep-ChAT$^+$ neurons.

To achieve that, pAAV_hSyn1-SIO-stGtACR2-FusionRed virus was injected into ipsilateral

Cg1, and optical fibers were implanted into the Cg1 regions of *Cr-Cre* mice **(Fig. 8A)**. The

inhibition of Cg1 CR$^+$ neurons were continuously conducted for two days and were delivered by

TTL control of a 473-nm laser **(Fig. 8B)**. The mice were perfused directly after the circuit

inhibition was terminated and brain slices of the targeted sections were stained against DCX,

ChAT and P-S6 antibodies. The coronal sections included in our studies showed higher neuronal

activity of subep-ChAT$^+$ neurons on the ipsilateral (inhibited) side compared to the activity of

subep-ChAT$^+$ neurons on the control side within the same mice brain **(Fig. 8C (C' & C'');**

upper) (-31.25 ± 9.03%) **(Fig. 8D)**. Adjacent to the subep-ChAT$^+$ neurons on the ipsilateral

(inhibited) side, the number of DCX$^+$ neuroblasts were significantly lower than their counterpart

on the control side **(Fig. 8C (C' & C''); lower)**. These results indicate that the (Cg1-subep-

ChAT$^+$) circuit regulates local neurogenesis in the ventral SVZ. Furthermore, the total number of

DCX$^+$ neuroblasts on the ipsilateral side was lower than their comparative neuroblasts on the

contralateral ones (-16.4 ± 3.2%) **(Fig. 8E & F)**. All previous findings proposed a crucial role for

(Cg1-subep-ChAT$^+$) circuit in managing neurogenesis of areas surrounding subep-ChAT$^+$

neurons in the ventral SVZ.

Subsequently, the inhibition effect of (Cg1-subep-ChAT$^+$) circuit on the LV-SVZ cellular

proliferative activity around subep-ChAT$^+$ neurons was investigated. To accomplish this,

pAAV_hSyn1-SIO-stGtACR2-FusionRed virus was injected into ipsilateral Cg1, and optical

fibers were implanted into the Cg1 regions of *Cr-Cre* mice **(Fig. 8A)**. The (Cg1-subep-ChAT$^+$)

circuit was continuously blocked for one day and EdU injected within the last 2-3 hours before

mice were sacrificed **(Fig. 8G).** The neuronal activity of subep-ChAT$^+$ neurons was examined in

the ipsilateral SVZ and compared to their counterpart in control SVZ within every studied group

of mice as described in the previous section **(Extended Fig. 8A)**. In the used mice for this

purpose, the subep-ChAT$^+$ neurons in the ipsilateral SVZ wholemounts have shown remarkably

lower activity than their contralateral counterparts (-38.1 ± 7.57%) **(Extended Fig. 8B)**. Upon

staining for GFAP and EdU markers, the number of spatially localized EdU$^+$/GFAP$^+$ cells above

subep-ChAT$^+$ neurons in the ipsilateral V-SVZ **(Extended Fig. 9A')** was noticeably less than in

the control V-SVZ **(Extended Fig. 9A'')** (-1.77 ± 0.55%) **(Fig. 9B)**. This suggests that (Cg1-

subep-ChAT$^+$) circuit modulates the proliferative activity of LV-NSCs. Furthermore, both

pEGFR and EdU markers were applied together to study the effect of circuit inhibition on the

proliferative activity of LV aNSCs and TAC in the V-SVZ and SVZ, respectively. The number

of EdU$^+$/pEGFR$^+$ cells that spatially resides over the subep-ChAT$^+$ neurons in the ipsilateral V-

SVZ was observably lower than in the contralateral V-SVZ **(Fig. 8H' & H''; upper)** (-2.5 ±

0.1%) **(Fig. 8I; left)**. Additionally, the noticed number of EdU$^+$/pEGFR$^+$ cells around subep-

ChAT$^+$ neurons in the ipsilateral SVZ was less than in control SVZ **(Fig. 8H' & H''; lower)** (-

2.7 ± 0.68%) **(Fig. 8I; right)**. Together, these findings suggest that (Cg1-subep-ChAT$^+$) circuit

has regulatory roles on the proliferative activity of NSCs and cellular division in the ventral LV-

SVZ.

Discussion

Postnatal and adult LV NSCs proliferation and SVZ neurogenesis are known to be

modulated by neural activity [18]. Here, we identified a novel circuit controlling SVZ

neurogenesis and LV-SVZ cell proliferation in the LV-SVZ. Using a Rabies tracing strategy, we

determined the source of presynaptic inputs to the subep-ChAT$^+$ neurons, which are

demonstrated to direct LV NSCs proliferation[25]. Our results showed that subep-ChAT^{+} neurons

have a different pattern of neural connectivity than the striatal cholinergic neurons which are

connected with their adjacent striatal cholinergic neurons[31]. Interestingly, we found that

glutamatergic inputs from a specific VGlut1^{+} neuronal population in the anterior cingulate cortex

area 1 (Cg1) that projects directly to subep-ChAT^{+} neurons. This is the first identification of a

distal cortical input that monosynaptically drives subep-ChAT^{+} neurons and regulate their

activity in the SVZ niche. In addition, we also identified a small population of local GABAergic

calretinin (CR[+]) interneurons that directly inhibit subep-ChAT^{+} neurons. Previously, GABA was

shown to be involved in various functions in the SVZ such as enhancing neuroblast maturation

[32], and preserving the postnatal/adult LV NSCs by inhibiting their proliferation and

differentiation , but the source of GABA was largely unclear [21,33]. We demonstrated here for the

first time GABAergic CR[+] interneurons are a local source of GABA. Future studies may reveal

the neuronal connectivity of these distinct local CR[+] interneurons and other roles in the LV-SVZ

cell proliferation and neurogenesis processes.

The subep-ChAT^{+} neurons predominantly reside in the ventral domain of the SVZ. I*n*

vivo optogenetics stimulation and inhibition of the defined (Cg1-subep-ChAT[+]) circuit is

sufficient to modulate SVZ neurogenesis surrounding the subep-ChAT^{+} neurons in the ventral

area. While this strongly suggests that subep-ChAT^{+} neurons release ACh locally in the ventral

SVZ, further research can be directed to understand their axon terminals in the whole SVZ niche.

In consequence, i*n vivo* optogenetics modulation of this circuit is also sufficient to regulate the

activity of LV NSCs proliferation and cellular division regionally in the ventral areas of LV-

SVZ. Future research may explore the molecular mechanisms of subep-ChAT[+] neuronal

regulation of the LV NSCs activity and cellular divisions in V-SVZ and SVZ, respectively.

Taken together, our results uncover a new neural circuit that allows direct cortical

regulation of the LV-SVZ neurogenesis. In humans, there is evidence for active neurogenesis at

the wall of the lateral ventricles that generate migratory neuroblasts for up to two years after

birth [34], but the circuit mechanism is unknown. The analogous process in rodents may shed light

on how the neurogenesis during postnatal brain development is influenced by synaptic inputs and

neural activity. At present, how cortical inputs to the SVZ are related to environmental factors

and behavior remains unclear, but our results promise to bridge neural circuit activity with

neurogenesis and open a new avenue of research into cortical activity dependent SVZ

neurogenesis in postnatal and adult animals.

Methods

Animals

All experiments were approved by the Institutional Animal Care and Use Committee at

Duke University. Mice were group housed on a standard 12 h light/dark cycle (lights on at 7

a.m.) with a controlled average ambient temperature of 21 °C and 45% humidity. The following

mouse lines were purchased from JAX*: C57BL/6J* (000664); *VGlut1-Cre* (023527); *ChAT-eGFP*

(007902); ChR2*(RCL-hChR2(H134R)/tdT)-D (Ai27)* (012567); *ChAT- Cre* (006410); *Cr-Cre*

(010774); *VGat-Cre* (016962); *Cb-Cre* (028532); *SST-Cre* (013044); *PV-Cre* (017320);

Ai9(RCL-tdT) (007905). We generated the *R26R-FLEX-TVA-2A-RabiesG-2A-tdTomato-FLEX*

(*R26F-RTT*) mouse line for rabies monosynaptic circuit tracing **(Fig. 2A)**.

Rabies virus retrograde tracing

An intraventricular approach via a Cre-dependent viral strategy was employed to avoid

labeling striatal ChAT^{+} neurons. In the beginning we used a monosynaptic circuit tracing with

Rabies strategy[35] by injecting the first virus into LV of *ChAT-Cre* mice to express TVA, Rabies-

G protein, and tdTomato in subep-ChAT^{+} neurons. The infected subep-ChAT^{+} neurons were

later targeted with the second EnvA G-deleted rabies virus to enable efficient mono-synaptic

tracing. It was extremely difficult to infect the same subep-ChAT^{+} neurons via two separated

viral injections. To overcome this obstacle, *R26R-FLEX-TVA-2A-RabiesG-2A-tdTomato-FLEX*

(R26F-RTT) mice was successfully generated and validated **(Fig. 2A and Extended Figs. 1, 2 &**

3). These mice allow for monosynaptic tracing of Cre-targeted neurons via a single EnvA G-

deleted Rabies-eGFP (Salk, USA) or EnvA rVSV-eGFP (EnvA/RABVG-eGFP) (Salk, USA)

viral injections.

Stereotaxic injections

Stereotaxic injections were performed as mice were kept deeply anesthetized in a

stereotaxic frame (David Kopf Instruments) with isoflurane. For circuit tracing, rabies virus, 300

nL of (EnvA G-deleted Rabies-eGFP or EnvA rVSV-eGFP (Salk, USA)) virus were injected into

striatum or lateral ventricle (LV) of P30 *ChAT-Cre; R26F-RTT* mice. Viruses were infused

slowly over 10 min into striatum as the following coordinates relative to Bregma (AP: +1, ML ±

2.0, DV: 2.2 from brain surface) or LV (AP: +0.8, ML ± 0.65, DV: 2.1 from brain surface) using

a microdriver with a 10 μL Hamilton syringe. For electrophysiological testing of Cg1 inputs to

the subep-ChAT^{+} neurons (optogenetic-light stimulation), adeno-associated viral viruses were

used for Cre-dependent expression of the excitatory channelrhodopsin. pAAV-CaMKIIa-

hChR2(E123A)-mCherry (addgene #35506) (300nl) virus or pAAV-CaMKIIa-

hChR2(E123T/T159C)-EYFP (addgene #35509) (300nl) virus were injected into P30 *C57BL/6J*

or P30 *ChAT-Cre; R26F-RTT* mice, respectively. Also, pAAV-Ef1a-DIO

hChR2(E123T/T159C)-mCherry (addgene #35510) (300nl) virus was injected into P30 *VGlut1-*

Cre; ChAT-eGFP and P28 *Cr-Cre; ChAT-eGFP* mice. Viruses were infused slowly over 10 min

into the Cg1 using the following coordinates relative to Bregma (AP: +0.8, ML ± 0.3, DV: 0.5

from brain surface). For pAAV retrograde tracing, pAAVrg-hSyn-DIO-mCherry (addgene

#50459-AAVrg) and pAAVrg-hSyn-mCherry (addgene # 114472-AAVrg) viruses (300 nl) was

injected into the LV P28 *Cr-Cre*, P30 *C57BL/6J* (P30) and P30 *VGlut1-Cre*. Viruses were

infused slowly over 10 min into the Cg1 as the following coordinates relative to Bregma (AP:

+0.8, ML ± 0.65, DV: 2.1 from brain surface) .For *in vivo* optogenetic testing of Cg1 inputs to

the subep-ChAT^{+} neurons, pAAV-Ef1a-DIO hChR2(E123T/T159C)-mCherry (addgene #35510)

(300nl) and pAAV_hSyn1-SIO-stGtACR2-FusionRed (addgene #105677) (300nl) viruses were

injected into P28 *Cr-Cre* mice. Viruses were infused slowly over 10 min into the Cg1 (AP: +0.8,

ML ± 0.25, DV: 0.5 from brain surface).

All viruses were infused slowly for over 10 min using a Nanoject (Drummond Scientific)

connected to a glass pipette. The injection pipette was left in place for 10 min post-injection

before it was retracted.

SVZ wholemount preparation and whole-cell patch-clamp recording

For electrophysiology experiments both male and female mice (4- to 10-week-old) were

used. They were anesthetized with isofluorane, transcardially perfused and then ventricular wall

were dissected as SVZ wholemounts in ice-cold NMDG artificial cerebrospinal fluid (ACSF;

containing 92 mM NMDG, 2.5 mM KCl, 1.2 mM NaH_2PO_4, 30 mM $NaHCO_3$, 20 mM HEPES,

2 mM glucose, 5 mM sodium ascorbate, 2 mM thiourea, 3 mM sodium pyruvate, 10 mM

$MgSO_4$, 0.5 mM $CaCl_2$), and bubbled with 5% CO_2/95% O_2. Tissues were then bubbled in same

solution at 37 °C for 15 min, transferred to bubbled, modified-HEPES ACSF at 23–25 °C (92

mM NaCl, 2.5 mM KCl, 1.2 mM NaH$_2$PO$_4$, 30 mM NaHCO$_3$, 20 mM HEPES, 2 mM glucose, 5

mM sodium ascorbate, 2 mM thiourea, 3 mM sodium pyruvate, 2 mM MgSO$_4$, 2 mM CaCl$_2$) for

at least 45 min before start experimentation. (For NMDG artificial cerebrospinal fluid and

modified-HEPES ACSF solutions, pH was adjusted to 7.5 and osmolarity to 290 mOsm).

Recordings were performed in submerged chamber, superfused with continuously bubbled

ACSF (125 mM NaCl, 2.5 mM KCl, 1.25 mM NaH$_2$PO$_4$, 26 mM NaHCO$_3$, 20 mM glucose, 2

mM CaCl$_2$, 1.3 mM MgCl$_2$) at 2.5–5 ml/min at 23–25 °C.

Patch electrodes with a resistance of 3–6 MΩ were pulled from borosilicate glass

capillaries using a horizontal puller (P-97, Sutter-instruments). All subep-ChAT$^+$ neurons used in

the electrophysiology recordings were identified by their distinctive morphology from striatal

cholinergic neurons, location (within 15-20 µM from LV surface), their non-spontaneous firing

activity, and their depolarized resting membrane potentials (~-45 mV). For measuring EPSCs

when holding at – 60 mV, the following internal solution was used: 130 mM potassium

gluconate, 2 mM NaCl, 4 mM MgCl$_2$, 20 mM HEPES, 4 mM Na$_2$ATP, 0.4 mM NaGTP and 0.5

mM EGTA, pH adjusted to 7.2 with KOH. A 40 µM picrotoxin ((124-87-8), Millipore Sigma)

was added in the bath solution in voltage-clamp mode. For measuring inward IPSCs when

holding at – 60 mV, we used internal solution containing K-Cl (high Chloride) KCl (135 Mm),

HEPES (10 mM), Na2ATP (2mM), NaGTP (0.2 mM), MgCl2 (2 mM), and EGTA (0.1 mM), pH

adjusted to 7.2 with KOH. To block glutamatergic transmission, 50 µM DL-AP5 (NMDA

antagonist) ((3693/50), R&D Systems) and 50 µM CNQX (AMPA antagonist) ((0190), Tocris)

were added in the bath solution in voltage-clamp mode. To block action potentials in the patched

cell, the sodium channel blocker QX-314 (10 mM, 552233, Millipore Sigma) was also added.

464 Signals were amplified with Multiclamp 700B (filtered at 10 kHz), digitized with Digidata

465 1440A (20 kHz), and recorded using pClamp 10 software (Axon). Light-activation of

466 channelrhodopsin was achieved using a 473-nm laser (X-Cite exacte) through a 40x objective

467 (Nikon). Action potentials, evoked EPSCs and evoked IPSCs were analyzed using AxoGraph

468 (AxoGraph). One or two subep-ChAT$^+$ neurons were recorded per wholemount SVZ. Up to

469 three neurons from each animal were recorded. Measurements were taken as an average of at

470 least five responses to obtain a data point.

471  For focal (targeted) photostimulation of local CR$^+$-IPSCs from subep-ChAT$^+$ neurons, we

472 used Polygon 400 Digital Micromirror Device (Mightex; multiwavelength pattered illuminator)

473 to control the temporal dynamics (the size, shape, intensity, and position) of light inputs. The

474 illumination consisted of a 100 ms light pulse (470-nm) at 50% intensity in the selected area (50

475 μm). The pulse was triggered using a TTL pulse from the Digidata to synchronize the

476 stimulation with electrophysiology. Photostimulation was performed for one fluorescent subep-

477 CR$^+$ neuron presents in the field of view resulted in postsynaptic effects in the recorded neuron.

478 **Immunofluorescence staining and imaging**

479  Preparation of brain tissue for IHC staining was described previously [25]. We used primary

480 antibodies to GFP (#GFP-1020, 1:400, AVES lab), Choline Acetyltransferase Antibody

481 (#AB144P, 1:250, Millipore sigma), tdTomato [16D7] (#EST203, 1:200, Kerafast), RFP (#600-

482 401-379, 1:250, Rockland), Calretinin (#ab702, 1:200, Abcam), calretinin (#6B3, 1:250, Swant),

483 Calretinin (#MCA-3G9, 1:250, EnCor Biotechnology), RFP (#ab62341, 1:200, Abcam),

484 Doublecortin (#AB2253, 1:250, Millipore), Doublecortin (#4604S, 1:200, Cell Signaling

485 Technology), Doublecortin (#MA5-17066, 1:200, ThermoFisher Scientific), EGFR (phospho

486 Y1068) (#ab5644, 1:250, Abcam), EGFR (phospho Y1068) [Y38] (#ab32430, 1:250, Abcam),

487 GFAP (#GFAP, 1:500, Aves Labs). In brief, for circuit tracing and *in vivo* optogenetic

488 experiments, after experiments mice were deeply anesthetized with isoflurane, perfused

489 transcardially with phosphate buffered saline (PBS), followed by 4% PFA in PBS. The perfused

490 brains were removed and postfixed overnight at 4°C in 4% PFA. The fixed brains were either cut

491 into 50 μm coronal sections by Precisionary Instruments VF-500-0Z vibrating microtome or the

492 SVZ wholemounts were dissected. The coronal slices or SVZ wholemounts were incubated in a

493 blocking solution containing 5% donkey serum and TBST for 100 min at room temperature. The

494 sections were then incubated at room temperature overnight in PBS containing 1% donkey serum

495 and antibodies. They were then washed with PBS and incubated with secondary antibodies,

496 Alexa-594 (1:1000, LifeTech) or Alexa-488 (1:1000, LifeTech) or Alexa-647 (1:1000, LifeTech)

497 for 2 h at room temperature, before washing with PBS. The stained sections were counterstained

498 with a 4′,6-diamidino-2-phenylindole solution (DAPI; (D9542) Sigma-Aldrich). After washing

499 four times with TBST, the sections were coverslipped with Fluoromount (Sigma) aqueous

500 mounting medium. Images were taken using Leica SP8 upright confocal microscope (Zeiss) with

501 10 ×, 20 × and 40 × objectives under the control of Zen software (Zeiss). All antibodies used

502 were validated as in previous publications [25,36] or by publications available on vendor websites

503 specific to each antibody.

504 ***In vivo* optogenetic stimulation and inhibition**

505 Cannula targeting the Cg1 region was implanted (AP: +0.9, ML ± 0.25, DV: 0.4 from brain

506 surface) after injecting pAAV-Ef1a-DIO hChR2(E123T/T159C)-mCherry and pAAV_hSyn1-

507 SIO-stGtACR2-FusionRed viruses as described in **Fig. 6A, 7A & 8A**, using implantable mono

508 fiber-optic fiber (200 μm, 0.22 NA, Doric). Protruding ferrule end of cannula was then connected

via fiber cord and a rotary coupling joint (Doric) was used to permit free movement. Three to

four weeks after viral infection, light-stimulation was delivered by TTL control (Master 8,

AMPI) of a 473-nm laser (IkeCool). For *In vivo* optogenetic stimulation, the ipsilateral Cg1

region was stimulated for 10 ms pulses (7-9 mW laser power) at 10 Hz, lasting 10 s, given once

every 1 min. For *In vivo* optogenetic inhibition, the Cg1 regions in ipsilateral and contralateral

sides were continuously stimulated using 3 mW laser power only to study and avoid overheating

in the local cortical areas.

For ChAT, DCX, EGFR (phospho Y1068) [Y38], GFAP, CR and RFP analyses, 50-μm

brain coronal sections were cut and collected by Precisionary Instruments VF-500-0Z vibrating

microtome or SVZ wholemounts were dissected. The sections or areas surrounding the

activated/inhibited subep-ChAT$^+$ neurons were selected for analyses.

***in vivo* EdU labeling**

The 5-ethynyl-2'-deoxyuridine (EdU) staining was conducted using EdU Cell

Proliferation Kit for Imaging (EdU *in vivo* Kits) (baseclick GmbH, Germany) according to the

manufacturer's protocol. EdU was prepared at 50mg/20mL in sterile PBS and used for pulse

labeling of adult mice by performing an intraperitoneal (IP) injection of 500 ul dissolved EdU

(50mg/kg). The mice were harvested as described in **Fig. 7 and 8**. The intended coronal slices or

SVZ wholemounts for EdU labeling were first stained for other antibodies, and then

counterstained with DAPI as described in the immunofluorescence staining section.

Subsequently, the sections were washed three times with 3% BSA in PBS. Then, incubated for

30 minutes in a reaction cocktail containing Deionized water, Reaction buffer, Catalyst solution,

Dye Azide and Buffer additive while protected from light. After the reaction cocktail was

removed, sections were washed three times with 3% BSA in PBS. They were then mounted in

vectashield mounting media (vector laboratories Inc, Burlingame, CA) and imaged by using

Leica SP8 upright confocal microscope (Zeiss) as above. All steps were carried out at room

temperature.

Quantification and Statistical Analysis

536 All data are expressed as mean ± SEM and all statistical analyses were performed using a

537 GraphPad Prism (version 8) program. Paired and unpaired t-tests were used for analysis of *in*

538 *vivo* optogenetics stimulation and inhibition studies. $p < 0.05$ was considered statistically

539 significant.

Acknowledgements

542 We thank Scott Soderling and Shawn Je for helpful comments on the manuscript, Transgenic and

543 Knockout Mouse Facility at Duke University for assistance with generation of R26R-FLEX-

544 TVA-2A-RabiesG-2A-tdTomato-FLEX mice, E. Adlaf, J. Erb, B. Asrican, D. Fromme, P. Paez-

545 Gonzalez, K. Abdi, G. Neves, S. Ramamoorthy, J. David and K. Woldemichael for project

546 assistance. This work was supported by NIH grant R01MH105416.

Contributions

549 M.M.N., C.T.K. and H.H.Y. conceived the project and participated in research design. M.M.N.

550 performed all experiments and analyzed data. R.R.K. helped with preparation of SVZ

551 wholemounts for electrophysiology experiments. M.M.N. and H.H.Y wrote the paper.

Reference

Lim, D. A. & Alvarez-Buylla, A. The Adult Ventricular-Subventricular Zone (V-SVZ) and Olfactory Bulb (OB) Neurogenesis. *Cold Spring Harb Perspect Biol* **8**, doi:10.1101/cshperspect.a018820 (2016).

Ihrie, R. A. & Alvarez-Buylla, A. Lake-front property: a unique germinal niche by the lateral ventricles of the adult brain. *Neuron* **70**, 674-686, doi:10.1016/j.neuron.2011.05.004 (2011).

Lazarini, F. & Lledo, P. M. Is adult neurogenesis essential for olfaction? *Trends Neurosci* **34**, 20-30, doi:10.1016/j.tins.2010.09.006 (2011).

Doetsch, F., Caille, I., Lim, D. A., Garcia-Verdugo, J. M. & Alvarez-Buylla, A. Subventricular zone astrocytes are neural stem cells in the adult mammalian brain. *Cell* **97**, 703-716, doi:10.1016/s0092-8674(00)80783-7 (1999).

Obernier, K. *et al.* Adult Neurogenesis Is Sustained by Symmetric Self-Renewal and Differentiation. *Cell Stem Cell* **22**, 221-234 e228, doi:10.1016/j.stem.2018.01.003 (2018).

Codega, P. *et al.* Prospective identification and purification of quiescent adult neural stem cells from their in vivo niche. *Neuron* **82**, 545-559, doi:10.1016/j.neuron.2014.02.039 (2014).

Lois, C., Garcia-Verdugo, J. M. & Alvarez-Buylla, A. Chain migration of neuronal precursors. *Science* **271**, 978-981, doi:10.1126/science.271.5251.978 (1996).

Ponti, G., Obernier, K. & Alvarez-Buylla, A. Lineage progression from stem cells to new neurons in the adult brain ventricular-subventricular zone. *Cell Cycle* **12**, 1649-1650, doi:10.4161/cc.24984 (2013).

Lois, C. & Alvarez-Buylla, A. Long-distance neuronal migration in the adult mammalian brain. *Science* **264**, 1145-1148, doi:10.1126/science.8178174 (1994).

Luskin, M. B. Restricted proliferation and migration of postnatally generated neurons derived from the forebrain subventricular zone. *Neuron* **11**, 173-189, doi:10.1016/0896-6273(93)90281-u (1993).

Petreanu, L. & Alvarez-Buylla, A. Maturation and death of adult-born olfactory bulb granule neurons: role of olfaction. *J Neurosci* **22**, 6106-6113, doi:20026588 (2002).

Imayoshi, I. *et al.* Roles of continuous neurogenesis in the structural and functional integrity of the adult forebrain. *Nat Neurosci* **11**, 1153-1161, doi:10.1038/nn.2185 (2008).

Livneh, Y., Adam, Y. & Mizrahi, A. Odor processing by adult-born neurons. *Neuron* **81**, 1097-1110, doi:10.1016/j.neuron.2014.01.007 (2014).

Sakamoto, M. *et al.* Continuous postnatal neurogenesis contributes to formation of the olfactory bulb neural circuits and flexible olfactory associative learning. *J Neurosci* **34**, 5788-5799, doi:10.1523/JNEUROSCI.0674-14.2014 (2014).

Mak, G. K. & Weiss, S. Paternal recognition of adult offspring mediated by newly generated CNS neurons. *Nat Neurosci* **13**, 753-758, doi:10.1038/nn.2550 (2010).

Sakamoto, M., Kageyama, R. & Imayoshi, I. The functional significance of newly born neurons integrated into olfactory bulb circuits. *Front Neurosci* **8**, 121, doi:10.3389/fnins.2014.00121 (2014).

Jones, K. S. & Connor, B. Intrinsic regulation of adult subventricular zone neural progenitor cells and the effect of brain injury. *Am J Stem Cells* **1**, 48-58 (2012).

Obernier, K. & Alvarez-Buylla, A. Neural stem cells: origin, heterogeneity and regulation in the adult mammalian brain. *Development* **146**, doi:10.1242/dev.156059 (2019).

Bovetti, S., Gribaudo, S., Puche, A. C., De Marchis, S. & Fasolo, A. From progenitors to integrated neurons: role of neurotransmitters in adult olfactory neurogenesis. *J Chem Neuroanat* **42**, 304-316, doi:10.1016/j.jchemneu.2011.05.006 (2011).

598	20	Young, S. Z., Taylor, M. M. & Bordey, A. Neurotransmitters couple brain activity to
599		subventricular zone neurogenesis. *Eur J Neurosci* **33**, 1123-1132, doi:10.1111/j.1460-
600		9568.2011.07611.x (2011).
601	21	Liu, X., Wang, Q., Haydar, T. F. & Bordey, A. Nonsynaptic GABA signaling in postnatal
602		subventricular zone controls proliferation of GFAP-expressing progenitors. *Nat Neurosci* **8**, 1179-
603		1187, doi:10.1038/nn1522 (2005).
604	22	Hoglinger, G. U. *et al.* Dopamine depletion impairs precursor cell proliferation in Parkinson
605		disease. *Nat Neurosci* **7**, 726-735, doi:10.1038/nn1265 (2004).
606	23	Brezun, J. M. & Daszuta, A. Depletion in serotonin decreases neurogenesis in the dentate gyrus
607		and the subventricular zone of adult rats. *Neuroscience* **89**, 999-1002, doi:10.1016/s0306-
608		4522(98)00693-9 (1999).
609	24	Banasr, M., Hery, M., Printemps, R. & Daszuta, A. Serotonin-induced increases in adult cell
610		proliferation and neurogenesis are mediated through different and common 5-HT receptor
611		subtypes in the dentate gyrus and the subventricular zone. *Neuropsychopharmacology* **29**, 450-
612		460, doi:10.1038/sj.npp.1300320 (2004).
613	25	Paez-Gonzalez, P., Asrican, B., Rodriguez, E. & Kuo, C. T. Identification of distinct ChAT(+)
614		neurons and activity-dependent control of postnatal SVZ neurogenesis. *Nat Neurosci* **17**, 934-
615		942, doi:10.1038/nn.3734 (2014).
616	26	Oldenburg, I. A. & Ding, J. B. Cholinergic modulation of synaptic integration and dendritic
617		excitability in the striatum. *Current opinion in neurobiology* **21**, 425-432 (2011).
618	27	Fremeau, R. T., Jr. *et al.* The expression of vesicular glutamate transporters defines two classes
619		of excitatory synapse. *Neuron* **31**, 247-260, doi:10.1016/s0896-6273(01)00344-0 (2001).
620	28	Madisen, L. *et al.* A toolbox of Cre-dependent optogenetic transgenic mice for light-induced
621		activation and silencing. *Nature neuroscience* **15**, 793-802, doi:nn.3078 [pii]
622		10.1038/nn.3078 (2012).
623	29	Lim, S. A., Kang, U. J. & McGehee, D. S. Striatal cholinergic interneuron regulation and circuit
624		effects. *Front Synaptic Neurosci* **6**, 22, doi:10.3389/fnsyn.2014.00022 (2014).
625	30	Beier, K. T., Saunders, A. B., Oldenburg, I. A., Sabatini, B. L. & Cepko, C. L. Vesicular stomatitis
626		virus with the rabies virus glycoprotein directs retrograde transsynaptic transport among
627		neurons in vivo. *Front Neural Circuits* **7**, 11, doi:10.3389/fncir.2013.00011 (2013).
628	31	Schafer, M. K., Eiden, L. E. & Weihe, E. Cholinergic neurons and terminal fields revealed by
629		immunohistochemistry for the vesicular acetylcholine transporter. I. Central nervous system.
630		*Neuroscience* **84**, 331-359, doi:10.1016/s0306-4522(97)00516-2 (1998).
631	32	Young, S. Z. *et al.* NKCC1 knockdown decreases neuron production through GABA(A)-regulated
632		neural progenitor proliferation and delays dendrite development. *J Neurosci* **32**, 13630-13638,
633		doi:10.1523/JNEUROSCI.2864-12.2012 (2012).
634	33	Alfonso, J., Le Magueresse, C., Zuccotti, A., Khodosevich, K. & Monyer, H. Diazepam binding
635		inhibitor promotes progenitor proliferation in the postnatal SVZ by reducing GABA signaling. *Cell
636		Stem Cell* **10**, 76-87, doi:10.1016/j.stem.2011.11.011 (2012).
637	34	Sanai, N. *et al.* Corridors of migrating neurons in the human brain and their decline during
638		infancy. *Nature* **478**, 382-386, doi:10.1038/nature10487 (2011).
639	35	Callaway, E. M. & Luo, L. Monosynaptic Circuit Tracing with Glycoprotein-Deleted Rabies
640		Viruses. *J Neurosci* **35**, 8979-8985, doi:10.1523/JNEUROSCI.0409-15.2015 (2015).
641	36	Benner, E. J. *et al.* Protective astrogenesis from the SVZ niche after injury is controlled by Notch
642		modulator Thbs4. *Nature* **497**, 369-373, doi:10.1038/nature12069 (2013).

Figure 1 Glutamatergic and GABAergic inputs to subep-ChAT$^+$ neurons.

(A) Illustration of electrophysiological recording of excitatory inputs (VGlut1$^+$) to the subep-ChAT$^+$ neuron from the wholemount of P30-50 *VGlut1-Cre; ChAT-eGFP; Ai27* mice.

(B) Representative trace from whole cell current clamp recording of evoked action potential (APs) from subep-ChAT$^+$ neurons upon blue (473nm) light stimulation for 5s (Left). Blue bar = duration of light stimulation.

Average frequency (Right). P < 0.0001, t_9 = 17.7, n = 10. Data collected from four *VGlut1-Cre; ChAT-eGFP; Ai27* mice. Each dot represents data from one subep-ChAT$^+$ neuron.

(C) Representative trace of EPSCs that were obtained in whole-cell voltage-clamp recordings from subep-ChAT$^+$ neurons after photostimulation for 500ms (black), and the application of AMPA and NMDA receptors antagonists; CNQX and AP-5, respectively (red) (left). Blue bar = duration of light stimulation.

Average evoked EPSC amplitude and latency (Right). P < 0.0001, t_9 = 29.1, n = 10 and P < 0.0001, t_9 = 44.2, n = 10, respectively. Data collected from four *VGlut-Cre; ChAT-eGFP; Ai27* mice. Each dot represents the amplitude (upper) and latency (lower) of a single Excitatory Postsynaptic Current (EPSC) recorded from subep-ChAT$^+$ neuron.

(D) Illustration of electrophysiological recording of inhibitory inputs (VGat$^+$) to the subep-ChAT$^+$ neuron from P30-50 *VGat-ChR2-eYFP; ChAT-Cre; Ai9* mice.

(E) Representative trace of electrophysiological recordings of IPSCs that were obtained in whole-
cell recordings from subep-ChAT$^+$ neurons upon blue (473nm) light stimulation for 500ms (black),
and the application of GABA$_A$R antagonist; picrotoxin (blue) (left). Blue bar = duration of light
stimulation. Each dot represents the amplitude (upper) and latency (lower) of a single Inhibitory
Postsynaptic Current (IPSC) recorded from subep-ChAT$^+$ neuron.

Average evoked IPSC amplitude and latency (Right). P value < 0.0001, $t_9 = 30.4$, n = 10 and P
value < 0.0001, $t_9 = 16.1$, n = 10, respectively. Data collected from four *VGat-ChR2-eYFP; ChAT-*
cre; Ai9 mice.

(F) Schematic representation of presynaptic (excitatory and inhibitory) inputs to the subep-ChAT$^+$
neuron.
All error bars indicate SEM

Figure 2 Rabies virus infection of subep-ChAT$^+$ neurons.

(A) Schematic representation of *R26R-FLEX-TVA-2A-RabiesG-2A-tdTomato-FLEX* (R26F-RTT) mice before and after Cre recombinase.

(B) Schematic of EnvA G-deleted Rabies-eGFP virus injection into the brain of *-Cre; R26F-RTT* mice.

(C) Experimental representation of EnvA G-deleted Rabies-eGFP virus injection into lateral ventricle (LV) of P30 *Chat-Cre; R26F-RTT* mice. Mice were sacrificed on the day 7 after injection.

(D-E) Immunofluorescence staining for GFP (green) and ChAT (red) in SVZ and striatum of ipsilateral (injected) side of *Chat-Cre; R26F-RTT* mice (7 days post-injection).

(F) Experimental representation of EnvA G-deleted Rabies-eGFP virus injection into LV of P30
Chat-Cre; R26F-RTT mice. Mice were sacrificed on the day 14 after injection.

(G-H) Immunofluorescence staining for GFP (green) and ChAT (red) in SVZ and striatum of the
ipsilateral (injected) side of *Chat-Cre; R26F-RTT* mice (14 days post-injection).

(I) Schematic representation of intra cholinergic connections of striatal cholinergic and subep-
ChAT^{+} neurons.

Figure 3 Local calretinin-positive (CR⁺) GABAergic interneurons provide inhibitory inputs to subep-ChAT⁺ neurons.

(A) Experimental design of EnvA G-deleted Rabies-eGFP virus injection into LV of P30 *Chat-Cre; R26F-RTT* mice.

(B-C) Immunofluorescence staining for GFP (green), Calretinin (purple) and DAPI (blue) in ipsilateral SVZ wholemount (injected) of P37 *Chat-Cre; R26F-RTT* mice.

(D) illustration of electrophysiological recording of calretinin-positive (CR$^+$) inhibitory inputs to the subep-ChAT$^+$ neurons from SVZ wholemount of P30-50 *Cr-Cre; ChAT-eGFP; Ai27* mice.

(E) Representative trace of evoked IPSCs from subep-ChAT$^+$ neurons upon photostimulation for 500ms (black) and after application of GABA$_A$R antagonist picrotoxin (green) (Left). Blue bar = duration of light stimulation.

Average current amplitude and latency of the IPSCs upon stimulation (Right). $P < 0.0001$, $t_9 = 19.1$, n = 10 and $P < 0.0001$, $t_9 = 46.9$, n = 10, respectively. Data collected from five *Cr-Cre; ChAT-eGFP; Ai27* mice. Each dot represents the amplitude (left) and latency (right) of a single Inhibitory Postsynaptic Current (EPSC) recorded from subep-ChAT$^+$ neuron.

(F) Representation of subep-ChAT$^+$ neuron recording from SVZ wholemount of P30-50 *Cr-Cre; ChAT-eGFP; Ai27* mice. Left; subep-ChAT$^+$ neuron (473nm light). Middle; Subep-CR$^+$ neurons (590nm light). Right; bright field image showing both subep-ChAT$^+$ and -CR$^+$ neurons.

(G) Representative trace of evoked IPSCs in subep-ChAT$^+$ neurons following photostimulation for 100ms (Left). Blue bar = duration of light stimulation.

Average current amplitude and latency of the IPSCs upon stimulation (Right). $P < 0.0001$, $t_7 = 12.25$, n = 8 and $P < 0.0001$, $t_9 = 32.4$, n = 8, respectively. Data collected from three *Cr-Cre; ChAT-eGFP; Ai27* mice. Each dot represents the amplitude (left) and latency (right) of an Inhibitory Postsynaptic Current (EPSC) recorded from subep-ChAT$^+$ neuron.

(H-I) Immunofluorescence staining for CR (purple) and ChAT (green) in the SVZ wholemount (H) and coronal section (I) of P28 *C57BL/6J* mice.

All error bars indicate SEM.

Figure 4 Anterior cingulate cortex projects directly to subep-ChAT⁺ neurons.

(A) Schematic illustration of EnvA rVSV-eGFP (EnvA/RABVG-eGFP) virus injection into LV of
P30 *Chat-Cre; R26F-RTT* mice.

(B) A sample of immunofluorescence staining for GFP (green) in anterior cingulate cortex areas 1 and 2 (Cg1 and Cg2) (left) and Cg1 region (right) of ipsilateral (injected) side of P33 *Chat-Cre; R26F-RTT* mice.

(C) Schematic representation of AAV-CaMKII-hChR2(E123A)-mCherry virus injection into Cg1 region of P30 *C57BL/6J* mice.

(D) A sample of immunofluorescence staining for ChAT (green) and infected projections (red) of the ipsilateral Cg1 neurons (injected side) in striatum and SVZ (left), and SVZ (right) of P50 *C57BL/6J* mice in panel C.

(E) Experimental design of pAAV-Ef1a-DIO hChR2(E123T/T159C)-EYFP virus injection into Cg1 region of P30 *Chat-Cre; Ai9* mice. Mice were used for recording 3-4 weeks post-injection.

(F) Representative trace of whole cell current clamp recoding of evoked action potentials (APs) from subep-ChAT$^+$ neurons upon photo-stimulation for 5s (Left). Blue bar = duration of light stimulation.
Average spike frequency (Right). P < 0.0001, t_7 = 7.6, n = 8. Data collected from five *Chat-Cre; Ai9* mice. Each dot represents data from one subep-ChAT$^+$ neuron.

(G) Representative trace of EPSCs were obtained in whole-cell recordings from subep-ChAT$^+$ neurons upon 473nm light stimulation for 500ms (black), and after blocking with AMPA and NMDA receptors antagonists (CNQX and AP-5, respectively) (red) (Left). Blue bar = duration of light stimulation.
Average EPSC amplitude and latency (Right). P value < 0.0001, t_9 = 36.16, n = 10 and P value < 0.0001, t_9 = 40.5, n = 10, respectively. Data collected from five *Chat-Cre; Ai9* mice. Each dot represents the amplitude (left) and latency (right) of a single Excitatory Postsynaptic Current (EPSC) recorded from subep-ChAT$^+$ neuron.

(H) Experimental design of pAAV-Ef1a-DIO hChR2(E123T/T159C)-mCherry virus injection into Cg1 region of P30 *VGlut1-Cre; ChAT-eGFP* mice. Mice were used for recording at 3-4 weeks post-injection.

(I) Representative trace of EPSCs were obtained in whole-cell voltage-clamp recordings from subep-ChAT$^+$ neurons upon 473nm light stimulation for 500ms (black), and after blocking with AMPA and NMDA receptors antagonists (CNQX and AP-5, respectively) (red) (Left). Blue bar = duration of light stimulation.
Average evoked EPSC amplitude and latency (Right). P value < 0.0001, t_9 = 23.3, n = 10 and P value < 0.0001, t_9 = 61.2, n = 10, respectively. Data collected from four *VGlut1-Cre; ChAT-eGFP* mice. Each dot represents the amplitude (left) and latency (right) of a single Excitatory Post Synaptic Current (EPSC) recorded from subep-ChAT$^+$ neuron.

All error bars indicate SEM.

Figure 5 Presynaptic cingulate neurons that excite subep-ChAT$^+$ neurons are calretinin-positive (CR$^+$).

(A) Schematic illustration of the experimental design of pAAVrg-hSyn-DIO-mCherry virus injection into LV of P28 *Cr-Cre* mice.

(B) Immunofluorescence staining for RFP (red) in ipsilateral (injected side) anterior cingulate cortex areas 1 and 2 (Cg1 and Cg2) (left) and ipsilateral (injected side) Cg1 region (right) of P50 *Cr-Cre* mice.

(C) Illustration of electrophysiological recording of calretinin-positive (CR$^+$) excitatory inputs to subep-ChAT$^+$ neurons from SVZ wholemount of P30-50 *Cr-Cre; ChAT-eGFP; Ai27* mice.

(D) Representative trace of EPSCs were obtained in whole-cell recordings from subep-ChAT$^+$ neurons upon 473nm light stimulation for 500ms (black), and after blocking with AMPA and NMDA receptors antagonists (CNQX and AP-5, respectively) (red) (Left). Blue bar = duration of light stimulation.
Average EPSC amplitude and latency (Right). P < 0.0001, t_9 = 19.6, n = 10 and P < 0.0001, t_9 = 27, n = 10, respectively. Data collected from five *Cr-Cre; ChAT-eGFP; Ai27* mice. Each dot represents the amplitude (left) and latency (right) of a single Excitatory Postsynaptic Current (EPSC) recorded from subep-ChAT$^+$ neuron.

(E) Experimental design of AAV-EF1a-DIO-hChR2(E123T/T159C)-mCherry virus injection into Cg1 of P28 *Cr-Cre; ChAT-eGFP* mice. Mice were used for recording 3-4 weeks post-injection.

(F) Representative trace of EPSCs evoked specifically by Cg1 CR$^+$ input. EPSCs were recorded from subep-ChAT$^+$ neurons of injected mice in panel E upon 473nm light stimulation for 500ms (black). EPSCs were blocked by CNQX and AP-5 (red) (Left).
An average current amplitude and latency (Right). P < 0.0001, t_7 = 15.3, n = 8 and P < 0.0001, t_7 = 24.9, n = 8, respectively. Data collected from four *Cr-Cre; ChAT-eGFP* mice. Each dot represents the amplitude (left) and latency (right) of a single Excitatory Postsynaptic Current (EPSC) recorded from subep-ChAT$^+$ neuron.

(G) Schematic summary of (Cg1-Subep-ChAT$^+$) circuit regulation of LV NSCs.

All error bars indicate SEM.

809
810
811 **Figure 6** *In vivo* **optogenetic (Cg1-subep-ChAT⁺) circuit stimulation modulates neurogenesis**
812 **in the SVZ niche.**

(A) Schematic representation of *in vivo* optogenetic stimulation experiment in panel B. Immunofluorescence staining for RFP (red) in the ipsilateral (injected) Cg1 of *Cr-Cre* mice.

(B) Experimental design of *in vivo* optogenetic stimulation for three days post AAV-EF1a-DIO-hChR2(E123T/T159C)-mCherry virus injection into ipsilateral Cg1 and optical fibers implantation into Cg1 regions of P28 *Cr-Cre* mice

(C) DCX (grey) and ChAT (green) immunofluorescence staining of (C') ipsilateral (upper images; activation) vs. (C'') contralateral (lower images; control) sides of SVZ from stimulated coronal sections of mice in panel B. Identical settings from same brain section were used for imaging. Red arrows represent DCX$^+$ neuroblasts around subep-ChAT$^+$ neurons.

(D) P-S6 intensity analysis of subep-ChAT$^+$ neurons in ipsilateral side (activation) vs. contralateral side (control). P<0.0004, n=8, Paired t-test. Data collected from three stimulated *Cr-Cre* mice. Each dot represents a subep-ChAT$^+$ neuron.

(E) DCX (grey) immunofluorescence staining of ipsilateral SVZ (left image; activation) vs. contralateral SVZ (right image; control) from stimulated coronal section of mice in panel C. Identical settings from same brain section were used for imaging. Blue arrows display DCX$^+$ neuroblasts in dorsal (upper) and ventral (lower) domains. Red arrow displays DCX$^+$ neuroblasts adjacent to subep-ChAT$^+$ neurons in ventral domain of the stimulated SVZ (left) and control SVZ (right).

(F) Analysis of DCX$^+$ neuroblasts in stimulated coronal sections of ipsilateral side (activation) vs. contralateral side (control). P<0.0001, n=9, Paired t-test. Data collected from four stimulated *Cr-Cre* mice. Each dot represents a total DCX$^+$ cells per coronal section.

(G) DCX (green) immunofluorescence staining of ipsilateral (upper images; activation) vs. contralateral (lower images; control) of SVZ wholemount from stimulated mice in panel B. Identical settings from same brain wholemounts were used for imaging. Yellow dotted circle represents an area where subep-ChAT$^+$ neurons mainly reside in the ventral region of SVZ.

(H) Analysis of DCX intensity in ipsilateral SVZ wholemount (activation) vs. contralateral SVZ wholemount (control). P = 0.766 (ns), n=4, Paired t-test. Data collected from four stimulated *Cr-Cre* mice. Each dot represents a total DCX protein per SVZ wholemount.

All error bars indicate SEM.

842
843
Figure 7 *In vivo* optogenetic (Cg1-subep-ChAT⁺) circuit stimulation modulates cellular
proliferation in the SVZ niche.

(A) Experimental design of *in vivo* optogenetic stimulation for one day post AAV-EF1a-DIO-
hChR2(E123T/T159C)-mCherry virus injection into ipsilateral Cg1 and optical fibers
implantation into Cg1 regions of P28 *Cr-Cre* mice

(B) Schematic representation of the cellular composition and organization of the LV-SVZ.

(C) EdU (purple), ChAT (green) and GFAP (grey) immunofluorescence staining of (C') ipsilateral
SVZ wholemount (activation) (upper images; V-SVZ and lower images; SVZ) vs. (C'')
contralateral SVZ wholemount (control) (upper images; V-SVZ and lower images; SVZ) from
stimulated mice in panel A. Identical settings from same brain section were used for imaging. Blue
arrows display EdU$^+$/GFAP$^+$ cells surrounding subep-ChAT$^+$ neurons in V-SVZ. Red arrows
display EdU$^+$/GFAP$^-$ cells surrounding subep-ChAT$^+$ neurons in V-SVZ and SVZ.

(D) Analysis of EdU$^+$/GFAP$^+$ cells in ipsilateral (activation) V-SVZ vs. contralateral (control) V-
SVZ of SVZ wholemounts. P<0.0001, n=13, Unpaired t-test. Data collected from four stimulated
Cr-Cre mice. Each dot represents total EdU$^+$/GFAP$^+$ cells surrounding a subep-ChAT$^+$ neuron.

(E) EdU (purple), ChAT (green) and pEGFR (grey) immunofluorescence staining of (E')
ipsilateral SVZ wholemount (activation) (upper images; V-SVZ and lower images; SVZ) vs. (E'')
contralateral SVZ wholemount (control) (upper images; V-SVZ and lower images; SVZ) from
stimulated mice in panel A. Identical settings from same brain section were used for imaging. Blue
arrows display EdU$^+$/pEGFR$^+$ cells surrounding subep-ChAT$^+$ neurons in V-SVZ and SVZ. Red
arrows display EdU$^+$/pEGFR$^-$ cells surrounding subep-ChAT$^+$ neurons in V-SVZ.

(F) Left: Analysis of EdU$^+$/pEGFR$^+$ cells in ipsilateral (activation) V-SVZ vs. contralateral
(control) V-SVZ of SVZ wholemounts. P = 0.0001, n=13, Unpaired t-test. Data collected from
four stimulated *Cr-Cre* mice. Each dot represents total EdU$^+$/pEGFR$^+$ cells surrounding a subep-
ChAT$^+$ neuron.

Right: Analysis of EdU$^+$/pEGFR$^+$ cells in ipsilateral (activation) SVZ vs. contralateral (control)
SVZ of SVZ wholemounts. P = 0.0002, n=13, Unpaired t-test. Data collected from four stimulated
Cr-Cre mice. Each dot represents total EdU$^+$/pEGFR$^+$ cells surrounding a subep-ChAT$^+$ neuron.

All error bars indicate SEM.

874
875
876 **Figure 8** *In vivo* **optogenetic (Cg1-subep-ChAT$^+$) circuit inhibition modulates neurogenesis**
877 **and cellular proliferation in the SVZ niche.**

878 (A) Schematic representation of *in vivo* optogenetic inhibition experiment in panel B.
879 Immunofluorescence staining for RFP (red) in ipsilateral (injected) Cg1 of *Cr-Cre* mice.

(B) Experimental design of *in vivo* optogenetic inhibition for two days post AAV_hSyn1-SIO-stGtACR2-FusionRed virus injection into ipsilateral Cg1 and optical fibers implantation into Cg1 regions of P28 *Cr-Cre* mice.

(C) DCX (grey) and ChAT (green) immunofluorescence staining of (C') ipsilateral (upper images; inhibition) vs. (C'') contralateral (lower images; control) sides of SVZ from stimulated coronal sections of mice in panel B. Identical settings from same brain section were used for imaging. Red arrows represent DCX$^+$ neuroblasts around subep-ChAT$^+$ neurons.

(D) P-S6 intensity analysis of subep-ChAT$^+$ neurons in ipsilateral side (inhibition) vs. contralateral side (control). P = 0.0038, n=8, Paired t-test. Data collected from three stimulated *Cr-Cre* mice. Each dot represents a Subep-ChAT$^+$ neuron.

(E) DCX (grey) immunofluorescence staining of ipsilateral SVZ (left image; inhibition) vs. contralateral SVZ (right image; control) from stimulated coronal section of mice in panel C. Identical settings from same brain section were used for imaging. Red arrow displays DCX$^+$ cells adjacent to subep-ChAT$^+$ neurons in ventral domain of the stimulated and control sides.

(F) Analysis of DCX$^+$ cells in stimulated coronal sections of ipsilateral side (inhibition) vs. contralateral side (control). P<0.0001, n=9, Paired t-test. Data collected from four stimulated *Cr-Cre* mice. Each dot represents a total DCX$^+$ cells per coronal section.

(G) Experimental design of *in vivo* optogenetic inhibition for one day post AAV_hSyn1-SIO-stGtACR2-FusionRed virus injection into ipsilateral Cg1 and optical fiber implantation into Cg1 regions of P28 *Cr-Cre* mice.

(H) EdU (purple), ChAT (green) and pEGFR (grey) immunofluorescence staining of (H') ipsilateral SVZ wholemount (inhibition) (upper images; V-SVZ and lower images; SVZ) vs. (H'') contralateral SVZ wholemount (control) (upper images; V-SVZ and lower images; SVZ) from stimulated mice in panel G. Identical settings from same brain section were used for imaging. Blue arrows display EdU$^+$/pEGFR$^+$ cells surrounding subep-ChAT$^+$ neurons in V-SVZ and SVZ. Red arrows display EdU$^+$/pEGFR$^-$ cells surrounding subep-ChAT$^+$ neurons in V-SVZ and SVZ.

(I) Left: Analysis of EdU$^+$/pEGFR$^+$ cells in ipsilateral (inhibition) V-SVZ vs. contralateral (control) V-SVZ of SVZ wholemounts. P = 0.039, n=13, Unpaired t-test. Data collected from four stimulated *Cr-Cre* mice. Each dot represents the total EdU$^+$/pEGFR$^+$ cells surrounding a subep-ChAT$^+$ neuron.

Right: Analysis of EdU$^+$/pEGFR$^+$ cells in ipsilateral (inhibition) SVZ vs. contralateral (control) SVZ of SVZ wholemounts. P = 0.0006, n=13, Unpaired t-test. Data collected from four stimulated *Cr-Cre* mice. Each dot represents total EdU$^+$/pEGFR$^+$ cells surrounding a subep-ChAT$^+$ neuron.

All error bars indicate SEM.

Figure 9 Schematic summary of presynaptic inputs into subep-ChAT$^+$ neuron within LV-SVZ niche.