

1 Single-deletion-mutant, third-generation rabies viral vectors allow 2 nontoxic retrograde targeting of projection neurons with greatly 3 increased efficiency

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10 11 SUMMARY

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13 Rabies viral vectors have become important components of the systems neuroscience
14 toolkit, allowing both direct retrograde targeting of projection neurons and monosynaptic
15 tracing of inputs to defined postsynaptic populations, but the rapid cytotoxicity of first-
16 generation (ΔG) vectors limits their use to short-term experiments. We recently introduced
17 second-generation, double-deletion-mutant (ΔGL) rabies viral vectors, showing that they
18 efficiently retrogradely infect projection neurons and express recombinases effectively but
19 with little to no detectable toxicity; more recently, we have shown that ΔGL viruses can be
20 used for monosynaptic tracing with far lower cytotoxicity than the first-generation system.
21 Here we introduce third-generation (ΔL) rabies viral vectors, which, like first-generation
22 vectors, have only a single gene deleted from their genomes (in this case the viral
23 polymerase gene L) but which appear to be as nontoxic as second-generation ones: using
24 longitudinal structural and functional two-photon imaging in mouse visual cortex *in vivo*, we
25 found that they did not kill labeled neurons or noticeably perturb their response properties
26 over the entire months-long courses of imaging. Although third-generation vectors are
27 therefore phenotypically very similar to second-generation ones, we show that they have the
28 major advantage of growing to much higher titers, and this key difference results in 25% -
29 525% increased numbers of retrogradely labeled neurons *in vivo*. These ΔL rabies viral
30 vectors therefore constitute a new state of the art for minimally perturbative, pathway-
31 specific expression of recombinases and transactivators in mammalian neurons selected on
32 the basis of their axonal projections. Because replication of deletion-mutant rabies viruses
33 within complementing cells is precisely the process that underlies monosynaptic tracing,
34 the higher replication efficiency of this new class of rabies viral vectors furthermore
35 suggests the potential to provide the foundation of an improved nontoxic monosynaptic
36 tracing system.

37 38 INTRODUCTION

39
40 Since their introduction to neuroscience in 2007 (Wickersham et al., 2007a; Wickersham et al., 2007b),
41 recombinant rabies viral vectors have become widely-adopted tools in neuroscience, allowing "monosynaptic
42 tracing" of direct inputs to genetically-targeted starting postsynaptic neuronal populations (Jin et al., 2021b;
43 Wall et al., 2010; Wickersham et al., 2007b) as well as simple retrograde targeting of projection neurons
44 when injected at the sites of these projection neurons' axonal arborizations (Chatterjee et al., 2018;
45 Wickersham et al., 2007a). These vectors are now used in a large number of laboratories worldwide and
46 have contributed to many high-impact studies of a wide variety of neural systems (Foster et al., 2021;
47 Miyamichi et al., 2011; Reardon et al., 2016; Schwarz et al., 2015; Siu et al., 2021; Smith et al., 2021;
48 Stephenson-Jones et al., 2016; Wu et al., 2021; Yao et al., 2021).

49 Because first generation (" ΔG ") rabies viral vectors (which have only the glycoprotein gene G deleted
50 from their genomes) are cytotoxic (Chatterjee et al., 2018; Jin et al., 2021a; Jin et al., 2021b; Wickersham et
51 al., 2007a), we recently introduced second-generation, " ΔGL " rabies viral vectors, which have both the
52 glycoprotein gene G and the viral polymerase gene L (for "large" protein) deleted from their genomes
53 (Chatterjee et al., 2018). Because the viral polymerase is absolutely required for transcription of all genes

54 from the rabies viral genome as well as for replication of the viral genome itself (Albertini et al., 2011; Finke
55 and Conzelmann, 2005; Horwitz et al., 2020; Morin et al., 2013; Ogino and Green, 2019; Te Velthuis et al.,
56 2021), this additional deletion, by design, reduces gene expression to a minimal level (provided by the few
57 starting copies of the polymerase protein that are copackaged in each viral particle) that appears to be
58 completely harmless to the "infected" cells. Because transgene expression is reduced by the same degree,
59 we inserted the genes for Cre and Flpo recombinase, of which even low levels of expression are sufficient
60 to cause neuroscientifically-useful downstream effects such as expression of fluorophores or calcium
61 indicators in labeled cells (Chatterjee et al., 2018). We originally showed that these Δ GL vectors are useful
62 tools for retrograde targeting of projection neurons (Chatterjee et al., 2018), and they have since been used
63 as such for applications including optogenetics and transcriptomic profiling (Ren et al., 2021; Roy et al., 2021;
64 Tasic et al., 2018). More recently, we have also shown that Δ GL vectors can be complemented *in vivo* by
65 expression of both G and L *in trans*, yielding a second-generation monosynaptic tracing system with far lower
66 cytotoxicity than the first-generation version (Jin et al., 2021a).

67 Here we show that deletion of L alone appears to make rabies viral vectors as nontoxic as Δ GL ones,
68 with labeled neurons surviving for at least months with apparently unperturbed visual response properties.
69 We find that these Δ L vectors have a major growth advantage over Δ GL ones in cell culture, attaining much
70 higher titers in complementing cells in culture. This higher replication efficiency translates into the practical
71 advantage of retrogradely labeling many more projection neurons when injected into these neurons' target
72 sites *in vivo*.

73

74 RESULTS

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76 Construction and characterization of Δ L rabies virus

77 We began by constructing rabies viral vectors with only the polymerase gene deleted and characterized their
78 gene expression levels and growth dynamics in cell culture (Figure 1). Beginning with the genome plasmid
79 of a Δ GL virus (Chatterjee et al., 2018), we reinserted the native glycoprotein gene in its original location,
80 followed by the gene for Cre recombinase (codon-optimized for mouse (Koresawa et al., 2000)) in an
81 additional transcriptional unit, then produced infectious virus by standard techniques (see Methods). We then
82 compared the gene expression levels of the resulting virus, RV Δ L-Cre, to those of first- and second-
83 generation versions (RV Δ G-Cre and RV Δ GL-Cre, respectively) in cell culture (HEK 293T/17 cells) using
84 immunostaining for Cre as well as for the viral nucleoprotein, the highest-expressed rabies viral protein.

85 As shown in Figure 1A-D, whereas the first-generation (Δ G) virus expressed high levels of
86 nucleoprotein (which accumulated in cytoplasmic inclusions) and Cre (which localized to the nuclei), the Δ GL
87 and Δ L viruses had very low expression levels of both Cre and nucleoprotein, with the amount of label for
88 these proteins appearing much more similar to that seen in uninfected control cells than in cells infected with
89 the Δ G virus. We also found similarly low transgene expression levels for Δ L and Δ GL viruses expressing
90 EGFP (Figure S1). However, just as we found previously for Δ GL viruses (Chatterjee et al., 2018), the Cre
91 expressed by RV Δ L-Cre was sufficient to result in bright labeling of Cre reporter cells (bottom row in panels
92 A-D).

93 These results led us to predict that Δ L viruses would be as nontoxic as Δ GL ones, because of their
94 similarly low expression levels, and also that they would be similarly able to recombine reporter alleles *in*
95 *vivo* in order to allow downstream expression of useful transgene products such as fluorophores, activity
96 indicators, or opsins.

97 It remained to be seen, however, whether Δ L viruses would have any particular advantage over Δ GL
98 ones for purposes of retrogradely targeting neurons. Specifically, if they could not be produced at significantly
99 higher titers, they could be expected to label similar numbers of projection neurons, making Δ L vectors a
100 mere curiosity of purely academic interest and with no relevance to neuroscientists. However, if they could
101 be grown to much higher titers than Δ GL vectors, that could be expected to translate to the ability to
102 retrogradely label many more projection neurons, a desirable characteristic indeed for a tool for retrograde
103 targeting.

104 To examine this, we directly compared the ability of Δ L virus to replicate in complementing cells with
105 that of Δ GL and Δ G viruses (Figure 1E-F). We infected cell lines expressing L, G, or both with the three
106 different generations of virus, at two different multiplicities of infection (MOI, measured in infectious units per
107 cell): either very low (MOI = 0.01, "multi-step growth curves" (Gomme et al., 2010; Wang and Bushman,
108 2006)) or high (MOI = 1, "single-step growth curves". Following a one-hour incubation in the presence of the

109 viruses, we washed the cells twice with DPBS and applied fresh medium, then collected supernatant samples
110 every 24 hours for five days after infection, then titered the samples on reporter cells.

111 As seen in Figure 1E-F, the results were clear: whereas the Δ GL virus (on cells expressing both G
112 and L) never accumulated to titers higher than 2.37×10^6 iu/mL in either experiment, the Δ L virus grew to
113 maximal titers of 6.51×10^6 iu/mL (at MOI of 0.01) and 1.96×10^7 iu/mL (at MOI of 1) on the same cell line
114 (expressing both G and L) and considerably higher (maximal titers of 1.31×10^7 iu/mL at MOI=0.01 and 2.60×10^7
115 iu/mL at MOI=1) on cells expressing L alone. The Δ G virus grew to similarly high (or slightly higher, in the
116 MOI=1 case) titers to the Δ L one: 6.83×10^6 iu/mL at MOI=0.01 and 3.03×10^7 iu/mL at MOI=1, suggesting that
117 single-deletion-mutant rabies viruses may in general be easier to make at high titers than viruses with multiple
118 deleted genes. In non-complementing cells, by contrast, no such replication of any of these viruses (Δ L, Δ GL,
119 or Δ G) occurred (Figure S2).

120 These findings that a Δ L rabies virus could be grown to 11-fold higher titer than a matched Δ GL one
121 led us to predict that Δ L viruses would be superior tools for retrograde targeting *in vivo*, because their much
122 higher titers would result in retrograde infection of many more projection neurons.

123

124 **Retrograde targeting *in vivo***

125 To test this prediction, we made matched preparations of Δ GL and Δ L viruses expressing either Cre or Flpo
126 (mouse-codon-optimized Flp recombinase (Raymond and Soriano, 2007)), then injected each of the four
127 viruses in the somatosensory thalami of reporter mice (Ai14 (Madisen et al., 2010) for the Cre viruses, Ai65F
128 (Daigle et al., 2018) for the Flpo ones; both lines express tdTomato following recombination by the respective
129 recombinase). We sacrificed the mice at either 7 days or 4 weeks after injection, sectioned and imaged the
130 brains by confocal microscopy, and counted the numbers of retrogradely labeled cells in cortex. Figure 2
131 shows the results.

132 As seen in panels 2B-E, for both recombinases, and at both timepoints, the Δ L viruses significantly
133 outperformed the Δ GL ones. For the Flpo viruses, the difference was dramatic: at the 1-week timepoint, the
134 Δ L virus labeled 24 times as many cells as the Δ GL one (although this difference was not statistically
135 significant due to high variance in the Δ L cohort: single factor ANOVA, $p=0.275$, $n = 8$ mice each group); by
136 the 4-week timepoint, the Δ L-Flpo virus had labeled 6.25 times as many cells as the Δ GL counterpart, a
137 difference that was extremely significant (single factor ANOVA, $p=3.21 \times 10^{-4}$, $n = 8$ mice each group). For the
138 Cre viruses, the difference was smaller, presumably due a ceiling effect (see Discussion), but still highly
139 significant: at 1 week, the Δ L-Cre virus had labeled 1.40 times as many cells as Δ GL-Cre (single factor
140 ANOVA, $p=4.20 \times 10^{-3}$; $n = 4$ mice each group); at the 4-week timepoint, the Δ L-Cre virus had labeled 1.25
141 times as many cells as Δ GL-Cre (single factor ANOVA, $p=7.38 \times 10^{-4}$, $n = 4$ mice per group).

142 We also made some injections of RV Δ L-Cre, in thalami of Ai14 mice, with the much longer survival
143 times of 4 or 6 months (Fig 2F-G). The results at both of these longer survival times appeared very similar to
144 those at the shorter ones. Consistent with extensive prior literature on corticothalamic neurons (Alitto and
145 Usrey, 2003; Rockland, 2021; Rouiller and Welker, 2000) and with our previous results with corticothalamic
146 injections of Δ G and Δ GL viruses (Chatterjee et al., 2018; Wickersham et al., 2007a), the cells labeled in
147 cortex by both viruses at all timepoints were pyramidal neurons in layer 6, with a few in layer 5. Furthermore,
148 labeled neurons all appeared morphologically normal even months after injection, with the fine processes of
149 axons and dendrites, including individual spines (rightmost images in 2F-G) clearly visible and without
150 blebbing or other obvious abnormalities.

151 As a further test of the flexibility of Δ L vectors, we made a version expression the tetracycline
152 transactivator (tTA) and injected it in the thalamus of Ai63 reporter mice (in which TRE-tight drives tdTomato
153 expression) (Daigle et al., 2018). As seen in Figure S3, thousands of cortical thalamic cells were found
154 retrogradely labeled at both 1-week and 4-week survival times, with no significant difference between the
155 numbers at the two timepoints (single factor ANOVA, $p = 0.772$, $n = 4$ mice per group).

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157 **Longitudinal structural two-photon imaging *in vivo***

158 Because examining only postmortem tissue can be very misleading when attempting to determine whether
159 a virus is nontoxic (see Jin et al. '21 (Jin et al., 2021a) for a detailed case study), we conducted longitudinal
160 two-photon imaging of RV Δ L-labeled neurons *in vivo* (Figure 3). We injected either RV Δ GL-Cre (Chatterjee
161 et al., 2018) or RV Δ L-Cre in primary visual cortex (V1) of Ai14 reporter mice, then imaged the resulting
162 tdTomato-expressing neurons at or near the injection site beginning 7 days after injection and continuing
163 every 7 or 14 days until 16 weeks postinjection. As seen in Figure 3, the results using the two viruses were
164 very similar. For both Δ GL and Δ L viruses, the numbers of visibly labeled neurons increased significantly

165 between 1 week and 4 weeks postinjection (Figure 3D, G), by 56.27% for Δ GL and by 67.77% for Δ L (paired
166 t-tests, $p = 1.319\text{E-}04$ for Δ GL, $p = 1.003\text{E-}05$ for Δ L, $n = 8$ FOVs for each virus). Also for both viruses, the
167 numbers of visibly labeled neurons remained nearly completely constant from the 4-week timepoint onward
168 through all remaining imaging sessions (Figure 3E, H) (with the number of labeled cells at the 4-week
169 timepoint being not significantly different than that at the 12-week timepoint for the Δ L virus (paired t-test, p
170 $= 0.1327$, $n = 8$ FOVs) and slightly (0.5%) lower for the Δ GL virus (paired t-test, $p = 0.0056$, $n = 8$ FOVs).
171 See File S3 for all counts and statistical comparisons; see also Video S1 for a rendering of a group of Δ L-
172 labeled neurons at 2 weeks and again at 10 weeks.
173

174 Longitudinal functional two-photon imaging *in vivo*

175 We went on to examine the functional properties of RV Δ L-labeled neurons *in vivo*. As for the structural
176 imaging (see above), we injected RV Δ L-Cre in the primary visual cortices of reporter mice, in this case mice
177 that express the calcium indicator GCaMP6s (Chen et al., 2013) after Cre recombination (Figure 4).
178 Beginning one week later, we began imaged the calcium signals in the labeled neurons in a series of imaging
179 sessions that continued until 16 weeks postinjection, in the awake mice viewed visual stimuli consisting of
180 drifting gratings of different orientations and frequencies. Just as we found previously for Δ GL viruses
181 (Chatterjee et al., 2018; Jin et al., 2021b), we found no signs of dysfunction in cells labeled by the third-
182 generation vector for as long as we followed them (Figure 4; see also Figures S4 and S5. See File S4 for all
183 counts and statistical comparisons and Video S2 for an example of calcium responses in a group of cortical
184 neurons 16 weeks after injection of RV Δ L-Cre).
185

186 DISCUSSION

187
188 Here we have shown that deletion of just the polymerase gene renders rabies viral vectors nontoxic, like
189 second-generation (Δ GL) vectors, but makes them much more efficient at replicating within complementing
190 cells in culture. This ability to be grown to much higher titers results in significantly increased transduction of
191 projection neurons within a given pathway. This more comprehensive access to projection neurons will
192 increase the yield and efficacy of systems neuroscience experiments that depend on the retrograde targeting
193 approach.

194 In the corticothalamic pathway that we have examined here, the advantage of a Δ L vector over the
195 Δ GL equivalent was clearest in the case of the Flpo-expressing versions, with the Δ L vector labeling 6.25
196 times as many neurons as the Δ GL one did at four weeks postinjection. This ratio is of the same order of
197 magnitude as the ratio of the titers of the injected Flpo viruses (14.4: see Methods). By contrast, for the Cre-
198 expressing versions, the advantage of the Δ L vector over the Δ GL one was more modest, labeling 1.25 times
199 as many cells, even though the ratio of the titers of these Cre vectors was even higher (20.5). Because the
200 absolute numbers of retrogradely labeled neurons, as well as the titers, were much higher for the Cre viruses
201 than for the corresponding Flpo ones, we presume that the smaller advantage of the Δ L version seen in this
202 case was because of a ceiling effect, with the Δ GL-Cre virus already labeling most of the available neurons
203 in this pathway.

204 One could certainly argue that the much higher titers that we are easily able to obtain with Δ L vectors
205 could also, in theory, potentially be achieved with Δ GL vectors, if enough effort were put into generating and
206 testing producer cell lines expressing both G and L in order to find one that expressed the two genes at just
207 the right ratio and levels. In practice, however, this hypothetical future research effort does not detract from
208 the fact that the best currently-existing preparations of Δ L rabies viral vectors label many more cells than do
209 Δ GL ones, making them the better choice for retrograde targeting applications.

210 We note that, although here we have only demonstrated the use of Δ L rabies viral vectors in mice,
211 they are also highly likely to work in a wide variety of mammalian species, because, apart from their shorter
212 RNA genomes, the structural properties of second- and third-generation rabies viral particles are identical to
213 those of first-generation ones, which have been successfully used in diverse mammalian species including
214 rats (Cruz et al., 2021), cats (Connolly et al., 2012; Liu et al., 2013), ferrets (Hasse et al., 2019), and
215 macaques (Bragg et al., 2017; Briggs et al., 2016; Lyon et al., 2010; Nassi and Callaway, 2006, 2007; Nassi
216 et al., 2006; Siu et al., 2021; Yarch et al., 2017) (and even in fish (Dohaku et al., 2019; Satou et al., 2021;
217 Zhu et al., 2009) and frogs (Faulkner et al., 2021)).

218 Our findings here that Δ L rabies viruses have extremely low expression levels and do not replicate
219 within (or spread beyond, *in vivo*) non-complementing cells are entirely consistent with similar findings in cell
220 culture in a recent report on an L-deficient rabies virus encoding firefly luciferase (Nakagawa et al., 2017).

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A note about safety: our results strongly suggest that ΔL rabies viruses are unable to replicate in the absence of complementation and moreover are harmless to any cells that they transduce. However, a mixture of ΔL and ΔG viruses could pose a safety risk, because such viruses will be mutually complementary. Care must therefore be taken to avoid contamination between ΔL and ΔG constructs – either packaged viruses or the genome plasmids used to make them – which would have the potential to create a self-complementing replication-competent mixture (see Hidaka et al. (Hidaka et al., 2018) for an example of such a self-complementing mixture).

Finally, we have recently shown (Jin et al., 2021b) that second-generation (ΔGL) rabies viral vectors can spread transsynaptically when complemented by provision of both G and L *in trans*. That is, complementation of an L-deficient rabies virus (in that case, a G- and L-deficient virus that is also complemented by G) allows it to spread beyond initially infected cells *in vivo*. It is therefore reasonable to infer that provision of L *in trans* should allow third-generation, ΔL rabies viral vectors to spread beyond initially infected cells, given that we have shown here that such complementation in cell culture allows ΔL viruses to replicate very efficiently. We have also shown here, with the longitudinal two-photon imaging of labeled neurons, that ΔL viruses do not spread beyond initially infected cells *in vivo* in the absence of complementation. Collectively, our results therefore suggest the outlines of a third-generation monosynaptic tracing system based on ΔL vectors complemented with L expression *in trans*. However, genetic targeting of a ΔL vector to specific starting cell types might appear elusive: in the first- and second-generation systems (Jin et al., 2021b; Wickersham et al., 2007b), this targeting is achieved by packaging the rabies viral particles with an avian retroviral envelope protein (EnvA) instead of its own envelope glycoprotein, so that they can only infect cells that have been engineered to express EnvA's cognate receptor. On the face of it, this pseudotyping strategy requires that G be deleted from the rabies viral genome, because expression of G by the virus within the EnvA-expressing producer cells would result in the production of virions with membranes populated by a mixture of EnvA and the rabies viral glycoprotein. If this challenge could be overcome, our present findings that ΔL viruses replicate more readily in complementing cells, which is the fundamental process central to monosynaptic tracing (Wickersham et al., 2007b), suggest that a third-generation monosynaptic tracing system could be more efficient than the second-generation one.

METHODS

All experiments involving animals were conducted according to NIH guidelines and approved by the MIT Committee for Animal Care. Mice were housed 1-5 per cage under a normal light/dark cycle for all experiments.

Cloning

The third-generation rabies viral vector genome plasmids pRV ΔL -5Cre, pRV ΔL -5Flpo, and pRV ΔL -5tTA (Addgene 182964, 182965, and 182966) (the "5" denoting the position of the transgene relative to the other genes in the viral genome) was made by replacing the mCre gene in pRV ΔGL -4Cre (Chatterjee et al., 2018) (Addgene 98039) with the SAD B19 glycoprotein gene from pCAG-B19G (Chatterjee et al., 2018) (Addgene 59921) and either the mCre, Flpo (from pRV ΔG -4Flpo (Addgene 98040)), or tTA (from pAAV-syn-FLEX-splitTVA-EGFP-tTA (Liu et al., 2017) (Addgene 100798)) gene, separated by endogenous rabies viral transcriptional stop and start signals, using seamless cloning (InFusion (Takara) or HiFi (NEB)).

The piggyBac vector pB-TREtight-EGFP (Addgene 182967) was made by cloning the TRE-tight element from pAAV-TREtight-mTagBFP2-B19G (Liu et al., 2017) and the EGFP gene into pB-CAG-TEVp-IRES-mCherry (Addgene 174377) in place of the CAG-TEVp-IRES-mCherry sequences using HiFi seamless cloning (NEB).

The piggyBac plasmid pB-CAG-B19G-IRES-EGFP-WPRE-BGHpA (Addgene 178517) was made by cloning the CAG promoter from pCAG-B19G (Addgene 59921), the SAD B19 L gene, the EMCV IRES (Gallardo et al., 1997), the mCherry (Shaner et al., 2004) gene, and the woodchuck post-transcriptional regulatory element and bovine growth hormone polyadenylation signal from pCSC-SP-PW-GFP (Addgene 12337), into PB-CMV-MCS-EF1-Puro (System Biosciences #PB510B-1).

Cell lines

The BHK-B19G3 cell line, expressing the SAD B19 strain rabies virus glycoprotein gene, was made by resorting BHK-B19G2 cells (Wickersham et al., 2010) on a BD FACS Aria cell sorter and retaining the

277 brightest 2% of EGFP-positive cells as well as the next-brightest 18%. Following the sort, both populations
278 were expanded and refrozen, then thawed and tested for their efficacy at supporting replication of ΔG virus;
279 the second-brightest population ("BHK-B19G3_2") was found to result in higher titers and is referred to
280 here as BHK-B19G3.

281 The BHK-B19L cell line, expressing the SAD B19 strain rabies virus polymerase gene, was made
282 by transfecting BHK-21 cells (ATCC CCL-10) with pCAG-hypBase (Jin et al., 2021a) and pB-CAG-B19L-
283 IRES-mCherry-WPRE-BGHpA (Jin et al., 2021b) using Lipofectamine 2000 (Thermo Fisher 11668019),
284 then expanding the cells and sorting on a FACS Aria sorter (BD) to collect the brightest 5%, as well as the
285 next brightest 5%, of mCherry-expressing cells. The two collected populations were expanded and
286 refrozen, then thawed and tested for their efficacy at supporting replication of ΔL virus; the second-brightest
287 population ("BHK-B19L_2") was found to result in higher titers and is referred to here as BHK-B19L.

288 The BHK-B19L-G cell line, expressing the SAD B19 strain rabies virus polymerase and glycoprotein
289 genes, was made by transfecting BHK-B19L cells (see above) with pCAG-hypBase and pB-CAG-B19G-
290 IRES-EGFP-WPRE-BGHpA (see above), then expanding and sorting on a BD FACS Aria, keeping the
291 brightest 5%, as well as the next brightest 5%, of EGFP-expressing cells which also expressed mCherry.
292 The sorted cells were expanded and refrozen, then thawed and tested for their efficacy at supporting
293 replication of ΔGL virus; the brightest population ("BHK-B19L-G_1") was found to result in higher titers and
294 is referred to here as BHK-B19L-G.

295 The 293T-TREtight-EGFP cell line for titering tTA-expressing viruses was made by transfecting
296 HEK 293T/17 cells with pCAG-hypBase and pB-TREtight-EGFP (described above), then expanded and
297 sorted on a BD FACS Aria, excluding the brightest 2% of EGFP cells, and keeping four of the next brightest
298 EGFP cell populations. The sorted cells were expanded, frozen, and then thawed for testing their efficacy
299 at titering ΔL -tTA virus. The fourth-brightest tranche of cells was used for subsequent titering of ΔL -tTA
300 virus.

301

302 Rabies virus production and titering

303 The first-generation vector RV ΔG -4Cre, the second-generation vectors RV ΔGL -4Cre and RV ΔGL -4Flpo,
304 and the third-generation vectors RV ΔL -5Cre, RV ΔL -5Flpo, and RV ΔL -5tTA were rescued as described
305 previously (Chatterjee et al., 2018) using genome plasmids pRV ΔGL -4Cre, pRV ΔGL -4Flpo, pRV ΔL -5Cre,
306 pRV ΔL -5Flpo, and pRV ΔL -5tTA, respectively. For simplicity, these viruses are referred to in this manuscript
307 as RV ΔG -Cre, RV ΔGL -Cre, RV ΔGL -Flpo, RV ΔL -Cre, RV ΔL -Flpo, and RV ΔL -tTA, omitting the numbers
308 denoting the positions of the transgenes within the viral genomes. Rescue supernatants were collected and
309 filtered as described (Wickersham and Sullivan, 2015), titered on the reporter cell lines 293T-FLEX BC (for
310 Cre viruses) or 293T-F14F15S-BC (for Flpo viruses) (Jin et al., 2021a) as described (Wickersham et al.,
311 2010), then used to infect BHK-B19G3, BHK-B19L-G, or BHK-B19L cells (see above) at multiplicities of
312 infection ranging from 0.1 to 1. Supernatants from these "P1" plates were collected and titered as
313 described (Wickersham and Sullivan, 2015); in some cases, these were used for a similar second passage
314 ("P2"). Purification and concentration of either P1 or P2 supernatants was as described (Wickersham et al.,
315 2010), with supernatants treated with benzonase (Sigma 71206) (25 minute incubation at 37°C with 30
316 units/ml at) before ultracentrifugation. Concentrated viruses were aliquoted and frozen at -80°C. Rabies
317 viruses were titered on reporter cells (293T-FLEX-BC for Cre viruses, 293T-F14F15S-BC for Flpo viruses,
318 293T-TREtight-EGFP (see above) for RV ΔL -tTA) as described (Wickersham et al., 2010), using a LUNA-II
319 cell counter (Logos Biosystems) instead of a hemocytometer for counting cells, and in some cases using
320 two-fold (as opposed to ten-fold) dilution series for more precise comparisons of titers.

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322 Immunostaining, imaging, and flow cytometry of cultured cells

323 For anti-nucleoprotein and anti-Cre staining (for Figure 1): HEK 293T/17 (ATCC 11268) cells were plated
324 on poly-L-lysine-coated coverslips in 24-well plates, then infected the following day with serial dilutions of
325 RV ΔG -4Cre (Chatterjee et al., 2018), RV ΔGL -4Cre (Chatterjee et al., 2018), or RV ΔL -5Cre. Three days
326 after infection, cells were fixed with 2% paraformaldehyde, washed repeatedly with
327 blocking/permeabilization buffer (0.1% Triton-X (Sigma) and 1% bovine serum albumin (Sigma) in PBS),
328 then labeled with a 1:100 dilution of anti-nucleoprotein monoclonal antibody blend (Light Diagnostics
329 Rabies DFA Reagent, EMD Millipore 5100) as well as a 1:250 dilution of rabbit anti-Cre polyclonal antibody
330 (Millipore Sigma 69050) followed by a 1:200 dilution of Alexa Fluor 594-conjugated donkey anti-rabbit
331 secondary (Jackson Immuno 711-585-152).

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333 For anti-EGFP staining (for Figure S1), HEK cells were plated as above, then infected the following day
334 with serial dilutions of RV Δ G-4EGFP (Wickersham et al., 2010), RV Δ GL-4EGFP (Chatterjee et al., 2018),
335 or RV Δ L-5EGFP, with immunostaining three days postinfection, using a 1:1000 dilution of chicken anti-GFP
336 polyclonal antibody (Aves Labs, GFP-1020) and a 1:500 dilution of Alexa Fluor 594-conjugated donkey
337 anti-chicken secondary antibody (Jackson Immuno 703-585-155).

338 Immunostained cells on coverslips were mounted on slides using Prolong Diamond Antifade
339 mounting medium (Thermo P36970) and imaged on a Zeiss LSM 900 confocal microscope using a 20x
340 objective.

341 For matched flow cytometric analysis of immunostained cells, cells were plated in 24-well plates
342 without poly-L-lysine-coated coverslips but otherwise immunostained as described above, then analyzed
343 on an LSR II flow cytometer (BD) using FACS Diva software (BD). Histograms displayed in Figure 1 were
344 smoothed using the FACS Diva "Smooth histogram" setting.

345 346 **Viral growth analysis**

347 For determining growth curves, BHK-B19G3, BHK-B19L-G, and BHK-B19L cells (see above) were plated
348 in 10 cm plates coated in poly-L-lysine in normal medium (10% fetal bovine serum (VWR 16777-014) and
349 antibiotic-antimycotic (Thermo 15240096) in DMEM (Thermo 11995073)) (Wickersham et al., 2010). The
350 following day, cells were infected with RV Δ G-4Cre(B19G), RV Δ GL-4Cre(B19G), or RV Δ L-5Cre(B19G) at
351 an MOI of either 1 (for single-step growth curves) or 0.01 (for multi-step growth curves), with viruses diluted
352 in normal medium at a total volume of 2 ml per plate, with each condition in triplicate. Following a one-hour
353 incubation, the virus-containing medium was aspirated, plates were washed twice in DPBS (Thermo
354 14190144), and 12 ml fresh medium was added to each plate before they were returned to the incubator.
355 Every 24 hours for the following five days, 200 μ l of supernatant was collected from each plate; these
356 supernatant samples were filter-sterilized using a 96-well 0.45 μ m PVDF filter plate (Millipore MSHVN4510),
357 then frozen at -80°C before all samples were thawed and titered on HEK 293T-FLEX-BC cells as described
358 above.

359 360 **Mouse strains**

361 The Cre-dependent tdTomato reporter line Ai14 (Madisen et al., 2010) was purchased from Jackson
362 Laboratory (catalog # 007914). The Flp-dependent tdTomato reporter line Ai65F was obtained by crossing
363 the Cre- and Flp-dependent tdTomato double-reporter line Ai65D (Madisen et al., 2015) (Jackson
364 Laboratory 021875) to the Cre deleter line Meox2-Cre (Tallquist and Soriano, 2000) (Jackson Laboratory
365 003755), then breeding out the Meox2-Cre allele. An equivalent Ai65F line, made using a different Cre
366 deleter line, was described in Daigle et al. '18 (Daigle et al., 2018) and is now available from Jackson
367 Laboratory (catalog # 032864). The tTA-dependent tdTomato reporter line Ai63 (Daigle et al., 2018) was a
368 generous gift from Hongkui Zeng and Tanya Daigle. Mice used for the functional two-photon imaging
369 experiments were crosses of the Cre- and tTA-dependent GCaMP6s line Ai94D (Jackson Laboratory
370 024104) with the Cre-dependent tTA line ROSA:LNL:tTA (Wang et al., 2008) (Jackson Laboratory 011008).
371 All mice were maintained in a C57BL/6J (Jackson Laboratory 000664) background.

372 For experiments, adult mice of both sexes were used, of the following mouse strains. For retrograde
373 targeting using Cre-expressing viruses (Figure 2) and structural two-photon imaging (Figure 3): Ai14
374 heterozygotes. For retrograde targeting using Flpo-expressing viruses (Figure 2): Ai65F heterozygotes. For
375 retrograde targeting using RV Δ L-tTA (Figure S3): Ai63 heterozygotes. For functional two-photon imaging
376 (Figures 4, S4, and S5): Ai94D x ROSA:LNL:tTA double homozygotes.

377 378 **Stereotaxic injections**

379 200 nl of rabies virus was injected into either somatosensory thalamus (VPM/Po, for figure 2) or primary
380 visual cortex (for two-photon experiments) of anesthetized adult mice using a stereotaxic instrument
381 (Stoelting Co., 51925) and a custom injection apparatus consisting of a hydraulic manipulator (Narishige,
382 MO-10) with headstage coupled via custom adaptors to a wire plunger advanced through pulled glass
383 capillaries (Drummond, Wiretrol II) back-filled with mineral oil and front-filled with viral vector solution (Lavin
384 et al., 2019). We have described this injection system in detail previously. Injection coordinates for VPM/Po
385 were: anteroposterior (AP) = -1.82 mm with respect to (w.r.t.) bregma, lateromedial (LM) = +1.54 mm w.r.t
386 bregma, dorsoventral (DV) = -3.15 mm w.r.t the brain surface; injection coordinates for V1 cortex were: AP
387 = -2.70 mm w.r.t. bregma, LM = 2.50 mm w.r.t. bregma, DV = -0.26 mm w.r.t the brain surface.

388

389 For mice to be used for two-photon imaging, a 3 mm craniotomy was opened over primary visual
390 cortex (V1). Glass windows composed of a 3mm-diameter glass coverslip (Warner Instruments CS-3R)
391 glued (Optical Adhesive 61, Norland Products) to a 5mm-diameter glass coverslip (Warner Instruments
392 CS-5R) were affixed over the craniotomy with Metabond (Parkell) after virus injection.

393 For the Δ GL vs. Δ L experiments (Figure 2), the four viruses were produced in parallel for direct
394 comparison, and RV Δ L-Cre (6.16E+10 i.u./ml) or RV Δ GL-Cre (3.01E+09 i.u./ml) was injected into Ai14
395 (het) mice, and RV Δ L-Flpo (1.61E+09 i.u./ml) or RV Δ GL-4Flpo (1.12E+08 i.u./ml) was injected into Ai65F
396 (het) mice. For the 4-month and 6-month experiments for Figure 2, RV Δ L-Cre (1.66E+10 i.u./ml) was
397 injected into Ai14 (het) mice. For Figure S3, RV Δ L-tTA (3.63E+10 iu/ml) was injected into Ai63 (het) mice.

398 For two-photon structural experiments (Figure 3), RV Δ GL-Cre (1.19E+10 iu/ml) or RV Δ L-Cre
399 (1.66E+10 iu/ml diluted to 1.19E+10 iu/ml for matching to RV Δ GL-Cre) was injected into Ai14 (het) mice.
400 For two-photon functional experiments in Figure 4, RV Δ L-Cre (2.61E+10 iu/ml) was injected into
401 homo/homo Ai94D x ROSA:LNL:tTA mice.

402

403 **Perfusions, histology, and confocal imaging**

404 1 week to 6 months (see main text) after injection of rabies virus, anesthetized mice were transcardially
405 perfused with 4% paraformaldehyde. Brains were postfixed overnight in 4% paraformaldehyde in PBS on a
406 shaker at 4°C and cut into 50 μ m coronal sections on a vibrating microtome (Leica, VT-1000S). Sections
407 were collected sequentially into 6 tubes containing cryoprotectant, so that each tube contained every sixth
408 section, then frozen at -20°C. Sections to be imaged were washed to remove cryoprotectant, then mounted
409 with Prolong Diamond Antifade mounting medium (Thermo Fisher P36970) and imaged on a confocal
410 microscope (Zeiss, LSM 900). To ensure that the confocal images included in the figures are
411 representative of each group, the images were taken after the counts were conducted, and the mouse with
412 the next higher number of labeled neurons than the average number for its group was selected for confocal
413 imaging.

414

415 **Quantification of retrograde targeting**

416 Coronal sections between 0.43mm anterior and 4.07mm posterior to bregma were imaged with an
417 epifluorescence microscope for cell counting (Zeiss, Imager.Z2). Due to the high density of retrogradely
418 labeled tdTomato neurons in the cortex at the injection site (VPM/Po), cells were counted using the
419 Analyze Particle function in ImageJ (size in micron²: 20-400; circularity: 0.20-1.00). Only one of the six
420 series of sections (i.e., every sixth section: see above) was counted for each mouse. P-values for all
421 comparisons were obtained using single-factor ANOVAs.

422

423 **Structural two-photon imaging and image analysis**

424 Beginning seven days after injection of each rabies virus and recurring at the subsequent indicated
425 timepoints (see main text) up to a maximum of 16 weeks following rabies virus injection, fields of view
426 (FOVs) were imaged on a Prairie/Bruker Ultima IV In Vivo two-photon microscope driven by a Spectra
427 Physics Mai-Tai Deep See laser with a mode locked Ti:sapphire laser emitting at a wavelength of 1020 nm
428 for excitation of tdTomato. In order to distinguish individual labeled neurons, FOVs were chosen some
429 distance away from the area of brightest tdTomato labeling. Two well-separated areas were chosen in
430 each mouse. For each imaging session, mice were reanesthetized and mounted via their headplates to a
431 custom frame, with ointment applied to protect their eyes and with a handwarmer maintaining body
432 temperature. Imaging parameters were as follows: image size 512 X 512 pixels (282.6 μ m x 282.6 μ m),
433 0.782 Hz frame rate, dwell time 4.0 μ s, 2x optical zoom, Z-stack step size 1 μ m. Image acquisition was
434 controlled with Prairie View 5.4 software. Laser power exiting the 20x water-immersion objective (Zeiss, W
435 plan-apochromat, NA 1.0) varied between 20 and 65 mW depending on focal plane depth (Pockels cell
436 value was automatically increased from 450 at the top section of each stack to 750 at the bottom section).
437 For the example images of labeled cells, maximum intensity projections (stacks of 100-200 μ m) were made
438 with ImageJ software. Cell counting was automated using the "Analyze Particles" function in ImageJ. Plots
439 of cell counts were made with Prism 9 (GraphPad Software, San Diego, California).

440

441 **Functional two-photon imaging and image analysis**

442 For functional two-photon imaging of RV Δ L-Cre-labeled cells, FOVs were slightly offset from the regions of
443 brightest GCaMP6s label in left-hemisphere V1 in order to allow separate identification of individual cells.
444 This imaging was performed using the same microscope (5.356-Hz frame rate, 1024X 128 pixels, 565.1

445 $\mu\text{m} \times 565.1 \mu\text{m}$, dwell time 0.8 μs , 1x optical zoom, scan angle 45 degree) with the same objective and
446 laser (at 920 nm) as for the structural imaging experiments. Laser power at the objective ranged from 10 to
447 65 mW. Calcium imaging data were acquired in supragranular layers (100 to 200 μm deep). Surface
448 vasculature provided coarse fiducial markers for finding the same FOVs in different imaging sessions.
449 For these experiments, mice were awake and head-fixed. No behavioral training or reward was given.
450 Visual stimuli were generated in Matlab (R2015R version) with custom software based on Psychtoolbox
451 (<http://psychtoolbox.org>) and shown on the same LCD screen as in the widefield mapping experiments.
452 Each condition consisted of 2 s of a full-field sine wave grating drifting in one direction, presented at 80%
453 contrast with spatial frequency of 0.04 cycles/ degree, followed by 2 s of uniform mean luminance (gray).
454 All permutations of 12 directions (30° steps) and 5 temporal frequencies (1, 2, 4, 8 and 15 Hz) were shown,
455 in randomized order. The complete set was repeated 10 times, for a total stimulation period of 40 min per
456 FOV per session. Cells were then manually segmented, and single-cell fluorescence traces were extracted
457 by averaging the fluorescence of all pixels masking the soma, using ImageJ (version 2-0-0-rc-69) software.
458 The mean $\Delta F/F$ over the full 2 s of each stimulus condition was used to calculate orientation tuning curves,
459 with background fluorescence (F) in $\Delta F/F$ taken as the value of the trace immediately preceding a
460 condition, averaged over all conditions. The raw calcium traces from cells within individual FOVs (not
461 across FOVs, given different imaging conditions across animals and time points) were sorted by mean
462 fluorescence. Randomly colored ROI view images were created by suite2p (<https://www.suite2p.org>). For
463 'tuned' cells in Figure 4 panels F and G, the counts are based on all imaged neurons' individual tuning
464 curves, plotted in MATLAB; any cell showing response to a preferred orientation (including narrowly tuned
465 neurons and broadly tuned neurons) at any temporal frequency (1Hz, 2Hz, 4Hz, 8Hz, or 15Hz) was
466 counted manually as a tuned cell.

467 **RESOURCE AVAILABILITY**

468
469
470 All cell counts and statistical analyses are provided in Supplemental Information. The novel plasmids
471 described in this paper have been deposited with Addgene with the accession numbers given in Methods.

472 **SUPPLEMENTAL INFORMATION**

473
474
475 Supplemental information can be found online.
476 A preprint version of this paper is available on bioRxiv.

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478
479
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485 **AUTHOR CONTRIBUTIONS**

486
487
488 L.J., N.E.L., M.M., Y.H., and M.Z. cloned constructs; H.A.S. produced viruses with assistance from L.J. and
489 M.Z.; H.A.S. conducted cell culture assays and immunocytochemistry; L.J., N.E.L., and T.K.L. performed
490 surgeries; L.J. and M.Z. performed histology and confocal imaging; L.J. performed two-photon imaging; L.J.,
491 M.Z., T.K.L., and N.E.L. managed mouse breeding; I.W. planned and supervised all work; I.W. and L.J. wrote
492 the manuscript with input from the other authors.

493 **DECLARATION OF INTERESTS**

494
495
496 I.R.W. is a consultant for Monosynaptix, LLC, advising on design of neuroscientific experiments.

497 **REFERENCES**

498
499

- 500 Albertini, A.A., Ruigrok, R.W., and Blondel, D. (2011). Rabies virus transcription and replication. *Advances*
501 *in virus research* 79, 1-22.
- 502 Alitto, H.J., and Usrey, W.M. (2003). Corticothalamic feedback and sensory processing. *Curr Opin Neurobiol*
503 13, 440-445.
- 504 Bragg, E.M., Fairless, E.A., Liu, S., and Briggs, F. (2017). Morphology of visual sector thalamic reticular
505 neurons in the macaque monkey suggests retinotopically specialized, parallel stream-mixed input to the
506 lateral geniculate nucleus. *J Comp Neurol* 525, 1273-1290.
- 507 Briggs, F., Kiley, C.W., Callaway, E.M., and Usrey, W.M. (2016). Morphological Substrates for Parallel
508 Streams of Corticogeniculate Feedback Originating in Both V1 and V2 of the Macaque Monkey. *Neuron* 90,
509 388-399.
- 510 Chatterjee, S., Sullivan, H.A., MacLennan, B.J., Xu, R., Hou, Y., Lavin, T.K., Lea, N.E., Michalski, J.E.,
511 Babcock, K.R., Dietrich, S., *et al.* (2018). Nontoxic, double-deletion-mutant rabies viral vectors for retrograde
512 targeting of projection neurons. *Nat Neurosci* 21, 638-646.
- 513 Chen, T.W., Wardill, T.J., Sun, Y., Pulver, S.R., Renninger, S.L., Baohan, A., Schreiter, E.R., Kerr, R.A.,
514 Orger, M.B., Jayaraman, V., *et al.* (2013). Ultrasensitive fluorescent proteins for imaging neuronal activity.
515 *Nature* 499, 295-300.
- 516 Connolly, J.D., Hashemi-Nezhad, M., and Lyon, D.C. (2012). Parallel feedback pathways in visual cortex of
517 cats revealed through a modified rabies virus. *J Comp Neurol* 520, 988-1004.
- 518 Cruz, A.M., Kim, T.H., and Smith, R.J. (2021). Monosynaptic Retrograde Tracing From Prelimbic Neuron
519 Subpopulations Projecting to Either Nucleus Accumbens Core or Rostromedial Tegmental Nucleus. *Front*
520 *Neural Circuits* 15, 639733.
- 521 Daigle, T.L., Madisen, L., Hage, T.A., Valley, M.T., Knoblich, U., Larsen, R.S., Takeno, M.M., Huang, L., Gu,
522 H., Larsen, R., *et al.* (2018). A Suite of Transgenic Driver and Reporter Mouse Lines with Enhanced Brain-
523 Cell-Type Targeting and Functionality. *Cell* 174, 465-480 e422.
- 524 Dohaku, R., Yamaguchi, M., Yamamoto, N., Shimizu, T., Osakada, F., and Hibi, M. (2019). Tracing of
525 Afferent Connections in the Zebrafish Cerebellum Using Recombinant Rabies Virus. *Front Neural Circuits*
526 13, 30.
- 527 Faulkner, R.L., Wall, N.R., Callaway, E.M., and Cline, H.T. (2021). Application of Recombinant Rabies Virus
528 to *Xenopus* Tadpole Brain. *eNeuro*.
- 529 Finke, S., and Conzelmann, K.K. (2005). Replication strategies of rabies virus. *Virus Res*.
- 530 Foster, N.N., Barry, J., Korobkova, L., Garcia, L., Gao, L., Becerra, M., Sherafat, Y., Peng, B., Li, X., Choi,
531 J.H., *et al.* (2021). The mouse cortico-basal ganglia-thalamic network. *Nature* 598, 188-194.
- 532 Gallardo, H.F., Tan, C., and Sadelain, M. (1997). The internal ribosomal entry site of the
533 encephalomyocarditis virus enables reliable coexpression of two transgenes in human primary T
534 lymphocytes. *Gene Ther* 4, 1115-1119.
- 535 Gomme, E.A., Faul, E.J., Flomenberg, P., McGettigan, J.P., and Schnell, M.J. (2010). Characterization of a
536 single-cycle rabies virus-based vaccine vector. *Journal of virology* 84, 2820-2831.
- 537 Hasse, J.M., Bragg, E.M., Murphy, A.J., and Briggs, F. (2019). Morphological heterogeneity among
538 corticogeniculate neurons in ferrets: quantification and comparison with a previous report in macaque
539 monkeys. *J Comp Neurol* 527, 546-557.
- 540 Hidaka, Y., Lim, C.K., Takayama-Ito, M., Park, C.H., Kimitsuki, K., Shiwa, N., Inoue, K.I., and Itou, T. (2018).
541 Segmentation of the rabies virus genome. *Virus Res* 252, 68-75.
- 542 Horwitz, J.A., Jenni, S., Harrison, S.C., and Whelan, S.P.J. (2020). Structure of a rabies virus polymerase
543 complex from electron cryo-microscopy. *Proc Natl Acad Sci U S A* 117, 2099-2107.
- 544 Jin, L., Matsuyama, M., Sullivan, H.A., Zhu, M., Lavin, T.K., Hou, Y., Lea, N.E., Pruner, M.T., Fernández,
545 M.L.D., and Wickersham, I.R. (2021a). Rabies virus with a destabilization domain added to its nucleoprotein
546 spreads between neurons only if the domain is removed. *bioRxiv*, 550640.
- 547 Jin, L., Sullivan, H.A., Zhu, M., Lavin, T.K., Matsuyama, M., Lea, N.E., Xu, R., Hou, Y., Rutigliani, L., Pruner,
548 M., *et al.* (2021b). Long-term labeling and imaging of synaptically-connected neuronal networks *in*
549 *vivo* using nontoxic, double-deletion-mutant rabies viruses. *bioRxiv*, 2021.2012.2004.471186.
- 550 Koresawa, Y., Miyagawa, S., Ikawa, M., Matsunami, K., Yamada, M., Shirakura, R., and Okabe, M. (2000).
551 Synthesis of a new Cre recombinase gene based on optimal codon usage for mammalian systems. *J*
552 *Biochem* 127, 367-372.
- 553 Lavin, T.K., Jin, L., and Wickersham, I.R. (2019). Monosynaptic tracing: a step-by-step protocol. *J Chem*
554 *Neuroanat*, 101661.

- 555 Liu, K., Kim, J., Kim, D.W., Zhang, Y.S., Bao, H., Denaxa, M., Lim, S.A., Kim, E., Liu, C., Wickersham, I.R.,
556 *et al.* (2017). Lhx6-positive GABA-releasing neurons of the zona incerta promote sleep. *Nature* 548, 582-
557 587.
- 558 Liu, Y.J., Ehrenguber, M.U., Negwer, M., Shao, H.J., Cetin, A.H., and Lyon, D.C. (2013). Tracing inputs to
559 inhibitory or excitatory neurons of mouse and cat visual cortex with a targeted rabies virus. *Curr Biol* 23,
560 1746-1755.
- 561 Lyon, D.C., Nassi, J.J., and Callaway, E.M. (2010). A disynaptic relay from superior colliculus to dorsal
562 stream visual cortex in macaque monkey. *Neuron* 65, 270-279.
- 563 Madisen, L., Garner, A.R., Shimaoka, D., Chuong, A.S., Klapoetke, N.C., Li, L., van der Bourg, A., Niino, Y.,
564 Egolf, L., Monetti, C., *et al.* (2015). Transgenic mice for intersectional targeting of neural sensors and
565 effectors with high specificity and performance. *Neuron* 85, 942-958.
- 566 Madisen, L., Zwingman, T.A., Sunkin, S.M., Oh, S.W., Zariwala, H.A., Gu, H., Ng, L.L., Palmiter, R.D.,
567 Hawrylycz, M.J., Jones, A.R., *et al.* (2010). A robust and high-throughput Cre reporting and characterization
568 system for the whole mouse brain. *Nature neuroscience* 13, 133-140.
- 569 Miyamichi, K., Amat, F., Moussavi, F., Wang, C., Wickersham, I., Wall, N.R., Taniguchi, H., Tasic, B., Huang,
570 Z.J., He, Z., *et al.* (2011). Cortical representations of olfactory input by trans-synaptic tracing. *Nature* 472,
571 191-196.
- 572 Morin, B., Kranzusch, P.J., Rahmeh, A.A., and Whelan, S.P. (2013). The polymerase of negative-stranded
573 RNA viruses. *Current opinion in virology* 3, 103-110.
- 574 Nakagawa, K., Kobayashi, Y., Ito, N., Suzuki, Y., Okada, K., Makino, M., Goto, H., Takahashi, T., and
575 Sugiyama, M. (2017). Molecular Function Analysis of Rabies Virus RNA Polymerase L Protein by Using an
576 L Gene-Deficient Virus. *J Virol* 91.
- 577 Nassi, J.J., and Callaway, E.M. (2006). Multiple circuits relaying primate parallel visual pathways to the
578 middle temporal area. *J Neurosci* 26, 12789-12798.
- 579 Nassi, J.J., and Callaway, E.M. (2007). Specialized circuits from primary visual cortex to V2 and area MT.
580 *Neuron* 55, 799-808.
- 581 Nassi, J.J., Lyon, D.C., and Callaway, E.M. (2006). The parvocellular LGN provides a robust disynaptic input
582 to the visual motion area MT. *Neuron* 50, 319-327.
- 583 Ogino, T., and Green, T.J. (2019). Transcriptional Control and mRNA Capping by the GDP
584 Polyribonucleotidyltransferase Domain of the Rabies Virus Large Protein. *Viruses* 11.
- 585 Raymond, C.S., and Soriano, P. (2007). High-efficiency FLP and PhiC31 site-specific recombination in
586 mammalian cells. *PLoS One* 2, e162.
- 587 Reardon, T.R., Murray, A.J., Turi, G.F., Wirblich, C., Croce, K.R., Schnell, M.J., Jessell, T.M., and Losonczy,
588 A. (2016). Rabies Virus CVS-N2c(DeltaG) Strain Enhances Retrograde Synaptic Transfer and Neuronal
589 Viability. *Neuron* 89, 711-724.
- 590 Ren, W., Centeno, M.V., Wei, X., Wickersham, I., Martina, M., Apkarian, A.V., and Surmeier, D.J. (2021).
591 Adaptive alterations in the mesoaccumbal network after peripheral nerve injury. *Pain* 162, 895-906.
- 592 Rockland, K.S. (2021). A Closer Look at Corticothalamic "Loops". *Front Neural Circuits* 15, 632668.
- 593 Rouiller, E.M., and Welker, E. (2000). A comparative analysis of the morphology of corticothalamic
594 projections in mammals. *Brain Res Bull* 53, 727-741.
- 595 Roy, D.S., Zhang, Y., Aida, T., Choi, S., Chen, Q., Hou, Y., Lea, N.E., Skaggs, K.M., Quay, J.C., Liew, M.,
596 *et al.* (2021). Anterior thalamic dysfunction underlies cognitive deficits in a subset of neuropsychiatric disease
597 models. *Neuron* 109, 2590-2603 e2513.
- 598 Satou, C., Neve, R.L., Oyibo, H.K., Zmarz, P., Huang, K.-H., Bouldoires, E.A., Mori, T., Higashijima, S.-i.,
599 Keller, G.B., and Friedrich, R.W. (2021). A viral toolbox for conditional and transneuronal gene expression in
600 zebrafish. *bioRxiv*, 2021.2003.2025.436574.
- 601 Schwarz, L.A., Miyamichi, K., Gao, X.J., Beier, K.T., Weissbourd, B., DeLoach, K.E., Ren, J., Ibanes, S.,
602 Malenka, R.C., Kremer, E.J., *et al.* (2015). Viral-genetic tracing of the input-output organization of a central
603 noradrenaline circuit. *Nature*.
- 604 Shaner, N.C., Campbell, R.E., Steinbach, P.A., Giepmans, B.N., Palmer, A.E., and Tsien, R.Y. (2004).
605 Improved monomeric red, orange and yellow fluorescent proteins derived from *Discosoma* sp. red fluorescent
606 protein. *Nat Biotechnol* 22, 1567-1572.
- 607 Siu, C., Balsor, J., Merlin, S., Federer, F., and Angelucci, A. (2021). A direct interareal feedback-to-
608 feedforward circuit in primate visual cortex. *Nat Commun* 12, 4911.
- 609 Smith, M.L., Asada, N., and Malenka, R.C. (2021). Anterior cingulate inputs to nucleus accumbens control
610 the social transfer of pain and analgesia. *Science* 371, 153-159.

611 Stephenson-Jones, M., Yu, K., Ahrens, S., Tucciarone, J.M., van Huijstee, A.N., Mejia, L.A., Penzo, M.A.,
612 Tai, L.H., Wilbrecht, L., and Li, B. (2016). A basal ganglia circuit for evaluating action outcomes. *Nature* 539,
613 289-293.

614 Tallquist, M.D., and Soriano, P. (2000). Epiblast-restricted Cre expression in MORE mice: a tool to distinguish
615 embryonic vs. extra-embryonic gene function. *Genesis* 26, 113-115.

616 Tasic, B., Yao, Z., Graybiel, L.T., Smith, K.A., Nguyen, T.N., Bertagnolli, D., Goldy, J., Garren, E., Economo,
617 M.N., Viswanathan, S., *et al.* (2018). Shared and distinct transcriptomic cell types across neocortical areas.
618 *Nature* 563, 72-78.

619 Te Velthuis, A.J.W., Grimes, J.M., and Fodor, E. (2021). Structural insights into RNA polymerases of
620 negative-sense RNA viruses. *Nat Rev Microbiol* 19, 303-318.

621 Wall, N.R., Wickersham, I.R., Cetin, A., De La Parra, M., and Callaway, E.M. (2010). Monosynaptic circuit
622 tracing in vivo through Cre-dependent targeting and complementation of modified rabies virus. *P Natl Acad*
623 *Sci USA*.

624 Wang, G.P., and Bushman, F.D. (2006). A statistical method for comparing viral growth curves. *J Virol*
625 *Methods* 135, 118-123.

626 Wang, L., Sharma, K., Deng, H.X., Siddique, T., Grisotti, G., Liu, E., and Roos, R.P. (2008). Restricted
627 expression of mutant SOD1 in spinal motor neurons and interneurons induces motor neuron pathology.
628 *Neurobiol Dis* 29, 400-408.

629 Wickersham, I.R., Finke, S., Conzelmann, K.K., and Callaway, E.M. (2007a). Retrograde neuronal tracing
630 with a deletion-mutant rabies virus. *Nature Methods* 4, 47-49.

631 Wickersham, I.R., Lyon, D.C., Barnard, R.J., Mori, T., Finke, S., Conzelmann, K.K., Young, J.A., and
632 Callaway, E.M. (2007b). Monosynaptic restriction of transsynaptic tracing from single, genetically targeted
633 neurons. *Neuron* 53, 639-647.

634 Wickersham, I.R., and Sullivan, H.A. (2015). Rabies viral vectors for monosynaptic tracing and targeted
635 transgene expression in neurons. *Cold Spring Harb Protoc* 2015, 375-385.

636 Wickersham, I.R., Sullivan, H.A., and Seung, H.S. (2010). Production of glycoprotein-deleted rabies viruses
637 for monosynaptic tracing and high-level gene expression in neurons. *Nature protocols* 5, 595-606.

638 Wu, X., Morishita, W., Beier, K.T., Heifets, B.D., and Malenka, R.C. (2021). 5-HT modulation of a medial
639 septal circuit tunes social memory stability. *Nature* 599, 96-101.

640 Yao, S., Wang, Q., Hirokawa, K.E., Ouellette, B., Ahmed, R., Bomben, J., Brouner, K., Casal, L., Caldejon,
641 S., Cho, A., *et al.* (2021). A whole-brain monosynaptic input connectome to neuron classes in mouse visual
642 cortex. *bioRxiv*, 2021.2009.2029.459010.

643 Yarch, J., Federer, F., and Angelucci, A. (2017). Local Circuits of V1 Layer 4B Neurons Projecting to V2
644 Thick Stripes Define Distinct Cell Classes and Avoid Cytochrome Oxidase Blobs. *J Neurosci* 37, 422-436.

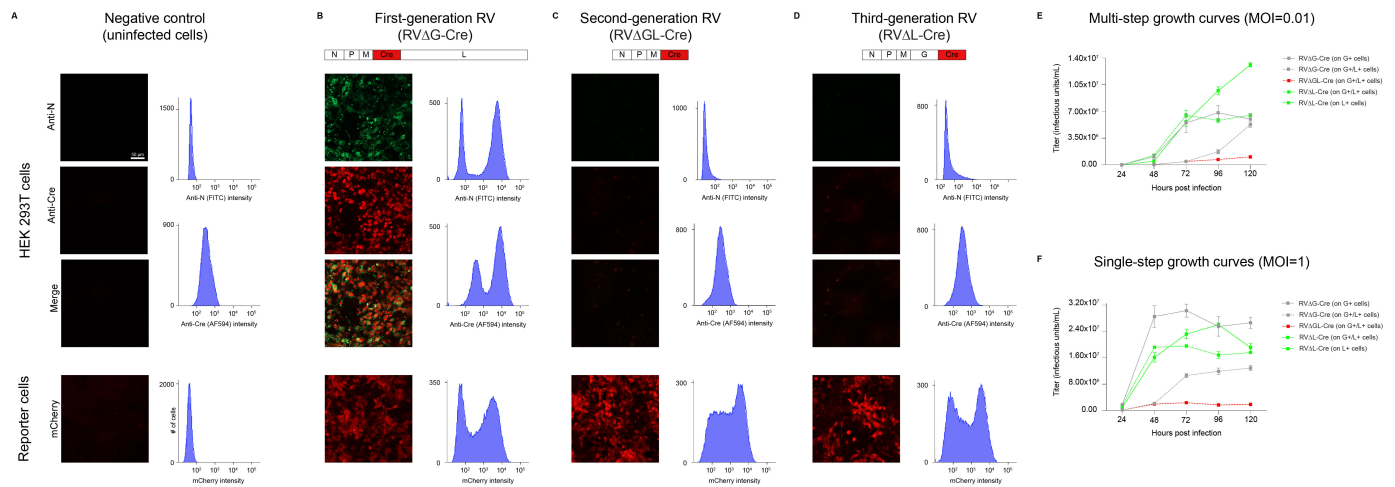
645 Zhu, P., Narita, Y., Bundschuh, S.T., Fajardo, O., Scharer, Y.P., Chattopadhyaya, B., Bouldoires, E.A.,
646 Stepien, A.E., Deisseroth, K., Arber, S., *et al.* (2009). Optogenetic Dissection of Neuronal Circuits in Zebrafish
647 using Viral Gene Transfer and the Tet System. *Front Neural Circuits* 3, 21.

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FIGURES



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Figure 1. Rabies virus with just the polymerase gene deleted (ΔL) is phenotypically similar to double-deletion-mutant (ΔGL) virus but replicates to much higher titers within complementing cells.

(A-D) Deletion of just the polymerase gene L reduces transgene expression to levels that are very low but still sufficient to support reporter allele recombination in Cre reporter cells.

(A) Negative controls (uninfected cells). Top: Uninfected HEK 293T cells stained for rabies virus nucleoprotein (green) and for Cre (red). Histograms to right of panels show flow cytometric quantification of baseline fluorescence of uninfected cells in these channels. Bottom: Uninfected reporter cells which express mCherry following Cre recombination. Little signal is seen in these negative controls.

(B) Cells infected with a first-generation (ΔG) vector expressing Cre. Both Cre and N are expressed at very high levels, and infected Cre reporter cells brightly express mCherry (note that dilutions at which roughly half of cells were infected were chosen for this figure).

(C) Consistent with our previous findings (Chatterjee et al., 2018), expression of both nucleoprotein and Cre from a second-generation (ΔGL) vector is drastically reduced with respect to the first-generation vector, with expression levels comparable to those seen in negative controls. Despite this, the low Cre levels are still high enough to activate mCherry expression in reporter cells.

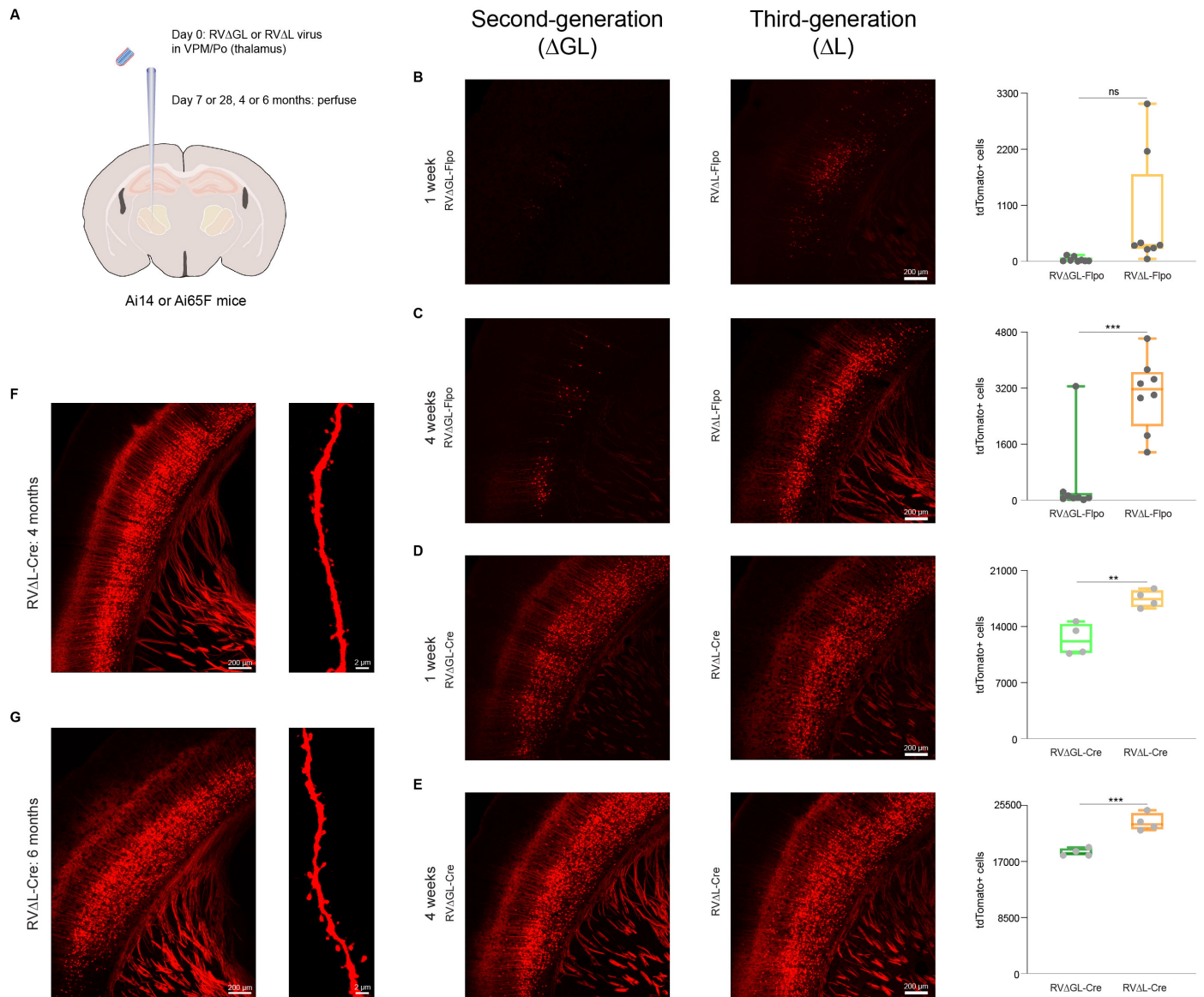
(D) A third-generation (ΔL) vector expresses nucleoprotein and Cre at similarly very low levels, but again Cre expression is nonetheless high enough to successfully activate mCherry expression in reporter cells.

(E-F) Third-generation (ΔL) vectors grow to much high titers in cultured cells than second-generation (ΔGL) ones do.

(E) Viral titers in supernatants of complementing cells (expressing L, G, or both) infected with ΔL , ΔGL , or ΔG viruses at a multiplicity of infection (MOI) of 0.01 ("multi-step growth curves"), with supernatants collected every 24 hours for five days. Whereas a ΔGL virus only achieves $1.05E+06$ infectious units (i.u./ml) over the duration of the experiment, the ΔL virus grows to 6.2-fold higher on the same cell line, and 12.5-fold higher on a line expressing L alone. The highest ΔL titers obtained in this experiment were significantly higher than the highest obtained with a first-generation (ΔG) virus (single-factor ANOVA, $p = 3.24E-03$, $n = 3$ replicates per condition).

(F) Similarly, at a MOI of 1 ("single-step" growth curves), the ΔGL virus titer peaks at $2.37E+06$ i.u./ml, whereas the peak titer of the ΔL virus is $2.60E+07$ i.u./ml, 11.0-fold higher than that of the ΔGL virus and not significantly different from that of the ΔG virus (single-factor ANOVA, $p = 0.105$, $n = 3$ replicates per condition). Graphs in (E-F) show means \pm s.e.m.

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686 **Figure 2. Third-generation (Δ L) rabies viral vectors retrogradely label many more projection**
687 **neurons *in vivo* than do second-generation (Δ GL) ones and leave cells morphologically normal**
688 **for at least six months.**

689 (A) Design of experiments retrogradely targeting corticothalamic cells in reporter mice. Either second-
690 generation vector RV Δ GL-Fipo or RV Δ GL-Cre, or third-generation vector RV Δ L-Fipo or RV Δ L-Cre, was
691 injected into somatosensory thalamus (VPM/Po) of either Ai65F (Fipo reporter) or Ai14 (Cre reporter). Mice
692 were perfused 1 week (b, d), 4 weeks (c, e), 4 months (f), or 6 months later (g).

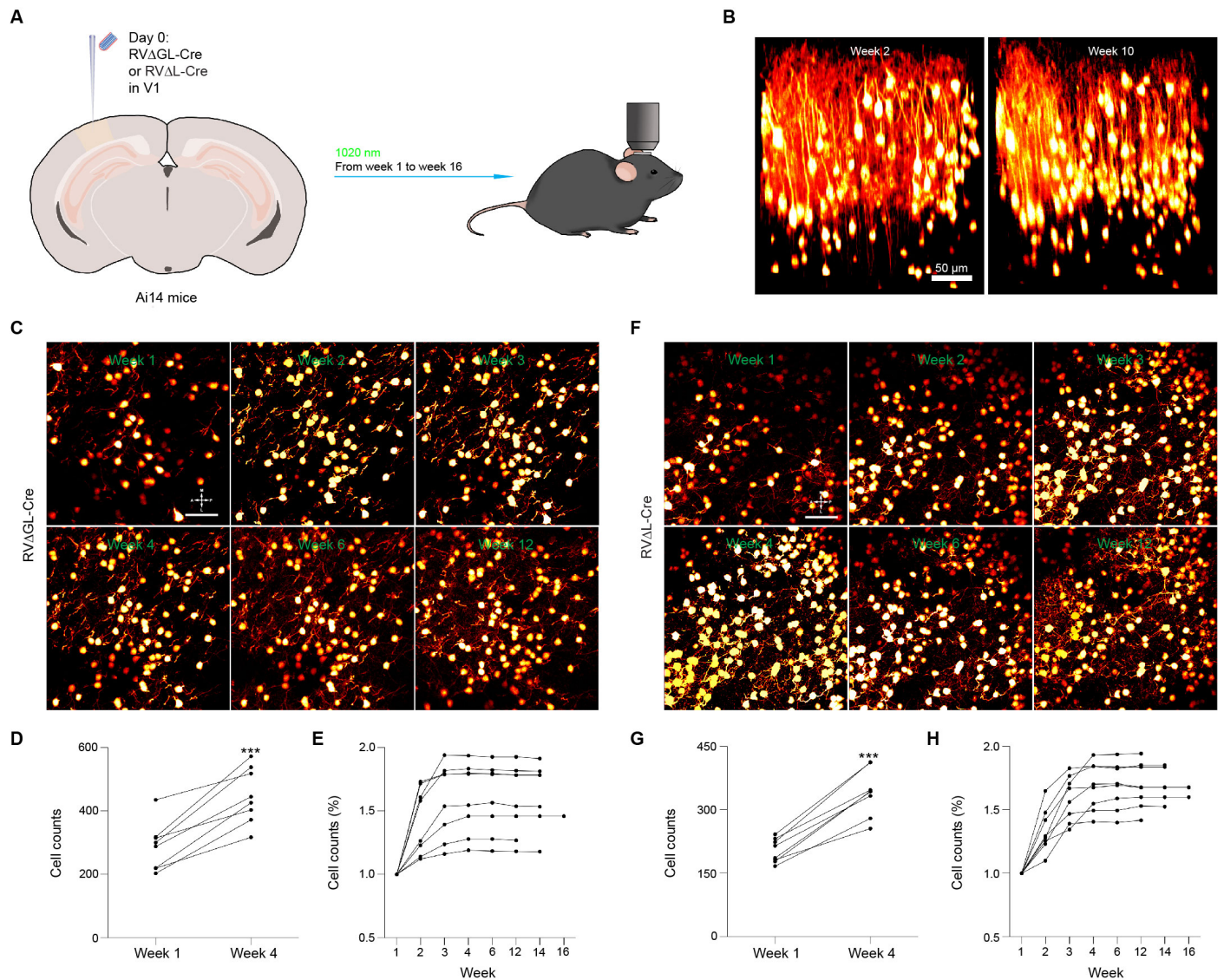
693 (B-E) Efficacy comparison of Fipo- and Cre-expressing Δ GL and Δ L vectors.

694 (B) Corticothalamic neurons in S1 of Ai65F mice labeled with RV Δ GL-Fipo (left) or RV Δ L-Fipo (center) at 1
695 week postinjection. Scale bar: 200 μ m, applies to both images. Counts of labeled cortical neurons
696 are shown at right (each data point is the total number in one series consisting of every sixth 50 μ m
697 section from a given brain - see Methods). The Δ L virus labeled 24 times as many cortical neurons
698 than the Δ GL virus did, although the difference in this case is not significant due to high variance
699 (single-factor ANOVA, $p = 0.0608$, $n = 8$ mice per group).

700 (C) Corticothalamic neurons in S1 of Ai65F mice labeled with RV Δ GL-Fipo (left) or RV Δ L-Fipo (center) at 4
701 weeks postinjection. Scale bar: 200 μ m, applies to both images. Counts of labeled cortical neurons
702 are shown at right. The Δ L virus labeled 6.25 times as many cortical neurons than the Δ GL virus
703 did, an extremely significant difference (single-factor ANOVA, $p = 0.000321$, $n = 8$ mice per group).

704 (D) Corticothalamic neurons in S1 of Ai14 mice labeled with RV Δ GL-Cre (left) or RV Δ L-Cre (center) at 1
705 week postinjection. Scale bar: 200 μ m, applies to both images. Counts of labeled cortical neurons are

706 shown at right. The ΔL virus labeled 1.4 times as many cortical neurons than the ΔGL virus did, a highly
707 significant difference (single-factor ANOVA, $p = 0.00420$, $n = 8$ mice per group).
708 (E) Corticothalamic neurons in S1 of Ai14 mice labeled with RV ΔGL -Cre (left) or RV ΔL -Cre (center) at 4
709 weeks postinjection. Scale bar: 200 μm , applies to both images. Counts of labeled cortical neurons are
710 shown at right. The ΔL virus labeled 1.25 times as many cortical neurons than the ΔGL virus did, an
711 extremely significant difference (single-factor ANOVA, $p = 0.000738$, $n = 8$ mice each group).
712 (F) Corticothalamic neurons in S1 of Ai14 mice labeled with RV ΔL -Cre at 4 months postinjection. Cells
713 appear morphologically completely normal, with no blebbing or decomposition of processes. Scale bars:
714 200 μm (left image) and 2 μm (right image).
715 (G) Corticothalamic neurons in S1 of Ai14 mice labeled with RV ΔL -Cre at 6 months postinjection. Cells still
716 appear morphologically completely normal. Scale bars: 200 μm (left image) and 2 μm (right image).



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719 **Figure 3. Neurons labeled by Δ L rabies virus survive for at least 16 weeks.**

720 (A) Experimental design for longitudinal structural two-photon imaging *in vivo*. Second-generation (Δ GL) or
721 third-generation (Δ L) virus expressing Cre was injected in primary visual cortex of reporter mice, then the
722 injection sites were imaged repeatedly for the following 16 weeks.

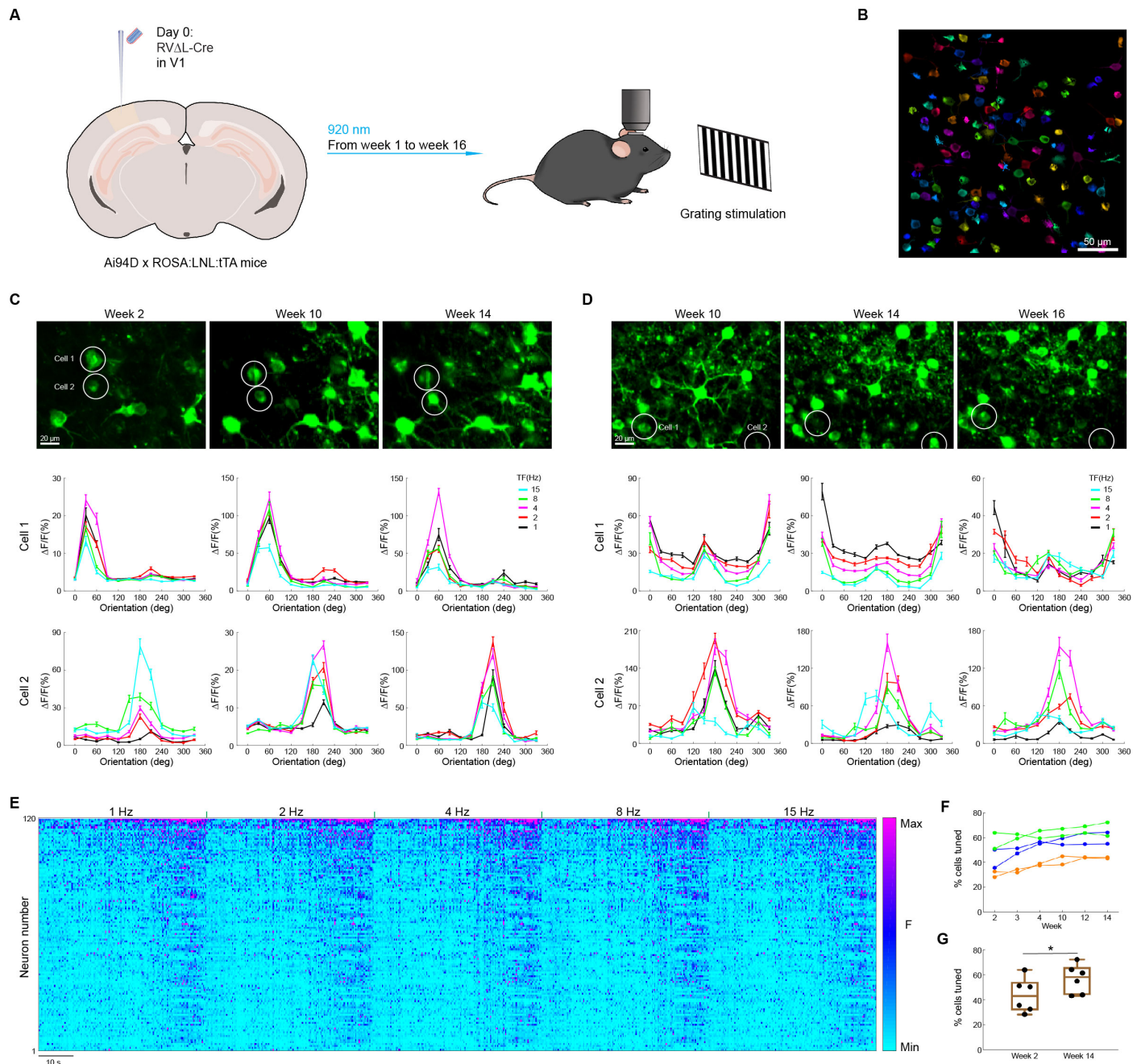
723 (B) Example renderings of the same volume of cortex labeled by RV Δ L-Cre and imaged with a two-photon
724 microscope at two different timepoints, 2 weeks (left) and 10 weeks (right). Every labeled neuron visible at
725 2 weeks is still present at 10 weeks. Scale bar: 50 μ m. See also Video S1.

726 (C) & (F), Example two-photon images of single fields of view (FOV) of cortex labeled by either the second-
727 generation vector RV Δ GL-Cre (C) or the third-generation vector RV Δ L-Cre (F), imaged at different
728 timepoints, from 1 week (top left) to 12 weeks (bottom right). All labeled neurons visible at earlier
729 timepoints are still present at later ones, for both viruses. Scale bars: 50 μ m, apply to all images.

730 (D) & (G), Absolute numbers of cells visibly labeled by RV Δ GL-Cre (D) or RV Δ L-Cre (G) for all structural
731 FOVs in the study, at the 1-week and 4-week timepoints. Numbers of visibly labeled cells increased by
732 56.27% for Δ GL and by 67.77% for Δ L, as we found previously for second-generation vectors (Chatterjee
733 et al., 2018)), suggesting accumulation and persistent activity of recombinase on this timescale. These
734 increases were both extremely significant (one-tailed paired t-tests, $p = 0.000132$ (Δ GL) and 0.00001003
735 (Δ L), $n = 8$ FOVs each virus), but there was no significant difference between the increases seen for the
736 two viruses (two-tailed unpaired t-test, $p = 0.5187$, $n = 8$ FOVs per group).

737 (E) & (H), Percentages of cells visibly labeled by RV Δ GL-Cre (E) and RV Δ L-Cre (H) over time, relative to
738 the numbers visible at 1 week after rabies injection; each connected set of dots represents numbers seen
739 in a given FOV at the different time points. For both viruses, the numbers of labeled neurons remain nearly

740 constant from the 4-week timepoint onward, as we found previously (Chatterjee et al., 2018) for Δ GL virus.
741 Imaging was discontinued for some mice at weeks 12 or 14 due to cloudiness of the optical windows.



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Figure 4. Δ L rabies virus does not appear to perturb neurons' visual response properties for at least 16 weeks.

746 (A) Experimental design for longitudinal functional two-photon imaging *in vivo*. Δ L virus expressing Cre was
747 injected in primary visual cortex of reporter mice expressing GCaMP6s (Chen et al., 2013) after Cre
748 recombination, then the injection sites were imaged while the awake mice were presented with drifting
749 grating stimuli of different orientations and temporal frequencies, repeatedly for 16 weeks following virus
750 injection.

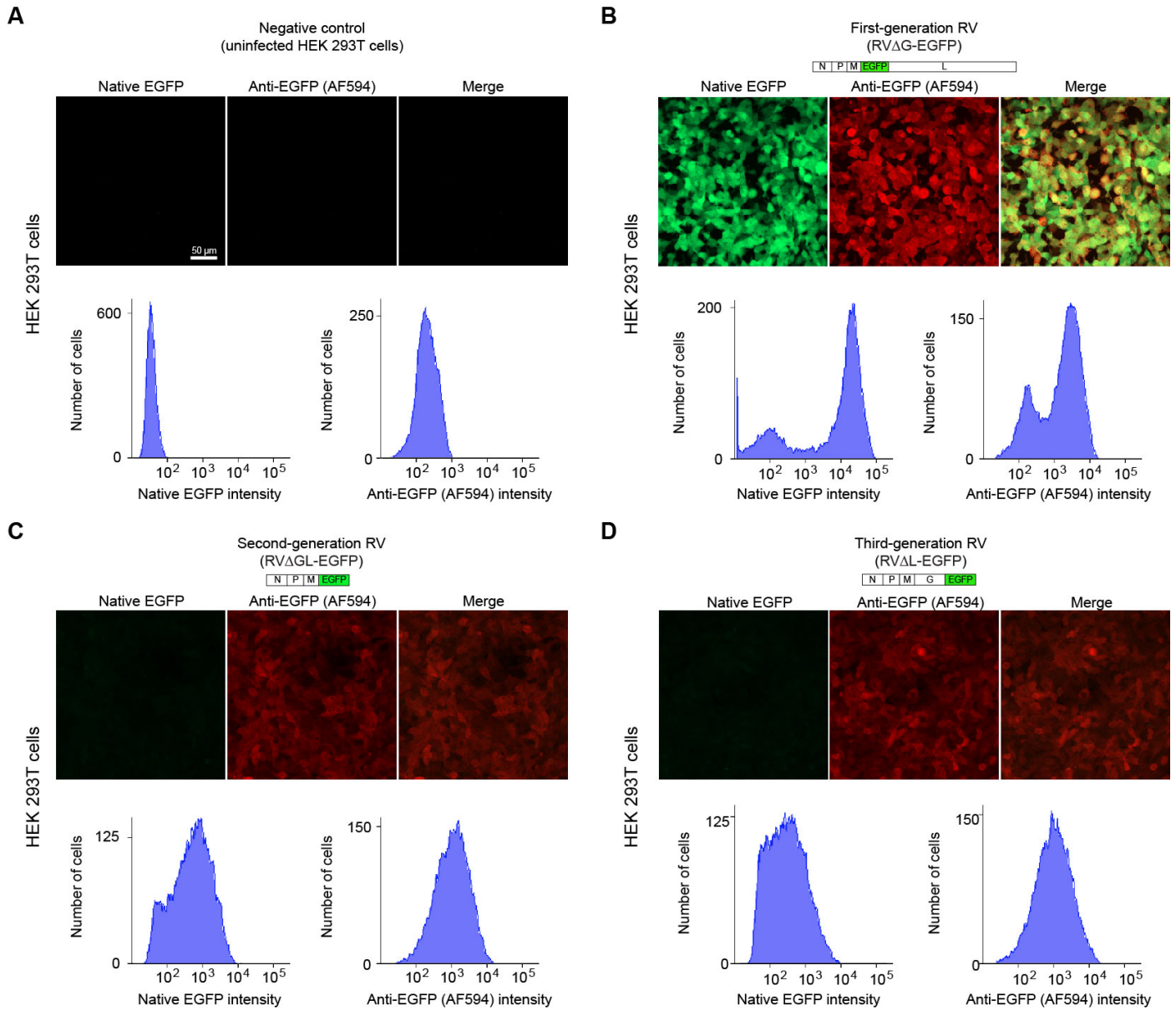
751 (B) Example FOV from a GCaMP6s imaging session 16 weeks after RV injection. Individual analyzed cells
752 are randomly pseudocolored. This is the same FOV as shown in Video S2. Scale bar: 50 μ m.

753 (C-D) Long-term stability of orientation and temporal frequency tuning in RV Δ L-Cre-labeled neurons. The
754 top rows show maximum intensity projections of the imaged GCaMP6s signal in two different FOVs at
755 three different timepoints for each FOV. Scale bars: 20 μ m, apply to all images. Visual response tuning
756 curves of the two circled cells in each FOV at the corresponding timepoint, obtained with drifting gratings
757 presented at 12 directions of motion and 5 temporal frequencies (TF) (mean Δ F/F \pm s.e.m., averaged over
758 10 repeats), are shown under each image. More examples from the same FOV are shown in Figure S5.

759 (E) Single-cell fluorescence time courses for 120 cells at the 12-week timepoint, showing activity over all
760 five temporal frequencies (mean $\Delta F/F$, averaged over 10 repeats). Cells are ranked in descending order of
761 total activity. Scale bar: 10 s.
762 (F) Percentages of labeled cells that were visually tuned (see Methods), from 6 different FOVs in 3 mice
763 imaged over 14 weeks. Connected sets of dots in a given color indicate data from a single mouse (data
764 from 2 FOVs are shown per mouse).
765 (G) Comparison of the percentages of labeled cells that were visually tuned at 2 weeks and 14 weeks. The
766 percentages increased moderately but significantly between the two timepoints, from 60% to 68% (paired
767 two-sample t-test, $p = 0.0178$, $n = 6$).
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SUPPLEMENTAL INFORMATION

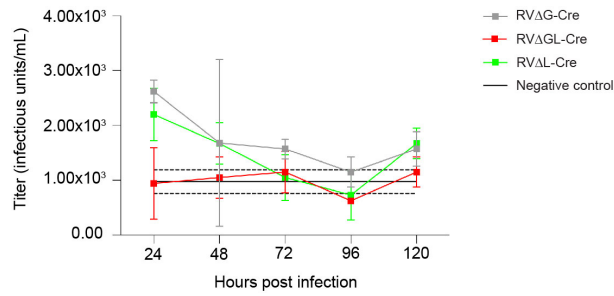


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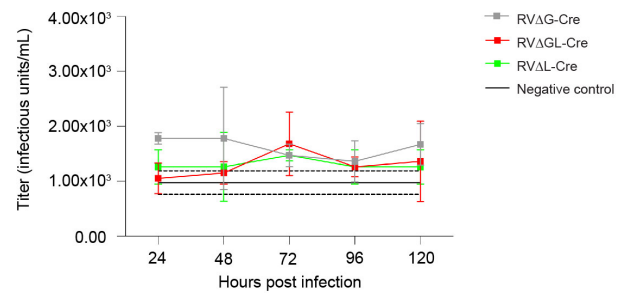
Figure S1. Δ L and Δ GL viruses express EGFP at similarly low levels, Related to Figure 1

Confocal images and flow cytometric histograms showing native and immunostained EGFP signal in uninfected cells (A) and cells infected with first-generation (Δ G) virus (B), second-generation (Δ GL) virus (C), or third-generation virus (D) expressing EGFP. Scale bar: 50 μ m, applies to all images.

A Multi-step growth curves (MOI=0.01)



B Single-step growth curves (MOI=1)



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782 Figure S2. ΔG , ΔGL , and ΔL viruses do not propagate in non-complementing cells, Related to Figure
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784 Viral titers in supernatants of BHK-21 cells not expressing any rabies viral genes, infected with ΔL , ΔGL , or
785 ΔG viruses at a multiplicity of infection (MOI) of 0.01 ("multi-step growth curves", panel A) or 1 ("single-
786 step" growth curves, panel B), with supernatants collected every 24 hours for five days. Graphs show mean
787 \pm s.e.m. .Black lines show negative control "titers" calculated from uninfected reporter cells (mean \pm s.e.m.
788 of 10 samples). Note that the titers in these graphs are 3-4 orders of magnitude lower than those obtained
789 on complementing cells (Figure 1).
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File S1. Titers and statistics for growth dynamics experiments, Related to Figure 1

See following pages.

FACS data from Single-step growth curves (Infectious units/mL)	24hr	48hr	72hr	96hr	120hr
(1) RVΔGL-Cre (on B19L-G_01 cells)	1.89E+05	1.99E+06	2.39E+06	1.76E+06	1.94E+06
(2) RVΔGL-Cre (on B19L-G_01 cells)	1.48E+05	1.97E+06	2.46E+06	1.77E+06	1.92E+06
(3) RVΔGL-Cre (on B19L-G_01 cells)	1.55E+05	1.89E+06	2.25E+06	1.55E+06	1.73E+06
(7) RVΔL-Cre (on B19L_02 cells)	8.60E+05	1.84E+07	2.60E+07	2.71E+07	2.10E+07
(8) RVΔL-Cre (on B19L_02 cells)	7.05E+05	1.35E+07	2.11E+07	2.54E+07	1.92E+07
(9) RVΔL-Cre (on B19L_02 cells)	1.17E+06	1.65E+07	2.25E+07	2.56E+07	1.70E+07
(13) RVΔL-Cre (on B19L-G_01 cells)	1.31E+06	1.92E+07	2.02E+07	1.87E+07	1.79E+07
(14) RVΔL-Cre (on B19L-G_01 cells)	1.88E+06	1.96E+07	1.92E+07	1.67E+07	1.79E+07
(15) RVΔL-Cre (on B19L-G_01 cells)	1.61E+06	1.89E+07	1.94E+07	1.51E+07	1.70E+07
(19) RVΔG-Cre (on B19G3-2 cells)	2.13E+06	3.12E+07	3.22E+07	3.02E+07	2.71E+07
(20) RVΔG-Cre (on B19G3-2 cells)	1.89E+06	3.22E+07	3.24E+07	2.63E+07	2.92E+07
(21) RVΔG-Cre (on B19G3-2 cells)	1.39E+06	2.19E+07	2.64E+07	2.01E+07	2.35E+07
(25) RVΔG-Cre (on B19L-G_01 cells)	1.71E+05	2.04E+06	1.14E+07	1.36E+07	1.37E+07
(26) RVΔG-Cre (on B19L-G_01 cells)	1.84E+05	2.40E+06	1.09E+07	1.18E+07	1.36E+07
(27) RVΔG-Cre (on B19L-G_01 cells)	1.65E+05	2.01E+06	9.61E+06	1.02E+07	1.14E+07

FACS data from Multi-step growth curves (Infectious units/mL)	24hr	48hr	72hr	96hr	120hr
(4) RVΔGL-Cre (on B19L-G_01 cells)	6.22E+03	4.36E+04	5.17E+05	7.09E+05	1.21E+06
(5) RVΔGL-Cre (on B19L-G_01 cells)	7.77E+03	4.39E+04	4.47E+05	7.78E+05	9.81E+05
(6) RVΔGL-Cre (on B19L-G_01 cells)	5.91E+03	6.31E+04	4.30E+05	6.82E+05	9.70E+05
(10) RVΔL-Cre (on B19L_02 cells)	6.84E+03	5.48E+05	5.50E+06	9.25E+06	1.26E+07
(11) RVΔL-Cre (on B19L_02 cells)	5.28E+03	2.96E+05	4.67E+06	9.18E+06	1.34E+07
(12) RVΔL-Cre (on B19L_02 cells)	5.59E+03	6.57E+05	6.63E+06	1.07E+07	1.34E+07
(16) RVΔL-Cre (on B19L-G_01 cells)	8.71E+03	1.31E+06	7.69E+06	5.80E+06	6.84E+06
(17) RVΔL-Cre (on B19L-G_01 cells)	4.04E+03	1.38E+06	6.40E+06	6.33E+06	6.57E+06
(18) RVΔL-Cre (on B19L-G_01 cells)	5.91E+03	1.11E+06	5.39E+06	5.40E+06	6.11E+06
(22) RVΔG-Cre (on B19G3-2 cells)	3.31E+04	1.70E+06	7.96E+06	8.63E+06	7.39E+06
(23) RVΔG-Cre (on B19G3-2 cells)	2.24E+04	6.82E+05	4.13E+06	5.36E+06	4.99E+06
(24) RVΔG-Cre (on B19G3-2 cells)	2.43E+04	8.56E+05	4.41E+06	6.52E+06	5.61E+06
(28) RVΔG-Cre (on B19L-G_01 cells)	6.84E+03	6.25E+04	5.56E+05	2.23E+06	5.91E+06
(29) RVΔG-Cre (on B19L-G_01 cells)	5.28E+03	3.23E+04	3.33E+05	1.27E+06	4.70E+06
(30) RVΔG-Cre (on B19L-G_01 cells)	8.40E+03	4.24E+04	4.88E+05	1.67E+06	5.21E+06

Average titers for Single-step growth curves (Infectious units/mL)	24hr	48hr	72hr	96hr	120hr
(1-3) RVΔGL-Cre (on B19L-G_01 cells)	1.64E+05	1.95E+06	2.37E+06	1.69E+06	1.86E+06
(7-9) RVΔL-Cre (on B19L_02 cells)	9.12E+05	1.61E+07	2.32E+07	2.60E+07	1.91E+07
(13-15) RVΔL-Cre (on B19L-G_01 cells)	1.60E+06	1.92E+07	1.96E+07	1.68E+07	1.76E+07
(19-21) RVΔG-Cre (on B19G3-2 cells)	1.81E+06	2.85E+07	3.03E+07	2.55E+07	2.66E+07
(25-27) RVΔG-Cre (on B19L-G_01 cells)	1.73E+05	2.15E+06	1.06E+07	1.19E+07	1.29E+07

Standard error of the mean for Single-step growth curves (Infectious units/mL)	24hr	48hr	72hr	96hr	120hr
(1-3) RVΔGL-Cre (on B19L-G_01 cells)	1.27E+04	3.12E+04	6.11E+04	7.25E+04	6.73E+04
(7-9) RVΔL-Cre (on B19L_02 cells)	1.37E+05	1.42E+06	1.45E+06	5.43E+05	1.16E+06
(13-15) RVΔL-Cre (on B19L-G_01 cells)	1.64E+05	2.12E+05	3.21E+05	1.04E+06	3.02E+05
(19-21) RVΔG-Cre (on B19G3-2 cells)	2.18E+05	3.29E+06	1.99E+06	2.94E+06	1.66E+06
(25-27) RVΔG-Cre (on B19L-G_01 cells)	5.71E+03	1.24E+05	5.38E+05	9.85E+05	7.57E+05

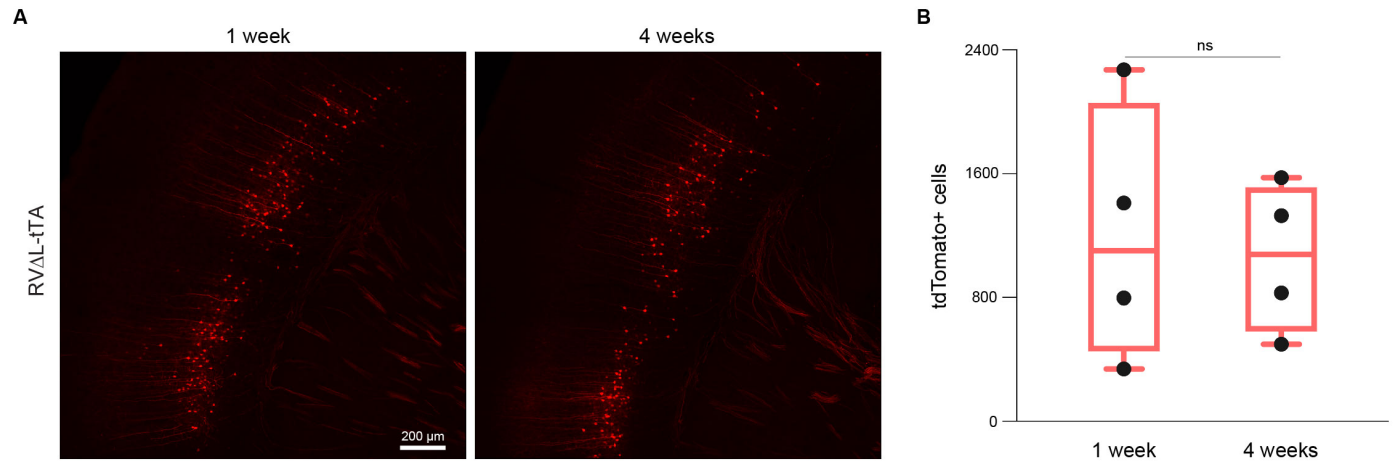
Average titers for Multi-step growth curves (Infectious units/mL)	24hr	48hr	72hr	96hr	120hr
(4-6) RVΔGL-Cre (on B19L-G_01 cells)	6.63E+03	5.02E+04	4.65E+05	7.23E+05	1.05E+06
(10-12) RVΔL-Cre (on B19L_02 cells)	5.91E+03	5.00E+05	5.60E+06	9.72E+06	1.31E+07
(16-18) RVΔL-Cre (on B19L-G_01 cells)	6.22E+03	1.27E+06	6.49E+06	5.84E+06	6.51E+06
(22-24) RVΔG-Cre (on B19G3-2 cells)	2.66E+04	1.08E+06	5.50E+06	6.83E+06	6.00E+06
(28-30) RVΔG-Cre (on B19L-G_01 cells)	6.84E+03	4.57E+04	4.59E+05	1.72E+06	5.27E+06

Standard error of the mean for Multi-step growth curves	24hr	48hr	72hr	96hr	120hr
(4-6) RVΔGL-Cre (on B19L-G_01 cells)	5.78E+02	6.45E+03	2.69E+04	2.87E+04	7.74E+04
(10-12) RVΔL-Cre (on B19L_02 cells)	4.75E+02	1.07E+05	5.70E+05	5.06E+05	2.77E+05
(16-18) RVΔL-Cre (on B19L-G_01 cells)	1.36E+03	8.07E+04	6.63E+05	2.71E+05	2.13E+05
(22-24) RVΔG-Cre (on B19G3-2 cells)	3.28E+03	3.16E+05	1.23E+06	9.57E+05	7.18E+05
(28-30) RVΔG-Cre (on B19L-G_01 cells)	8.98E+02	8.87E+03	6.60E+04	2.80E+05	3.51E+05

Conditions with the highest average titers for Single-step growth curves	Infectious units/mL
(1-3) RVΔGL-Cre (on B19L-G_01 cells): 72hr	2.39E+06
	2.46E+06
	2.25E+06
(7-9) RVΔL-Cre (on B19L_02 cells): 96hr	2.71E+07
	2.54E+07
	2.56E+07
(19-21) RVΔG-Cre (on B19G3-2 cells): 72hr	3.22E+07
	3.24E+07
	2.64E+07

Column1	Column2
average "titer" of BC-FLEX cells:	
3.63E+03	
Individual BC-FLEX negative "titers"	
NEG_01	5.28E+03
NEG_02	5.91E+03
NEG_03	3.11E+03
NEG_04	2.80E+03
NEG_05	2.17E+03
NEG_06	2.49E+03

Conditions with the highest average titers for Multi-step growth curves	Infectious units/mL
(4-6) RVΔGL-Cre (on B19L-G_01 cells): 120hr	1.21E+06
	9.81E+05
	9.70E+05
(10-12) RVΔL-Cre (on B19L_02 cells): 120hr	1.26E+07
	1.34E+07
	1.34E+07
(22-24) RVΔG-Cre (on B19G3-2 cells): 96hr	8.63E+06
	5.36E+06
	6.52E+06



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Figure S3. Retrograde targeting with third-generation (Δ L) rabies virus expressing the tetracycline transactivator, Related to Figure 2

(A) Corticothalamic neurons retrogradely labeled by a Δ L virus expressing tTA injected in the somatosensory thalamus of Ai63 reporter mice (Daigle et al., 2018) (tdTomato driven by TRE-tight) 1 week (left image) or 4 weeks (right image) prior to perfusion. Scale bar: 200 μ m, applies to both images.

(B) Counts of total labeled cortical neurons across every sixth section (see Methods) of each mouse brain. Numbers are not significantly different between the two time points (single factor ANOVA, $p = 0.772$, $n = 4$ mice per group).

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File S2. Cell counts and statistics for retrograde targeting experiments, Related to Figure 2

See following pages.

Condition	Mouse number	Section number	Cell counts
RVΔGL-Flpo: 1 week	21021505LJ	10	5
RVΔGL-Flpo: 1 week	21021505LJ	11	8
RVΔGL-Flpo: 1 week	21021505LJ	12	9
RVΔGL-Flpo: 1 week	21021505LJ	total:	22
RVΔGL-Flpo: 1 week	21021506LJ	9	5
RVΔGL-Flpo: 1 week	21021506LJ	10	6
RVΔGL-Flpo: 1 week	21021506LJ	11	4
RVΔGL-Flpo: 1 week	21021506LJ	12	3
RVΔGL-Flpo: 1 week	21021506LJ	total:	18
RVΔGL-Flpo: 1 week	21021507LJ	5	1
RVΔGL-Flpo: 1 week	21021507LJ	6	27
RVΔGL-Flpo: 1 week	21021507LJ	7	29
RVΔGL-Flpo: 1 week	21021507LJ	8	25
RVΔGL-Flpo: 1 week	21021507LJ	9	20
RVΔGL-Flpo: 1 week	21021507LJ	10	10
RVΔGL-Flpo: 1 week	21021507LJ	11	4
RVΔGL-Flpo: 1 week	21021507LJ	12	3
RVΔGL-Flpo: 1 week	21021507LJ	13	2
RVΔGL-Flpo: 1 week	21021507LJ	total:	121
RVΔGL-Flpo: 1 week	21021508LJ	11	1
RVΔGL-Flpo: 1 week	21021508LJ	15	1
RVΔGL-Flpo: 1 week	21021508LJ	total:	2
RVΔGL-Flpo: 1 week	21052505LJ	9	1
RVΔGL-Flpo: 1 week	21052505LJ	10	5
RVΔGL-Flpo: 1 week	21052505LJ	11	1
RVΔGL-Flpo: 1 week	21052505LJ	total:	7
RVΔGL-Flpo: 1 week	21052506LJ	11	4
RVΔGL-Flpo: 1 week	21052506LJ	12	4
RVΔGL-Flpo: 1 week	21052506LJ	total:	8
RVΔGL-Flpo: 1 week	21052507LJ	6	2
RVΔGL-Flpo: 1 week	21052507LJ	7	20
RVΔGL-Flpo: 1 week	21052507LJ	8	45
RVΔGL-Flpo: 1 week	21052507LJ	9	26
RVΔGL-Flpo: 1 week	21052507LJ	10	5
RVΔGL-Flpo: 1 week	21052507LJ	total:	98
RVΔGL-Flpo: 1 week	21052508LJ	8	1
RVΔGL-Flpo: 1 week	21052508LJ	10	1
RVΔGL-Flpo: 1 week	21052508LJ	11	1
RVΔGL-Flpo: 1 week	21052508LJ	12	1
RVΔGL-Flpo: 1 week	21052508LJ	14	2
RVΔGL-Flpo: 1 week	21052508LJ	total:	6
RVΔL-Flpo: 1 week	21021805LJ	1	57
RVΔL-Flpo: 1 week	21021805LJ	2	20
RVΔL-Flpo: 1 week	21021805LJ	3	11
RVΔL-Flpo: 1 week	21021805LJ	4	115
RVΔL-Flpo: 1 week	21021805LJ	5	270
RVΔL-Flpo: 1 week	21021805LJ	6	399
RVΔL-Flpo: 1 week	21021805LJ	7	428
RVΔL-Flpo: 1 week	21021805LJ	8	542
RVΔL-Flpo: 1 week	21021805LJ	9	416
RVΔL-Flpo: 1 week	21021805LJ	10	368
RVΔL-Flpo: 1 week	21021805LJ	11	240
RVΔL-Flpo: 1 week	21021805LJ	12	106
RVΔL-Flpo: 1 week	21021805LJ	13	42

RVΔL-Flpo: 1 week	21021805LJ	14	52
RVΔL-Flpo: 1 week	21021805LJ	15	27
RVΔL-Flpo: 1 week	21021805LJ	total:	3093
RVΔL-Flpo: 1 week	21021806LJ	1	1
RVΔL-Flpo: 1 week	21021806LJ	4	5
RVΔL-Flpo: 1 week	21021806LJ	5	3
RVΔL-Flpo: 1 week	21021806LJ	6	4
RVΔL-Flpo: 1 week	21021806LJ	7	16
RVΔL-Flpo: 1 week	21021806LJ	8	35
RVΔL-Flpo: 1 week	21021806LJ	9	93
RVΔL-Flpo: 1 week	21021806LJ	10	74
RVΔL-Flpo: 1 week	21021806LJ	11	40
RVΔL-Flpo: 1 week	21021806LJ	12	24
RVΔL-Flpo: 1 week	21021806LJ	13	11
RVΔL-Flpo: 1 week	21021806LJ	14	7
RVΔL-Flpo: 1 week	21021806LJ	15	7
RVΔL-Flpo: 1 week	21021806LJ	total:	320
RVΔL-Flpo: 1 week	21021807LJ	5	1
RVΔL-Flpo: 1 week	21021807LJ	6	2
RVΔL-Flpo: 1 week	21021807LJ	7	6
RVΔL-Flpo: 1 week	21021807LJ	8	8
RVΔL-Flpo: 1 week	21021807LJ	9	35
RVΔL-Flpo: 1 week	21021807LJ	10	64
RVΔL-Flpo: 1 week	21021807LJ	11	80
RVΔL-Flpo: 1 week	21021807LJ	12	55
RVΔL-Flpo: 1 week	21021807LJ	13	34
RVΔL-Flpo: 1 week	21021807LJ	14	13
RVΔL-Flpo: 1 week	21021807LJ	15	13
RVΔL-Flpo: 1 week	21021807LJ	total:	311
RVΔL-Flpo: 1 week	21021808LJ	6	7
RVΔL-Flpo: 1 week	21021808LJ	7	13
RVΔL-Flpo: 1 week	21021808LJ	8	48
RVΔL-Flpo: 1 week	21021808LJ	9	74
RVΔL-Flpo: 1 week	21021808LJ	10	57
RVΔL-Flpo: 1 week	21021808LJ	11	20
RVΔL-Flpo: 1 week	21021808LJ	12	13
RVΔL-Flpo: 1 week	21021808LJ	13	13
RVΔL-Flpo: 1 week	21021808LJ	14	14
RVΔL-Flpo: 1 week	21021808LJ	15	4
RVΔL-Flpo: 1 week	21021808LJ	total:	263
RVΔL-Flpo: 1 week	21052607LJ	4	3
RVΔL-Flpo: 1 week	21052607LJ	5	2
RVΔL-Flpo: 1 week	21052607LJ	6	4
RVΔL-Flpo: 1 week	21052607LJ	7	11
RVΔL-Flpo: 1 week	21052607LJ	8	10
RVΔL-Flpo: 1 week	21052607LJ	9	4
RVΔL-Flpo: 1 week	21052607LJ	11	6
RVΔL-Flpo: 1 week	21052607LJ	12	7
RVΔL-Flpo: 1 week	21052607LJ	total:	47
RVΔL-Flpo: 1 week	21052608LJ	1	105
RVΔL-Flpo: 1 week	21052608LJ	2	116
RVΔL-Flpo: 1 week	21052608LJ	3	110
RVΔL-Flpo: 1 week	21052608LJ	4	192
RVΔL-Flpo: 1 week	21052608LJ	5	395
RVΔL-Flpo: 1 week	21052608LJ	6	389

RVΔL-Flpo: 1 week	21052608LJ	7	421
RVΔL-Flpo: 1 week	21052608LJ	8	340
RVΔL-Flpo: 1 week	21052608LJ	9	72
RVΔL-Flpo: 1 week	21052608LJ	10	14
RVΔL-Flpo: 1 week	21052608LJ	11	5
RVΔL-Flpo: 1 week	21052608LJ	12	2
RVΔL-Flpo: 1 week	21052608LJ	total:	2161
RVΔL-Flpo: 1 week	21060302LJ	1	8
RVΔL-Flpo: 1 week	21060302LJ	2	10
RVΔL-Flpo: 1 week	21060302LJ	3	17
RVΔL-Flpo: 1 week	21060302LJ	4	24
RVΔL-Flpo: 1 week	21060302LJ	5	22
RVΔL-Flpo: 1 week	21060302LJ	6	29
RVΔL-Flpo: 1 week	21060302LJ	7	34
RVΔL-Flpo: 1 week	21060302LJ	8	59
RVΔL-Flpo: 1 week	21060302LJ	9	78
RVΔL-Flpo: 1 week	21060302LJ	10	46
RVΔL-Flpo: 1 week	21060302LJ	11	16
RVΔL-Flpo: 1 week	21060302LJ	12	12
RVΔL-Flpo: 1 week	21060302LJ	13	6
RVΔL-Flpo: 1 week	21060302LJ	14	2
RVΔL-Flpo: 1 week	21060302LJ	15	2
RVΔL-Flpo: 1 week	21060302LJ	total:	365
RVΔL-Flpo: 1 week	21060303LJ	1	7
RVΔL-Flpo: 1 week	21060303LJ	2	9
RVΔL-Flpo: 1 week	21060303LJ	3	10
RVΔL-Flpo: 1 week	21060303LJ	4	16
RVΔL-Flpo: 1 week	21060303LJ	5	22
RVΔL-Flpo: 1 week	21060303LJ	6	27
RVΔL-Flpo: 1 week	21060303LJ	7	25
RVΔL-Flpo: 1 week	21060303LJ	8	27
RVΔL-Flpo: 1 week	21060303LJ	9	36
RVΔL-Flpo: 1 week	21060303LJ	10	30
RVΔL-Flpo: 1 week	21060303LJ	11	15
RVΔL-Flpo: 1 week	21060303LJ	12	7
RVΔL-Flpo: 1 week	21060303LJ	14	6
RVΔL-Flpo: 1 week	21060303LJ	15	1
RVΔL-Flpo: 1 week	21060303LJ	total:	238
RVΔGL-Flpo: 4 weeks	21021501LJ	7	2
RVΔGL-Flpo: 4 weeks	21021501LJ	8	12
RVΔGL-Flpo: 4 weeks	21021501LJ	9	25
RVΔGL-Flpo: 4 weeks	21021501LJ	10	26
RVΔGL-Flpo: 4 weeks	21021501LJ	11	7
RVΔGL-Flpo: 4 weeks	21021501LJ	12	3
RVΔGL-Flpo: 4 weeks	21021501LJ	13	3
RVΔGL-Flpo: 4 weeks	21021501LJ	14	1
RVΔGL-Flpo: 4 weeks	21021501LJ	15	1
RVΔGL-Flpo: 4 weeks	21021501LJ	total:	80
RVΔGL-Flpo: 4 weeks	21021502LJ	9	2
RVΔGL-Flpo: 4 weeks	21021502LJ	10	25
RVΔGL-Flpo: 4 weeks	21021502LJ	11	26
RVΔGL-Flpo: 4 weeks	21021502LJ	12	5
RVΔGL-Flpo: 4 weeks	21021502LJ	13	4
RVΔGL-Flpo: 4 weeks	21021502LJ	total:	62
RVΔGL-Flpo: 4 weeks	21021503LJ	1	88

RVΔGL-Flpo: 4 weeks	21021503LJ	2	129
RVΔGL-Flpo: 4 weeks	21021503LJ	3	199
RVΔGL-Flpo: 4 weeks	21021503LJ	4	254
RVΔGL-Flpo: 4 weeks	21021503LJ	5	312
RVΔGL-Flpo: 4 weeks	21021503LJ	6	396
RVΔGL-Flpo: 4 weeks	21021503LJ	7	351
RVΔGL-Flpo: 4 weeks	21021503LJ	8	391
RVΔGL-Flpo: 4 weeks	21021503LJ	9	359
RVΔGL-Flpo: 4 weeks	21021503LJ	10	295
RVΔGL-Flpo: 4 weeks	21021503LJ	11	133
RVΔGL-Flpo: 4 weeks	21021503LJ	12	216
RVΔGL-Flpo: 4 weeks	21021503LJ	13	59
RVΔGL-Flpo: 4 weeks	21021503LJ	14	40
RVΔGL-Flpo: 4 weeks	21021503LJ	15	32
RVΔGL-Flpo: 4 weeks	21021503LJ	total:	3254
RVΔGL-Flpo: 4 weeks	21021504LJ	3	1
RVΔGL-Flpo: 4 weeks	21021504LJ	4	4
RVΔGL-Flpo: 4 weeks	21021504LJ	5	22
RVΔGL-Flpo: 4 weeks	21021504LJ	6	24
RVΔGL-Flpo: 4 weeks	21021504LJ	7	56
RVΔGL-Flpo: 4 weeks	21021504LJ	8	78
RVΔGL-Flpo: 4 weeks	21021504LJ	9	32
RVΔGL-Flpo: 4 weeks	21021504LJ	10	18
RVΔGL-Flpo: 4 weeks	21021504LJ	11	1
RVΔGL-Flpo: 4 weeks	21021504LJ	12	1
RVΔGL-Flpo: 4 weeks	21021504LJ	total:	237
RVΔGL-Flpo: 4 weeks	21052501LJ	2	1
RVΔGL-Flpo: 4 weeks	21052501LJ	4	1
RVΔGL-Flpo: 4 weeks	21052501LJ	7	2
RVΔGL-Flpo: 4 weeks	21052501LJ	8	6
RVΔGL-Flpo: 4 weeks	21052501LJ	9	9
RVΔGL-Flpo: 4 weeks	21052501LJ	10	9
RVΔGL-Flpo: 4 weeks	21052501LJ	11	2
RVΔGL-Flpo: 4 weeks	21052501LJ	12	6
RVΔGL-Flpo: 4 weeks	21052501LJ	13	4
RVΔGL-Flpo: 4 weeks	21052501LJ	14	1
RVΔGL-Flpo: 4 weeks	21052501LJ	15	2
RVΔGL-Flpo: 4 weeks	21052501LJ	total:	43
RVΔGL-Flpo: 4 weeks	21052502LJ	6	2
RVΔGL-Flpo: 4 weeks	21052502LJ	7	1
RVΔGL-Flpo: 4 weeks	21052502LJ	8	4
RVΔGL-Flpo: 4 weeks	21052502LJ	11	1
RVΔGL-Flpo: 4 weeks	21052502LJ	14	3
RVΔGL-Flpo: 4 weeks	21052502LJ	15	3
RVΔGL-Flpo: 4 weeks	21052502LJ	total:	14
RVΔGL-Flpo: 4 weeks	21052503LJ	6	2
RVΔGL-Flpo: 4 weeks	21052503LJ	8	1
RVΔGL-Flpo: 4 weeks	21052503LJ	9	9
RVΔGL-Flpo: 4 weeks	21052503LJ	10	45
RVΔGL-Flpo: 4 weeks	21052503LJ	11	33
RVΔGL-Flpo: 4 weeks	21052503LJ	12	17
RVΔGL-Flpo: 4 weeks	21052503LJ	13	10
RVΔGL-Flpo: 4 weeks	21052503LJ	14	7
RVΔGL-Flpo: 4 weeks	21052503LJ	15	6
RVΔGL-Flpo: 4 weeks	21052503LJ	total:	130

RVΔGL-Flpo: 4 weeks	21052504LJ	7	2
RVΔGL-Flpo: 4 weeks	21052504LJ	8	13
RVΔGL-Flpo: 4 weeks	21052504LJ	9	18
RVΔGL-Flpo: 4 weeks	21052504LJ	10	16
RVΔGL-Flpo: 4 weeks	21052504LJ	11	8
RVΔGL-Flpo: 4 weeks	21052504LJ	12	3
RVΔGL-Flpo: 4 weeks	21052504LJ	13	3
RVΔGL-Flpo: 4 weeks	21052504LJ	14	1
RVΔGL-Flpo: 4 weeks	21052504LJ	15	1
RVΔGL-Flpo: 4 weeks	21052504LJ	total:	65
RVΔL-Flpo: 4 weeks	21021801LJ	1	36
RVΔL-Flpo: 4 weeks	21021801LJ	2	18
RVΔL-Flpo: 4 weeks	21021801LJ	3	2
RVΔL-Flpo: 4 weeks	21021801LJ	4	28
RVΔL-Flpo: 4 weeks	21021801LJ	5	180
RVΔL-Flpo: 4 weeks	21021801LJ	6	415
RVΔL-Flpo: 4 weeks	21021801LJ	7	622
RVΔL-Flpo: 4 weeks	21021801LJ	8	649
RVΔL-Flpo: 4 weeks	21021801LJ	9	739
RVΔL-Flpo: 4 weeks	21021801LJ	10	577
RVΔL-Flpo: 4 weeks	21021801LJ	11	431
RVΔL-Flpo: 4 weeks	21021801LJ	12	211
RVΔL-Flpo: 4 weeks	21021801LJ	13	166
RVΔL-Flpo: 4 weeks	21021801LJ	14	170
RVΔL-Flpo: 4 weeks	21021801LJ	15	370
RVΔL-Flpo: 4 weeks	21021801LJ	total:	4614
RVΔL-Flpo: 4 weeks	21021802LJ	1	3
RVΔL-Flpo: 4 weeks	21021802LJ	2	6
RVΔL-Flpo: 4 weeks	21021802LJ	3	8
RVΔL-Flpo: 4 weeks	21021802LJ	5	32
RVΔL-Flpo: 4 weeks	21021802LJ	6	224
RVΔL-Flpo: 4 weeks	21021802LJ	7	279
RVΔL-Flpo: 4 weeks	21021802LJ	8	287
RVΔL-Flpo: 4 weeks	21021802LJ	9	457
RVΔL-Flpo: 4 weeks	21021802LJ	10	501
RVΔL-Flpo: 4 weeks	21021802LJ	11	491
RVΔL-Flpo: 4 weeks	21021802LJ	12	211
RVΔL-Flpo: 4 weeks	21021802LJ	13	287
RVΔL-Flpo: 4 weeks	21021802LJ	14	220
RVΔL-Flpo: 4 weeks	21021802LJ	total:	3006
RVΔL-Flpo: 4 weeks	21021803LJ	2	6
RVΔL-Flpo: 4 weeks	21021803LJ	3	9
RVΔL-Flpo: 4 weeks	21021803LJ	4	48
RVΔL-Flpo: 4 weeks	21021803LJ	5	177
RVΔL-Flpo: 4 weeks	21021803LJ	6	247
RVΔL-Flpo: 4 weeks	21021803LJ	7	339
RVΔL-Flpo: 4 weeks	21021803LJ	8	316
RVΔL-Flpo: 4 weeks	21021803LJ	9	322
RVΔL-Flpo: 4 weeks	21021803LJ	10	205
RVΔL-Flpo: 4 weeks	21021803LJ	11	78
RVΔL-Flpo: 4 weeks	21021803LJ	12	46
RVΔL-Flpo: 4 weeks	21021803LJ	13	41
RVΔL-Flpo: 4 weeks	21021803LJ	14	13
RVΔL-Flpo: 4 weeks	21021803LJ	total:	1847
RVΔL-Flpo: 4 weeks	21021804LJ	1	6

RVΔL-Flpo: 4 weeks	21021804LJ	2	14
RVΔL-Flpo: 4 weeks	21021804LJ	3	2
RVΔL-Flpo: 4 weeks	21021804LJ	4	57
RVΔL-Flpo: 4 weeks	21021804LJ	5	359
RVΔL-Flpo: 4 weeks	21021804LJ	6	414
RVΔL-Flpo: 4 weeks	21021804LJ	7	469
RVΔL-Flpo: 4 weeks	21021804LJ	8	576
RVΔL-Flpo: 4 weeks	21021804LJ	9	533
RVΔL-Flpo: 4 weeks	21021804LJ	10	383
RVΔL-Flpo: 4 weeks	21021804LJ	11	283
RVΔL-Flpo: 4 weeks	21021804LJ	12	114
RVΔL-Flpo: 4 weeks	21021804LJ	13	34
RVΔL-Flpo: 4 weeks	21021804LJ	14	87
RVΔL-Flpo: 4 weeks	21021804LJ	total:	3331
RVΔL-Flpo: 4 weeks	21052601LJ	2	1
RVΔL-Flpo: 4 weeks	21052601LJ	3	1
RVΔL-Flpo: 4 weeks	21052601LJ	5	16
RVΔL-Flpo: 4 weeks	21052601LJ	6	10
RVΔL-Flpo: 4 weeks	21052601LJ	7	28
RVΔL-Flpo: 4 weeks	21052601LJ	8	92
RVΔL-Flpo: 4 weeks	21052601LJ	9	161
RVΔL-Flpo: 4 weeks	21052601LJ	10	232
RVΔL-Flpo: 4 weeks	21052601LJ	11	280
RVΔL-Flpo: 4 weeks	21052601LJ	12	201
RVΔL-Flpo: 4 weeks	21052601LJ	13	143
RVΔL-Flpo: 4 weeks	21052601LJ	14	106
RVΔL-Flpo: 4 weeks	21052601LJ	15	98
RVΔL-Flpo: 4 weeks	21052601LJ	total:	1369
RVΔL-Flpo: 4 weeks	21052602LJ	1	49
RVΔL-Flpo: 4 weeks	21052602LJ	2	12
RVΔL-Flpo: 4 weeks	21052602LJ	3	17
RVΔL-Flpo: 4 weeks	21052602LJ	4	51
RVΔL-Flpo: 4 weeks	21052602LJ	5	163
RVΔL-Flpo: 4 weeks	21052602LJ	6	418
RVΔL-Flpo: 4 weeks	21052602LJ	7	489
RVΔL-Flpo: 4 weeks	21052602LJ	8	526
RVΔL-Flpo: 4 weeks	21052602LJ	9	510
RVΔL-Flpo: 4 weeks	21052602LJ	10	319
RVΔL-Flpo: 4 weeks	21052602LJ	11	179
RVΔL-Flpo: 4 weeks	21052602LJ	12	72
RVΔL-Flpo: 4 weeks	21052602LJ	13	48
RVΔL-Flpo: 4 weeks	21052602LJ	14	32
RVΔL-Flpo: 4 weeks	21052602LJ	15	34
RVΔL-Flpo: 4 weeks	21052602LJ	total:	2919
RVΔL-Flpo: 4 weeks	21052603LJ	1	38
RVΔL-Flpo: 4 weeks	21052603LJ	2	33
RVΔL-Flpo: 4 weeks	21052603LJ	3	16
RVΔL-Flpo: 4 weeks	21052603LJ	4	44
RVΔL-Flpo: 4 weeks	21052603LJ	5	302
RVΔL-Flpo: 4 weeks	21052603LJ	6	648
RVΔL-Flpo: 4 weeks	21052603LJ	7	636
RVΔL-Flpo: 4 weeks	21052603LJ	8	666
RVΔL-Flpo: 4 weeks	21052603LJ	9	582
RVΔL-Flpo: 4 weeks	21052603LJ	10	409
RVΔL-Flpo: 4 weeks	21052603LJ	11	207

RVΔL-Flpo: 4 weeks	21052603LJ	12	79
RVΔL-Flpo: 4 weeks	21052603LJ	13	26
RVΔL-Flpo: 4 weeks	21052603LJ	14	25
RVΔL-Flpo: 4 weeks	21052603LJ	15	19
RVΔL-Flpo: 4 weeks	21052603LJ	total:	3730
RVΔL-Flpo: 4 weeks	21052606LJ	1	43
RVΔL-Flpo: 4 weeks	21052606LJ	2	23
RVΔL-Flpo: 4 weeks	21052606LJ	3	14
RVΔL-Flpo: 4 weeks	21052606LJ	4	32
RVΔL-Flpo: 4 weeks	21052606LJ	5	220
RVΔL-Flpo: 4 weeks	21052606LJ	6	503
RVΔL-Flpo: 4 weeks	21052606LJ	7	616
RVΔL-Flpo: 4 weeks	21052606LJ	8	622
RVΔL-Flpo: 4 weeks	21052606LJ	9	560
RVΔL-Flpo: 4 weeks	21052606LJ	10	403
RVΔL-Flpo: 4 weeks	21052606LJ	11	179
RVΔL-Flpo: 4 weeks	21052606LJ	12	82
RVΔL-Flpo: 4 weeks	21052606LJ	13	56
RVΔL-Flpo: 4 weeks	21052606LJ	14	68
RVΔL-Flpo: 4 weeks	21052606LJ	15	33
RVΔL-Flpo: 4 weeks	21052606LJ	total:	3454
RVΔGL-Cre: 1 week	21021703LJ	1	422
RVΔGL-Cre: 1 week	21021703LJ	2	395
RVΔGL-Cre: 1 week	21021703LJ	3	606
RVΔGL-Cre: 1 week	21021703LJ	4	543
RVΔGL-Cre: 1 week	21021703LJ	5	637
RVΔGL-Cre: 1 week	21021703LJ	6	848
RVΔGL-Cre: 1 week	21021703LJ	7	1145
RVΔGL-Cre: 1 week	21021703LJ	8	1183
RVΔGL-Cre: 1 week	21021703LJ	9	1292
RVΔGL-Cre: 1 week	21021703LJ	10	1311
RVΔGL-Cre: 1 week	21021703LJ	11	1377
RVΔGL-Cre: 1 week	21021703LJ	12	1251
RVΔGL-Cre: 1 week	21021703LJ	13	998
RVΔGL-Cre: 1 week	21021703LJ	14	835
RVΔGL-Cre: 1 week	21021703LJ	15	641
RVΔGL-Cre: 1 week	21021703LJ	total:	13484
RVΔGL-Cre: 1 week	21021704LJ	1	918
RVΔGL-Cre: 1 week	21021704LJ	2	975
RVΔGL-Cre: 1 week	21021704LJ	3	1144
RVΔGL-Cre: 1 week	21021704LJ	4	1404
RVΔGL-Cre: 1 week	21021704LJ	5	1491
RVΔGL-Cre: 1 week	21021704LJ	6	1377
RVΔGL-Cre: 1 week	21021704LJ	7	1231
RVΔGL-Cre: 1 week	21021704LJ	8	1337
RVΔGL-Cre: 1 week	21021704LJ	9	1135
RVΔGL-Cre: 1 week	21021704LJ	10	1058
RVΔGL-Cre: 1 week	21021704LJ	11	711
RVΔGL-Cre: 1 week	21021704LJ	12	568
RVΔGL-Cre: 1 week	21021704LJ	13	443
RVΔGL-Cre: 1 week	21021704LJ	14	432
RVΔGL-Cre: 1 week	21021704LJ	15	411
RVΔGL-Cre: 1 week	21021704LJ	total:	14635
RVΔGL-Cre: 1 week	21021707LJ	1	497
RVΔGL-Cre: 1 week	21021707LJ	2	568

RVΔGL-Cre: 1 week	21021707LJ	3	454
RVΔGL-Cre: 1 week	21021707LJ	4	454
RVΔGL-Cre: 1 week	21021707LJ	5	583
RVΔGL-Cre: 1 week	21021707LJ	6	873
RVΔGL-Cre: 1 week	21021707LJ	7	973
RVΔGL-Cre: 1 week	21021707LJ	8	1133
RVΔGL-Cre: 1 week	21021707LJ	9	1220
RVΔGL-Cre: 1 week	21021707LJ	10	1176
RVΔGL-Cre: 1 week	21021707LJ	11	994
RVΔGL-Cre: 1 week	21021707LJ	12	645
RVΔGL-Cre: 1 week	21021707LJ	13	467
RVΔGL-Cre: 1 week	21021707LJ	14	330
RVΔGL-Cre: 1 week	21021707LJ	15	280
RVΔGL-Cre: 1 week	21021707LJ	total:	10647
RVΔGL-Cre: 1 week	21021708LJ	1	271
RVΔGL-Cre: 1 week	21021708LJ	2	303
RVΔGL-Cre: 1 week	21021708LJ	3	386
RVΔGL-Cre: 1 week	21021708LJ	4	634
RVΔGL-Cre: 1 week	21021708LJ	5	805
RVΔGL-Cre: 1 week	21021708LJ	6	931
RVΔGL-Cre: 1 week	21021708LJ	7	1029
RVΔGL-Cre: 1 week	21021708LJ	8	1186
RVΔGL-Cre: 1 week	21021708LJ	9	1227
RVΔGL-Cre: 1 week	21021708LJ	10	1102
RVΔGL-Cre: 1 week	21021708LJ	11	1019
RVΔGL-Cre: 1 week	21021708LJ	12	666
RVΔGL-Cre: 1 week	21021708LJ	13	431
RVΔGL-Cre: 1 week	21021708LJ	14	369
RVΔGL-Cre: 1 week	21021708LJ	15	486
RVΔGL-Cre: 1 week	21021708LJ	total:	10845
RVΔL-Cre: 1 week	21021905LJ	1	1161
RVΔL-Cre: 1 week	21021905LJ	2	1093
RVΔL-Cre: 1 week	21021905LJ	3	1072
RVΔL-Cre: 1 week	21021905LJ	4	1079
RVΔL-Cre: 1 week	21021905LJ	5	1318
RVΔL-Cre: 1 week	21021905LJ	6	1417
RVΔL-Cre: 1 week	21021905LJ	7	1541
RVΔL-Cre: 1 week	21021905LJ	8	1510
RVΔL-Cre: 1 week	21021905LJ	9	1500
RVΔL-Cre: 1 week	21021905LJ	10	1549
RVΔL-Cre: 1 week	21021905LJ	11	1381
RVΔL-Cre: 1 week	21021905LJ	12	1109
RVΔL-Cre: 1 week	21021905LJ	13	532
RVΔL-Cre: 1 week	21021905LJ	14	380
RVΔL-Cre: 1 week	21021905LJ	15	265
RVΔL-Cre: 1 week	21021905LJ	total:	16907
RVΔL-Cre: 1 week	21021906LJ	1	985
RVΔL-Cre: 1 week	21021906LJ	2	1012
RVΔL-Cre: 1 week	21021906LJ	3	1020
RVΔL-Cre: 1 week	21021906LJ	4	1060
RVΔL-Cre: 1 week	21021906LJ	5	1082
RVΔL-Cre: 1 week	21021906LJ	6	1515
RVΔL-Cre: 1 week	21021906LJ	7	1808
RVΔL-Cre: 1 week	21021906LJ	8	1675
RVΔL-Cre: 1 week	21021906LJ	9	1538

RVΔL-Cre: 1 week	21021906LJ	10	1728
RVΔL-Cre: 1 week	21021906LJ	11	955
RVΔL-Cre: 1 week	21021906LJ	12	1018
RVΔL-Cre: 1 week	21021906LJ	13	993
RVΔL-Cre: 1 week	21021906LJ	14	701
RVΔL-Cre: 1 week	21021906LJ	15	483
RVΔL-Cre: 1 week	21021906LJ	16	350
RVΔL-Cre: 1 week	21021906LJ	total:	17923
RVΔL-Cre: 1 week	21021907LJ	1	1207
RVΔL-Cre: 1 week	21021907LJ	2	1083
RVΔL-Cre: 1 week	21021907LJ	3	1152
RVΔL-Cre: 1 week	21021907LJ	4	1225
RVΔL-Cre: 1 week	21021907LJ	5	1336
RVΔL-Cre: 1 week	21021907LJ	6	1558
RVΔL-Cre: 1 week	21021907LJ	7	1498
RVΔL-Cre: 1 week	21021907LJ	8	1709
RVΔL-Cre: 1 week	21021907LJ	9	1547
RVΔL-Cre: 1 week	21021907LJ	10	1718
RVΔL-Cre: 1 week	21021907LJ	11	1503
RVΔL-Cre: 1 week	21021907LJ	12	1249
RVΔL-Cre: 1 week	21021907LJ	13	845
RVΔL-Cre: 1 week	21021907LJ	14	626
RVΔL-Cre: 1 week	21021907LJ	15	495
RVΔL-Cre: 1 week	21021907LJ	total:	18751
RVΔL-Cre: 1 week	21021908LJ	1	1011
RVΔL-Cre: 1 week	21021908LJ	2	892
RVΔL-Cre: 1 week	21021908LJ	3	896
RVΔL-Cre: 1 week	21021908LJ	4	957
RVΔL-Cre: 1 week	21021908LJ	5	1243
RVΔL-Cre: 1 week	21021908LJ	6	1448
RVΔL-Cre: 1 week	21021908LJ	7	1568
RVΔL-Cre: 1 week	21021908LJ	8	1459
RVΔL-Cre: 1 week	21021908LJ	9	1573
RVΔL-Cre: 1 week	21021908LJ	10	1313
RVΔL-Cre: 1 week	21021908LJ	11	1146
RVΔL-Cre: 1 week	21021908LJ	12	1046
RVΔL-Cre: 1 week	21021908LJ	13	774
RVΔL-Cre: 1 week	21021908LJ	14	483
RVΔL-Cre: 1 week	21021908LJ	15	448
RVΔL-Cre: 1 week	21021908LJ	total:	16257
RVΔGL-Cre: 4 weeks	21021701LJ	1	1314
RVΔGL-Cre: 4 weeks	21021701LJ	2	1246
RVΔGL-Cre: 4 weeks	21021701LJ	3	1351
RVΔGL-Cre: 4 weeks	21021701LJ	4	1601
RVΔGL-Cre: 4 weeks	21021701LJ	5	1668
RVΔGL-Cre: 4 weeks	21021701LJ	6	1618
RVΔGL-Cre: 4 weeks	21021701LJ	7	1593
RVΔGL-Cre: 4 weeks	21021701LJ	8	1757
RVΔGL-Cre: 4 weeks	21021701LJ	9	1730
RVΔGL-Cre: 4 weeks	21021701LJ	10	1600
RVΔGL-Cre: 4 weeks	21021701LJ	11	1268
RVΔGL-Cre: 4 weeks	21021701LJ	12	931
RVΔGL-Cre: 4 weeks	21021701LJ	13	652
RVΔGL-Cre: 4 weeks	21021701LJ	14	491
RVΔGL-Cre: 4 weeks	21021701LJ	15	336

RVΔGL-Cre: 4 weeks	21021701LJ	total:	19156
RVΔGL-Cre: 4 weeks	21021702LJ	1	957
RVΔGL-Cre: 4 weeks	21021702LJ	2	1175
RVΔGL-Cre: 4 weeks	21021702LJ	3	1414
RVΔGL-Cre: 4 weeks	21021702LJ	4	1470
RVΔGL-Cre: 4 weeks	21021702LJ	5	1481
RVΔGL-Cre: 4 weeks	21021702LJ	6	1654
RVΔGL-Cre: 4 weeks	21021702LJ	7	1616
RVΔGL-Cre: 4 weeks	21021702LJ	8	1601
RVΔGL-Cre: 4 weeks	21021702LJ	9	1717
RVΔGL-Cre: 4 weeks	21021702LJ	10	1449
RVΔGL-Cre: 4 weeks	21021702LJ	11	1132
RVΔGL-Cre: 4 weeks	21021702LJ	12	871
RVΔGL-Cre: 4 weeks	21021702LJ	13	587
RVΔGL-Cre: 4 weeks	21021702LJ	14	456
RVΔGL-Cre: 4 weeks	21021702LJ	15	355
RVΔGL-Cre: 4 weeks	21021702LJ	total:	17935
RVΔGL-Cre: 4 weeks	21021705LJ	1	892
RVΔGL-Cre: 4 weeks	21021705LJ	2	824
RVΔGL-Cre: 4 weeks	21021705LJ	3	879
RVΔGL-Cre: 4 weeks	21021705LJ	4	1094
RVΔGL-Cre: 4 weeks	21021705LJ	5	1182
RVΔGL-Cre: 4 weeks	21021705LJ	6	1318
RVΔGL-Cre: 4 weeks	21021705LJ	7	1482
RVΔGL-Cre: 4 weeks	21021705LJ	8	1637
RVΔGL-Cre: 4 weeks	21021705LJ	9	1666
RVΔGL-Cre: 4 weeks	21021705LJ	10	1796
RVΔGL-Cre: 4 weeks	21021705LJ	11	1636
RVΔGL-Cre: 4 weeks	21021705LJ	12	1415
RVΔGL-Cre: 4 weeks	21021705LJ	13	1121
RVΔGL-Cre: 4 weeks	21021705LJ	14	894
RVΔGL-Cre: 4 weeks	21021705LJ	15	703
RVΔGL-Cre: 4 weeks	21021705LJ	total:	18539
RVΔGL-Cre: 4 weeks	21021706LJ	1	756
RVΔGL-Cre: 4 weeks	21021706LJ	2	817
RVΔGL-Cre: 4 weeks	21021706LJ	3	750
RVΔGL-Cre: 4 weeks	21021706LJ	4	798
RVΔGL-Cre: 4 weeks	21021706LJ	5	920
RVΔGL-Cre: 4 weeks	21021706LJ	6	1337
RVΔGL-Cre: 4 weeks	21021706LJ	7	1485
RVΔGL-Cre: 4 weeks	21021706LJ	8	1490
RVΔGL-Cre: 4 weeks	21021706LJ	9	1641
RVΔGL-Cre: 4 weeks	21021706LJ	10	2004
RVΔGL-Cre: 4 weeks	21021706LJ	11	1955
RVΔGL-Cre: 4 weeks	21021706LJ	12	1453
RVΔGL-Cre: 4 weeks	21021706LJ	13	1122
RVΔGL-Cre: 4 weeks	21021706LJ	14	769
RVΔGL-Cre: 4 weeks	21021706LJ	15	641
RVΔGL-Cre: 4 weeks	21021706LJ	total:	17938
RVΔL-Cre: 4 weeks	21021901LJ	1	943
RVΔL-Cre: 4 weeks	21021901LJ	2	1003
RVΔL-Cre: 4 weeks	21021901LJ	3	1036
RVΔL-Cre: 4 weeks	21021901LJ	4	1195
RVΔL-Cre: 4 weeks	21021901LJ	5	1402
RVΔL-Cre: 4 weeks	21021901LJ	6	1739

RVΔL-Cre: 4 weeks	21021901LJ	7	1793
RVΔL-Cre: 4 weeks	21021901LJ	8	1913
RVΔL-Cre: 4 weeks	21021901LJ	9	2026
RVΔL-Cre: 4 weeks	21021901LJ	10	1913
RVΔL-Cre: 4 weeks	21021901LJ	11	1929
RVΔL-Cre: 4 weeks	21021901LJ	12	1668
RVΔL-Cre: 4 weeks	21021901LJ	13	1247
RVΔL-Cre: 4 weeks	21021901LJ	14	1062
RVΔL-Cre: 4 weeks	21021901LJ	15	874
RVΔL-Cre: 4 weeks	21021901LJ	total:	21743
RVΔL-Cre: 4 weeks	21021902LJ	1	1357
RVΔL-Cre: 4 weeks	21021902LJ	2	1416
RVΔL-Cre: 4 weeks	21021902LJ	3	1287
RVΔL-Cre: 4 weeks	21021902LJ	4	1308
RVΔL-Cre: 4 weeks	21021902LJ	5	1583
RVΔL-Cre: 4 weeks	21021902LJ	6	1850
RVΔL-Cre: 4 weeks	21021902LJ	7	2020
RVΔL-Cre: 4 weeks	21021902LJ	8	2238
RVΔL-Cre: 4 weeks	21021902LJ	9	2092
RVΔL-Cre: 4 weeks	21021902LJ	10	2321
RVΔL-Cre: 4 weeks	21021902LJ	11	2241
RVΔL-Cre: 4 weeks	21021902LJ	12	1858
RVΔL-Cre: 4 weeks	21021902LJ	13	1326
RVΔL-Cre: 4 weeks	21021902LJ	14	1083
RVΔL-Cre: 4 weeks	21021902LJ	15	793
RVΔL-Cre: 4 weeks	21021902LJ	total:	24773
RVΔL-Cre: 4 weeks	21021903LJ	1	1543
RVΔL-Cre: 4 weeks	21021903LJ	2	1406
RVΔL-Cre: 4 weeks	21021903LJ	3	1303
RVΔL-Cre: 4 weeks	21021903LJ	4	1489
RVΔL-Cre: 4 weeks	21021903LJ	5	1825
RVΔL-Cre: 4 weeks	21021903LJ	6	1946
RVΔL-Cre: 4 weeks	21021903LJ	7	1924
RVΔL-Cre: 4 weeks	21021903LJ	8	2142
RVΔL-Cre: 4 weeks	21021903LJ	9	2077
RVΔL-Cre: 4 weeks	21021903LJ	10	1954
RVΔL-Cre: 4 weeks	21021903LJ	11	1672
RVΔL-Cre: 4 weeks	21021903LJ	12	1411
RVΔL-Cre: 4 weeks	21021903LJ	13	1020
RVΔL-Cre: 4 weeks	21021903LJ	14	687
RVΔL-Cre: 4 weeks	21021903LJ	15	627
RVΔL-Cre: 4 weeks	21021903LJ	total:	23026
RVΔL-Cre: 4 weeks	21021904LJ	1	1592
RVΔL-Cre: 4 weeks	21021904LJ	2	1466
RVΔL-Cre: 4 weeks	21021904LJ	3	1355
RVΔL-Cre: 4 weeks	21021904LJ	4	1599
RVΔL-Cre: 4 weeks	21021904LJ	5	1736
RVΔL-Cre: 4 weeks	21021904LJ	6	1862
RVΔL-Cre: 4 weeks	21021904LJ	7	1794
RVΔL-Cre: 4 weeks	21021904LJ	8	1930
RVΔL-Cre: 4 weeks	21021904LJ	9	2047
RVΔL-Cre: 4 weeks	21021904LJ	10	1584
RVΔL-Cre: 4 weeks	21021904LJ	11	1755
RVΔL-Cre: 4 weeks	21021904LJ	12	1431
RVΔL-Cre: 4 weeks	21021904LJ	13	741

RVΔL-Cre: 4 weeks	21021904LJ	14	755
RVΔL-Cre: 4 weeks	21021904LJ	15	625
RVΔL-Cre: 4 weeks	21021904LJ	total:	22272
RVΔL-5tTA: 1 week	21042901LJ	3	2
RVΔL-5tTA: 1 week	21042901LJ	6	9
RVΔL-5tTA: 1 week	21042901LJ	7	4
RVΔL-5tTA: 1 week	21042901LJ	8	26
RVΔL-5tTA: 1 week	21042901LJ	9	32
RVΔL-5tTA: 1 week	21042901LJ	10	54
RVΔL-5tTA: 1 week	21042901LJ	11	69
RVΔL-5tTA: 1 week	21042901LJ	12	56
RVΔL-5tTA: 1 week	21042901LJ	13	51
RVΔL-5tTA: 1 week	21042901LJ	14	14
RVΔL-5tTA: 1 week	21042901LJ	15	22
RVΔL-5tTA: 1 week	21042901LJ	total:	339
RVΔL-5tTA: 1 week	21042902LJ	1	4
RVΔL-5tTA: 1 week	21042902LJ	2	11
RVΔL-5tTA: 1 week	21042902LJ	3	14
RVΔL-5tTA: 1 week	21042902LJ	4	4
RVΔL-5tTA: 1 week	21042902LJ	5	39
RVΔL-5tTA: 1 week	21042902LJ	6	108
RVΔL-5tTA: 1 week	21042902LJ	7	213
RVΔL-5tTA: 1 week	21042902LJ	8	255
RVΔL-5tTA: 1 week	21042902LJ	9	296
RVΔL-5tTA: 1 week	21042902LJ	10	219
RVΔL-5tTA: 1 week	21042902LJ	11	134
RVΔL-5tTA: 1 week	21042902LJ	12	64
RVΔL-5tTA: 1 week	21042902LJ	13	29
RVΔL-5tTA: 1 week	21042902LJ	14	14
RVΔL-5tTA: 1 week	21042902LJ	15	7
RVΔL-5tTA: 1 week	21042902LJ	total:	1411
RVΔL-5tTA: 1 week	21042903LJ	2	3
RVΔL-5tTA: 1 week	21042903LJ	3	7
RVΔL-5tTA: 1 week	21042903LJ	4	6
RVΔL-5tTA: 1 week	21042903LJ	5	4
RVΔL-5tTA: 1 week	21042903LJ	6	27
RVΔL-5tTA: 1 week	21042903LJ	7	40
RVΔL-5tTA: 1 week	21042903LJ	8	81
RVΔL-5tTA: 1 week	21042903LJ	9	115
RVΔL-5tTA: 1 week	21042903LJ	10	165
RVΔL-5tTA: 1 week	21042903LJ	11	120
RVΔL-5tTA: 1 week	21042903LJ	12	84
RVΔL-5tTA: 1 week	21042903LJ	13	67
RVΔL-5tTA: 1 week	21042903LJ	14	44
RVΔL-5tTA: 1 week	21042903LJ	15	33
RVΔL-5tTA: 1 week	21042903LJ	total:	796
RVΔL-5tTA: 1 week	21060301LJ	1	85
RVΔL-5tTA: 1 week	21060301LJ	2	117
RVΔL-5tTA: 1 week	21060301LJ	3	154
RVΔL-5tTA: 1 week	21060301LJ	4	59
RVΔL-5tTA: 1 week	21060301LJ	5	207
RVΔL-5tTA: 1 week	21060301LJ	6	325
RVΔL-5tTA: 1 week	21060301LJ	7	314
RVΔL-5tTA: 1 week	21060301LJ	8	303
RVΔL-5tTA: 1 week	21060301LJ	9	288

RVΔL-5tTA: 1 week	21060301J	10	185
RVΔL-5tTA: 1 week	21060301J	11	141
RVΔL-5tTA: 1 week	21060301J	12	47
RVΔL-5tTA: 1 week	21060301J	13	19
RVΔL-5tTA: 1 week	21060301J	14	14
RVΔL-5tTA: 1 week	21060301J	15	13
RVΔL-5tTA: 1 week	21060301J	total:	2271
RVΔL-5tTA: 4 weeks	21042905J	1	4
RVΔL-5tTA: 4 weeks	21042905J	2	14
RVΔL-5tTA: 4 weeks	21042905J	3	6
RVΔL-5tTA: 4 weeks	21042905J	4	21
RVΔL-5tTA: 4 weeks	21042905J	5	15
RVΔL-5tTA: 4 weeks	21042905J	6	41
RVΔL-5tTA: 4 weeks	21042905J	7	42
RVΔL-5tTA: 4 weeks	21042905J	8	101
RVΔL-5tTA: 4 weeks	21042905J	9	156
RVΔL-5tTA: 4 weeks	21042905J	10	109
RVΔL-5tTA: 4 weeks	21042905J	11	94
RVΔL-5tTA: 4 weeks	21042905J	12	24
RVΔL-5tTA: 4 weeks	21042905J	13	87
RVΔL-5tTA: 4 weeks	21042905J	14	44
RVΔL-5tTA: 4 weeks	21042905J	15	71
RVΔL-5tTA: 4 weeks	21042905J	total:	829
RVΔL-5tTA: 4 weeks	21042906J	1	9
RVΔL-5tTA: 4 weeks	21042906J	2	17
RVΔL-5tTA: 4 weeks	21042906J	3	28
RVΔL-5tTA: 4 weeks	21042906J	4	23
RVΔL-5tTA: 4 weeks	21042906J	5	24
RVΔL-5tTA: 4 weeks	21042906J	6	25
RVΔL-5tTA: 4 weeks	21042906J	7	93
RVΔL-5tTA: 4 weeks	21042906J	8	200
RVΔL-5tTA: 4 weeks	21042906J	9	305
RVΔL-5tTA: 4 weeks	21042906J	10	233
RVΔL-5tTA: 4 weeks	21042906J	11	200
RVΔL-5tTA: 4 weeks	21042906J	12	141
RVΔL-5tTA: 4 weeks	21042906J	13	74
RVΔL-5tTA: 4 weeks	21042906J	14	108
RVΔL-5tTA: 4 weeks	21042906J	15	94
RVΔL-5tTA: 4 weeks	21042906J	total:	1574
RVΔL-5tTA: 4 weeks	21042907J	2	2
RVΔL-5tTA: 4 weeks	21042907J	4	3
RVΔL-5tTA: 4 weeks	21042907J	5	5
RVΔL-5tTA: 4 weeks	21042907J	6	24
RVΔL-5tTA: 4 weeks	21042907J	7	32
RVΔL-5tTA: 4 weeks	21042907J	8	51
RVΔL-5tTA: 4 weeks	21042907J	9	100
RVΔL-5tTA: 4 weeks	21042907J	10	74
RVΔL-5tTA: 4 weeks	21042907J	11	69
RVΔL-5tTA: 4 weeks	21042907J	12	54
RVΔL-5tTA: 4 weeks	21042907J	13	26
RVΔL-5tTA: 4 weeks	21042907J	14	31
RVΔL-5tTA: 4 weeks	21042907J	15	28
RVΔL-5tTA: 4 weeks	21042907J	total:	499
RVΔL-5tTA: 4 weeks	21042908J	1	64
RVΔL-5tTA: 4 weeks	21042908J	2	67

RVΔL-5tTA: 4 weeks	21042908LJ	3	66
RVΔL-5tTA: 4 weeks	21042908LJ	4	79
RVΔL-5tTA: 4 weeks	21042908LJ	5	151
RVΔL-5tTA: 4 weeks	21042908LJ	6	149
RVΔL-5tTA: 4 weeks	21042908LJ	7	166
RVΔL-5tTA: 4 weeks	21042908LJ	8	171
RVΔL-5tTA: 4 weeks	21042908LJ	9	184
RVΔL-5tTA: 4 weeks	21042908LJ	10	57
RVΔL-5tTA: 4 weeks	21042908LJ	11	48
RVΔL-5tTA: 4 weeks	21042908LJ	12	29
RVΔL-5tTA: 4 weeks	21042908LJ	13	14
RVΔL-5tTA: 4 weeks	21042908LJ	14	27
RVΔL-5tTA: 4 weeks	21042908LJ	15	57
RVΔL-5tTA: 4 weeks	21042908LJ	total:	1329

Condition	Mouse number	Cell counts
RVΔGL-Flpo: 1 week	21021505LJ	22
RVΔGL-Flpo: 1 week	21021506LJ	18
RVΔGL-Flpo: 1 week	21021507LJ	121
RVΔGL-Flpo: 1 week	21021508LJ	2
RVΔGL-Flpo: 1 week	21052505LJ	7
RVΔGL-Flpo: 1 week	21052506LJ	8
RVΔGL-Flpo: 1 week	21052507LJ	98
RVΔGL-Flpo: 1 week	21052508LJ	6
RVΔGL-Flpo: 1 week	Mean:	35.25
RVΔL-Flpo: 1 week	21021805LJ	3093
RVΔL-Flpo: 1 week	21021806LJ	320
RVΔL-Flpo: 1 week	21021807LJ	311
RVΔL-Flpo: 1 week	21021808LJ	263
RVΔL-Flpo: 1 week	21052607LJ	47
RVΔL-Flpo: 1 week	21052608LJ	2161
RVΔL-Flpo: 1 week	21060302LJ	365
RVΔL-Flpo: 1 week	21060303LJ	238
RVΔL-Flpo: 1 week	Mean:	849.75
RVΔGL-Flpo: 4 weeks	21021501LJ	80
RVΔGL-Flpo: 4 weeks	21021502LJ	62
RVΔGL-Flpo: 4 weeks	21021503LJ	3254
RVΔGL-Flpo: 4 weeks	21021504LJ	237
RVΔGL-Flpo: 4 weeks	21052501LJ	43
RVΔGL-Flpo: 4 weeks	21052502LJ	14
RVΔGL-Flpo: 4 weeks	21052503LJ	130
RVΔGL-Flpo: 4 weeks	21052504LJ	65
RVΔGL-Flpo: 4 weeks	Mean:	485.625
RVΔL-Flpo: 4 weeks	21021801LJ	4614
RVΔL-Flpo: 4 weeks	21021802LJ	3006
RVΔL-Flpo: 4 weeks	21021803LJ	1847
RVΔL-Flpo: 4 weeks	21021804LJ	3331
RVΔL-Flpo: 4 weeks	21052601LJ	1369
RVΔL-Flpo: 4 weeks	21052602LJ	2919
RVΔL-Flpo: 4 weeks	21052603LJ	3730
RVΔL-Flpo: 4 weeks	21052606LJ	3454
RVΔL-Flpo: 4 weeks	Mean:	3033.75
RVΔGL-Cre: 1 week	21021703LJ	13484
RVΔGL-Cre: 1 week	21021704LJ	14635
RVΔGL-Cre: 1 week	21021707LJ	10647
RVΔGL-Cre: 1 week	21021708LJ	10845
RVΔGL-Cre: 1 week	Mean:	12402.75
RVΔL-Cre: 1 week	21021905LJ	16907
RVΔL-Cre: 1 week	21021906LJ	17923
RVΔL-Cre: 1 week	21021907LJ	18751
RVΔL-Cre: 1 week	21021908LJ	16257
RVΔL-Cre: 1 week	Mean:	17459.5
RVΔGL-Cre: 4 weeks	21021701LJ	19156
RVΔGL-Cre: 4 weeks	21021702LJ	17935
RVΔGL-Cre: 4 weeks	21021705LJ	18539
RVΔGL-Cre: 4 weeks	21021706LJ	17938
RVΔGL-Cre: 4 weeks	Mean:	18392
RVΔL-Cre: 4 weeks	21021901LJ	21743
RVΔL-Cre: 4 weeks	21021902LJ	24773
RVΔL-Cre: 4 weeks	21021903LJ	23026
RVΔL-Cre: 4 weeks	21021904LJ	22272
RVΔL-Cre: 4 weeks	Mean:	22953.5
RVΔL-5tTA: 1 week	21042901LJ	339
RVΔL-5tTA: 1 week	21042902LJ	1411
RVΔL-5tTA: 1 week	21042903LJ	796
RVΔL-5tTA: 1 week	21060301LJ	2271
RVΔL-5tTA: 1 week	Mean:	1204.25
RVΔL-5tTA: 4 weeks	21042905LJ	829
RVΔL-5tTA: 4 weeks	21042906LJ	1574
RVΔL-5tTA: 4 weeks	21042907LJ	499
RVΔL-5tTA: 4 weeks	21042908LJ	1329
RVΔL-5tTA: 4 weeks	Mean:	1057.75

RVΔGL-Flpo 4 weeks	RVΔL-Flpo 4 weeks
80	4614
62	3006
3254	1847
237	3331
43	1369
14	2919
130	3730
65	3454

Anova: Single Factor

SUMMARY

Groups	Count	Sum	Average	Variance
RVΔGL-Flpo 4 weeks	8	3885	485.625	1255920.84
RVΔL-Flpo 4 weeks	8	24270	3033.75	1062946.79

ANOVA

Source of Variation	SS	df	MS	F	P-value	F crit
Between Groups	25971764.06	1	25971764.1	22.4003852	0.00032063	4.60010994
Within Groups	16232073.38	14	1159433.81			
Total	42203837.44	15				

RVΔGL-Flpo 1 week	RVΔGL-Flpo 4 weeks
22	80
18	62
121	3254
2	237
7	43
8	14
98	130
6	65

Anova: Single Factor

SUMMARY

Groups	Count	Sum	Average	Variance
RVΔGL-Flpo 1 week	8	282	35.25	2180.78571
RVΔGL-Flpo 4 weeks	8	3885	485.625	1255920.84

ANOVA

Source of Variation	SS	df	MS	F	P-value	F crit
Between Groups	811350.5625	1	811350.563	1.28980131	0.27515341	4.60010994
Within Groups	8806711.375	14	629050.813			
Total	9618061.938	15				

RVΔGL-Flpo 4 weeks	RVΔL-Flpo 1 week
80	3093
62	320
3254	311
237	263

Anova: Single Factor

SUMMARY

Groups	Count	Sum	Average	Variance
RVΔGL-Flpo 4 weeks	4	3633	908.25	2451752.25
RVΔL-Flpo 1 week	4	3987	996.75	1953632.25

ANOVA

Source of Variation	SS	df	MS	F	P-value	F crit
Between Groups	15664.5	1	15664.5	0.00711152	0.93553752	5.98737761
Within Groups	13216153.5	6	2202692.25			
Total	13231818	7				

RVΔL-Flpo 4 weeks	RVΔGL-Flpo 1 week
4614	22
3006	18
1847	121
3331	2

Anova: Single Factor

SUMMARY

Groups	Count	Sum	Average	Variance
RVΔL-Flpo 4 weeks	4	12798	3199.5	1294933.67
RVΔGL-Flpo 1 week	4	163	40.75	2936.91667

ANOVA

Source of Variation	SS	df	MS	F	P-value	F crit
Between Groups	19955403.13	1	19955403.1	30.7509907	0.00145233	5.98737761
Within Groups	3893611.75	6	648935.292			
Total	23849014.88	7				

RVΔL-Flpo 1 week	RVΔL-Flpo 4 weeks
3093	4614
320	3006
311	1847
263	3331
47	1369
2161	2919
365	3730
238	3454

Anova: Single Factor

SUMMARY

<i>Groups</i>	<i>Count</i>	<i>Sum</i>	<i>Average</i>	<i>Variance</i>
RVΔL-Flpo 1 week	8	6798	849.75	1274333.93
RVΔL-Flpo 4 weeks	8	24270	3033.75	1062946.79

ANOVA

<i>Source of Variation</i>	<i>SS</i>	<i>df</i>	<i>MS</i>	<i>F</i>	<i>P-value</i>	<i>F crit</i>
Between Groups	19079424	1	19079424	16.3261724	0.00121549	4.60010994
Within Groups	16360965	14	1168640.36			
Total	35440389	15				

RVΔGL-Flpo 1 week	RVΔL-Flpo 1 week
22	3093
18	320
121	311
2	263
7	47
8	2161
98	365
6	238

Anova: Single Factor

SUMMARY

<i>Groups</i>	<i>Count</i>	<i>Sum</i>	<i>Average</i>	<i>Variance</i>
RVΔGL-Flpo 1 week	8	282	35.25	2180.78571
RVΔL-Flpo 1 week	8	6798	849.75	1274333.93

ANOVA

<i>Source of Variation</i>	<i>SS</i>	<i>df</i>	<i>MS</i>	<i>F</i>	<i>P-value</i>	<i>F crit</i>
Between Groups	2653641	1	2653641	4.1576348	0.06079134	4.60010994
Within Groups	8935603	14	638257.357			
Total	11589244	15				

RVΔGL-Cre 4 weeks	RVΔL-Cre 4 weeks
19156	21743
17935	24773
18539	23026
17938	22272

Anova: Single Factor

SUMMARY

<i>Groups</i>	<i>Count</i>	<i>Sum</i>	<i>Average</i>	<i>Variance</i>
RVΔGL-Cre 4 weeks	4	73568	18392	340090
RVΔL-Cre 4 weeks	4	91814	22953.5	1748529.67

ANOVA

Source of Variation	SS	df	MS	F	P-value	F crit
Between Groups	41614564.5	1	41614564.5	39.8488678	0.00073769	5.98737761
Within Groups	6265859	6	1044309.83			
Total	47880423.5	7				

RVΔGL-Cre 4 weeks RVΔGL-Cre 1 week

19156	13484
17935	14635
18539	10647
17938	10845

Anova: Single Factor

SUMMARY

Groups	Count	Sum	Average	Variance
RVΔGL-Cre 4 weeks	4	73568	18392	340090
RVΔGL-Cre 1 week	4	49611	12402.75	3887094.92

ANOVA

Source of Variation	SS	df	MS	F	P-value	F crit
Between Groups	71742231.13	1	71742231.1	33.943266	0.00112486	5.98737761
Within Groups	12681554.75	6	2113592.46			
Total	84423785.88	7				

RVΔGL-Cre 4 weeks RVΔL-Cre 1 week

19156	16907
17935	17923
18539	18751
17938	16257

Anova: Single Factor

SUMMARY

Groups	Count	Sum	Average	Variance
RVΔGL-Cre 4 weeks	4	73568	18392	340090
RVΔL-Cre 1 week	4	69838	17459.5	1211355.67

ANOVA

Source of Variation	SS	df	MS	F	P-value	F crit
Between Groups	1739112.5	1	1739112.5	2.24192511	0.1849584	5.98737761
Within Groups	4654337	6	775722.833			
Total	6393449.5	7				

RVΔL-Cre 4 weeks	RVΔGL-Cre 1 week
21743	13484
24773	14635
23026	10647
22272	10845

Anova: Single Factor

SUMMARY

<i>Groups</i>	<i>Count</i>	<i>Sum</i>	<i>Average</i>	<i>Variance</i>
RVΔL-Cre 4 weeks	4	91814	22953.5	1748529.67
RVΔGL-Cre 1 week	4	49611	12402.75	3887094.92

ANOVA

<i>Source of Variation</i>	<i>SS</i>	<i>df</i>	<i>MS</i>	<i>F</i>	<i>P-value</i>	<i>F crit</i>
Between Groups	222636651.1	1	222636651	79.010462	0.00011291	5.98737761
Within Groups	16906873.75	6	2817812.29			
Total	239543524.9	7				

RVΔL-Cre 4 weeks	RVΔL-Cre 1 week
21743	16907
24773	17923
23026	18751
22272	16257

Anova: Single Factor

SUMMARY

<i>Groups</i>	<i>Count</i>	<i>Sum</i>	<i>Average</i>	<i>Variance</i>
RVΔL-Cre 4 weeks	4	91814	22953.5	1748529.67
RVΔL-Cre 1 week	4	69838	17459.5	1211355.67

ANOVA

<i>Source of Variation</i>	<i>SS</i>	<i>df</i>	<i>MS</i>	<i>F</i>	<i>P-value</i>	<i>F crit</i>
Between Groups	60368072	1	60368072	40.790818	0.00069327	5.98737761
Within Groups	8879656	6	1479942.67			
Total	69247728	7				

RVΔGL-Cre 1 week	RVΔL-Cre 1 week
13484	16907
14635	17923
10647	18751
10845	16257

Anova: Single Factor

SUMMARY

<i>Groups</i>	<i>Count</i>	<i>Sum</i>	<i>Average</i>	<i>Variance</i>
RVΔGL-Cre 1 week	4	49611	12402.75	3887094.92
RVΔL-Cre 1 week	4	69838	17459.5	1211355.67

ANOVA

<i>Source of Variation</i>	<i>SS</i>	<i>df</i>	<i>MS</i>	<i>F</i>	<i>P-value</i>	<i>F crit</i>
Between Groups	51141441.13	1	51141441.1	20.061562	0.00419681	5.98737761
Within Groups	15295351.75	6	2549225.29			
Total	66436792.88	7				

RVΔL-5tTA 1 week	RVΔL-5tTA 4 weeks
339	829
1411	1574
796	499
2271	1329

Anova: Single Factor

SUMMARY

<i>Groups</i>	<i>Count</i>	<i>Sum</i>	<i>Average</i>	<i>Variance</i>
RVΔL-5tTA 1 week	4	4817	1204.25	698675.583
RVΔL-5tTA 4 weeks	4	4231	1057.75	234872.917

ANOVA

<i>Source of Variation</i>	<i>SS</i>	<i>df</i>	<i>MS</i>	<i>F</i>	<i>P-value</i>	<i>F crit</i>
Between Groups	42924.5	1	42924.5	0.09195987	0.77193993	5.98737761
Within Groups	2800645.5	6	466774.25			
Total	2843570	7				

Comparison	p-value
RVΔGL-Flpo 4 weeks vs RVΔL-Flpo 4 weeks	0.000320625
RVΔGL-Flpo 4 weeks vs RVΔGL-Flpo 1 week	0.27515341
RVΔGL-Flpo 4 weeks vs RVΔL-Flpo 1 week	0.93553752
RVΔL-Flpo 4 weeks vs RVΔGL-Flpo 1 week	0.001452333
RVΔL-Flpo 4 weeks vs RVΔL-Flpo 1 week	0.001215493
RVΔGL-Flpo 1 week vs RVΔL-Flpo 1 week	0.060791343
RVΔGL-Cre 4 weeks vs RVΔL-Cre 4 weeks	0.000737695
RVΔGL-Cre 4 weeks vs RVΔGL-Cre 1 week	0.001124862
RVΔGL-Cre 4 weeks vs RVΔL-Cre 1 weeks	0.184958403
RVΔL-Cre 4 weeks vs RVΔGL-Cre 1 week	0.000112909
RVΔL-Cre 4 weeks vs RVΔL-Cre 1 week	0.000693266
RVΔGL-Cre 1 week vs RVΔL-Cre 1 week	0.004196815
RVΔL-5tTA 1 week vs RVΔL-5tTA 4 weeks	0.77193993

812 **Video S1. RV Δ L-Cre-labeled cortical neurons at 2 weeks vs 10 weeks postinjection, Related to**
813 **Figure 3**
814 3-dimensional rendering of the same cortical volume shown in Figure 3, panel B.
815
816 *See external file.*
817
818

819 **File S3. Cell counts and statistics for structural two-photon imaging experiments, Related to Figure**
820 **3**
821
822 *See following pages.*
823
824

RVΔGL-Cre: cell counts	Animal-1 FOV-1	Animal-1 FOV-2	Animal-2 FOV-1	Animal-2 FOV-2	Animal-3 FOV-1	Animal-3 FOV-2	Animal-4 FOV-1	Animal-4 FOV-2
Week-1	300	315	219	318	435	288	203	220
Week-2	519	359	269	546	488	364	321	354
Week-3	537	390	305	569	505	443	369	427
Week-4	538	403	317	572	518	445	372	426
Week-6	537	403	319	571	515	451	370	424
Week-12	535	400	319	567	514	443	369	424
Week-14			319	567	513	442	368	421
Week-16			319					

RVΔGL-Cre: percentage to week-1	Animal-1 FOV-1	Animal-1 FOV-2	Animal-2 FOV-1	Animal-2 FOV-2	Animal-3 FOV-1	Animal-3 FOV-2	Animal-4 FOV-1	Animal-4 FOV-2
Week-1	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00
Week-2	1.73	1.14	1.23	1.72	1.12	1.26	1.58	1.61
Week-3	1.79	1.24	1.39	1.79	1.16	1.54	1.82	1.94
Week-4	1.79	1.28	1.45	1.80	1.19	1.55	1.83	1.94
Week-6	1.79	1.28	1.46	1.80	1.18	1.57	1.82	1.93
Week-12	1.78	1.27	1.46	1.78	1.18	1.54	1.82	1.93
Week-14			1.46	1.78	1.18	1.53	1.81	1.91
Week-16			1.46					

RVΔGL-Cre: comparison between week-1 and week-4	Animal-1 FOV-1	Animal-1 FOV-2	Animal-2 FOV-1	Animal-2 FOV-2	Animal-3 FOV-1	Animal-3 FOV-2	Animal-4 FOV-1	Animal-4 FOV-2
Week-1	300	315	219	318	435	288	203	220
Week-4	538	403	317	572	518	445	372	426
Increase (Week-4 - Week-1)	238	88	98	254	83	157	169	206
% increase (Week-4 - Week-1)	79.33%	27.94%	44.75%	79.87%	19.08%	54.51%	83.25%	93.64%
Increase (Week-12 - Week-4)	-3	-3	2	-5	-4	-2	-3	-2
% increase (Week-12 - Week-4)	-0.56%	-0.74%	0.63%	-0.87%	-0.77%	-0.45%	-0.81%	-0.47%

t-Test: Paired Two Sample for Means		
	Week-1	Week-4
Mean	287.25	448.875
Variance	5712.5	7689.267857
Observations	8	8
Pearson Correlation		0.665513457
Hypothesized Mean Difference		0
df		7
t Stat		-6.75473154
P(T<=t) one-tail		0.000131934
t Critical one-tail		1.894578605
P(T<=t) two-tail		0.000263868
t Critical two-tail		2.364624252

t-Test: Paired Two Sample for Means		
	Week-4	Week-12
Mean	448.875	446.375
Variance	7689.267857	7407.410714
Observations	8	8
Pearson Correlation		0.99890399
Hypothesized Mean Difference		0
df		7
t Stat		3.415650255
P(T<=t) one-tail		0.005600716
t Critical one-tail		1.894578605
P(T<=t) two-tail		0.011201433
t Critical two-tail		2.364624252

% increase from 1 week to 4 weeks:								
RVΔGL-Cre	79.33%	27.94%	44.75%	79.87%	19.08%	54.51%	83.25%	93.64%
RVΔL-Cre	84.38%	84.41%	49.57%	70.25%	54.88%	67.66%	40.66%	93.26%

Unpaired t test	
Table Analyzed	% increase
P value	0.5187
P value summary	ns
Significantly different (P < 0.05)?	No
One- or two-tailed P value?	Two-tailed
t, df	t=0.6621, df=14
How big is the difference?	
Mean of RVΔGL-Cre	0.603
Mean of RVΔL-Cre	0.6813
Difference between means (B - A) ± SEM	0.07836 ± 0.1184
95% confidence interval	-0.1755 to 0.3322
R squared (eta squared)	0.03036
F test to compare variances	
F, DFn, Dfd	2.216, 7, 7
P value	0.3158
P value summary	ns
Significantly different (P < 0.05)?	No
Data analyzed	
Sample size, RVΔGL-Cre	8
Sample size, RVΔL-Cre	8

% increase from 4 weeks to 12 weeks:								
RVΔGL-Cre	-0.56%	-0.74%	0.63%	-0.87%	-0.77%	-0.45%	-0.81%	-0.47%
RVΔL-Cre	0.48%	-0.29%	2.31%	-1.46%	3.30%	0.00%	0.78%	0.58%

Unpaired t test	
P value	0.0451
P value summary	*
Significantly different (P < 0.05)?	Yes
One- or two-tailed P value?	Two-tailed
t, df	t=2.199, df=14
How big is the difference?	
Mean of RVΔGL-Cre	-0.005054
Mean of RVΔL-Cre	0.007135
Difference between means (B - A) ± SEM	.01219 ± 0.005542
95% confidence interval	0.003030 to 0.02407
R squared (eta squared)	0.2568
F test to compare variances	
F, DFn, Dfd	9.402, 7, 7
P value	0.0085
P value summary	**
Significantly different (P < 0.05)?	Yes
Data analyzed	
Sample size, RVΔGL-Cre	8
Sample size, RVΔL-Cre	8

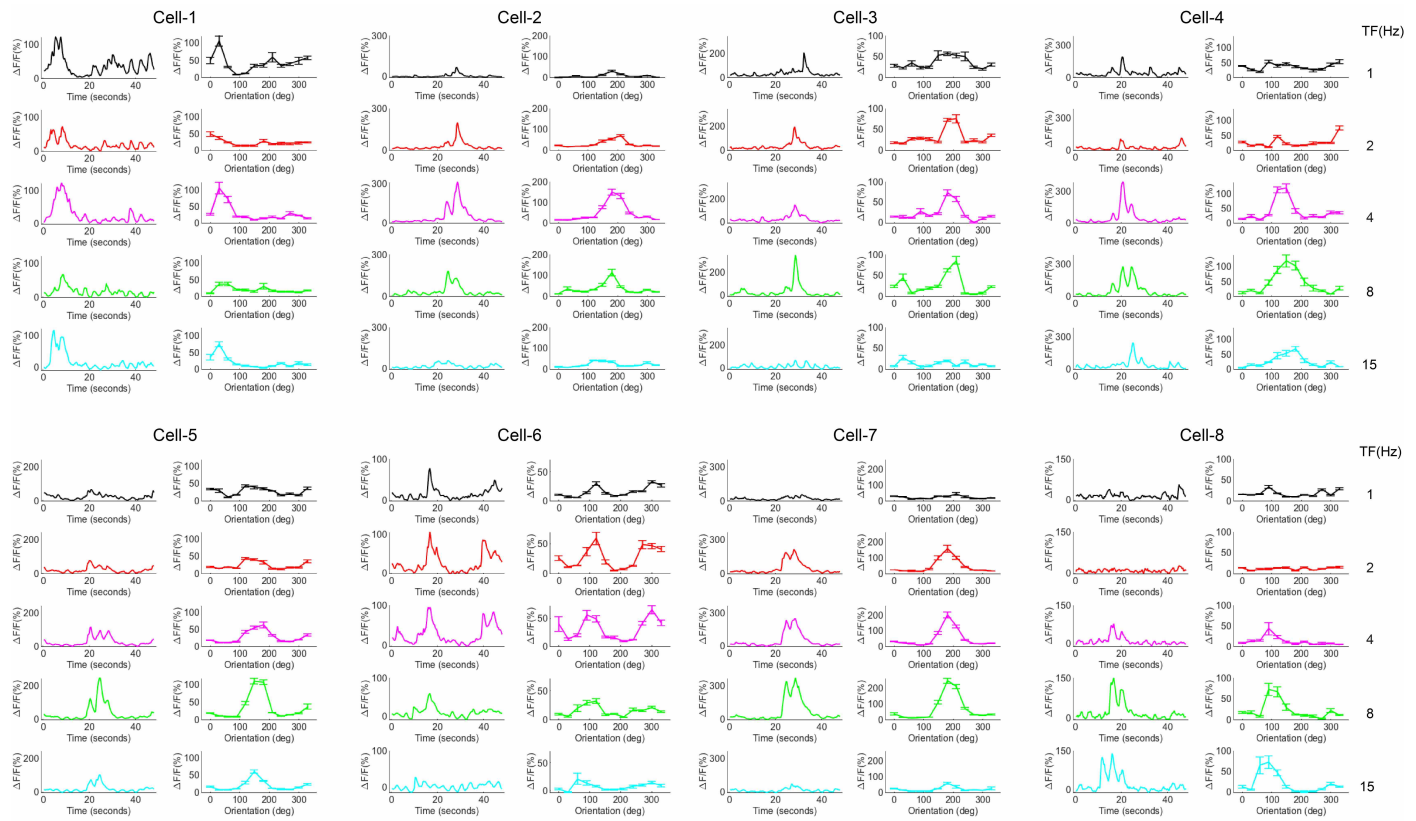
RVΔL-Cre: cell counts	Animal-1 FOV-1	Animal-1 FOV-2	Animal-2 FOV-1	Animal-2 FOV-2	Animal-3 FOV-1	Animal-3 FOV-2	Animal-4 FOV-1	Animal-4 FOV-2
Week-1	224	186	232	242	215	167	182	178
Week-2	331	307	300	298	270	237	200	228
Week-3	396	340	341	378	289	279	253	304
Week-4	413	343	347	412	333	280	256	344
Week-6	409	342	347	413	342	283	255	345
Week-12	415	342	355	406	344	280	258	346
Week-14	415	342	354	406	344	280		
Week-16				406	344	280		

RVΔL-Cre: percentage to week-1	Animal-1 FOV-1	Animal-1 FOV-2	Animal-2 FOV-1	Animal-2 FOV-2	Animal-3 FOV-1	Animal-3 FOV-2	Animal-4 FOV-1	Animal-4 FOV-2
Week-1	1.00	1.00	1.00	1.00	1.00	1.00	1.00	1.00
Week-2	1.48	1.65	1.29	1.23	1.26	1.42	1.10	1.28
Week-3	1.77	1.83	1.47	1.56	1.34	1.67	1.39	1.71
Week-4	1.84	1.84	1.50	1.70	1.55	1.68	1.41	1.93
Week-6	1.83	1.84	1.50	1.71	1.59	1.69	1.40	1.94
Week-12	1.85	1.84	1.53	1.68	1.60	1.68	1.42	1.94
Week-14	1.85	1.84	1.53	1.68	1.60	1.68		
Week-16				1.68	1.60	1.68		

RVΔL-Cre: comparison between week-1 and week-4	Animal-1 FOV-1	Animal-1 FOV-2	Animal-2 FOV-1	Animal-2 FOV-2	Animal-3 FOV-1	Animal-3 FOV-2	Animal-4 FOV-1	Animal-4 FOV-2
Week-1	224	186	232	242	215	167	182	178
Week-4	413	343	347	412	333	280	256	344
Increase (Week-4 - Week-1)	189	157	115	170	118	113	74	166
% increase (Week-4 - Week-1)	84.38%	84.41%	49.57%	70.25%	54.88%	67.66%	40.66%	93.26%
Increase (Week-12 - Week-4)	2	-1	8	-6	11	0	2	2
% increase (Week-12 - Week-4)	0.48%	-0.29%	2.31%	-1.46%	3.30%	0.00%	0.78%	0.58%

t-Test: Paired Two Sample for Means		
	Week-1	Week-4
Mean	203.25	341
Variance	799.6428571	3040.571429
Observations	8	8
Pearson Correlation	0.754099973	
Hypothesized Mean Difference	0	
df	7	
t Stat	-10.09862351	
P(T<=t) one-tail	1.00266E-05	
t Critical one-tail	1.894578605	
P(T<=t) two-tail	2.00531E-05	
t Critical two-tail	2.364624252	

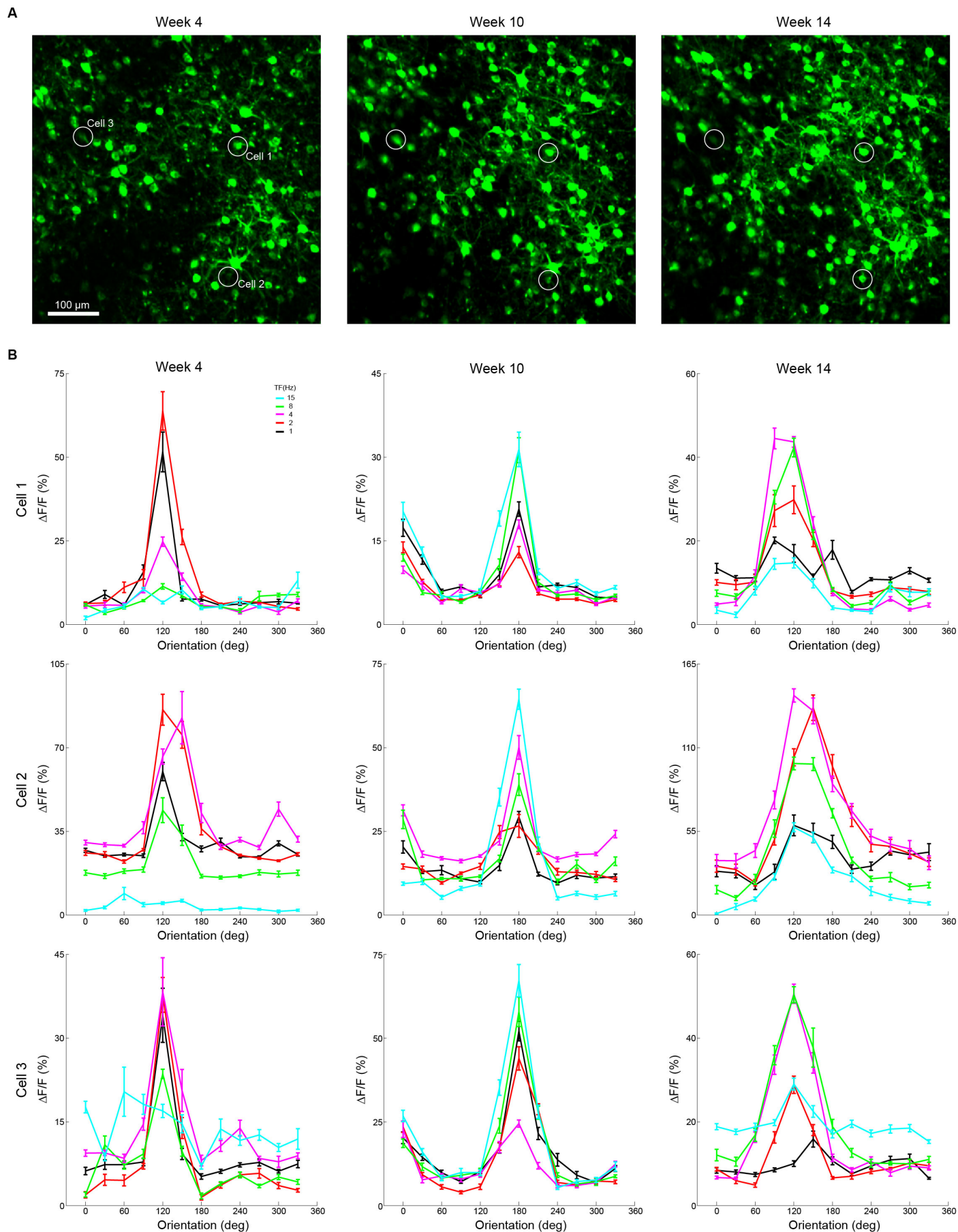
t-Test: Paired Two Sample for Means		
	Week-4	Week-12
Mean	341	343.25
Variance	3040.571429	2928.785714
Observations	8	8
Pearson Correlation	0.995543784	
Hypothesized Mean Difference	0	
df	7	
t Stat	-1.210419877	
P(T<=t) one-tail	0.13269902	
t Critical one-tail	1.894578605	
P(T<=t) two-tail	0.265398039	
t Critical two-tail	2.364624252	



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Figure S4. GCaMP6s signals and tuning curves of 8 example V1 neurons at 16 weeks postinjection, Related to Figure 4

These data were obtained with drifting gratings presented at 12 directions of motion and 5 temporal frequencies, repeated 10 times (tuning curve: mean $\Delta F/F \pm$ s.e.m; GCaMP6s signals: mean $\Delta F/F$) at two different FOVs at the 16-week timepoint.



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Figure S5. More examples showing long-term stability of orientation and temporal frequency tuning in RV Δ L-Cre labeled neurons, Related to Figure 4

836 (A) Maximum intensity projections of the same FOV at 4 weeks, 10 weeks, and 14 weeks postinjection.
837 Scale bar: 100 μm , applies to all images.
838 (B) Visual responses of the three circled cells in panel a measured at the three different timepoints. Data
839 were obtained with drifting gratings presented at 12 directions of motion and 5 temporal frequencies (mean
840 $\Delta F/F \pm \text{s.e.m.}$, averaged over 10 repeats).
841
842

843 **Video S2. GCaMP6s signals of visual cortical neurons 16 weeks after injection of RV Δ L-Cre, Related**
844 **to Figure 4**

845 Video shows responses to 10 repeats of drifting gratings at 12 directions of motion at one temporal
846 frequency (1 Hz) over a total of 480 seconds.

847

848 *See external file.*

849

850

851 **File S4. Tuned cell counts and statistics for functional two-photon imaging experiments, Related to**
852 **Figure 4**
853
854 *See following page.*