

1 **Increasing adult-born neurons protects mice from epilepsy.**

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50 **ABSTRACT**

51 Neurogenesis occurs in the adult brain in the hippocampal dentate gyrus, an
52 area that contains neurons which are vulnerable to insults and injury, such as severe
53 seizures. Previous studies showed that increasing adult neurogenesis reduced neuronal
54 damage after these seizures. Because the damage typically is followed by chronic life-
55 long seizures (epilepsy), we asked if increasing adult-born neurons would prevent
56 epilepsy. Adult-born neurons were selectively increased by deleting the pro-apoptotic
57 gene *Bax* from Nestin-expressing progenitors. Tamoxifen was administered at 6 weeks
58 of age to conditionally delete *Bax* in Nestin-CreER^{T2} *Bax*^{fl/fl} mice. Six weeks after
59 tamoxifen administration, severe seizures (status epilepticus; SE) were induced by
60 injection of the convulsant pilocarpine. After mice developed epilepsy, seizure frequency
61 was quantified for 3 weeks. Mice with increased adult-born neurons exhibited fewer
62 chronic seizures. Postictal depression was reduced also. These results were primarily in
63 female mice, possibly because they were the more affected by *Bax* deletion than males,
64 consistent with sex differences in *Bax*. The female mice with enhanced adult-born
65 neurons also showed less neuronal loss of hilar mossy cells and hilar somatostatin-
66 expressing neurons than wild type females or males, which is notable because these
67 two hilar cell types are implicated in epileptogenesis. The results suggest that selective
68 *Bax* deletion to increase adult-born neurons can reduce experimental epilepsy, and the
69 effect shows a striking sex difference. The results are surprising in light of past studies
70 showing that suppressing adult-born neurons can also reduce chronic seizures.

71 INTRODUCTION

72 It has been shown that neurogenesis occurs in the hippocampal dentate gyrus
73 (DG) during adult life of mammals (Taupin 2006; Gage et al. 2008; Altman 2011;
74 Kempermann 2012; Kazanis 2013). It is important to note that this idea was challenged
75 recently (Paredes et al. 2018; Sorrells et al. 2018) but afterwards more studies provided
76 support for the original idea (Boldrini et al. 2018; Kempermann et al. 2018; Tartt et al.
77 2018; Moreno-Jimenez et al. 2019; Tobin et al. 2019).

78 In the DG, adult-born neurons are born in the subgranular zone (SGZ; Altman
79 and Das 1965; Kaplan and Hinds 1977; Altman 2011). Upon maturation, newborn
80 neurons migrate to the granule cell layer (GCL; Cameron et al. 1993), develop almost
81 exclusively into GCs, and integrate into the DG circuitry like other GCs (Ramirez-Amaya
82 et al. 2006; Kempermann et al. 2015).

83 Prior studies suggest that the immature adult-born GCs can inhibit the other GCs
84 (Ash et al. 2023) especially when they are up to 6 weeks-old (Drew et al. 2016). By
85 inhibition of the GC population, young adult-born GCs could support DG functions that
86 require GCs to restrict action potential (AP) discharge, such as pattern separation
87 (Sahay et al. 2011a; Sahay et al. 2011b). Indeed suppressing adult neurogenesis in
88 mice appears to weaken pattern separation (Clelland et al. 2009; Nakashiba et al. 2012;
89 Niibori et al. 2012; Tronel et al. 2012) and increasing adult neurogenesis improves it
90 (Sahay et al. 2011a).

91 In addition, inhibition of the GC population by young adult-born GCs could limit
92 excessive excitation from glutamatergic input and protect the cells in the DG hilus and
93 hippocampus that are vulnerable to excitotoxicity. Thus, strong excitation of GCs can
94 cause excitotoxicity of hilar neurons, area CA1 pyramidal cells, and area CA3 pyramidal
95 cells (Scharfman and Schwartzkroin 1990b; a; Sloviter 1994; Scharfman 1999; Sloviter
96 et al. 2003). Indeed, increasing adult-born neurons protects hilar neurons, and CA3
97 from neuronal loss 3 days after severe seizures are induced by the convulsant
98 pilocarpine (Jain et al. 2019).

99 The seizures induced by kainic acid or pilocarpine are severe, continuous, and
100 last several hours, a condition called *status epilepticus* (SE). The neuronal injury in
101 hippocampus after SE has been suggested to be important because it is typically
102 followed by chronic seizures (epilepsy) in rodents and humans, and has been
103 suggested to cause the epilepsy (Falconer et al. 1964; Sloviter 1994; Cavalheiro et al.
104 1996; Herman 2002; Mathern et al. 2008; Dudek and Staley 2012; Dingledine et al.
105 2014). Chronic seizures involve the temporal lobe, so the type of epilepsy is called
106 temporal lobe epilepsy (TLE). In the current study we asked if increasing adult -born
107 neurons can protect from chronic seizures in an animal model of TLE. We used a very
108 common method to induce a TLE-like syndrome, which involves injection of the
109 muscarinic cholinergic agonist pilocarpine at a dose that elicits SE. Several weeks later,
110 spontaneous intermittent seizures begin and continue for the lifespan (Scorza et al.
111 2009; Botterill et al. 2019; Levesque et al. 2021; Whitebirch et al. 2022). Seizure
112 frequency, duration, and severity were measured by continuous video-EEG with 4
113 electrodes to monitor the hippocampus and cortex bilaterally.

114 It is known that SE increases adult neurogenesis (Parent and Kron 2012). SE
115 triggers a proliferation of progenitors in the week after SE (Parent et al. 1997). Although
116 many GCs that are born in the days after SE die in subsequent weeks by apoptosis,

117 some survive. Young neurons that arise after SE and migrate into the GCL may
118 suppress seizures by supporting inhibition of GCs because adult-born GCs in the
119 normal brain inhibit GCs when they are young (Drew et al. 2016; Ash et al. 2023). In
120 addition, after SE, the newborn GCs in the GCL can exhibit low excitability (Jakubs et
121 al. 2006). However, some neurons born after SE mismigrate to ectopic locations such
122 as the hilus (hilar ectopic GCs), where they can contribute to recurrent excitatory circuits
123 that promote seizures (Scharfman et al. 2000; Parent and Lowenstein 2002; Scharfman
124 2004; Scharfman and Hen 2007; Parent and Murphy 2008; Scharfman and McCloskey
125 2009; Zhan et al. 2010; Myers et al. 2013; Cho et al. 2015; Althaus et al. 2019; Zhou et
126 al. 2019). Since the hilar ectopic GCs are potential contributors to epileptogenesis, we
127 also studied whether enhancing adult -born neurons would alter the number of hilar
128 ectopic GCs.

129 Mossy cells are a major subset of glutamatergic hilar neurons which are
130 vulnerable to excitotoxicity after SE (Scharfman 1999; Sloviter et al. 2003). During SE,
131 mossy cells may contribute to the activity that ultimately leads to widespread neuronal
132 loss (Botterill et al., 2019). However, surviving mossy cells can be beneficial after SE
133 because they inhibit spontaneous chronic seizures in mice (Bui et al. 2018). Another
134 large subset of vulnerable hilar neurons co-express GABA and somatostatin (SOM;
135 Sloviter 1987; de Lanerolle et al. 1989; Freund et al. 1992; Sun et al. 2007) and
136 correspond to so-called HIPP cells (neurons with *h*ilar cell bodies and axons that project
137 to the terminal zone of the *p*erforant *p*ath; (Han et al. 1993)). HIPP cells are important
138 because they normally inhibit GCs and have the potential to prevent seizures.
139 Therefore, we studied mossy cells and SOM cells in the current study.

140 The results showed that increasing adult -born neurons protects mossy cells and
141 hilar SOM cells and reduces chronic seizures. Remarkably, the preservation of hilar
142 mossy cells and SOM cells, and the reduction in chronic seizures, was found in females
143 only. The sex difference may have been due to a greater ability to increase adult -born
144 neurons in females than males, consistent with sex differences in *Bax*- and caspase-
145 dependent cell death (Forger et al. 2004; Siegel and McCullough 2011).

146 The results are surprising because prior studies that suppressed neurogenesis
147 reduces chronic seizures. Therefore, taken together with the results presented here,
148 both increasing and suppressing adult-born neurons appear to reduce chronic seizures.
149 How could this be? Past studies suggested that suppressing adult-born neurons led to a
150 reduction in chronic seizures because there were fewer hilar ectopic granule cells. In
151 the current study, increasing adult-born neurons may have reduced chronic seizures for
152 another reason. Regardless, the present data suggest a novel and surprising series of
153 findings which, taken together with past studies, suggest that adult--born neurons can
154 be targeted in multiple ways to reduce chronic seizures in epilepsy.

155

156 RESULTS

157

158 I. Increasing adult-born neurons reduced the duration of pilocarpine-induced SE

159 A. General approach

160 The first experiment addressed the effect of increasing adult -born neurons on
161 pilocarpine-induced SE in Nestin-CreER^{T2}*Bax*^{fl/fl} mice (called "Cre+", below). To
162 produce Nestin-CreER^{T2}*Bax*^{fl/fl} mice, hemizygous Nestin-CreER^{T2} mice were bred with

163 homozygous *Bax^{fl/fl}* mice. Littermates of Cre+ mice that lacked Cre (called "Cre-", below)
164 were also treated with tamoxifen and were controls.

165 Fig. 1A1 shows the experimental timeline. Tamoxifen was injected s.c. once per
166 day for 5 days to delete *Bax* from Nestin-expressing progenitors. After 6 weeks, a time
167 sufficient for a substantial increase in adult-born neurons (Drew et al., 2016, Jain et al.,
168 2019), pilocarpine was injected s.c. to induce SE.

169 Fig. 1A2 shows the experimental timeline during the day of pilocarpine injection.
170 The location of electrodes for EEG are shown in Fig. 1B. Mice monitored with EEG were
171 implanted with electrodes 3 weeks before SE (see Methods). Examples of the EEG are
172 shown in Fig. 1C for Cre- and Cre+ mice and details are shown in Fig. 1- Supplemental
173 figure 1.

174

175 **B. Effects of increasing adult-born neurons on SE**

176 The latency to the first seizure after pilocarpine injection was measured for all
177 mice (with and without EEG electrodes; Fig. 1D) or just those that had EEG electrodes
178 (Fig. 1E). When mice with and without electrodes were pooled, the latency to the onset
179 of first seizure was similar in both genotypes (Cre-: 47.2 ± 4.8 min, n=27; Cre+: $45.3 \pm$
180 3.9 min, n=28; Student's t-test, $t(53)=0.3$, $p=0.761$; Fig. 1D1).

181 The total number of seizures was quantified until 2 hr after pilocarpine injection
182 because at that time diazepam was administered to decrease the severity SE. The total
183 number of seizures were similar in both genotypes (Cre-: 3.0 ± 0.2 seizures; Cre+: $2.8 \pm$
184 0.1 seizures; Student's t-test, $t(54)=0.9$, $p=0.377$; Fig. 1D2).

185 Interestingly, when sexes were separated, Cre+ females had a shorter latency to
186 the first seizure than all other groups (Fig. 1D3). Thus, a two-way ANOVA with genotype
187 (Cre- and Cre+) and sex (female and male) as main factors showed a main effect of sex
188 ($F(1,51)=4.31$; $p=0.043$) with Cre+ females exhibiting a shorter latency compared to
189 Cre+ males (Cre+ females, 34.3 ± 3.4 min, n=15; Cre+ males, 57.9 ± 5.6 min, n=13;
190 Bonferroni's test, $p=0.026$) but not other groups (Cre- female, 46.7 ± 9.3 min, n=12;
191 Cre- males, 47.4 ± 4.4 min, n=15; Bonferroni's tests, all $p > 0.344$; Fig. 1D3). There was
192 no effect of genotype ($F(1,49)=0.75$; $p=0.305$) or sex ($F(1,49)=0.62$; $p=0.436$) on the
193 total number of seizures by two-way ANOVA (Fig. 1D4).

194 When adult neurogenesis was suppressed by thymidine kinase activation in
195 GFAP-expressing progenitors, the severity of the first seizure was worse, meaning it
196 was often convulsive rather than non-convulsive (Iyengar et al., 2015). Therefore we
197 examined the severity of the first seizure. These analyses were conducted only with
198 mice implanted with electrodes because only with the EEG can one determine if a
199 seizure is non-convulsive. A non-convulsive seizure was defined as an EEG seizure
200 without movement. When sexes were pooled, the proportion of mice with a non-
201 convulsive first seizure was not different (Cre-: 18.2%, 2/11 mice; Cre+: 28.6%, 4/14
202 mice Chi-square test, $p>0.999$; Fig. 1E1). However, when sexes were separated, the
203 first seizure was non-convulsive in 60% of Cre+ females (3/5 mice) whereas only 25%
204 of Cre- females had a first seizure that was nonconvulsive (1/4 mice), 14% of Cre-
205 males (1/7 mice), and 11% of Cre+ males (1/9 mice; Fig. 1E3). Although the
206 percentages suggest differences, i.e., Cre+ females were protected from an initial
207 severe seizure, the differences were not significantly different (Fisher's exact test,
208 $p=0.166$; Fig. 1E3).

209 The duration of SE was shorter in Cre+ mice compared to Cre- mice (Cre-: 280.5
210 \pm 14.6 min, n=11; Cre+: 211.4 \pm 17.2 min, n=14; Student's t-test, t(23)=0.30, p=0.007;
211 Fig. 1E2). When sexes were separated, effects of genotype were modest. A two-way
212 ANOVA showed that the duration of SE was significantly affected by genotype
213 (F(1,21)=6.7; p=0.017) but not sex (F(1,21)=5.04; p=0.487). Cre+ males showed a trend
214 for a shorter SE duration compared to Cre- males (Cre- males: 280.1 \pm 22.8 min, n=7;
215 Cre+ males: 199.1 \pm 24.1 min, n=9; Bonferroni's test, p=0.078; Fig. 1E4). Cre+ females
216 had a mean SE duration that was shorter than Cre- females, but it was not a significant
217 difference (Cre- females: 281.0 \pm 11.8 min, n=4; Cre+ females: 233.4 \pm 20.5 min, n=5;
218 Bonferroni's test, p=0.485; Fig. 1E4). More females would have been useful, but the
219 incidence of SE in females was only 42.8% if they were implanted with EEG electrodes
220 (Fig. 1 – Supplemental figure 2). In contrast, the incidence of SE in the unimplanted
221 females was 100%, a significant difference by Fisher's exact test (p < 0.001; Fig. 1-
222 Supplemental figure 2). In males the incidence of SE was also significantly different in
223 implanted and unimplanted mice (implanted males, 70.4%; unimplanted males, 100%;
224 Fisher's exact test, p=0.016, Fig. 1- Supplemental figure 2).

225

226 **D. Power**

227 We also investigated power during SE (Fig. 1- Supplemental figure 3). The
228 baseline was measured, and then power was assessed for 5 hrs, at which time SE had
229 ended. Power was assessed in 20 min consecutive bins. Females were used in this
230 analysis (Cre- and Cre+). Two-way RMANOVA with genotype and time as main factors
231 showed no effect of genotype for any frequency range: delta (1-4 Hz, F(1,7)=1.61;
232 p=0.245); theta (4-8 Hz, F(1,7)=1.75; p=0.227); beta (8-30 Hz, F(1,7)=1.65; p=0.240);
233 low gamma (80 Hz, F(1,7)=0.29; p=0.174); high gamma (80-100 Hz, F(1,7)=0.17;
234 p=0.689). There was a significant effect of time for all bands (delta, p=0.003; theta,
235 p=0.002, beta, low gamma and high gamma, p < 0.001), which is consistent with the
236 declining power in SE with time.

237

238 **E. Role of diazepam**

239 Diazepam was administered earlier in females during SE than males, and this
240 could have influenced the results. However, the timing of SE was not significantly
241 different in females than males (Fig. 1 – Supplemental figure 4). Also, diazepam was
242 administered the same way in all Cre+ and Cre- females similarly but only the Cre+
243 females were protected as discussed below.

244 In summary, Cre+ mice did not show extensive differences in SE except for SE
245 duration, which was shorter.

246

247 **II. Increasing adult-born neurons decreased chronic seizures**

248 **A. Numbers and frequency of chronic seizures**

249 Continuous video-EEG was recorded for 3 weeks to capture chronic seizures
250 (Fig. 2A). Representative examples of chronic seizures are presented in Fig. 2B. All
251 chronic seizures were convulsive. First, we analyzed data with sexes pooled (Fig. 2C)
252 and the total number of chronic seizures were similar in the two genotypes (Cre-: 22.6 \pm
253 3.0 seizures, n=18; Cre+: 21.3 \pm 1.6 seizures, n=17; Student's t-test, t(33)=0.15,
254 p=0.882; Fig. 2C1). The frequency of chronic seizures were also similar among

255 genotypes (Cre-: 1.1 ± 0.14 seizures/day, $n=18$; Cre+: 1.0 ± 0.08 seizures/day, $n=17$;
256 Welch's t-test, $t(26)=0.37$, $p=0.717$; Fig. 2D1).

257 Data were then segregated based on sex and a two-way ANOVA was conducted
258 with genotype and sex as main factors. There was a main effect of genotype
259 ($F(1,32)=4.26$; $p=0.047$) and sex ($F(1,32)=12.46$; $p=0.001$) on the total number of
260 chronic seizures and a significant interaction between sex and genotype ($F(1,32)=8.54$;
261 $p=0.006$). Bonferroni's post-hoc tests showed that Cre+ females had ~half the chronic
262 seizures of Cre- females (Cre- female: 44.6 ± 10.2 seizures, $n=7$; Cre+ female: $22.6 \pm$
263 2.0 seizures, $n=9$; $p=0.004$; Fig. 2C). However, Cre+ males and Cre- males had a
264 similar number of chronic seizures (Cre- male: 16.1 ± 1.6 seizures, $n=12$; Cre+ male:
265 19.9 ± 2.7 seizures, $n=8$; $p>0.999$; Fig. 2C2).

266 Results for seizure frequency were similar to results comparing total numbers of
267 seizures. There was a main effect of genotype ($F(1,32)=4.18$; $p=0.049$) and sex ($F(1,$
268 $32)=11.96$; $p=0.002$) on chronic seizure frequency, and a significant interaction between
269 sex and genotype ($F(1,32)=8.29$; $p=0.007$). Cre+ female mice had approximately half
270 the seizures per day as Cre- females (Bonferroni's test, Cre- female: 2.1 ± 0.5
271 seizures/day; Cre+ female: 1.1 ± 0.1 seizures/day; $p=0.004$; Fig. 2D2).

272

273 **B. Additional analyses**

274 While reviewing the data for each mouse plotted in Fig. 2C2 and 2D2, one point
275 appeared spurious in the Cre- females, potentially influencing the comparison. The
276 seizures in this mouse were more than 2x the standard deviation of the mean. Although
277 not an outlier using the ROUT method (see Methods), we were curious if removing the
278 data of this mouse would lead to a difference in the statistical results. There was still a
279 main effect of sex ($F(1,31)=16.04$; $p=0.0004$) with a significant interaction between sex
280 and genotype ($F(1,31)=9.20$; $p=0.005$) and Cre+ females had significantly fewer
281 seizures than Cre- female mice ($p=0.020$; Fig. 2- Supplemental figure 1A1). Tests for
282 seizure frequency led to the same conclusions (Fig. 2- Supplemental figure 1A2). These
283 data suggest that spurious data point was not the reason for the results.

284 All mice were included in the analyses above, both those implanted and
285 unimplanted during SE. Mice which were unimplanted prior to SE were implanted at
286 approximately 2-3 weeks after pilocarpine to study chronic seizures. Because
287 implantation affected the incidence of SE (discussed above), we asked if chronic
288 seizures were different in implanted and unimplanted mice. The total number of chronic
289 seizures ($F(1,17)=1.33$, $p=0.265$) and seizure frequency ($F(1,17)=1.27$, $p=0.276$) were
290 similar, suggesting that implantation did not influence chronic seizures (Fig. 2-
291 Supplemental figure 1B).

292 There were no significant differences in mortality associated with SE or chronic
293 seizures. For quantification, we examined mortality during SE and the subsequent 3
294 days, 3 days until the end of the 3 week-long EEG recording period, or both (Fig. 2-
295 Supplemental figure 2A). Graphs of mouse numbers (Fig. 2- Supplemental figure 2B) or
296 percentages of mice (Fig. 2- Supplemental figure 2C) were similar: groups (Cre-
297 females, Cre+ females, Cre- males, Cre+ males) were not significantly different (Chi-Sq.
298 test, $p>0.999$).

299

300

301 C. Mean duration of individual chronic seizures

302 To evaluate the duration of individual seizures at the time mice were epileptic,
303 two measurements were made. First, durations of each seizure of a given mouse were
304 averaged, and then the averages for Cre- mice were compared to the averages for Cre+
305 mice (Fig. 2E1). There was no difference in the genotypes (Cre-: 46.8 ± 2.9 sec, $n=17$;
306 Cre+: 43.4 ± 2.5 sec, $n=16$; Student's t-test, $t(31)=0.89$, $p=0.379$; Fig. 2E1).

307 When separated by sex, a two-way ANOVA showed that female seizure
308 durations were shorter than males ($F(1,29)=12.42$; $p=0.001$). However, this was a sex
309 difference, not an effect of genotype ($F(1,29)=0.033$; $p=0.856$; Fig. 2E2), with Cre-
310 female seizure duration shorter than Cre- male seizure duration (Bonferroni post-hoc
311 test, $p=0.015$), and the same for Cre+ females compared to Cre+ males (Bonferroni
312 post-hoc test, $p=0.037$; Fig. 2E2). One reason for the sex difference could be related to
313 the greater incidence of postictal depression in females (see below), because that
314 suggests spreading depolarizations truncated the seizures in females but not males.

315 The second method to compare seizure durations compared the duration of
316 every seizure of every Cre- and Cre+ mouse. In the previous comparison (Fig. 2E1),
317 every mouse was a data point, whereas here every seizure was a data point (Fig. 2F1).
318 The data were similar between genotypes (Cre-: 41.9 ± 0.9 sec; Cre+: 36.8 ± 0.7 sec;
319 Mann-Whitney U test, U statistic, 18873, $p=0.079$; Fig. 2F1). When sexes were
320 separated, a Kruskal-Wallis test was significant (Kruskal-Wallis statistic, 69.30,
321 $p<0.001$). Post-hoc tests showed that Cre+ females had longer seizure durations than
322 Cre- females (Cre- female: 33.2 ± 0.7 sec; Cre+ female: 39.3 ± 0.6 sec, $p<0.001$; Fig.
323 2F2). Cre+ females may have had longer seizures because they were protected from
324 spreading depolarizations that truncated seizures in Cre- females. Seizure durations
325 were not significantly different in males (Cre- male: 51.4 ± 1.7 sec; Cre+ male: $44.0 \pm$
326 1.3 sec, $p=0.298$; Fig. 2F2).

327

328 D. Postictal depression

329 Postictal depression is a debilitating condition in humans where individuals suffer
330 fatigue, confusion and cognitive impairment after a seizure. In the EEG, it is exhibited by
331 a decrease in the EEG amplitude immediately after a seizure ends relative to baseline.
332 In recent years the advent of DC amplifiers made it possible to show that postictal
333 depression is often associated with spreading depolarization (Ssentongo et al. 2017), a
334 large depolarization shift that is accompanied by depolarization block. As action
335 potentials are blocked there are large decreases in input resistance leading to cessation
336 of synaptic responses. As ion pumps are activated to restore equilibrium, there is
337 recovery and the EEG returns to normal (Somjen 2001; Hartings et al. 2017; Herreras
338 and Makarova 2020; Lu and Scharfman 2021).

339 We found that males had little evidence of postictal depression but it was
340 common in females (Fig. 3), a sex difference that is consistent with greater spreading
341 depolarization in females (Eikermann-Haerter et al. 2009; Bolay et al. 2011; Kudo et al.
342 2023). As shown in Fig. 3A1, a male showed a robust spontaneous seizure (selected
343 from the 3 week-long recording period when mice are epileptic). However, the end of
344 the seizure did not exhibit a decrease in the amplitude of the EEG relative to baseline.
345 The EEG before and immediately after the seizure is expanded in Fig. 3A2 to show the
346 EEG amplitude is similar. In contrast, the seizure from the female in Fig. 3B1-2 shows a

347 large reduction in the EEG immediately after the seizure. For quantification, the mean
348 peak-to-trough amplitude of the EEG 25-30 sec before the seizure was compared to the
349 mean amplitude for the EEG during the maximal depression of the EEG after the
350 seizure. If the depression was more than half, the animal was said to have had postictal
351 depression.

352 When all chronic seizures were analyzed (n=274), the number of seizures with
353 postictal depression was significantly different in the four groups (Cre- females, Cre+
354 females, Cre- males, Cre+ males; Chi-square test, $p < 0.0001$; Fig. 3C-D). Cre+ females
355 showed less postictal depression compared to Cre- females (Fisher's exact test,
356 $p=0.009$; Fig. 3C). There was a sex difference, with females showing more postictal
357 depression (108/154 seizures, 70.5%) than males (17/120 seizures, 14.2%; $p < 0.0001$;
358 Fig. 3C-D).

359

360 E. Clusters of seizures

361 Next, we asked if the distribution of seizures during the 3 weeks of video-EEG
362 was affected by increasing adult-born neurons. In Fig. 4A, a plot of the day-to-day
363 variation in seizures is shown with each day of recording either black (if there were
364 seizures) or white (if there were no seizures).

365 The number of days with seizures were similar between genotypes (Cre-: $8.6 \pm$
366 0.6 days, n=18; Cre+: 8.4 ± 0.6 days, n=17; Student's t-test, $t(33)=0.23$, $p=0.822$; Fig.
367 4B1). The number of consecutive days without seizures, called the seizure-free interval,
368 was also similar between genotypes (Cre-: 6.4 ± 0.4 days, n=19; Cre+: 7.7 ± 0.6 days,
369 n=17; Student's t-test, $t(34)=1.65$, $p=0.107$; Fig. 4B2). When data were segregated
370 based on sex, a two-way ANOVA showed no effect of genotype ($F(1,32)=1.18$,
371 $p=0.286$) or sex on the number of days with seizures ($F(1,32)=0.86$, $p=0.361$; Fig. 4B3).
372 There also was no effect of genotype ($F(1,32)=2.86$, $p=0.100$) or sex $F(1,32)=0.53$,
373 $p=0.471$) on seizure-free interval (Fig. 4B4).

374 Clustering is commonly manifested by consecutive days with frequent seizures.
375 Clusters of seizures can have a substantial impact on the quality of life (Haut 2015;
376 Jafarpour et al. 2019) so they are important. In humans, clusters are defined as at least
377 3 seizures within 24 hr (Goffin et al. 2007; Jafarpour et al. 2019). Therefore, we defined
378 clusters as >1 consecutive day with ≥ 3 seizures/day (Fig. 4C). The duration of clusters
379 were similar between genotypes (Cre-: 3.8 ± 0.7 days, n=19; Cre+: 3.0 ± 0.3 days,
380 n=18; Mann-Whitney's U test, U statistic 159, $p=0.723$; Fig. 4D1). Next, we calculated
381 the number of days between clusters, which we call the intercluster interval. Genotypes
382 were similar (Cre-: 7.2 ± 0.8 days, n=13; Cre+: 9.2 ± 0.8 days, n=9; Student's t-test,
383 $t(20)=1.70$, $p=0.104$; Fig. 4D2).

384 Two-way ANOVA was then performed on the sex-separated data. For cluster
385 duration, there was no effect of genotype ($F(1,33)=3.36$, $p=0.076$) but there was a main
386 effect of sex ($F(1,33)=7.66$, $p=0.009$) and a significant interaction of genotype and sex
387 ($F(1,33)=.66$, $p=0.009$). Cre+ females had fewer days with ≥ 3 seizures than Cre-
388 females (Cre- females: 6.3 ± 1.4 days; Cre+ females: 3.0 ± 0.4 days; Bonferroni's test,
389 $p=0.009$; Fig. 4D3). These data suggest Cre+ females were protected from the peak of
390 a cluster, when seizures increase above 3/day.

391 There was no effect of genotype ($F(1,17)=2.72$, $p=0.117$) or sex ($F(1,17)=2.72$,
392 $p=0.117$) on the intercluster interval (Fig. 4D4). However, this result may have

393 underestimated effects because Cre⁺ females often had such a long interval that it was
394 not captured in the 3 week-long recording period. That led to fewer Cre⁺ females that
395 were included in the measurement of intercluster interval. In 5 out of 9 (i.e., 55%) Cre⁺
396 females, there was only one cluster in 3 weeks, so intercluster interval was too long to
397 capture. Of those mice where intercluster interval could be measured, Cre⁻ females had
398 an interval of 5.7 ± 1.0 days (n=7) and Cre⁺ females had a 9.0 ± 1.1 day interval (n=4).
399 That difference was not significant.

400 In summary, Cre⁺ females had fewer seizures, fewer days with ≥ 3 seizures,
401 reduced postictal depression, and appeared to have a long period between clusters of
402 seizures.

403

404 **III. Before and after epileptogenesis, Cre⁺ female mice exhibited more immature** 405 **neurons than Cre⁻ female mice but that was not true for male mice.**

406 **A. Prior to SE**

407 We first confirmed that prior to pilocarpine treatment, Cre⁺ mice had more young
408 adult-born neurons compared to Cre⁻ mice (Fig. 5A, Fig.5- Supplemental Fig. 1A-D). To
409 that end, we quantified the adult-born GCs associated with the GCL/SGZ in both Cre⁺
410 and Cre⁻ mice. DCX was used as a marker because it is highly expressed in immature
411 neurons (Brown et al. 2003; Couillard-Despres et al. 2005). The area of the GCL/SGZ
412 that exhibited DCX-ir was calculated and expressed as a percent of the total area of the
413 GCL/SGZ (Fig. 5C).

414 A two-way ANOVA with sex and genotype as factors showed a significant effect
415 of genotype ($F(1,9)=60.78$, $p<0.001$) but not sex ($F(1,9)=1.20$, $p=0.301$; Fig. 5D). Post-
416 hoc comparisons showed that Cre⁺ females had more DCX than Cre⁻ females
417 ($p=0.001$) and the same was true for males ($p=0.003$). Cre⁺ males had more DCX than
418 Cre⁻ females ($p<0.001$), and Cre⁺ females had more DCX than Cre⁻ males ($p=0.006$).
419 Cre⁻ females and males were not different ($p=0.774$). The results are consistent with
420 studies using the same methods which showed that Cre⁺ males have more DCX
421 compared to Cre⁻ males (Jain et al., 2019). Together the data suggest that Cre⁺ mice
422 had more young adult-born neurons than Cre⁻ mice immediately before SE.

423

424 **B. After epileptogenesis**

425 We also quantified DCX at the time when epilepsy had developed, after the 3
426 week-long EEG recording (Fig. 5B). Representative examples of DCX expression in the
427 GCL/SGZ are presented in Fig. 5F and Fig. 5E shows the area fraction of DCX in the
428 GCL/SGZ was significantly greater in Cre⁺ mice than Cre⁻ mice (Cre⁻: $3.1 \pm 0.4\%$,
429 $n=20$; Cre⁺: $4.2 \pm 0.3\%$, $n=17$; Student's t-test, $t(35)=2.13$, $p=0.041$; Fig. 5D1).
430 Therefore, Cre⁺ mice had increased DCX in the GCL/SGZ after chronic seizures had
431 developed.

432 To investigate a sex difference, a two-way ANOVA was conducted with genotype
433 and sex as main factors. There was a significant effect of genotype ($F(1,33)=12.62$,
434 $p=0.001$) and sex ($F(1,33)=11.68$, $p=0.002$), with Cre⁺ females having more DCX than
435 Cre⁻ females (Cre⁻ female: 1.8 ± 0.3 , $n=7$; Cre⁺ female: 3.8 ± 0.4 , $n=9$; Bonferroni's
436 test, $p=0.001$; Fig. 5E). In contrast, DCX levels were similar between Cre⁺ and Cre⁻
437 male mice ($p=0.498$, Fig. 5E). Therefore, elevated DCX occurred after chronic seizures
438 had developed in Cre⁺ mice but the effect was limited to females. Because Cre⁺

439 epileptic females had increased immature neurons relative to Cre- females at the time
440 of SE, and prior studies show that Cre+ females had less neuronal damage after SE
441 (Jain et al. 2019), female Cre+ mice might have had reduced chronic seizures because
442 of high numbers of immature neurons. However, the data do not prove a causal role.

443 It is notable that the Cre+ male mice did not show increased numbers of
444 immature neurons at the time of chronic seizures but Cre+ females did. It is possible
445 that there was a “ceiling” effect in DCX expression that would explain why male Cre+
446 mice did not have a significant increase in immature neurons relative to male Cre- mice.

447

448 **IV. Hilar ectopic granule cells**

449 Based on the literature showing that reducing hilar ectopic GCs decreases
450 chronic seizures after pilocarpine-induced SE (Cho et al., 2015), we hypothesized that
451 female Cre+ mice would have fewer hilar ectopic GCs than female Cre- mice. However,
452 that female Cre+ mice did not have fewer hilar ectopic GCs.

453 To quantify hilar ectopic GCs we used Prox1 as a marker. Prox1 is a common
454 marker of GCs in the GCL (Pleasure et al. 2000; Galeeva et al. 2007; Galichet et al.
455 2008; Steiner et al. 2008; Iwano et al. 2012), and the hilus (Scharfman et al. 2007;
456 Hester and Danzer 2013; Cho et al. 2015; Bermudez-Hernandez et al. 2017).

457 Cre+ mice had significantly more hilar Prox1 cells than Cre- mice (Cre-: $19.6 \pm$
458 1.9 cells, $n=18$; Cre+: 60.5 ± 7.9 cells, $n=18$; Student's t-test, $t(34)=5.76$, $p<0.001$; Fig.
459 6C1). A two-way ANOVA with genotype and sex as main factors showed no effect of
460 sex ($F(1,32)=0.28$, $p=0.595$) but a significant effect of genotype ($F(1,32)=0.23$,
461 $p<0.0001$) with more hilar Prox1 cells in female Cre+ than female Cre- mice (Cre-
462 female: 18.2 ± 3.3 cells, $n=7$; Cre+ female: 57.0 ± 8.3 cells, $n=9$; Bonferroni's test,
463 $p<0.009$; Fig. 6C2) and the same for males (Cre- male: 20.4 ± 2.4 cells, $n=11$; Cre+
464 male: 63.9 ± 14.0 cells, $n=9$; Bonferroni's test, $p=0.001$; Fig. 6C2).

465 In past studies, hilar ectopic GCs have been suggested to promote seizures
466 (Scharfman et al. 2000; Jung et al. 2006; Cho et al. 2015). Therefore, we asked if the
467 numbers of hilar ectopic GCs correlated with the numbers of chronic seizures. When
468 Cre- and Cre+ mice were compared (both sexes pooled), there was a correlation with
469 numbers of chronic seizures (Fig. 6D1) but it suggested that more hilar ectopic GCs
470 improved rather than worsened seizures. However, the correlation was only in Cre-
471 mice, and when sexes were separated there was no correlation (Fig. 6D3).

472 When seizure-free interval was examined with sexes pooled, there was a
473 correlation for Cre+ mice (Fig. 6D2) but not Cre- mice. Strangely, the correlations of
474 Cre+ mice with seizure-free interval (Fig. 6D2, D4) suggest ectopic GCs shorten the
475 seizure-free interval and therefore worsen epilepsy, opposite of the correlative data for
476 numbers of chronic seizures. In light of these inconsistent results it seems that hilar
477 ectopic granule cells had no consistent effect on chronic seizures.

478

479 **V. Increased adult-born neurons preserves mossy cells and hilar SOM** 480 **interneurons but has little effect on parvalbumin interneurons**

481 It has been suggested that epileptogenesis after a brain insult like SE is due to
482 the hippocampal damage caused by the insult (Cavalheiro et al. 1996; Herman 2002;
483 Mathern et al. 2008; Dudek and Staley 2012; Dingledine et al. 2014). Therefore, one of
484 the reasons why increasing adult-born neurons reduced chronic seizures could be that

485 it reduced neuronal damage after SE. Indeed, that has been shown (Jain et al., 2019).
486 Here we examined the loss of vulnerable hilar mossy cells and SOM cells because they
487 have been suggested to be critical (Sloviter 1987; Cavazos and Sutula 1990; Cavazos
488 et al. 1994; Henshall and Meldrum 2012; Huusko et al. 2015). We asked whether Cre+
489 mice had preserved mossy cells (Fig. 7A) and SOM neurons (Fig. 7B). For comparison,
490 we quantified the relatively seizure-resistant parvalbumin-expressing GABAergic
491 neurons (Fig. 7C). An antibody to GluR2/3 was used as a marker of mossy cells
492 (Leranth et al. 1996) and a SOM antibody for SOM cells (Leranth et al. 1990;
493 Savanthrapadian et al. 2014; Botterill et al. 2019).

494 The results showed that Cre+ mice had more GluR2/3-expressing hilar cells than
495 Cre- mice (Cre-: 10.0 ± 1.8 cells, n=10; Cre+: 17.0 ± 2.0 cells, n=13; Student's t-test,
496 $t(21)=2.46$, $p=0.022$; Fig. 7A1-3). We confirmed that the GluR2/3+ hilar cells were not
497 double-labeled with Prox1, suggesting they corresponded to mossy cells, not hilar
498 ectopic GCs (Supplementary Fig. 8A). To investigate sex differences, a two-way
499 ANOVA was conducted with genotype and sex as main factors. There was a significant
500 effect of genotype ($F(1,18)=4.95$, $p=0.039$) with Cre+ females having more GluR2/3
501 cells than Cre- females (Cre- female: 8.0 ± 2.0 cells, n=6; Cre+ female: 19.1 ± 2.7 cells,
502 n=8; Bonferroni's test, $p=0.011$; Fig. 7A4). GluR2/3-ir hilar cells were similar in males
503 (Cre- male: 13.0 ± 3.0 cells, n=4; Cre+ male: 13.6 ± 2.5 cells, n=5; Bonferroni's test,
504 $p=0.915$; Fig. 7A4). These results in dorsal DG also were obtained in ventral DG
505 (Supplementary Fig. 8B-C). The data suggest that having more GluR2/3-ir mossy cells
506 could be a mechanism that allowed Cre+ females to have reduced chronic seizures
507 compared to Cre- females. Equal numbers of GluR2/3 mossy cells in Cre+ and Cre-
508 males could relate to their lack of protection against chronic seizures.

509 Next, we measured SOM hilar cells in pooled data (females and males together).
510 These results were analogous to the data for GluR2/3, showing that Cre+ mice had
511 more hilar SOM cells than Cre- mice (Cre-: 2.1 ± 0.5 cells, n=9; Cre+: 4.6 ± 0.6 cells,
512 n=11; Student's t-test, $t(18)=2.95$, $p=0.008$; Fig. 7B1-3). When sexes were separated, a
513 two-way ANOVA showed a significant effect of genotype ($F(1,16)=5.14$, $p=0.038$) and
514 no effect of sex ($F(1,18)=0.94$, $p=0.346$). However, Cre+ females had more SOM cells
515 than Cre- females (Cre- female: 2.2 ± 0.6 cells, n=6; Cre+ female: 5.1 ± 0.6 cells, n=8;
516 Bonferroni's test, $p=0.019$; Fig. 7B4), although only in dorsal DG (Fig. 7B4) not ventral
517 DG (Supplementary Fig. 8C). Numbers of SOM cells were similar in males (Cre- male:
518 2.2 ± 0.7 cells, n=3; Cre+ male: 3.3 ± 1.8 cells, n=3; Bonferroni's test, $p=0.897$; Fig.
519 7B4) in both dorsal and ventral DG (Supplementary Fig. 8B-C). Therefore, the ability to
520 preserve more mossy cells and SOM hilar cells in Cre+ females could be a mechanism
521 by which Cre+ females were protected from chronic seizures.

522 Parvalbumin-ir cells were not significantly different between genotypes (Student's
523 test, $t(19)=1.76$, $p=0.095$; Fig. 7C1-3). A two-way ANOVA showed no effect of genotype
524 ($F(1,17)=3.10$, $p=0.096$) or sex ($F(1,17)=0.26$, $p=0.616$) on the numbers of parvalbumin
525 cells. The results were the same in dorsal and ventral DG (Supplementary Fig. 8B-C).
526 These data are consistent with the idea that loss of parvalbumin-expressing cells has
527 not been considered to play a substantial in epileptogenesis in the past (Sloviter 1987;
528 1994). However, it should be noted that subsequent research has shown that the topic
529 is complicated because parvalbumin expression may decline even if the cells do not die

530 (Andre et al. 2001; Sun et al. 2007) and data vary depending on the animal model (van
531 Vliet et al. 2004; Huusko et al. 2015).

532

533 **VI. Increased adult-born neurons decreased neuronal damage after SE**

534 In our previous study of Cre⁺ and Cre⁻ mice (Jain et al., 2019), tamoxifen was
535 administered at 6 weeks and SE was induced at 12 weeks (like the current study). We
536 examined neuronal loss 3 days after SE, when neuronal loss in the hilus and area CA3
537 is robust in wild type mice. There is also some neuronal loss in CA1 at 3 days but more
538 at 10 days after SE. We found less neuronal loss in Cre⁺ mice in these three areas
539 (Jain et al. 2019). In the current study we examined 10 days after SE (Fig. 8A-B)
540 because at this time delayed neuronal loss occurs, providing a better understanding of
541 CA1 and the subiculum because delayed cell death occurs there. The intent was to
542 determine if Cre⁺ mice exhibited less neuronal loss or not in CA1 and the subiculum.

543 To quantify Fluor Jade-C, ROIs were drawn digitally around the pyramidal cell
544 layers (Fig. 8B). As shown in Fig. 8C, there was less Fluor Jade-C staining in Cre⁺
545 female mice relative to Cre⁻ female mice in both CA1 and the subiculum. The area of
546 the ROI that showed Fluor Jade C-positive cells was calculated as area fraction and
547 expressed as % in Fig. 8D. For females, a two-way ANOVA with genotype and subfield
548 as factors showed a significant effect of genotype ($F(1,14)=11.21$, $p=0.005$) with a
549 smaller area fraction in Cre⁺ mice than Cre⁻ mice for subiculum ($p=0.045$) and but not
550 CA1 ($p=0.095$; Fig. 8D1). Males showed no significant differences either in genotype
551 ($F(1,8)=0.06$, $p=0.816$) or subfield ($F(1,8)=0.05$, $p=0.825$; Fig. 8D2). When genotypes
552 were pooled, females were not different than males either for CA1 or the subiculum
553 (two-way ANOVA, sex, $F(1,37)=0.466$, $p=0.499$; Fig. 8D3). Thus, Cre⁺ female mice
554 were protected from hilar, CA3, and CA1 damage at 3 days and were protected from
555 subicular damage at 10 days after SE.

556

557 **DISCUSSION**

558 This study showed that conditional deletion of *Bax* from Nestin-expressing
559 progenitors increased young adult-born neurons in the DG when studied 6 weeks after
560 deletion and using DCX as a marker of immature neurons. In a different set of mice,
561 pilocarpine was used to induce epileptogenesis. The chronic seizures, measured 4-7
562 weeks after pilocarpine, were reduced in frequency by about 50% in females. Therefore,
563 increasing young adult-born neurons before the epileptogenic insult can protect against
564 epilepsy. However, we do not know if the protective effect was due to the greater
565 number of new neurons before SE or other effects. Past data would suggest that
566 increased numbers of newborn neurons before SE leads to a reduced SE duration and
567 less neuronal damage in the days after SE. That would be likely to lessen the epilepsy
568 after SE. However, there may have been additional effects of larger numbers of
569 newborn neurons prior to SE.

570 Increasing young adult-born neurons has been shown to protect the
571 hippocampus from SE-induced neuronal loss (Jain et al., 2019) which is a major
572 contributor to epileptogenesis. Therefore, by protecting against SE-induced neuronal
573 loss the young adult-born neurons could have reduced the severity of epilepsy. Indeed,
574 we showed that the Cre⁺ female mice that had reduced chronic seizures had

575 preservation of hilar mossy cells and SOM cells, two populations that are lost in SE-
576 induced epilepsy and considered to contribute to epileptogenesis.

577 There were major surprises in the current study. First, the results are unexpected
578 because suppressing adult-born neurons was shown to reduce chronic seizures (Cho et
579 al., 2015). Here, increasing adult-born neurons did not have the opposite effect.
580 It was also unanticipated that only females with *Bax* deletion showed a significant
581 increase in young adult-born neurons and a significant reduction in chronic seizures.
582 The larger effect of increasing adult-born neurons in female mice may be attributable to
583 sex differences in *Bax* (discussed below). The other remarkable finding relates to hilar
584 ectopic GCs. These GCs have been suggested to promote epilepsy (Scharfman 2004;
585 Jung et al. 2006; Scharfman et al. 2007; Parent and Murphy 2008; Hester and Danzer
586 2013; Cho et al. 2015), but hilar ectopic GCs increased in females with reduced
587 seizures. The association of more hilar ectopic GCs with fewer chronic seizures was
588 unexpected.

589

590 **Effects of *Bax* deletion on SE**

591 In past studies, suppressing adult-born neurons made kainic acid-induced SE
592 worse, and pilocarpine-induced SE was also worse (Iyengar et al., 2015; Jain et al.,
593 2019). In the present study, SE was affected also. The duration of SE was reduced in
594 Cre+ mice. In the Cre+ females, the first seizure after pilocarpine injection was often
595 less severe, and power showed a tendency to be reduced during SE. Therefore, SE
596 might have been less severe in the Cre+ females, and this could have contributed to
597 reduced neuronal loss and chronic seizures.

598

599 **Chronic seizures**

600 It is remarkable that increasing adult-born neurons for 6 weeks was sufficient to
601 reduce seizures long-term. It is consistent with the idea that normally the young adult-
602 born neurons inhibit other GCs, which supports the DG gate function (Hsu 2007; Drew
603 et al. 2016). This gate has been suggested to be an inhibitory barrier to entry of
604 seizures from cortex into hippocampus (Coulter and Carlson 2007; Hsu 2007; Krook-
605 Magnuson et al. 2015). That entry is deleterious because seizures that pass from
606 entorhinal cortex to the GCs and then CA3 are likely to continue to CA1 and back to
607 cortex, causing reverberatory (long-lasting, severe) seizures. The reason for the
608 relatively ease of reverberation once past the DG gate is that the synapses between
609 GCs and CA3, CA1 and cortex are excitatory. The GCs have especially powerful
610 excitatory synapses on CA3 pyramidal cells (Henze et al. 2000; Scharfman and
611 MacLusky 2014), although these are normally mitigated by GABAergic circuitry (Acsady
612 et al. 1998).

613 These data are consistent with the demonstration that adult-born neurons protect
614 against other pathological conditions such as Alzheimer's disease (Choi et al. 2018;
615 Choi and Tanzi 2019). However, it is important to note that all effects are unlikely to be
616 mediated only by the DG. The olfactory bulb and other areas also have adult-born
617 neurons and they could contribute to epilepsy, especially those epilepsy syndromes with
618 mechanisms that are extrahippocampal.

619

620

621 **Clusters of seizures**

622 There were fewer days with > 3 seizures in Cre+ female mice which is another
623 way that Cre+ females were protected from chronic seizures. These findings are
624 valuable because clusters in humans have a significantly negative impact on health and
625 quality of life (Haut 2015; Jafarpour et al. 2019).

626 The results may have underestimated the effects on clusters because we did not
627 measure the interval between clusters in many Cre+ female mice. The reason is that the
628 interval between clusters increased in some mice so they only had one cluster in 3
629 weeks. Thus intercluster interval appeared to lengthen in Cre+ females but animals with
630 only one cluster had to be excluded. In the end, the results were not statistically
631 significant.

632

633 **Sex differences**

634 Females showed more of an effect of conditional *Bax* deletion than males. Insight
635 into this sex difference came when the same assessments were made before SE
636 Before SE, there was no sex difference. Cre+ females had more adult-born neurons
637 than Cre- females and Cre+ males had more than Cre- males. In addition, the levels of
638 DCX were similar in Cre+ females and Cre+ males.

639 However, after epilepsy developed, there was a sex difference. Cre- females had
640 less DCX than Cre- males. One explanation is that cell birth during epileptogenesis was
641 greater in males because it is in developing hippocampus (Sisk et al. 2016) and SE has
642 been suggested to rekindle developmental programs (Ben-Ari and Holmes 2006).
643 Another possibility is males had less programmed cell death during epileptogenesis.
644 Indeed during development, females have more apoptotic profiles than males and the
645 sex difference was blocked by *Bax* deletion (Forger et al. 2004). A final possibility is
646 that cell death during epileptogenesis is *Bax*-dependent in females but *Bax*-independent
647 in males. Support for this idea comes from studies of ischemic cell death, which is
648 caspase-dependent cell death in females but not males (Siegel and McCullough 2011).

649

650 **Hilar ectopic GCs**

651 In the normal brain, adult-born neurons in the DG are thought to arise mainly
652 from the SGZ and migrate to the GCL (Kempermann 2012). After SE, there is a surge in
653 proliferation in the SGZ and neurons either migrate correctly to the GCL or aberrantly in
654 the hilus (Parent et al. 1997).

655 These hilar ectopic GCs are thought to contribute to seizure generation in the
656 epileptic brain because they are innervated by residual CA3 neurons, and project to
657 GCs, making a major contribution to mossy fiber innervation of GCs in the inner
658 molecular layer (Scharfman et al. 2000; Kron et al. 2010; Pierce et al. 2011; Scharfman
659 and Pierce 2012; Althaus et al. 2016). When epileptiform activity occurs in CA3 in slices
660 of epileptic rats, CA3 evokes discharges in hilar ectopic GCs that in turn excite GCs in
661 the GCL (Scharfman et al. 2000). Consistent with the idea that hilar ectopic GCs
662 promote seizures, the numbers of hilar ectopic GCs are correlated with chronic seizure
663 frequency in rats (McCloskey et al. 2006) and mice (Hester and Danzer 2013).
664 Furthermore, suppressing hilar ectopic GC formation reduces chronic seizures (Jung et
665 al. 2006; Cho et al. 2015; Hosford et al. 2016).

666 Notably, this is the first study to our knowledge showing that increased hilar
667 ectopic GCs were found in mice that had reduced seizures. One potential explanation is
668 that SE-induced hippocampal damage was reduced in Cre+ females with high numbers
669 of hilar ectopic GCs. Therefore, the circuitry of the DG would be very different compared
670 to past studies of hilar ectopic GCs where neuronal loss was severe (Supplementary
671 Fig. 7). The presence of mossy cells is one way the circuitry would be different. MCs
672 normally support the young adult-born GCs that migrate to the GCL (Piatti and Schinder
673 2018). Mossy cells provide an important activator of newborn GCs when they are young
674 (Chancey et al. 2014). Mossy cells also innervate hilar ectopic GCs (Pierce et al. 2007).

675 Another possibility is that there was protection against chronic seizures in female
676 Cre+ mice by increasing adult-born neurons *in the GCL*. The reason to suggest this
677 possibility is that prior studies showed that young adult-born neurons in the GCL
678 primarily inhibit GCs in the normal brain (Drew et al. 2016) and are relatively quiescent
679 in the epileptic brain (Jakubs et al. 2006) .

680

681 **Additional considerations**

682 This study is limited by the possibilities of type II statistical errors in those
683 instances where we divided groups by genotype and sex, leading to comparisons of 3-5
684 mice/group. Another potential caveat is that female mice were selected regardless of
685 the stage of the estrous cycle.

686

687 **Conclusions**

688 In the past, suppressing adult neurogenesis before SE was followed by fewer
689 hilar ectopic GCs and reduced chronic seizures. Here, we show that the opposite -
690 enhancing adult-born neurons before SE and increased hilar ectopic GCs - do not
691 necessarily reduce seizures. We suggest instead that protection of the hilar neurons
692 from SE-induced excitotoxicity was critical to reducing seizures. The reason for the
693 suggestion is that the survival of hilar neurons would lead to persistence of the normal
694 inhibitory functions of hilar neurons, protecting against seizures. However, this is only a
695 suggestion at the present time because we do not have data to prove it. Additionally,
696 because protection was in females, sex differences are likely to have played an
697 important role. Regardless, the results show that enhancing-born neurons of young
698 adult-born neurons in Nestin-Cre+ mice had a striking effect in the pilocarpine model,
699 reducing chronic seizures in female mice.

700

701 **MATERIALS AND METHODS**

702

703 **I. General information**

704 Animal care and use was approved by the Nathan Kline Institute Institutional
705 Animal Care and Use Committee and met the regulations of the National Institute of
706 Health and the New York State Department of Health. Mice were housed in standard
707 mouse cages, with a 12 hr light/dark cycle and food (Laboratory rodent diet 5001; W.F.
708 Fisher & Sons) and water *ad libitum*. During gestation and until weaning, mice were fed
709 chow formulated for breeding (Formulab diet 5008; W.F. Fisher & Sons).

710

711

712 **II. Increasing adult-born neurons**

713 To enhance-born neurons, a method was used that depends on deletion of *Bax*,
714 the major regulator of programmed cell death in adult-born neurons (Sun et al. 2004;
715 Sahay et al. 2011a; Ikrar et al. 2013; Adlaf et al. 2017). Enhancement of-born neurons
716 was induced by conditional deletion of *Bax* from Nestin-expressing progenitors (Sahay
717 et al. 2011a). These mice were created by crossing mice that have *loxP* sites flanking
718 the pro-apoptotic gene *Bax* (*Bax^{fl/fl}*) with a Nestin-CreER^{T2} mouse line in which
719 tamoxifen-inducible Cre recombinase (CreER^{T2}) is expressed under the control of the
720 rat *Nestin* promoter (Sahay et al. 2011a). It was shown that after tamoxifen injection in
721 adult mice there is an increase in dentate gyrus neurogenesis based on studies of
722 bromo-deoxyuridine, Ki67, and doublecortin (Sahay et al. 2011a). The Nestin-
723 CreER^{T2}*Bax^{fl/fl}* mouse line was kindly provided by Drs. Amar Sahay and Rene Hen and
724 used and described previously by our group (Bermudez-Hernandez et al. 2017; Jain et
725 al. 2019). Although Nestin-Cre-ER^{T2} mouse lines have been criticized because they
726 can have leaky expression, the mouse line used in the present study did not (Sun et al.
727 2014), which we confirmed (Jain et al. 2019).

728 Starting at 6 weeks of age, mice were injected subcutaneously (s.c.) with
729 tamoxifen (dose 100 mg/kg, 1/day for 5 days; Cat# T5648, Sigma-Aldrich). Tamoxifen
730 was administered from a stock solution (20 mg/ml in corn oil, containing 10% absolute
731 alcohol; Cat# C8267, Sigma-Aldrich). Tamoxifen is light-sensitive so it was stored at 4°C
732 in an aluminum foil-wrapped container for the duration of treatment (5 days).

733

734 **III. Pilocarpine-induced SE**

735 Six weeks after the last dose of tamoxifen injection, mice were injected with
736 pilocarpine to induce SE. Methods were similar to those used previously (Jain et al.
737 2019). On the day of pilocarpine injection, there were 2 initial injections of pre-
738 treatments and then one injection of pilocarpine. The first injection of pre-treatments
739 was a solution of ethosuximide (150 mg/kg of 84 mg/ml in phosphate buffered saline,
740 s.c.; Cat# E;7138, Sigma-Aldrich). Ethosuximide was used because the background
741 strain, C57BL6/J, is susceptible to respiratory arrest during a severe seizure and
742 ethosuximide decreases the susceptibility (Iyengar et al. 2015). The second injection of
743 pre-treatments was a solution of scopolamine methyl nitrate (1 mg/kg of 0.2 mg/ml in
744 sterile 0.9% sodium chloride solution, s.c.; Cat# 2250, Sigma-Aldrich) and terbutaline
745 hemisulfate (1 mg/kg of 0.2 mg/ml in sterile 0.9% sodium chloride solution, s.c.; Cat#
746 T2528, Sigma-Aldrich). Scopolamine is a muscarinic cholinergic antagonist and when
747 injected as methyl nitrate it does not cross the blood brain barrier. Therefore,
748 scopolamine decreased peripheral cholinergic side effects of pilocarpine without
749 interfering with central actions of pilocarpine. Terbutaline was used to keep airways
750 patent during severe seizures, minimizing mortality. Ethosuximide had to be
751 administered separately because it precipitates when mixed with scopolamine and
752 terbutaline.

753 Thirty min after the pre-treatments, pilocarpine hydrochloride was injected (260-
754 280 mg/kg of 50 mg/ml in sterile 0.9% sodium chloride solution, s.c.; Cat# P6503;
755 Sigma-Aldrich). Different doses were used because different batches of pilocarpine had
756 different ability to elicit SE.

757 The severity of SE was decreased by administering the benzodiazepine
758 diazepam (10 mg/kg of 5 mg/ml stock solution, s.c.; NDC# 0409-3213-12, Hospira, Inc.)
759 2 hr after pilocarpine injection. In females, diazepam was injected earlier, 40 minutes
760 after the onset of first seizure, because in the first group of females in which diazepam
761 was injected 2 hr after pilocarpine, there was severe brain damage. While sedated with
762 diazepam, animals were injected with warm (31°C) lactated Ringer's solution (s.c.;
763 NDC# 07-893-1389, Aspen Veterinary Resources). At the end of the day, mice were
764 injected with ethosuximide using the same dose as before pilocarpine. For the next 3
765 days, chow was provided that was moistened with water. The cage was placed on a
766 heating blanket to maintain cage temperature at 31°C.

767

768 **IV. Stereotaxic surgery**

769 A. General information

770 Mice were anesthetized by isoflurane inhalation (3% isoflurane for induction and
771 1.75 - 2% isoflurane for maintenance during surgery; NDC# 07-893-1389, Patterson
772 Veterinary) and placed in a stereotaxic apparatus (David Kopf Instruments). Prior to
773 surgery, the analgesic Buprenex (Buprenorphine hydrochloride; NDC# 1296-0757-5;
774 Reckitt Benckheiser) was diluted in sterile saline (0.9% sodium chloride solution) to
775 yield a 0.03 mg/ml stock solution and 0.2 mg/kg was injected s.c. During surgery, mice
776 were placed on a heating blanket with a rectal probe for automatic maintenance of body
777 temperature at 31°C.

778

779 B. Implantation of EEG electrodes

780 Before electrode implantation, hair over the skull was shaved and then the scalp
781 was cleaned with 70% ethanol. A midline incision was made to expose the skull with a
782 sterile scalpel. To implant subdural screw electrodes (0.10" stainless steel screws, Cat#
783 8209, Pinnacle Technology), 6 holes were drilled over the exposed skull. The
784 coordinates were: right occipital cortex (anterior-posterior or AP -3.5 mm from Bregma,
785 medio-lateral or ML, 2.0 mm from the midline); left frontal cortex (Lt FC, AP -0.5 mm;
786 ML -1.5 mm); left hippocampus (AP -2.5 mm; ML -2.0 mm) and right hippocampus (AP
787 -2.5 mm; ML 2.0 mm). An additional screw was placed over the right olfactory bulb as
788 ground (AP 2.3 mm; ML 1.8 mm) and another screw over the cerebellum at the midline
789 as reference (relative to Lambda: AP -1.5 mm; ML -0.5 mm). Here, "ground" refers to
790 the earth ground and "reference" refers to the reference for all 4 screw electrode
791 recordings (Moyer et al. 2017). An 8-pin connector (Cat# ED85100-ND, Digi-Key
792 Corporation) was placed over the skull and secured with dental cement (Cat# 51459,
793 Dental Cement Kit; Stoelting Co.).

794 After surgery, mice were injected with 50 ml/kg warm (31°C) lactated Ringer's
795 solution (s.c.; NDC# 09355000476, Aspen Veterinary Resources). Mice were housed a
796 clean cage on a heating blanket for 24 hr. Moistened food pellets were placed at the
797 base of the cage to encourage food intake. Afterwards mice were housed individually
798 because group housing leads to disturbance of the implant by other mice in the cage.

799

800

801

802

803 **V. Continuous Video-EEG recording and analysis**

804 A. Video-EEG recording

805 Mice were allowed 3 weeks to recover from surgery. During this time, mice were
806 housed in the room where video-EEG equipment are placed so that mice could
807 acclimate to the recording environment. To record video-EEG, the pin connector on the
808 head of the mouse was attached to a preamplifier (Cat# 8406, Pinnacle Technology)
809 which was attached to a commutator (Cat# 8408, Mouse Swivel/Commutator, 4-
810 channel, Pinnacle Technology) to allow freedom of movement. Signals were acquired at
811 a 500 Hz sampling rate, and band-pass filtered at 1-100 Hz using Sirenia Acquisition
812 software (<https://www.pinnaclet.com>, RRID:SCR_016183). Video was captured with a
813 high-intensity infrared LED camera (Cat# PE-605EH, Pecham) and was synchronized to
814 the EEG record.

815 To monitor pilocarpine-induced SE, video-EEG was recorded for hour before ad
816 24 hr after pilocarpine injection. Approximately 5-6 weeks after pilocarpine-induced SE,
817 video-EEG was recorded to measure spontaneous recurrent seizures. Video-EEG was
818 recorded continuously for 3 weeks.

819

820 B. Video-EEG analysis

821 EEG was analyzed offline with Sirenia Seizure Pro, V2.0.7 (Pinnacle Technology,
822 RRID:SCR_016184). A seizure was defined as a period of rhythmic (>3 Hz) deflections
823 that were >2x the standard deviation of baseline mean and lasted at least 10 sec (Jain
824 et al. 2019). Seizures were rated as convulsive if an electrographic seizure was
825 accompanied by a behavioral convulsion (observed by video playback), defined as
826 stages 3-5 using the Racine scale (Racine 1972) where stage 3 is unilateral forelimb
827 clonus, stage 4 is bilateral forelimb clonus with rearing, and stage 5 is bilateral forelimb
828 clonus followed by rearing and falling. A seizure was defined as non-convulsive when
829 there was electrographic evidence of a seizure but there were no stage 3-5 behavior.

830 SE was defined as continuous seizures for >5 min (Chen and Wasterlain 2006)
831 and EEG amplitude in all 4 channels >3x the baseline mean. For mice without EEG, SE
832 was defined by stage 3-5 seizures that did not stop with a resumption of normal
833 behavior. Often stage 3-5 seizures heralded the onset of SE and then occurred
834 intermittently for hours. In between convulsive behavior mice had twitching of their body,
835 typically in a prone position.

836 SE duration was defined in light of the fact that the EEG did not return to normal
837 after the initial period of intense activity. Instead, intermittent spiking occurred for at
838 least 24 hrs, as we previously described (Jain et al. 2019) and has been described by
839 others (Mazzuferi et al. 2012; Bumanglag and Sloviter 2018; Smith et al. 2018). We
840 therefore chose a definition that captured the initial, intense activity. We defined the end
841 of this time as the point when the amplitude of the EEG deflections were reduced to
842 50% or less of the peak deflections during the initial hour of SE. Specifically, we
843 selected the time after the onset of SE when the EEG amplitude in at least 3 channels
844 had dropped to approximately 2 times the amplitude of the EEG during the first hour of
845 SE, and remained depressed for at least 10 min (Fig S2 in (Jain et al. 2019). Thus, the
846 duration of SE was defined as the time between the onset and this definition of the
847 "end" of SE.

848 To access the severity of chronic seizures, frequency and duration of seizures
849 were measured during the 3 weeks of EEG recording. Inter-cluster interval was defined
850 as the maximum number of days between two clusters.

851

852 **VI. Tissue processing**

853 **A. Perfusion-fixation and sectioning**

854 Mice were perfused after video-EEG recording. To perfuse, mice were deeply
855 anesthetized by isoflurane inhalation followed by urethane (250 mg/kg of 250 mg/ml in
856 0.9% sodium chloride, intraperitoneal, i.p.; Cat#U2500; Sigma-Aldrich). After loss of a
857 reflex to a tail pinch, and loss of a righting reflex, consistent with deep anesthesia, the
858 heart cavity was opened, and a 25-gauge needle inserted into the heart, followed by
859 perfusion with 10 ml saline (0.9% sodium chloride in double-distilled water (ddH₂O))
860 using a peristaltic pump (Minipuls 1; Gilson) followed by 30 ml of cold (4°C) 4%
861 paraformaldehyde (PFA; Cat# 19210, Electron Microscopy Sciences) in 0.1 M
862 phosphate buffer (PB; pH 7.4). The brains were removed immediately, hemisected, and
863 post-fixed for at least 24 hr in 4% PFA at 4°C. After post-fixation, one hemisphere was
864 cut in the coronal plane and the other in the horizontal plane (50 µm-thick sections)
865 using a vibratome (Cat# TPI-3000, Vibratome Co.). Sections were collected sequentially
866 to select sections that were from similar septotemporal levels. For dorsal hippocampus,
867 coronal sections were selected every 300 µm starting at the first section where the DG
868 blades are fully formed (between AP -1.94 and -2.06 mm). Horizontal sections were
869 chosen every 300 µm starting from the temporal pole at the place where the GCL is
870 clearly defined (between DV 0.84 and 1.08 mm). This scheme is diagrammed and
871 described in more detail in prior studies (Moretto et al. 2017).

872

873 **B. Doublecortin**

874 *1) Procedures for staining*

875 Doublecortin (DCX), a microtubule-associated protein (Gleeson et al. 1999), was
876 used to identify immature adult-born neurons (Brown et al. 2003; Couillard-Despres et
877 al. 2005), and was stained after antigen retrieval (Botterill et al. 2015). First, free floating
878 sections were washed in 0.1 M Tris buffer (TB, 3 × 5 min). Sections were then
879 incubated in sodium citrate (Cat# S4641, Sigma-Aldrich) buffer (2.94 mg/ml in ddH₂O,
880 pH 6.0 adjusted with HCl) in a preheated water bath at 85°C for 30 min. Sections were
881 washed with 0.1 M TB (3 × 5 min), blocked in 5% goat serum (Cat# S-1000,
882 RRID:AB_2336615, Vector Laboratories) in 0.1 M TB with 0.5% (v/v) Triton X-100 (Cat#
883 X-100, Sigma-Aldrich) and 1% (w/v) bovine serum albumin for 1 hr. Next, sections were
884 incubated overnight with primary antibody (1:1000 diluted in blocking serum,
885 monoclonal anti-rabbit DCX; Cat#4604S, Cell Signaling Technology) on a shaker
886 (Model# BDRAA115S, Stovall Life Science Inc.) at room temperature.

887 On the next day, sections were washed in 0.1 M TB (3 × 5 min), treated with
888 2.5% hydrogen peroxide (Cat# 216763, Sigma-Aldrich) for 30 min to block endogenous
889 peroxidase, and washed with 0.1 M TB (3 × 5 min). Next, sections were incubated in
890 secondary antibody (biotinylated goat anti-rabbit IgG, 1:500, Vector Laboratories) for 1
891 hr in 0.1 M TB, followed by washes with 0.1 M TB (3 × 5 min). Sections were then
892 incubated in avidin-biotin complex (1:500 in 0.1 M Tris buffer; Cat# PK-6100, Vector) for
893 2 hr, washed in 0.1 M TB (1 × 5 min) and then in 0.175 M sodium acetate (14.36 mg/ml

894 in ddH₂O, pH 6.8, adjusted with glacial acetic acid, 2 × 5 min; Cat# S8750, Sigma-
895 Aldrich). Sections were reacted in 0.5 mg/ml 3, 3'-diaminobenzidine (DAB; Cat# D5905,
896 Sigma-Aldrich) with 40 µg/ml ammonium chloride (Cat# A4514, Sigma-Aldrich), 3 µg/ml
897 glucose oxidase (Cat# G2133, Sigma-Aldrich), 2 mg/ml (D+)-glucose (Cat# G5767,
898 Sigma-Aldrich) and 25 mg/ml ammonium nickel sulfate (Cat# A1827, Sigma-Aldrich) in
899 0.175 M sodium acetate. Sections were washed in 0.175 M sodium acetate (2 × 5 min)
900 and 0.1 M TB (5 min), mounted on gelatin-coated slides (1% bovine gelatin; Cat#
901 G9391, Sigma-Aldrich), and dried overnight at room temperature.

902 On the next day, sections were dehydrated with increasing concentrations of
903 ethanol, cleared in Xylene (Cat# 534-56, Sigma-Aldrich), and coverslipped with
904 Permount (Cat# 17986-01, Electron Microscopy Sciences). Sections were viewed with a
905 brightfield microscope (Model BX51; Olympus of America) and images were captured
906 with a digital camera (Model Infinity3-6URC, Teledyne Lumenera).

907

908 2) DCX Analysis

909 DCX was quantified by first defining a region of interest (ROI) that included the
910 adult-born cells and the majority of their DCX-labeled dendrites: the SGZ, GCL, and
911 inner molecular layer. The SGZ was defined as a region that extended from the GCL
912 into the hilus for a width of 100 µm because this region included the vast majority of the
913 DCX immunoreactivity. The inner molecular layer was defined as the 100 µm
914 immediately above the GCL. Next, a threshold was selected where DCX-
915 immunoreactive (ir) cells were above, but the background was below threshold, as
916 described in more detail elsewhere (Jain et al. 2019).

917 This measurement is referred to as area fraction in the Results and expressed as
918 a percent. For a given animal, the area fraction was determined for 3 coronal sections in
919 the dorsal hippocampus between AP -1.94 to -2.06 mm and 3-4 horizontal sections in
920 the ventral hippocampus between DV 0.84 to 1.08 mm, with sections spaced 300 µm
921 apart. These area fractions were averaged so that a mean area fraction was defined for
922 each animal. For these and other analyses described below, the investigator was
923 blinded.

924

925 C. Prox-1

926 1) Procedures for staining

927 In normal rodent adult brain, prospero homeobox 1 (Prox1) is expressed in the
928 GCs (Pleasure et al. 2000) and in the hilus (Bermudez-Hernandez et al. 2017). To stain
929 for Prox1, free-floating sections were washed in 0.1 M TB pH 7.4, 3×5 min). Sections
930 were then incubated in 0.1 M TB with 0.25% Triton X-100 for 30 min followed by a 10
931 min-long wash in 0.1 M TB with 0.1% Triton X-100 (referred to as Tris A). Next, sections
932 were treated with 1% hydrogen peroxide in Tris A for 5 min followed by a 5 min-long
933 wash in Tris A. Sections were blocked in 10% normal horse serum (Cat# S-2000,
934 RRID:AB_2336617, Vector) in Tris A for 1 hr followed by a 10 min-long wash in Tris A
935 and then 0.1 M TB with 0.1% Triton X-100 and 0.005% bovine serum albumin (referred
936 to as Tris B). Next, sections were incubated overnight with primary antibody (goat anti-
937 human Prox1 polyclonal antibody, 1:2,000 diluted in Tris B, R and D systems) rotated
938 on a shaker (described above) at room temperature.

939 On the next day, sections were washed in Tris A then in Tris B (5 min each).
940 Sections were then incubated in secondary antibody (biotinylated anti-goat IgG made in
941 horse, 1:500, Vector Laboratories, see Table 2) for 1 hr in Tris B, followed by a wash
942 with Tris A (5 min) and then Tris B (5 min), blocked in avidin-biotin complex (1:500 in
943 Tris B) for 2 hr, and washed in 0.1 M TB (3 x 5 min). Sections were reacted in 0.5 mg/ml
944 3, 3'-diaminobenzidine (DAB) with 40 µg/ml ammonium chloride, 3 µg/ml glucose
945 oxidase, 2 mg/ml (D+)-glucose and 5mM nickel chloride (Cat# N6136, Sigma-Aldrich) in
946 0.1 M TB. Sections were washed in 0.1 M TB (3 x 5 min), mounted on 1% gelatin-
947 coated slides and dried overnight at room temperature. On the next day, sections were
948 dehydrated, cleared, and coverslipped (as described above). Sections were viewed and
949 images were captured as DCX above.

950

951 2) *Prox1 Analysis*

952 Prox1 was quantified in the hilus, defined based on zone 4 of Amaral (Amaral
953 1978). The definition of Amaral was modified to exclude 20 µm below the GCL
954 (Winawer et al. 2007). The GCL boundary was defined as the location where GCs
955 stopped being contiguous. Practically that meant there was no GC with more than a cell
956 body width of cell-free space around it. A cell body width was 10 µm (Claiborne et al.
957 1990; Amaral et al. 2007).

958 CA3c was included in the ROI but hilar Prox1 cells have not been detected in the
959 CA3c layer (Scharfman et al. 2000; Winawer et al. 2007). However, there are rare GCs
960 in CA3 according to one study (Szabadics et al. 2010).

961 In ImageJ, a ROI was traced in the image taken at 20x magnification and then a
962 threshold was selected where Prox1-immunoreactivity was above the background
963 threshold (Jain et al. 2019). Then Prox1 cells were counted using the Analyzed particle
964 plugin where a particle with an area $\geq 10 \mu\text{m}^2$ was counted. The following criteria were
965 used to define a hilar Prox1-ir cell (Bermudez-Hernandez et al. 2017): (1) the hilar cell
966 had sufficient Prox1-ir to reach a threshold equal to the average level of Prox1-ir of GCs
967 in the adjacent GC layer, (2) All hilar Prox-ir cells were complete, i.e., not cut at the
968 edge of the ROI. When hilar Prox1-ir cells were in clusters, although not many (typically
969 2–3 per 50 µm section), cells were counted manually. For each animal 3 coronal
970 sections in the dorsal hippocampus and 3-4 horizontal sections in the ventral
971 hippocampus, with sections spaced 300 µm apart were chosen.

972

973 D. Immunofluorescence

974 1) *Procedures for staining*

975 Free floating sections were washed (3x5 min) in 0.1 M TB followed by a 10-min
976 long wash in Tris A and another 10 min-long wash in Tris B. Sections were incubated in
977 blocking solution (5% normal goat serum or donkey serum in Tris B) for 1 hr at room
978 temperature. Next, primary antibodies for anti-rabbit GluR2/3, anti-goat Prox1, anti-
979 rabbit SOM and anti-mouse parvalbumin (Table 1) were diluted in blocking solution and
980 sections were incubated for 48 hr at 4°C. For SOM labelling, antigen retrieval was used.
981 Prior to the blocking step, sections were incubated in sodium citrate buffer (2.94 mg/ml
982 in ddH₂O, pH 6.0 adjusted with HCl) in a preheated water bath at 85°C for 30 min.

983 Next, sections were washed in Tris A and Tris B (10 min each) followed by 2 hr-
984 long incubation in secondary antibody (1:500 in Tris B, see Table 2). Sections were

985 washed in 0.1 M TB (3 x 5 min), and coverslipped with Citifluor™ AF1 mounting solution
986 (Cat# 17970-25, Vector Labs). Images were captured on a confocal microscope (Model
987 LSM 510 Meta; Carl Zeiss Microimaging).

988

989 2) Procedures for analysis

990 GluR2/3-, SOM- and parvalbumin- ir cells in the hilus and SGZ were counted
991 from 3 dorsal and 3 ventral sections. Sections were viewed at 40x of the confocal
992 microscope for manual counts. Because ectopic GCs express GluR2/3, sections were
993 co-labelled with Prox1. All co-labelled cells were considered as ectopic and excluded
994 from the GluR2/3- ir cell counting to measure mossy cells.

995

996 E. Fluorojade-C

997 1) Procedures for staining

998 Fluorojade-C (FJ) is a fluorescent dye that is the “gold standard” to stain
999 degenerating neurons (Schmued and Hopkins 2000; Schmued et al. 2005). First,
1000 sections were mounted on gelatin-coated slides (1% porcine gelatin in ddH2O; Cat#
1001 G1890, Sigma-Aldrich) and dried on a hot plate at 50–55°C for 1 hr. Then slides were
1002 placed in a staining rack and immersed in a 100% ethanol solution for 5 min, then in
1003 70% ethanol for 2 min, followed by a 1 min wash in ddH2O.

1004 Slides were then incubated in 0.06% potassium permanganate (Cat# P-279,
1005 Fisher Scientific) solution for 10 min on a shaker (described above) with gentle
1006 agitation, followed by washes in ddH2O (2 x 1 min). Slides were then incubated for 20
1007 min in a 0.0002% solution of FJ in ddH2O with 0.1% acetic acid in the dark. The stock
1008 solution of FJ was 0.01% in ddH2O and was stored at 4°C for up to 3 months. To
1009 prepare a working solution, 6 ml of stock solution was added to 294 mL of 0.1% acetic
1010 acid (Cat# UN2789, Fisher Scientific) in ddH2O and used within 10 min of preparation.
1011 Slides were subsequently protected from direct light. They were washed in ddH2O (3 x
1012 1 min) and dried overnight at room temperature. On the next day, slides were cleared in
1013 Xylene (2 x 3 min) and coverslipped with DPX mounting medium (Cat# 44581, Sigma-
1014 Aldrich). Sections were photographed with an epifluorescence microscope (Model
1015 BX51; Olympus of America) and images were captured with a digital camera (Model
1016 Infinity3-6URC, Teledyne Lumenera).

1017

1018 2) Procedures for analysis

1019 We measured the FJ in the cell layers of CA1 and CA3. Manual counting of FJ-
1020 positive (FJ+) cells was not possible in these cell layers because there could be so
1021 many FJ+ cells that were overlapping. Instead, FJ staining in cell layers was quantified
1022 by first outlining the cell layer as a ROI at 10x magnification in ImageJ as before (Jain et
1023 al. 2019).

1024 To outline the CA1 cell layer, the border with CA2 was defined as the point where
1025 the cell layer changed width, a sudden change that could be appreciated by the
1026 background in FJ-stained sections and confirmed by cresyl violet-stained sections. The
1027 border of CA1 and the subiculum was defined as the location where the normally
1028 compact CA1 cell layer suddenly dispersed. To outline CA3, the border with CA2 and
1029 CA3 was defined by the point where stratum lucidum of CA3 terminated. This location
1030 was distinct in its background in FJ-stained sections. The border of CA3 and the hilus

1031 was defined according to zone 4 of Amaral (Amaral 1978). This location was also
1032 possible to detect in FJ-stained sections because the background in the hilus was
1033 relatively dark compared to area CA3.

1034 After defining ROIs, a threshold fluorescence level was selected so that all cells
1035 that had very bright immunofluorescence were above threshold but other cells that were
1036 similar in fluorescence to background staining were not (Iyengar et al. 2015; Jain et al.
1037 2019). ImageJ was then used to calculate the area within the ROI and this
1038 measurement is referred to as area fraction in the Results and expressed as a percent.
1039 For a given animal, the area fraction was determined for three coronal sections in the
1040 dorsal hippocampus between AP -1.94 and -2.06 mm and 3–4 horizontal sections in
1041 the ventral hippocampus between DV 0.84 and 1.08 mm, with sections spaced $300\ \mu\text{m}$
1042 apart. These area fractions were averaged so that a mean area fraction was defined for
1043 each animal.

1044

1045 **VI. Statistical Analysis**

1046 Data are presented as the mean \pm standard error of the mean (SEM). Statistical
1047 analyses were performed using GraphPad Prism Software ([https://www.graphpad.com/
1048 scientific-software/prism/](https://www.graphpad.com/scientific-software/prism/), RRID: SCR_002798). Statistical significance was set at $p <$
1049 0.05 . Robust regression and Outlier removal (ROUT) method was used to remove
1050 outliers with ROUT coefficient Q set at 1%. Parametric tests were used when data fit a
1051 normal distribution, determined by the D'Agostino and Pearson or Shapiro-Wilk's
1052 normality tests, and there was homoscedasticity of variance (confirmed by a F-test). A
1053 Student's unpaired two-tailed t-test was used to assess differences between two
1054 groups. A Welch's test was used instead of a Student's t-test when there was
1055 heteroscedasticity of variance. One-way Analysis of Variance (ANOVA), two-way ANOVA,
1056 and three-way ANOVA were performed when there were multiple groups and were
1057 followed by Bonferroni's multiple comparison post-hoc test (Bonferroni's test). The main
1058 factors for two-way ANOVA were genotype and sex; region was added as another
1059 factor for three-way ANOVA. Interaction between factors is reported in the Results if it
1060 was significant. A Fisher's exact test was used for comparing proportions of binary data
1061 (yes/no). Pearson's Correlation was used to assess the association between 2
1062 variables.

1063 For data that did not follow a normal distribution, typically some data had a 0
1064 value. In these cases, non-parametric tests were selected. The Mann-Whitney U test
1065 was used to compare two groups, and a Kruskal-Wallis test followed by post-hoc
1066 Dunn's test was used for multiple groups comparison.

1067

1068

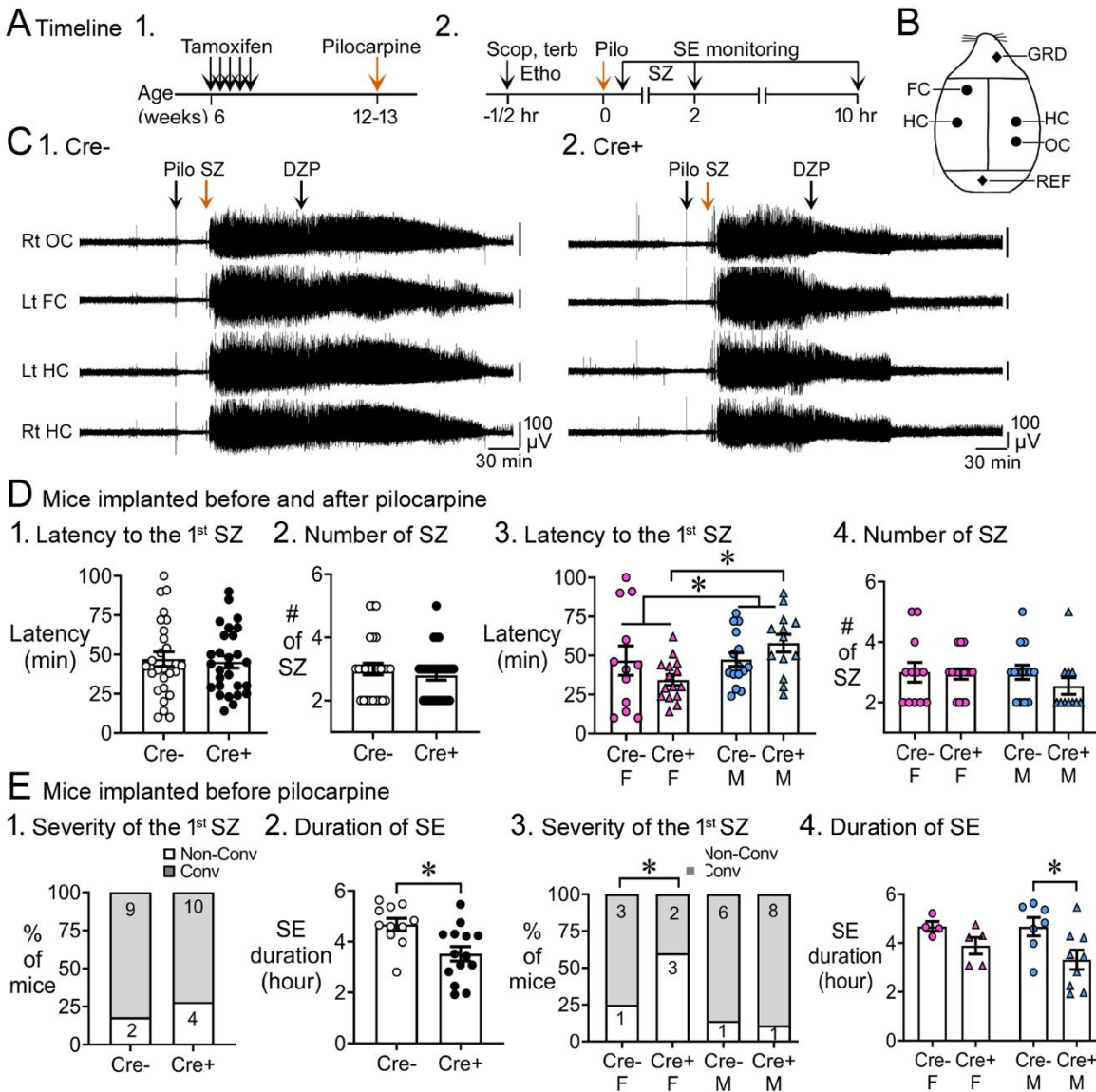
1069 **Table 1.**

Primary antibodies			Secondary antibodies		
Name	Dilution	Source, identifier	Name	Dilution	Source, identifier
anti-doublecortin (rabbit monoclonal)	1:1000	Cell Signaling Technology Cat# 4604S, RRID:AB_10693771	Biotinylated goat anti-rabbit IgG	1:500	Vector Laboratories Cat# BA-1000, RRID:AB_2313606
anti-human Prox1 (goat polyclonal)	1:2,000	R and D systems Cat# AF2727, RRID:AB_2170716	Biotinylated horse anti-goat IgG	1:500	Vector Laboratories Cat# BA-9500, RRID:AB_2336123
anti-GluR2/3 (rabbit polyclonal)	1:300	Millipore Cat# AB1506, RRID:AB_90710	Donkey anti-rabbit, Alexa Fluor 488	1:500	Thermo Fisher Scientific Cat# A-21206, RRID:AB_2535792
anti-Prox1 (goat polyclonal)	1:2000	R and D Systems Cat# AF2727, RRID:AB_2170716	Donkey anti-goat, Alexa Fluor 546	1:500	Thermo Fisher Scientific Cat# A-11056, RRID:AB_2534103
anti-somatostatin (rabbit polyclonal)	1:750	Peninsula Laboratories Cat# T-4103.0050, RRID:AB_518614	Goat anti-rabbit, Alexa Fluor 488	1:500	Thermo Fisher Scientific Cat# A-11034, RRID:AB_2576217
anti-parvalbumin (mouse monoclonal)	1:1000	Millipore Cat# MAB1572, RRID:AB_2174013	Goat anti-mouse, Alexa Fluor 568	1:500	Thermo Fisher Scientific Cat# A-11004, RRID:AB_2534072

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Figure 1. Pilocarpine-induced SE in Cre+ and Cre- mice.



1075
1076

A. The experimental timeline is shown.

1077

1. Tamoxifen was injected 1/day for 5 days in 6 week-old Nestin-CreER^{T2} *Bax^{fl/fl}* mice. Six weeks after the last tamoxifen injection, mice were injected with pilocarpine (Pilo) at a dose that induces status epilepticus (SE).

1078

2. On the day of pilocarpine injection, one group of mice without EEG electrodes were monitored for behavioral seizures for 2 hr after pilocarpine injection. Another group of mice were implanted with EEG electrodes 3 weeks prior to pilocarpine injection. In these mice, video-electroencephalogram (video-EEG) was used to monitor SE for 10 hr after pilocarpine injection.

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- B.** Locations to implant EEG electrodes are shown. Four circles represent recording sites: left frontal cortex (Lt FC), left hippocampus (Lt HC), right hippocampus (Rt HC) and right occipital cortex (Rt OC). Two diamonds represent ground (GRD) and

1088 reference (REF) electrodes. **D.** Pooled data for mice that were implanted with EEG
1089 electrodes and unimplanted mice. These data showed no significant genotypic
1090 differences but there was a sex difference.

- 1091 1. The latency to the onset of first seizure was similar in both genotypes (t-test,
1092 $p=0.761$). The seizure was a behavioral seizure \geq stage 3 of the Racine scale
1093 (unilateral forelimb jerking). For this figure and all others, detailed statistics are in
1094 the Results.
- 1095 2. The number of seizures in the first 2 hr after pilocarpine injection was similar in
1096 both genotypes (t-test, $p=0.377$).
- 1097 3. After separating males and females, females showed a shorter latency to the
1098 onset of the first seizure compared to males (two-way ANOVA, $p=0.043$); Cre+
1099 females had a shorter latency to the first seizure relative to Cre+ males
1100 (Bonferroni's test, $p=0.010$).
- 1101 4. The number of seizures in the first 2 hr after pilocarpine injection were similar in
1102 males and females (two-way ANOVA, $p=0.436$).

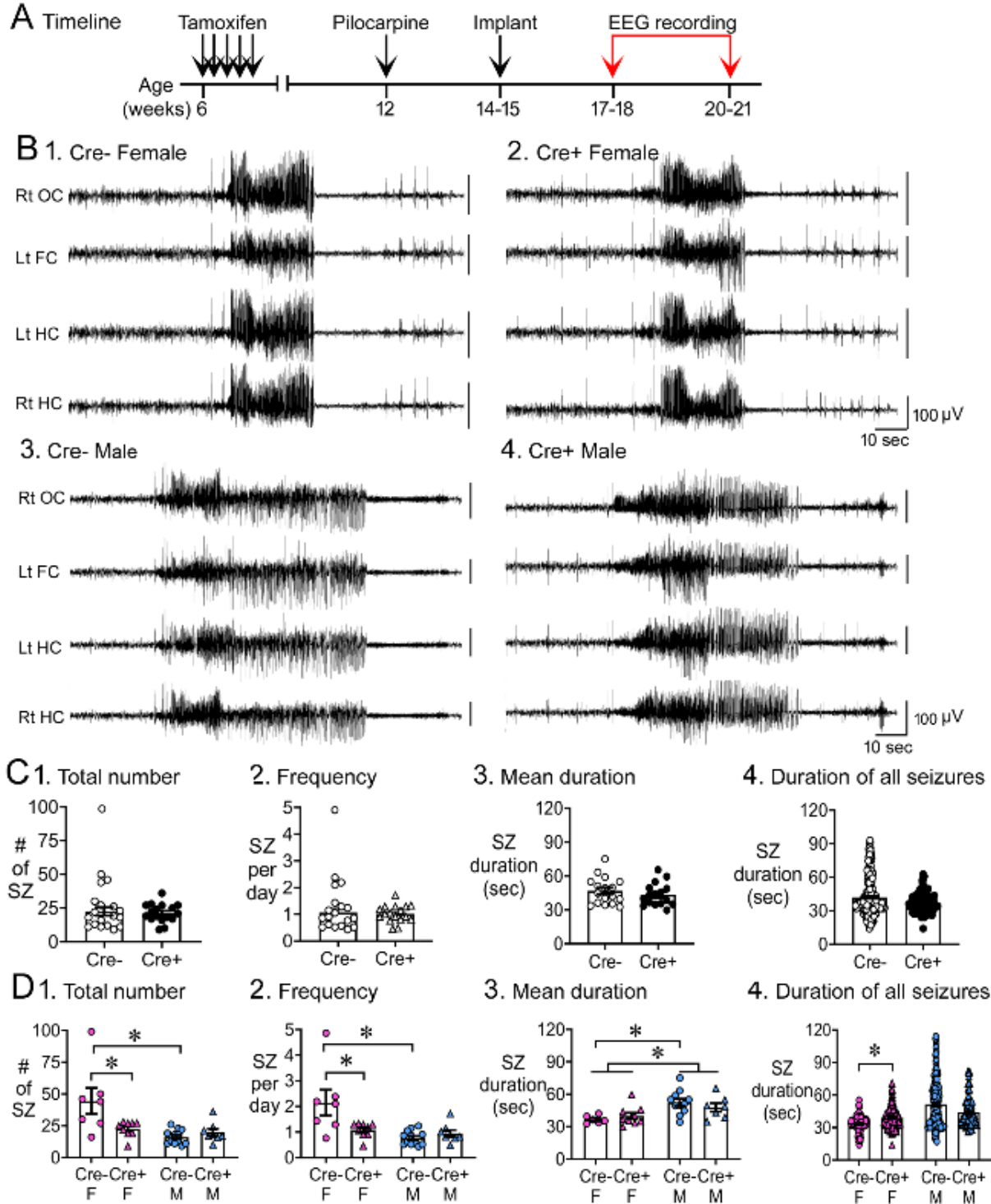
1103 **E.** Implanted mice. These data showed a significant protection of Cre+ mice on SE
1104 duration.

- 1105 1. The severity of the first seizure (non-convulsive or convulsive) was similar
1106 between genotypes (Chi-square test, $p=0.093$).
- 1107 2. Cre+ mice had a shorter duration of SE than Cre- mice (t-test, $p=0.007$).
- 1108 3. After separating males and females, the first seizure was mostly non-convulsive
1109 in Cre+ females compared to Cre- females (60% vs. 14%) but no groups were
1110 statistically different (Fisher's exact tests, $p>0.05$).
- 1111 4. Once sexes were separated, there was no effect of sex by two-way ANOVA but a
1112 trend in Cre+ males to have a shorter SE duration than Cre- males (Bonferroni's
1113 test, $p=0.078$).

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1115 **Figure 2. Reduced chronic seizures in Cre+ mice.**

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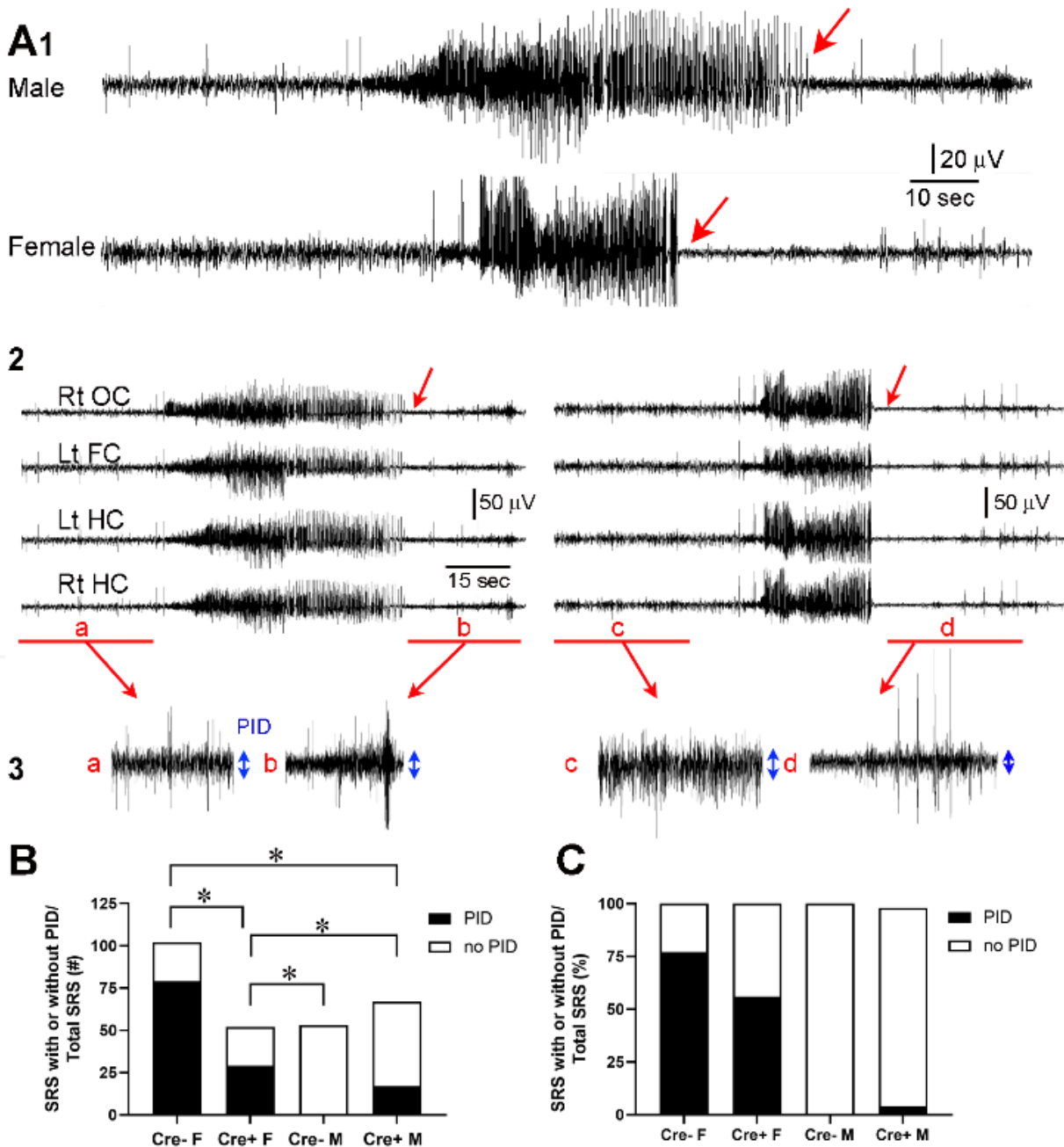
1122

A. The experimental timeline is shown. Six weeks after pilocarpine injection, continuous video-EEG was recorded for 3 weeks to capture chronic seizures. Mice that were unimplanted prior to SE were implanted at 2-3 weeks after pilocarpine injection.

B. Representative examples of 2 min-long EEG segments show a seizure in a Cre- (1, 3) and Cre+ (2, 4) mouse.

- 1123 **C. Numbers of chronic seizures.**
1124 1. Pooled data of females and males showed no significant effect of genotype on
1125 chronic seizure number. The total number of seizures during 3 weeks of
1126 recording were similar between genotypes (t-test, $p=0.882$).
1127 2. After separating data based on sex, females showed fewer seizures. Cre+
1128 females had fewer seizures than Cre- females (Bonferroni's test, $p=0.004$). There
1129 was a sex difference in control mice, with fewer seizures in Cre- males compared
1130 to Cre- females (Bonferroni's test, $p<0.001$).
1131 **D. Chronic seizure frequency.**
1132 1. Pooled data of females and males showed no significant effect of genotype on or
1133 chronic seizure frequency. The frequency of chronic seizures (number of
1134 seizures per day) were similar (Welch's t-test, $p=0.717$).
1135 2. Seizure frequency was reduced in Cre+ females compared to Cre- females
1136 (Bonferroni's test, $p=0.004$). There was a sex difference in control mice, with
1137 lower seizure frequency in Cre- males compared to Cre- females (Bonferroni's
1138 test, $p<0.001$).
1139 **E. Seizure duration per mouse.**
1140 1. Each data point is the mean seizure duration for a mouse. Pooled data of
1141 females and males showed no significant effect of genotype on seizure duration
1142 (t-test, $p=0.379$).
1143 2. There was a sex difference in seizure duration, with Cre- males having longer
1144 seizures than Cre- females (Bonferroni's test, $p=0.005$). Because females
1145 exhibited more postictal depression (see Fig. 3), corresponding to spreading
1146 depolarization (Ssentongo et al. 2017), the shorter female seizures may have
1147 been due to truncation of seizures by spreading depolarization.
1148 **F. Seizure durations for all seizures.**
1149 1. Every seizure is shown as a data point. The durations were similar for each
1150 genotype (Mann-Whitney U test, $p=0.079$).
1151 2. Cre+ females showed longer seizures than Cre- females (Dunn's test, $p<0.001$).
1152 Cre+ females may have had longer seizures because they were protected from
1153 spreading depolarization.
1154

1155 **Figure 3. Reduced postictal depression in Cre+ female mice.**



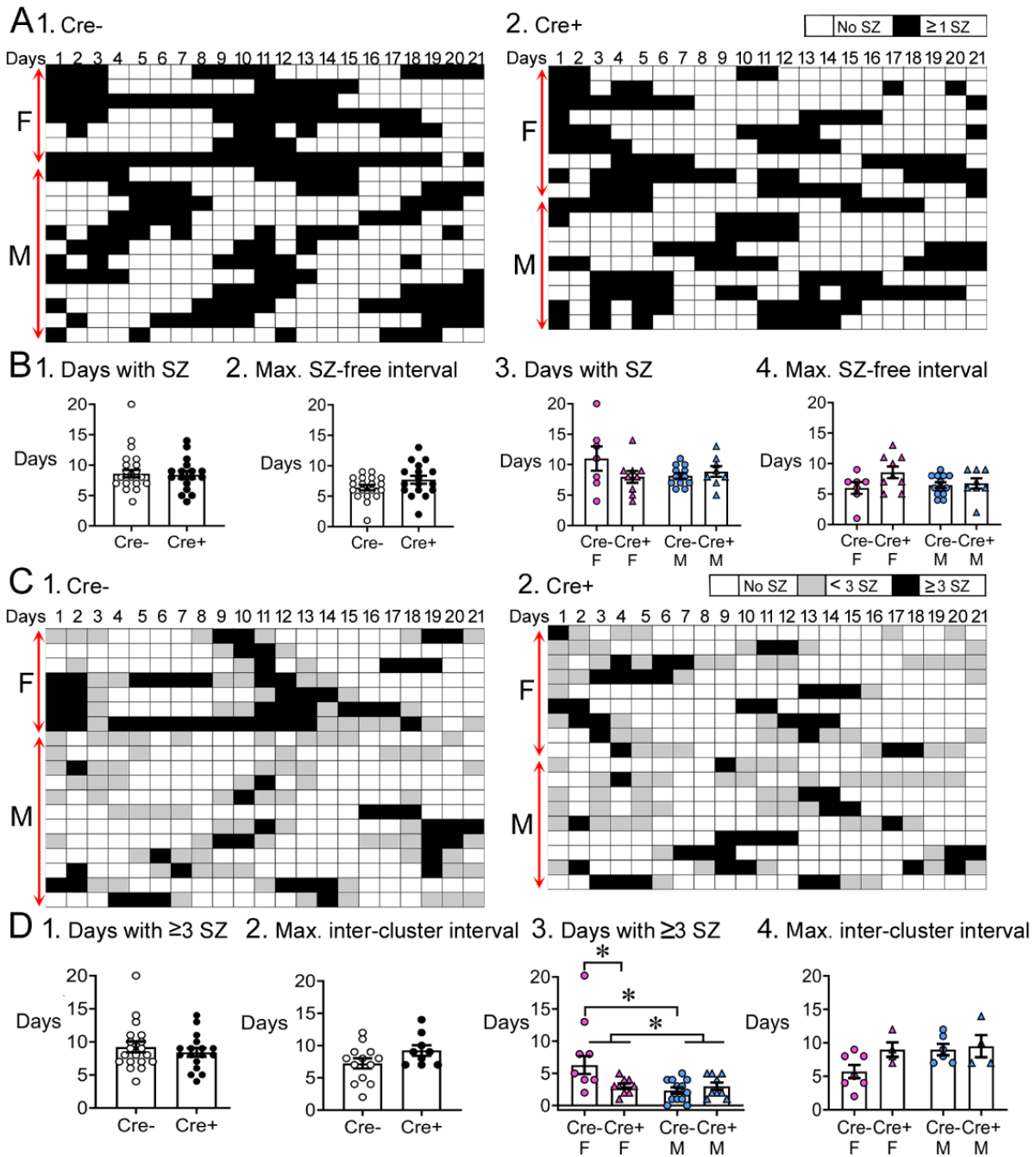
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- 1158 **A. 1.** A seizure of a male mouse and female mouse are shown to illustrate postictal
1159 depression starting at the end of the seizure (red arrow).
1160 **2.** All 4 channels are shown for the male (left) and female mouse (right). Rt OC,
1161 right occipital cortex; Lt FC, left frontal cortex; Lt HC, left hippocampus; Rt HC, right
1162 hippocampus. The red arrows point to the end of the seizure.
1163 **3.** The areas in A2 marked by the red bar are expanded. The blue double -sided
1164 arrows reflect the mean EEG amplitude before (a, c) and after the seizure (b,d).

- 1165 **B.** For all spontaneous recurrent seizures (SRS) in the 3 week-long recording period,
1166 there was a significant difference between groups, with number of SRS with PID
1167 reduced in Cre+ females compared to Cre- females (Fisher's exact test, all p
1168 <0.05). Males had very little postictal depression and there was no significant effect
1169 of genotype.
- 1170 **C.** The same data are plotted but the percentages are shown instead of the numbers of
1171 seizures.

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Figure 4. Temporal dynamics of chronic seizures.



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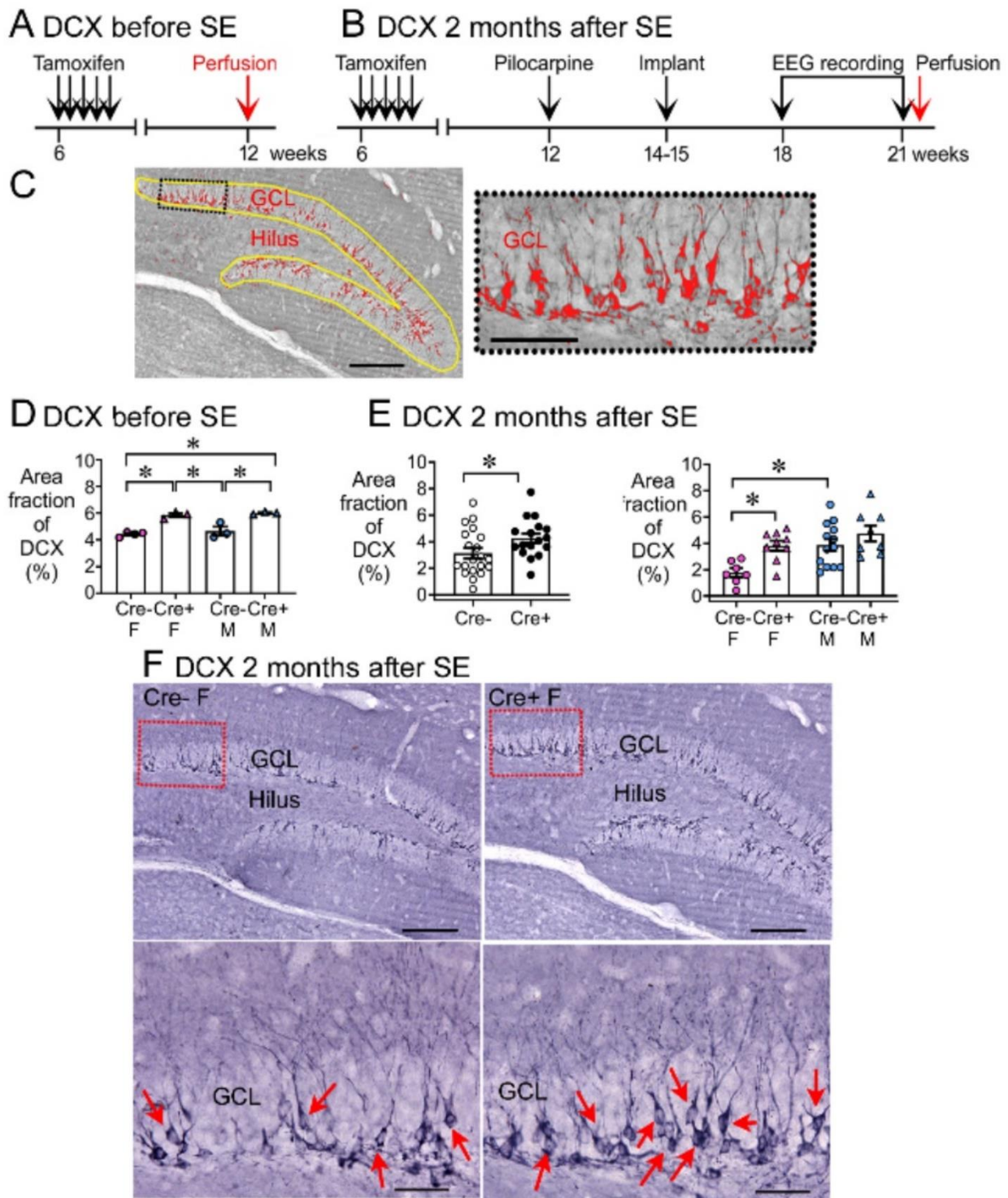
A. Each day of the 3 weeks-long EEG recording periods are shown. Each row is a different mouse. Days with seizures are coded as black boxes and days without seizures are white.

B.

1. The number of days with seizures were similar between genotypes (t-test, $p=0.822$).

- 1181 2. The maximum seizure-free interval was similar between genotypes (t-test,
1182 p=0.107).
- 1183 3. After separating females and males, two-way ANOVA showed no effect of
1184 genotype or sex on days with seizures.
- 1185 4. Two-way ANOVA showed no effect of genotype or sex on the maximum seizure-
1186 free interval.
- 1187 **C.** Then same data are shown but days with ≥ 3 seizures are black, days with < 3
1188 seizures as grey, and are white. Clusters of seizures are reflected by the
1189 consecutive black boxes.
- 1190 **D.**
- 1191 1. The cluster durations were similar between genotypes (Mann-Whitney's *U* test,
1192 p=0.723).
- 1193 2. The maximum inter-cluster interval was similar between genotypes (t-test,
1194 p=0.104).
- 1195 3. Cre+ females had significantly fewer clusters than Cre- females (two-way
1196 ANOVA followed by Bonferroni's test, p=0.009). There was a sex difference, with
1197 females having more clusters than males. Cre- females had more days with >3
1198 seizures than control males (Cre- females: 6.3 ± 1.4 days; Cre- males: 2.3 ± 0.5
1199 days; Bonferroni's test, $p < 0.001$).
- 1200 4. There was no significant effect of genotype or sex on the maximum inter-cluster
1201 interval. However, there was a trend for the inter-cluster interval to be longer in
1202 Cre+ females relative to than Cre- females.
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1204 **Figure 5. Increased DCX in Cre+ mice.**
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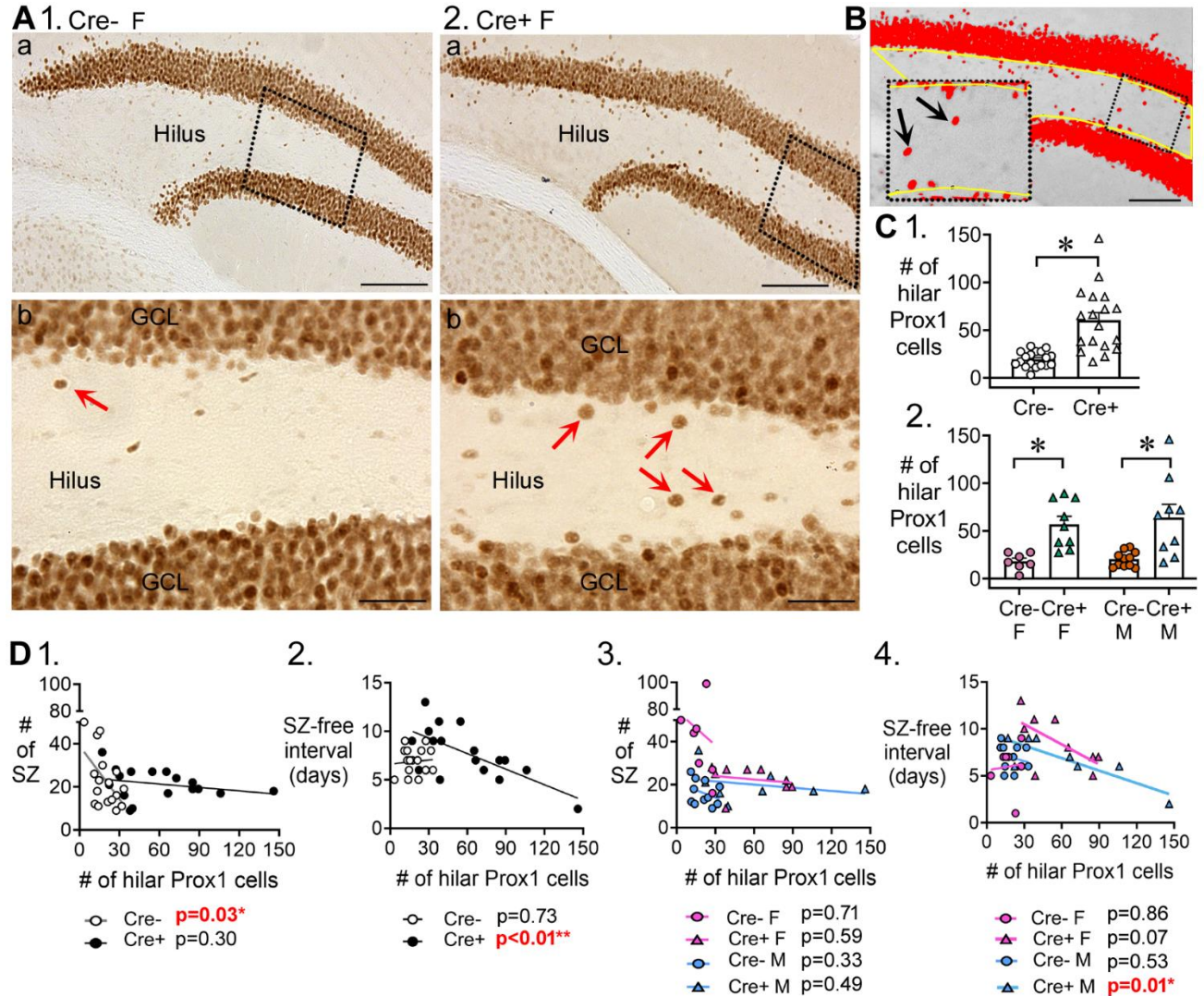
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A-B. The experimental timelines are shown. **A.** Mice were perfusion-fixed 6 weeks after tamoxifen injection, just before SE. Sections were then stained for DCX. **B.** Mice

1211 were tested 2 months after SE, after EEG recording. Then mice were perfused and
1212 staining was conducted for DCX.
1213 **C.** DCX quantification.
1214 DCX-ir within a region of interest (ROI; yellow lines) including the SGZ and GCL
1215 was thresholded. DCX-ir above the threshold is shown in red. Calibration, 100 μm
1216 (a); 50 μm (b). The inset is expanded to the right.
1217 **D.** The area of DCX-ir relative to the area of the ROI (referred to as area fraction) was
1218 greater in Cre⁺ mice compared to Cre⁻ mice. Two-way ANOVA followed by Tukey
1219 pot-hoc tests, all $p < 0.05$).
1220 **E.** Cre⁺ mice had increased DCX-ir relative to Cre⁻ mice 2 months after SE.
1221 1. Sexes were pooled. The area fraction of DCX-ir was greater in Cre⁺ than Cre⁻
1222 mice (t-test, $p = 0.041$).
1223 2. When sexes were separated, Cre⁺ females showed greater DCX-ir than Cre⁻
1224 females (two-way ANOVA followed by Bonferroni's test, $p = 0.015$). There was a
1225 sex difference, with Cre⁻ males showing more DCX-ir than Cre⁻ females
1226 (Bonferroni's test, $p = 0.007$). DCX-ir was similar in Cre⁻ and Cre⁺ males
1227 (Bonferroni's test, $p = 0.498$).
1228 **F.** Representative examples of DCX-ir 2 months after SE.
1229 1. Cre⁻ female mouse.
1230 2. Cre⁺ female mouse. The red boxes in a are expanded in b. Arrows point to DCX-
1231 ir cells. Calibration, 100 μm (a); 50 μm (b).
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Figure 6. Hilar Prox1-ir cells increased in Cre+ mice.



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A. Representative examples of hilar Prox1-ir in Cre- (1) and Cre+ (2) mice are shown.

The boxes in a are expanded in b. Arrows point to hilar Prox1-ir cells, corresponding to hilar ectopic GCs. Calibration, 100 μ m (a); 50 μ m (b).

B. Prox1-ir is shown, within a hilar ROI. The area of the ROI above the threshold, relative to the area of the ROI, is red. This area is called the area fraction, and was used to quantify hilar Prox1-ir. Calibration, 100 μ m.

C.

1. Cre+ mice had more hilar Prox1-ir cells than Cre- mice (t-test, $p < 0.001$).
2. When sexes were divided, Cre+ mice had more hilar Prox1-ir cells than Cre- mice in both female (two-way ANOVA followed by Bonferroni's test, $p < 0.001$) and male mice (Bonferroni's test, $p = 0.001$).

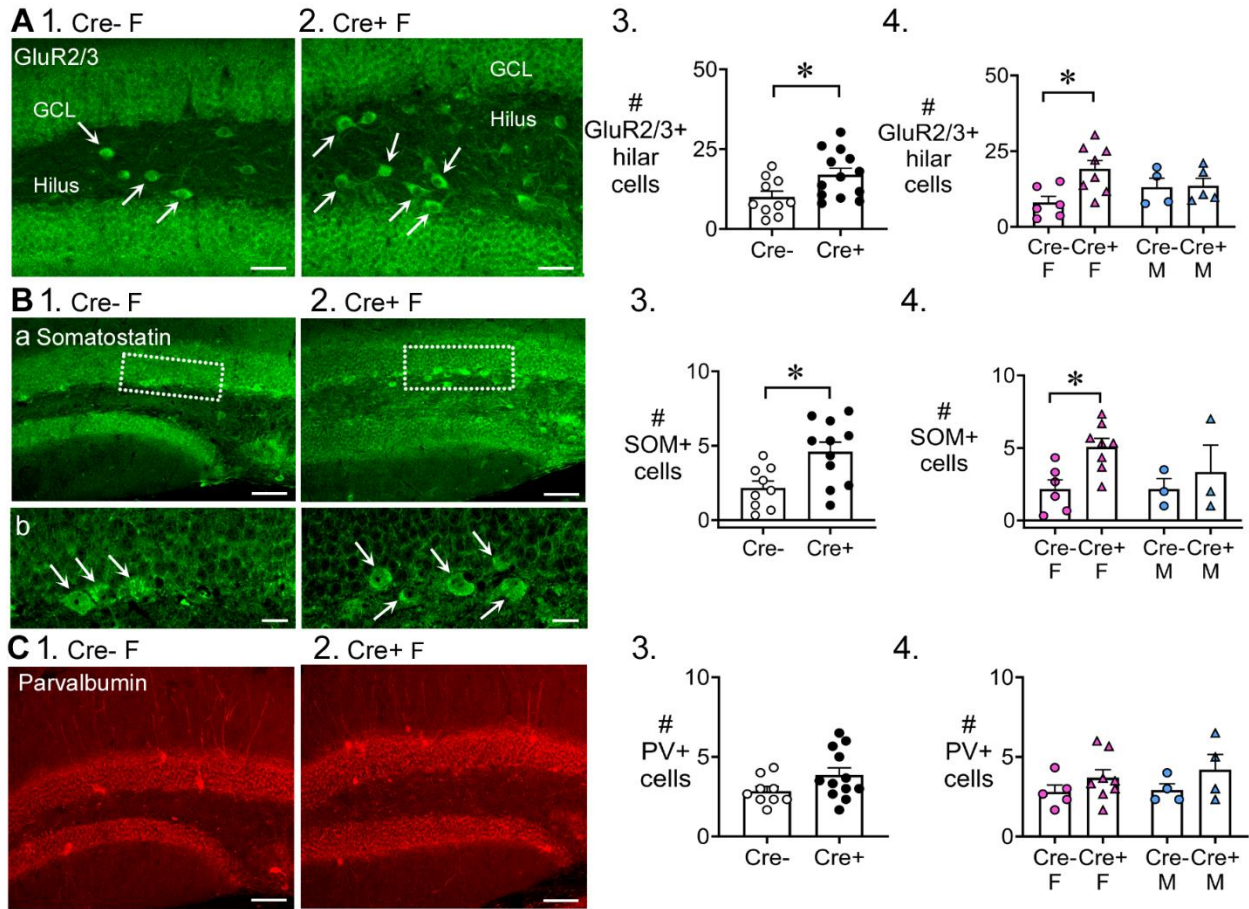
D. Correlations between hilar Prox1-ir cells and measurements of chronic seizures.

1. All Cre- and Cre- mice were compared regardless of sex. For the Cre- mice there was a significant inverse correlation between the # of Prox1-ir cells and # of

1250 chronic seizures ($R^2=0.296$). Thus, the more Prox1-ir cells there were, the fewer
1251 chronic seizures there were. However, that was not true for Cre+ mice ($R^2=0.072$).
1252 2. There was an inverse correlation between the number of hilar Prox1-ir cells and
1253 the seizure-free interval for Cre+ mice ($R^2=0.467$) but not Cre- mice ($R^2=0.008$).
1254 Thus, the more hilar Prox1-ir cells there were, the shorter the seizure-free periods
1255 were. However, this was not true for Cre- mice.
1256 3. When data were divided by genotype and sex there was no significant correlation
1257 between hilar Prox1-ir cells and # of seizures (Cre- F, $R^2=0.0035$; Cre+ F,
1258 $R^2=0.043$; Cre- M, $R^2=0.104$; Cre+ M, $R^2=0.083$).
1259 4. When data were divided by genotype and sex, there was a significant inverse
1260 correlation for the # of hilar Prox1-ir cells and seizure-free interval, but only for
1261 male Cre+ mice ($R^2=0.704$). Cre+ females showed a trend ($R^2=0.395$) and Cre-
1262 mice did not (Cre- F, $R^2=0.007$, Cre- M, $R^2=0.046$).
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Figure 7. Preserved mossy cells and hilar SOM cells in Cre+ female mice but not parvalbumin interneurons.



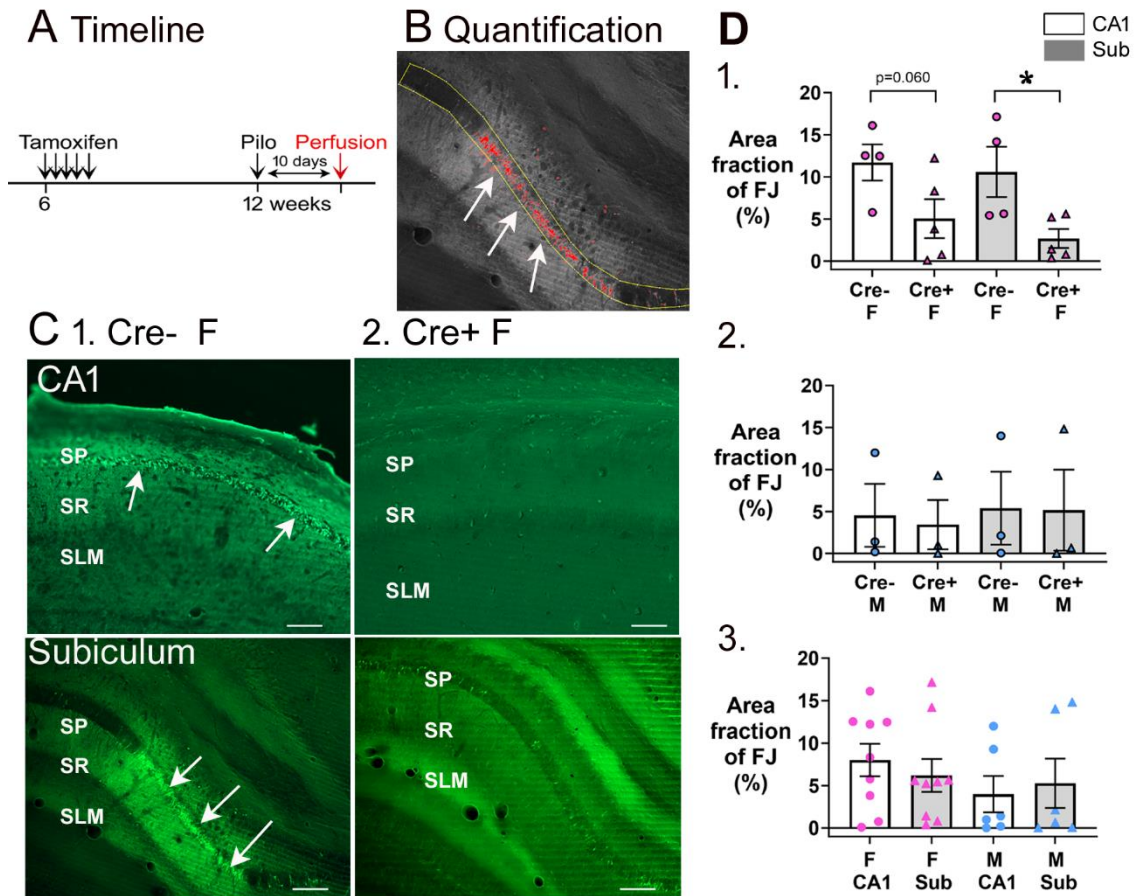
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- A.**
- 1-2. Representative examples of GluR2/3 labelling of Cre- (1) and Cre+ mice (2). Calibration, 50 μ m.
 3. Cre+ mice had more hilar GluR2/3-immunofluorescent (positive; +) cells than Cre- mice (t-test, $p=0.022$). Sexes were pooled.
 4. After separating females and males, Cre+ females showed more hilar GluR2/3+ cells than Cre- females (Bonferroni's test, $p=0.011$). Hilar GluR2/3+ cells were similar between genotypes in males (Bonferroni's test, $p=0.915$).
- B.**
- 1-2. Representative examples of SOM labelling in Cre- and Cre+ mice are shown. Calibration, 100 μ m (a); 20 μ m (b).
 3. In pooled data, Cre+ mice had more hilar SOM cells than Cre- mice (t-test, $p=0.008$).
 4. After separating females and males, Cre+ females showed more hilar SOM cells than Cre- females (Bonferroni's test, $p=0.019$). Hilar SOM cells were similar between genotypes in males (Bonferroni's test, $p=0.897$).
- C.**

- 1286 1-2. Representative examples of parvalbumin labelling in Cre- and Cre+ mice are
1287 shown. Calibration, 100 μ m.
1288 3. The number of parvalbumin+ cells in the DG were similar in Cre- and Cre+ mice
1289 in pooled data (t-test, $p=0.095$).
1290 4. There was no effect of genotype ($p=0.096$) or sex ($p=0.616$) on the number of DG
1291 parvalbumin+ cells.
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Figure 8. Cre+ female mice had less neuronal loss in hippocampus after SE.



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- A.** A timeline is shown to illustrate when mice were perfused to examine Fluorojade-C staining. All mice were perfused 10 days after SE, a time when delayed cell death occurs after SE, mainly in area CA1 and subiculum. Note that prior studies showed hilar and CA3 neurons, which exhibit more rapid cell death after SE, are protected from cell loss in Cre+ mice examined 3 days after SE (Jain et al., 2019). Also, there was protection of CA1 at 3 days (Jain et al., 2019).
- B.** Quantification. Fluorojade-C was thresholded using ImageJ and the pyramidal cell layer outlined in yellow. The fraction above threshold relative to the entire ROI (area fraction) was calculated (see Methods)
2. The Fluorojade-C area fraction was greater in Cre- mice than Cre+ mice. Statistical comparisons showed a trend for CA1 of Cre- mice to exhibit more Fluorojade-C than Cre+ mice (Mann-Whitney *U* test, $p=0.060$). Cre- mice had a significantly greater area fraction in the subiculum than Cre+ mice (Mann-Whitney *U* test, $p=0.032$).
- C.** Examples of Fluorojade-C staining in CA1 (top) and subiculum (bottom) of Cre+ female (1) and Cre- female (2) mice. SO, stratum oriens; SP, stratum pyramidale; SR, stratum radiatum; SLM, stratum lacunosum-moleculare. Arrows point to numerous Fluorojade-C-stained neurons in Cre- mice but not Cre+ mice. Calibration, 200 μ m.

1315 **D.** 1. Comparisons of female mice by two-way ANOVA showed an effect of
1316 genotype ($F(1,15)=11.97$, $p=0.004$) with less Fluorojade C in Cre+ mice for CA1
1317 ($p=0.016$) and subiculum $p=0.016$).
1318 2. Comparisons of male mice showed no significant effect of genotype on
1319 Fluorojade C in either CA1 or the subiculum ($F(1,8)=0.002$, $p=0.965$; CA1,
1320 $p=0.828$, subiculum, $p=0.973$, respectively).
1321 3. When genotypes were pooled, female mice did not have significantly more
1322 damage than males (two-way ANOVA, sex ($F(1,34)=3.16$, $p=0.085$) and there
1323 was no effect of subfield ($F(1,34)=0.0016$, $p=0.968$).
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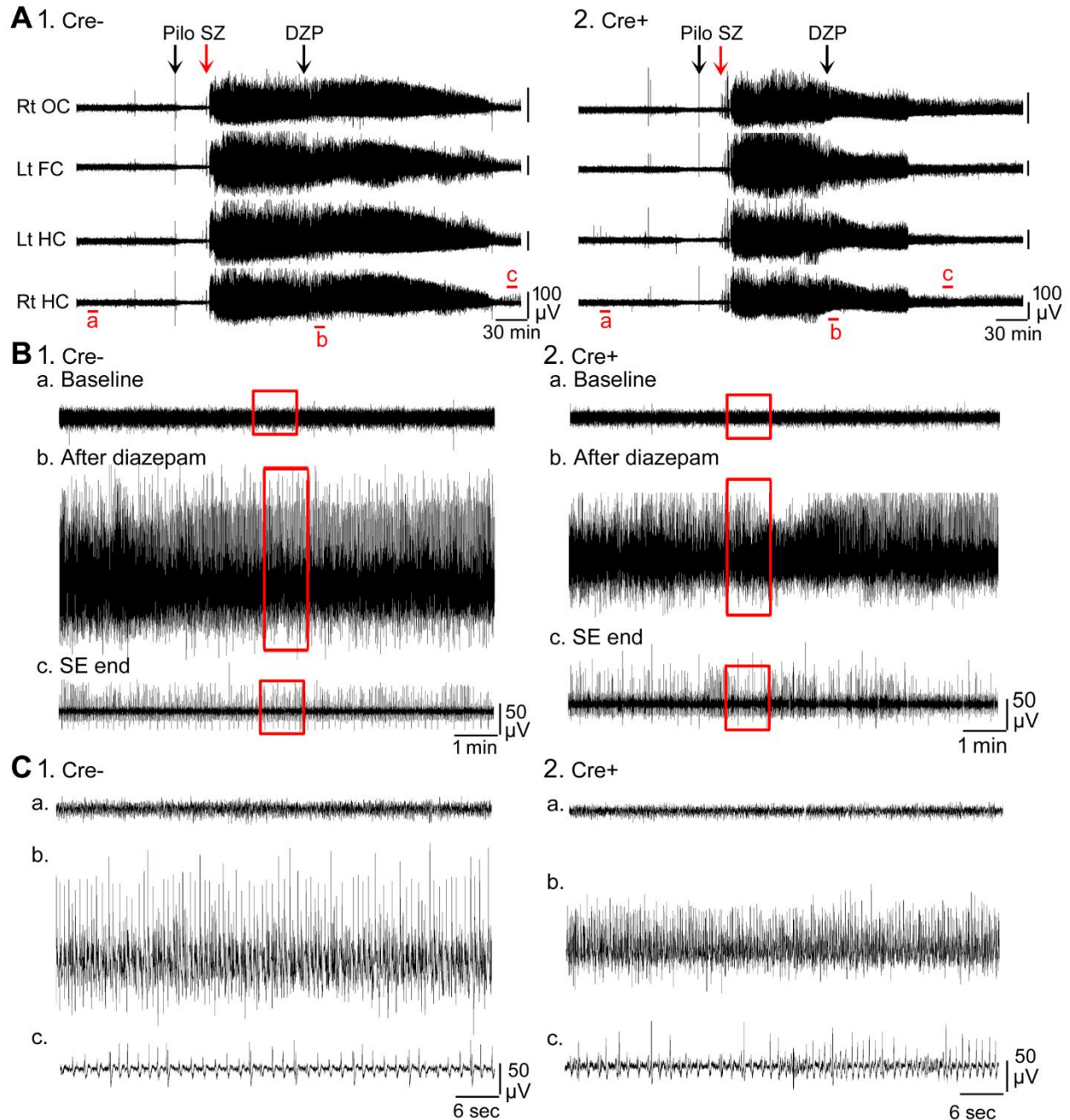
1326 **SUPPLEMENTARY FIGURES**

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1328 **Fig. 1.- Supplemental Fig. 1.**

1329 **Examples of EEG during SE.**

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1333 **A.** Representative examples of a 10 hr-long EEG recording are shown for Cre- (1) and

1334 Cre+ (2) mice. These records are the same as in Fig. 1.

1335 **B.** 10 min-long EEG recording segments from the left hippocampus A are shown with

1336 higher temporal gain.

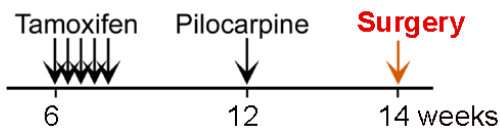
1337 1. Cre- mouse.

- 1338 a. Part of the baseline is shown. The area surrounded by the red box is
1339 expanded in C1a.
- 1340 b. The time when DZP was injected is shown. The area surrounded by the red
1341 box is expanded in C1b.
- 1342 c. The time following the seizure is shown. The area surrounded by the red box is
1343 expanded in C1c.
- 1344 2. Cre+ mouse.
- 1345 a. Part of the baseline is shown. The area surrounded by the red box is
1346 expanded in C2a. Note the baselines are similar in the two mice, suggesting
1347 no effect of genotype.
- 1348 b. The time when DZP was injected is shown. The area surrounded by the red
1349 box is expanded in C2b. Note there was a reduction in EEG amplitude in the
1350 Cre+ mouse.
- 1351 c. The time at the end of SE is shown. The area surrounded by the red box is
1352 expanded in C2c. Note that these two mice were similar after SE ended.
- 1353 **C.** The areas surrounded by the red boxes in B are expanded. The traces are 1 min-
1354 long.
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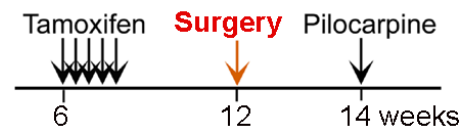
1356 **Fig. 1.- Supplemental Fig. 2.**
 1357 **Incidence of SE in the unimplanted and implanted mice.**
 1358

A Timeline

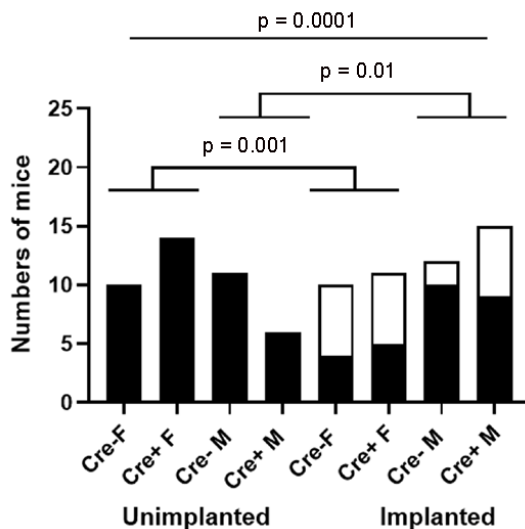
1. Unimplanted



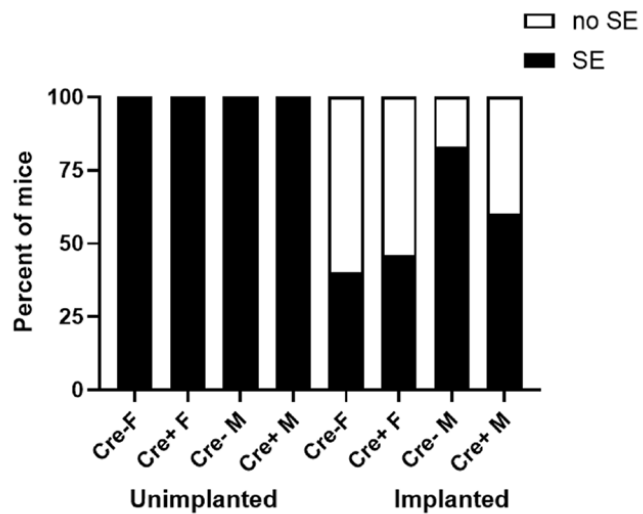
2. Implanted



B Numbers of mice

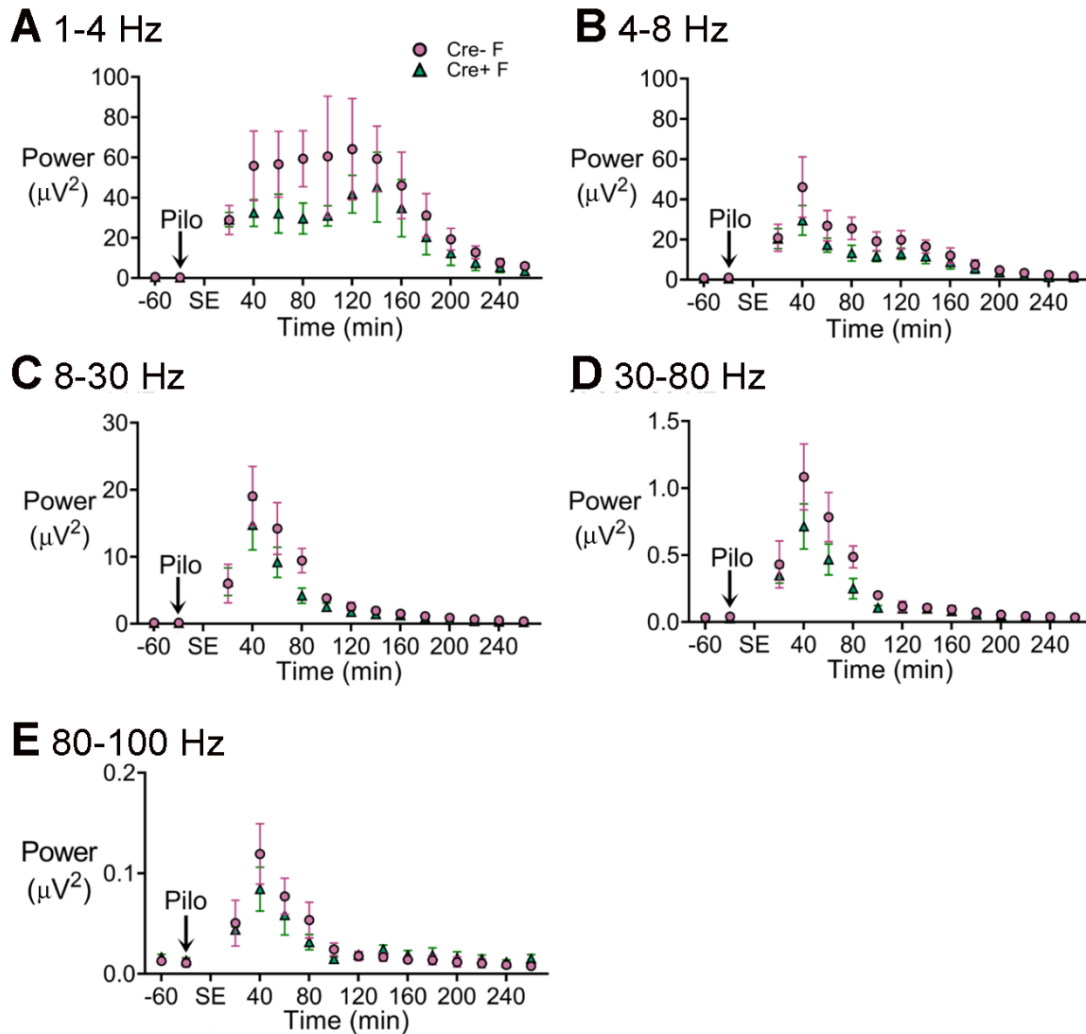


C Percentages of mice



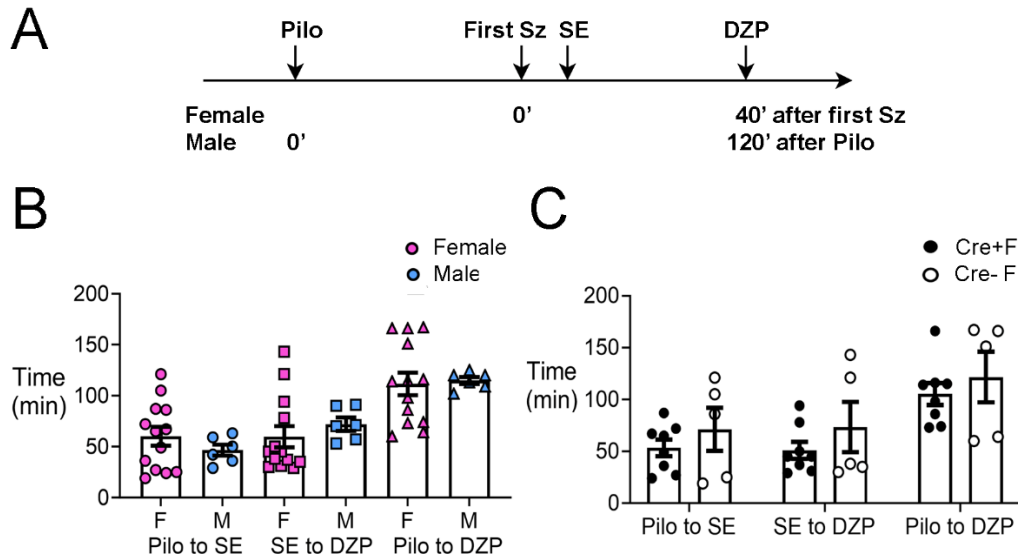
- 1359 **A.** The experimental timelines of pilocarpine injection and surgery.
 1360 1. Unimplanted mice.
 1361 2. Implanted mice. Only the timing of surgery with respect to pilocarpine injection
 1362 was different.
 1363 **B.** The incidence of SE in unimplanted and implanted mice is shown based on numbers
 1364 of mice. The incidence of SE was significantly higher in the unimplanted mice
 1365 (Fisher's exact test, $p < 0.0001$). Genotype had no effect on the incidence of SE.
 1366 **C.** The incidence of SE is shown as percentages.
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1381 **Fig. 1.- Supplemental Fig. 3.**
1382 **Power during SE in Cre+ female and Cre- female mice.**
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1384 **A.** Power was calculated for consecutive 20 min-long bins before and during SE. Power
1385 in the 1-4 Hz band was decreased during SE in female Cre+ mice (blue triangles)
1386 relative to Cre- mice (red circles) but it was not statistically significant.
1387
1388 **B.** Power in the 4-8 Hz band.
1389 **C.** Power in 8-30 Hz range.
1390 **D.** Power in 30-80 Hz range
1391 **E.** Power between 80 and 100 Hz.
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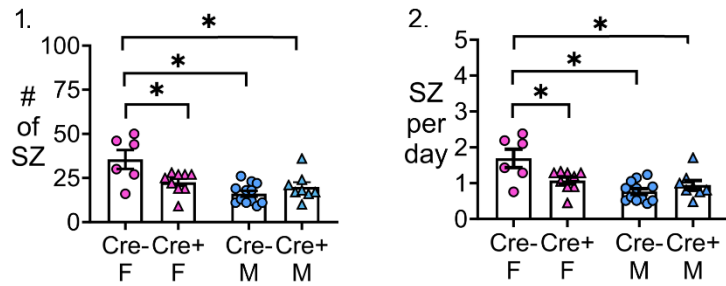
1396 **Fig. 1- Supplemental Fig. 4.**
1397 **The latency to SE, interval between SE and diazepam administration, and interval**
1398 **between pilocarpine and diazepam injections were not significantly different in**
1399 **experimental groups.**



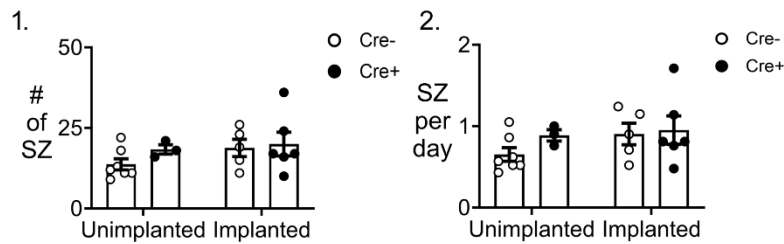
1400 **A.** A timeline of experimental procedures is shown for the day of pilocarpine-induced
1401 SE. Pilocarpine was injected and the first seizure stage 3 or greater was noted. The
1402 onset of SE was noted also. For females, diazepam (DZP) was injected 45 min after
1403 the first seizure (Sz). Males were administered DZP 2 hrs after pilocarpine (Pilo). The
1404 reason for the difference is that it made the latencies to SE, interval between SE and
1405 DZP injection, and interval between pilocarpine and DZP injections similar.
1406 **B.** The mean \pm SEM is shown for the time from pilocarpine to SE, SE to DZP injection,
1407 and pilocarpine to DZP injection. There were no sex differences: a two-way ANOVA
1408 with sex and type of measurement as main factors showed no effect of sex
1409 ($F(1,45)=0.004$, $p=0.949$). However, there were differences between the types of
1410 measurements ($F(2,45)=11.89$, $p<0.0001$).
1411 **C.** When Cre+ and Cre- females were compared, there was no effect of genotype
1412 ($F(1,33)=2.33$, $p=0.136$) but there was a significant effect of the type of measurement
1413 ($F(2,33)=7.66$, $p=0.002$).
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1417 **Fig. 2.- Supplemental Fig. 1.**
 1418 **Additional analyses of chronic seizures.**

A After outlier removal



B Similar results for unimplanted and implanted mice



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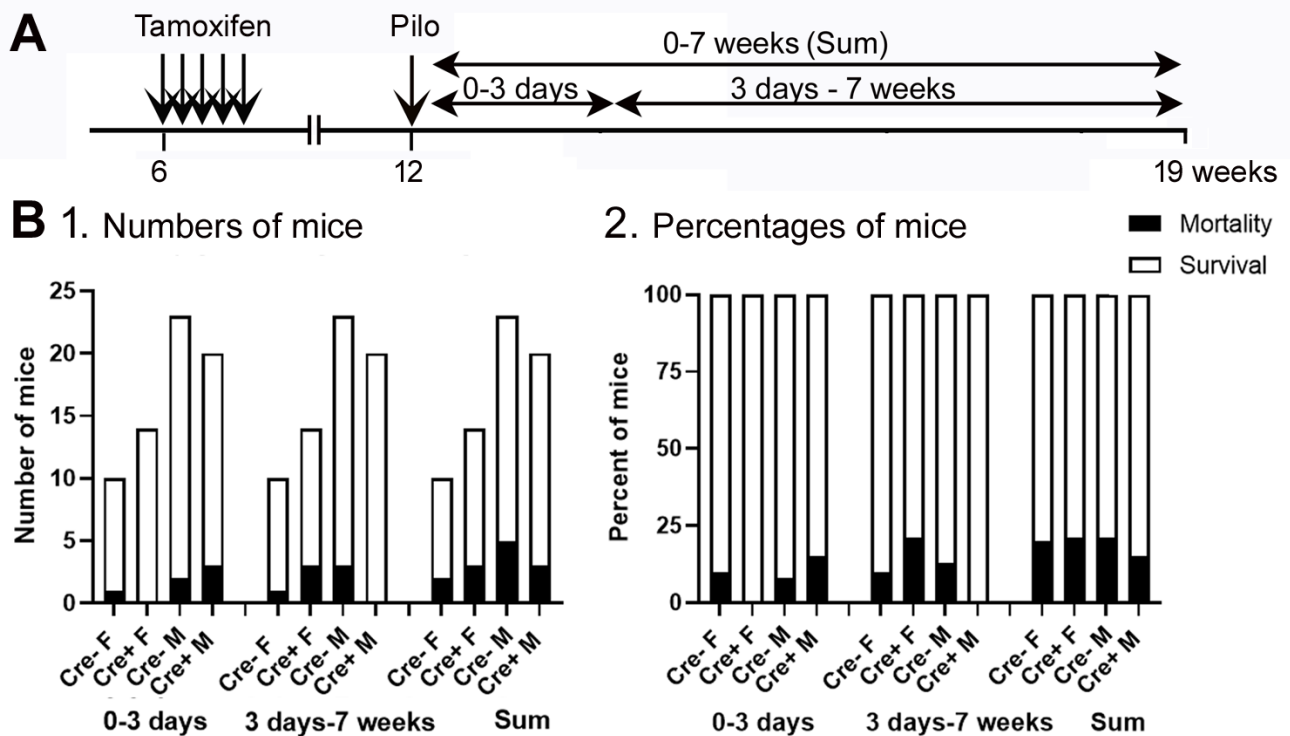
1437 **A. Similar results after outlier removal.**

- 1438 1. A two-way ANOVA with sex and genotype as factors showed a significant effect of
 1439 sex on the # of seizures ($F(1,31)=16.04$, $p=0.0004$) and an interaction
 1440 ($F(1,31)=9.20$, $p=0.005$) with no main effect of genotype ($F(1,31)=1.75$, $p=0.107$).
 1441 Cre- females had significantly more seizures than all other groups (post-hoc tests,
 1442 Cre- females vs. Cre+ females, $p=0.020$; vs. Cre- males, $p=0.0002$; vs. Cre+
 1443 males, $p=0.005$). Cre- males and Cre+ males were not different ($p=0.722$).
 1444 2. A two-way ANOVA with sex and genotype as factors showed a significant effect of
 1445 sex ($F(1,31)=15.84$, $p=0.0004$) on seizure frequency and an interaction
 1446 ($F(1,31)=9.16$, $p=0.005$) although no main effect of genotype ($F(1,31)=2.72$,
 1447 $p=0.109$). Cre- females had significantly more seizures than all other groups (post-
 1448 hoc tests, Cre- females vs. Cre+ females, $p=0.021$; vs. Cre- males, $p=0.0002$; vs.
 1449 Cre+ males, $p=0.005$). Cre- males were not different from Cre+ males ($p=0.720$).

1450 **B. Results were independent of the time when EEG electrodes were implanted.**

- 1451 1. The number of chronic seizures were similar between mice that were implanted
 1452 before and after SE by two-way ANOVA with implant status and genotype as
 1453 factors (implant status, $F(1,17)=1.33$, $p=0.265$; genotype, $F(1,17)=0.88$, $p=0.334$).
 1454 Sexes were pooled.
 1455 2. The frequency of seizures was similar between mice implanted before or after SE
 1456 (implant status, $F(1,17)=1.27$, $p=0.276$; genotype, $F(1,17)=1.00$, $p=0.330$). Sexes
 1457 were pooled.
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1460 **Fig. 2.- Supplemental Fig. 2.**
 1461 **Mortality was not significantly affected by genotype or sex.**
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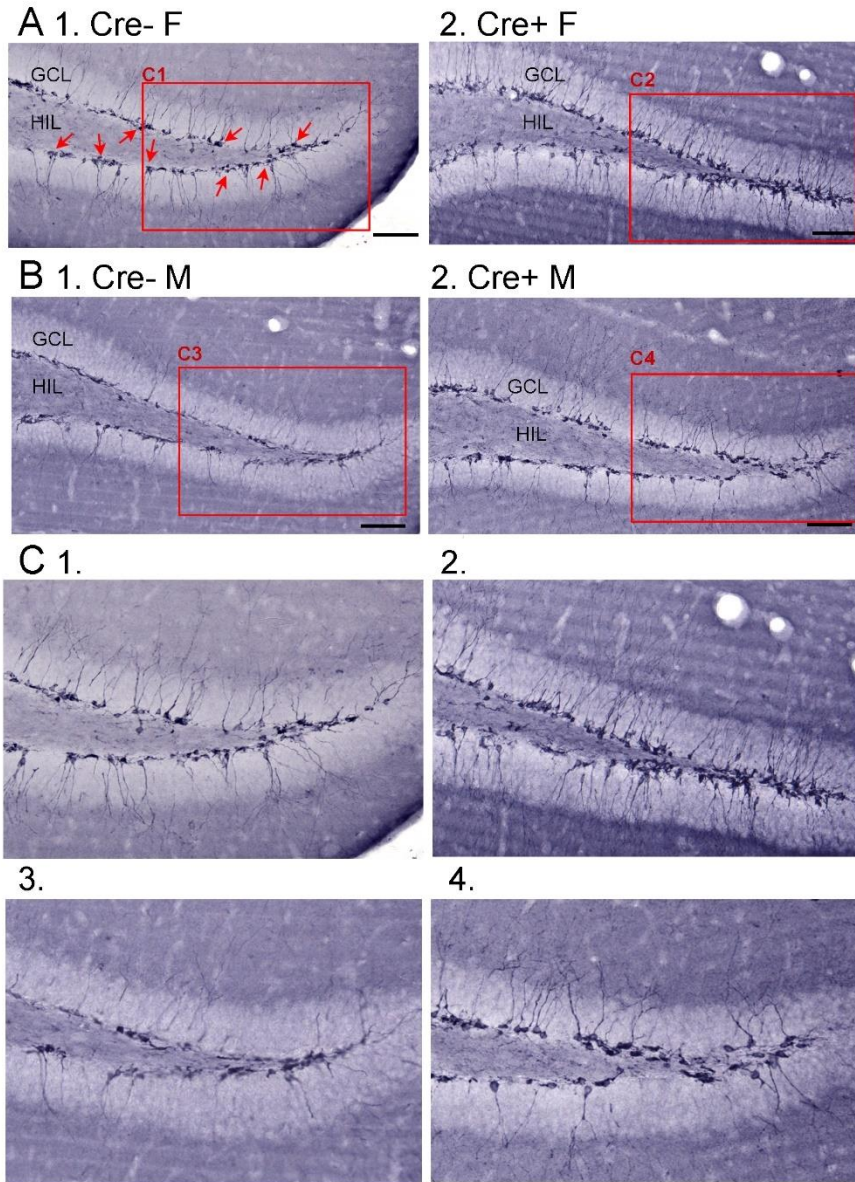


1463 **A.** The timeline of measurements of mortality is shown. Mice were categorized as dying
 1464 during SE or within 3 days of SE (0-3 days), 3 days to 7 weeks, or both (Sum). Mice
 1465 are included whether they were implanted with EEG electrodes before SE or
 1466 implanted 3 weeks after SE.
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1469 **B.**
 1470 1. The numbers of mice that died were not significantly different between genotypes
 1471 or sexes (Fisher's Exact test, all $p > 0.05$).
 1472 2. The percent of mice that died is shown.
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Fig. 5.- Supplemental Fig. 1.
DCX in Cre- and Cre+ mice before SE.



1512 **A.** Mice were administered tamoxifen for 5 days at 6 weeks of age as for other experiments. Six
1513 weeks after tamoxifen, mice were perfusion-fixed and staining was conducted using an
1514 antibody to DCX.

1515 1. Representative image from a Cre- female (F) of dorsal dentate gyrus in coronal section
1516 shows many DCX-ir cells in the SGZ and GCL (red arrows). The area surrounded by the
1517 red box is expanded in C1. Calibration, 100 μ m.

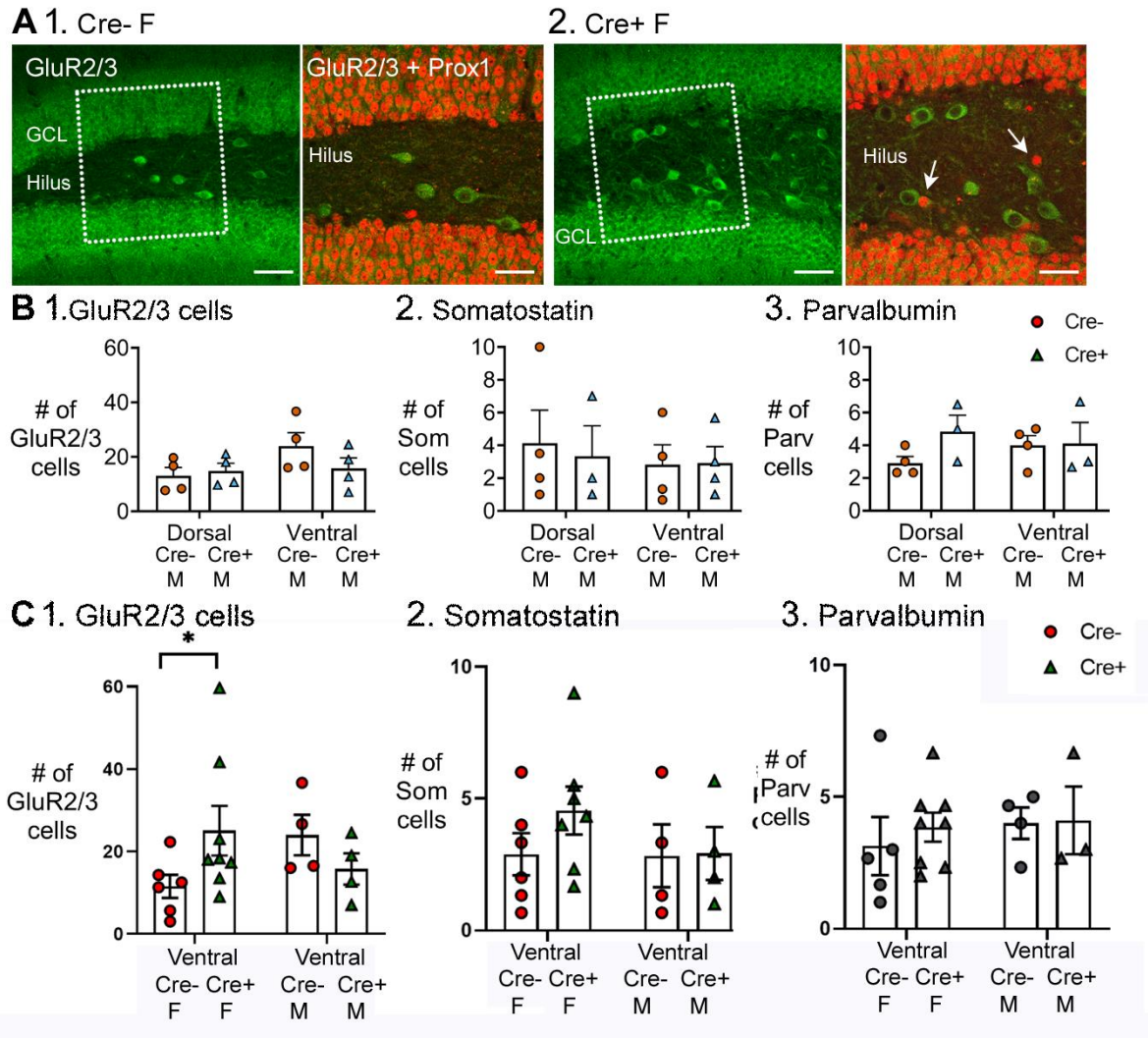
1518 2. Example of DCX-ir from a Cre+ female mouse shows more DCX-ir, reflecting more
1519 immature neurons. The area surrounded by the red box is expanded in C2. Calibration,
1520 100 μ m.

1521 **B.**

1522 1. Example from a Cre- male (M). The area surrounded by the red box is expanded in C3.

1523 2. Example of a Cre+ M showing more DCX-ir than the Cre- M. The area surrounded by the
1524 red box is expanded in C4. Calibration, 100 μ m.
1525 **C.** Expanded insets from A-B. 1. Cre- F; 2. Cre+ F, 3. Cre- M, 4. Cre+ M.
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1528 **Fig. 7.- Supplemental Fig. 1.**
 1529 **Additional analyses of GluR2/3, SOM and parvalbumin-expressing cells.**



- 1530 **A. GluR2/3 hilar cells lacked Prox1 expression.**
 1531 1. Cre- female mouse. Left: several GluR2/3+ cells (green) are located in the hilus
 1532 within the box (marked by dotted white lines). Calibration, 70 μ m. Right: The area
 1533 within the box in the left panel is expanded. The merged image of GluR2/3+
 1534 (green) and Prox1+ (red) cells shows no double labeling. Calibration, 40 μ m.
 1535 2. Cre + female mouse. Similar results are shown as for the Cre- mouse. White
 1536 arrows mark ectopic GCs. Calibrations are the same as for the Cre- mouse.
- 1537 **B. A comparison of dorsal and ventral measurements for Cre- and Cre+ male mice**
 1538 show no significant genotype effects.
 1539 1. GluR2/3. A two-way ANOVA showed no effect of dorsal or ventral location
 1540 ($F(1,13)=3.38$; $p=0.089$) or genotype ($F(1,13)=1.158$; $p=0.302$).
 1541 2. SOM. A two-way ANOVA showed no effect of dorsal or ventral location
 1542 ($F(1,10)=0.172$; $p=0.687$) or genotype ($F(1,10)=0.014$; $p=0.908$).
 1543 3. Parvalbumin. A two-way ANOVA showed no effect of dorsal or ventral location
 1544 ($F(1,13)=0.358$; $p=0.560$) or genotype ($F(1,13)=1.068$ $p=0.320$).

- 1545 **C. A comparison of ventral measurements for both Cre- and Cre+ female and male**
1546 **mice.**
- 1547 1. There were significantly more GluR2/3+ hilar cells in Cre+ female mice compared
1548 to Cre- female mice, like the dorsal hippocampus (Fig. 7). Thus, GluR2/3+ hilar
1549 cells were spared in Cre+ females in dorsal and ventral hippocampus. A two-way
1550 ANOVA showed o effect of sex ($F(1,18)=0.744$; $p=0.400$) or genotype
1551 $F(1,18)=0.386$; $p=0.542$) but there was a significant interaction $F(1,18)=5.433$;
1552 $p=0.0316$, and post-hoc tests showed that Cre+ females had significantly more
1553 GluR2/3+ cells than Cre- females ($p=0.045$).
 - 1554 2. There were no significant differences among groups for SOM+ cells. Thus, there
1555 was an effect in dorsal (Fig. 7) but not ventral hippocampus. Thus, SOM cells
1556 were spared in Cre+ females dorsally but not ventrally. A two-way ANOVA
1557 showed no effect of sex ($F(1,17)=0.718$; $p=0.408$) or genotype $F(1,17)=0.769$;
1558 $P=0.393$).
 - 1559 3. There were no significant differences in numbers of parvalbumin+ cells, like
1560 dorsal hippocampus (Fig. 7). Thus, parvalbumin cells were similar regardless of
1561 genotype in dorsal and ventral hippocampus. A two-way ANOVA showed no
1562 significant effect of sex ($F(1,16)=0.401$; $p=0.536$) or genotype ($F(1,16)=0.221$;
1563 $p=0.645$).
- 1564

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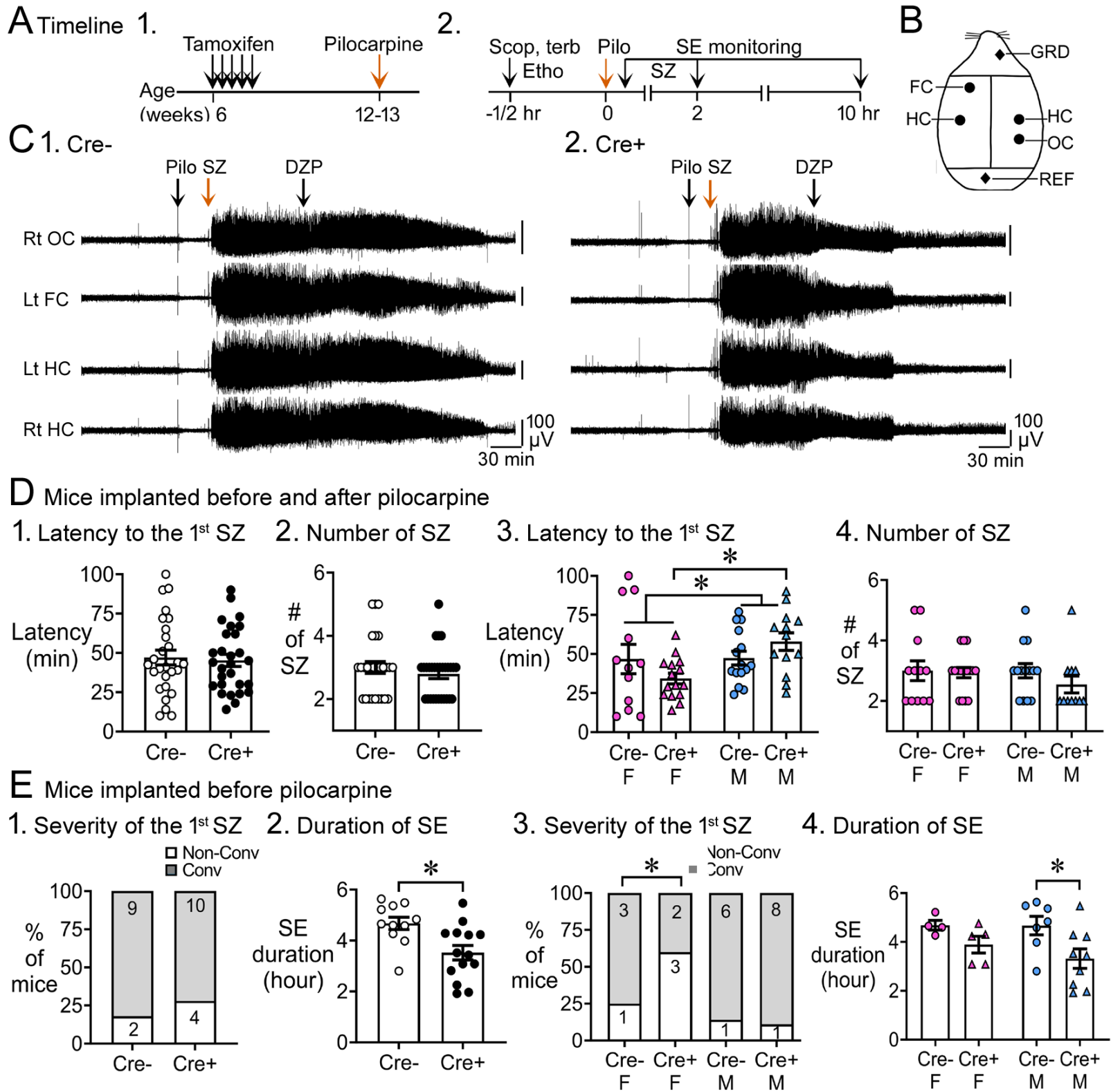
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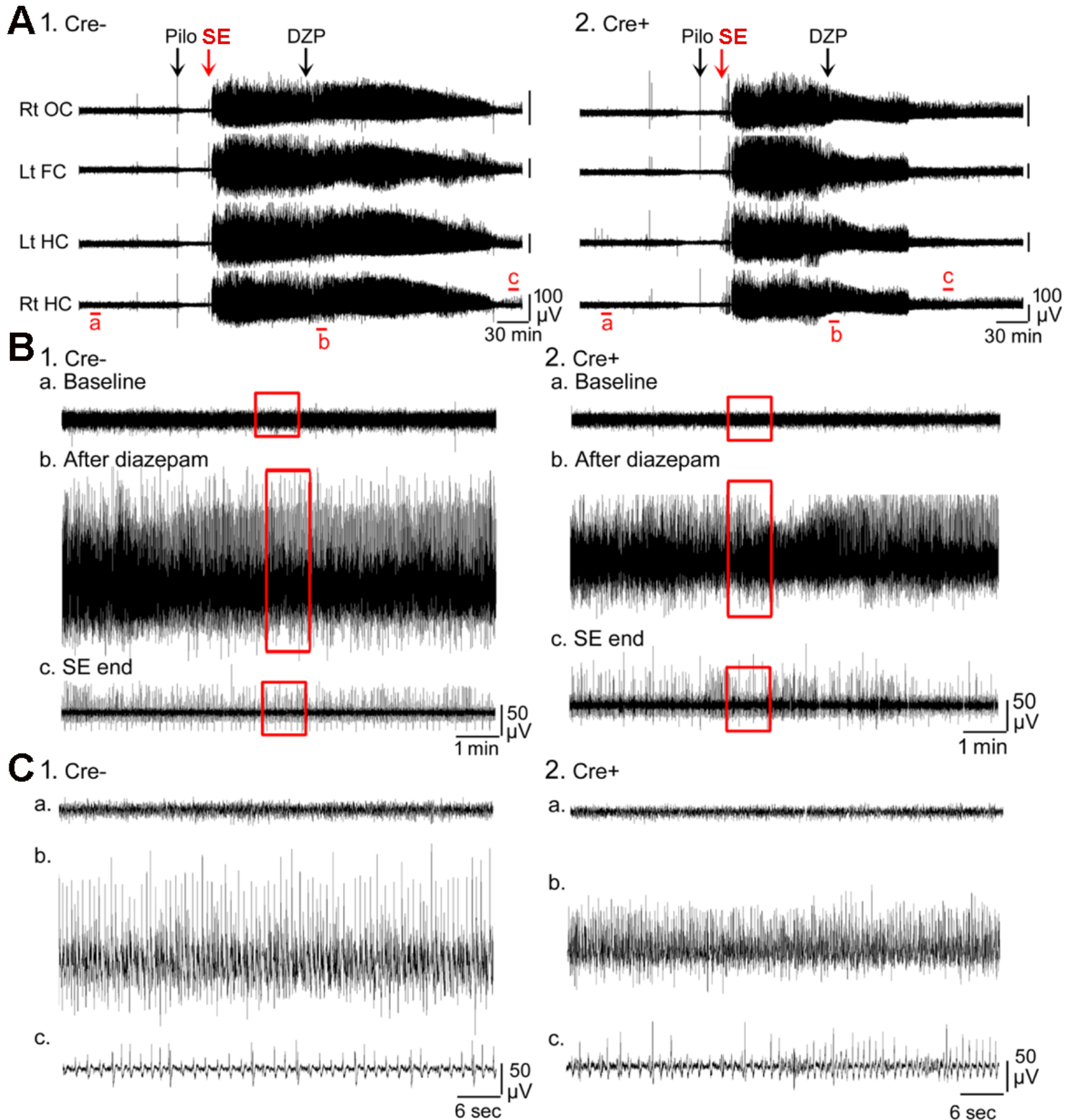
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1975

1976

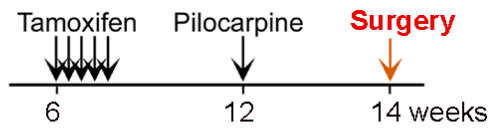
1977





A Timeline

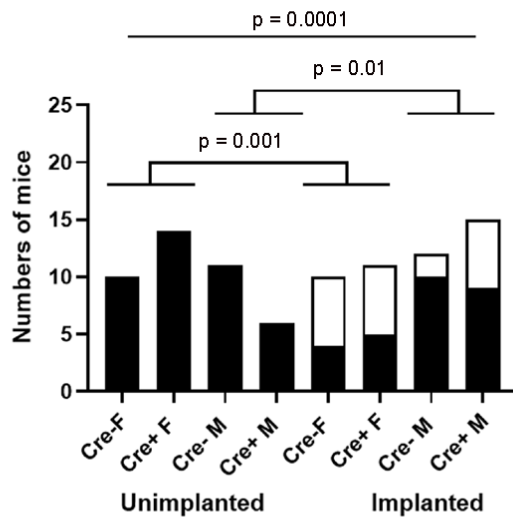
1. Unimplanted



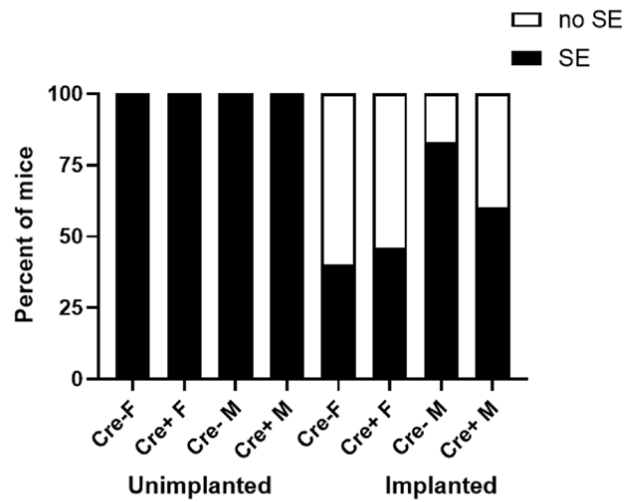
2. Implanted



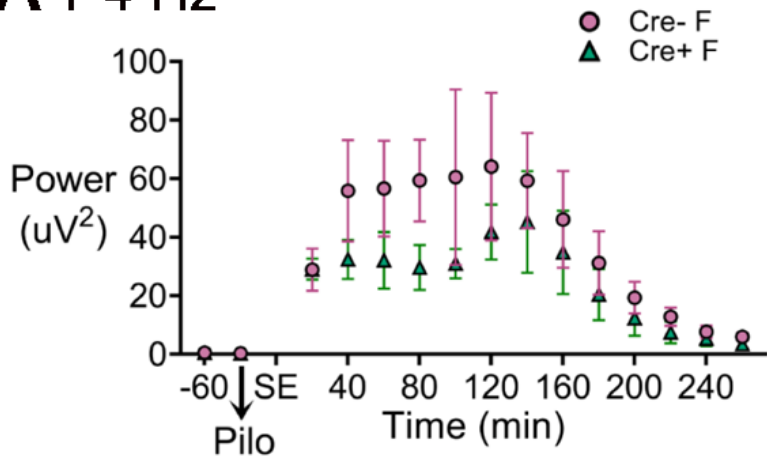
B Numbers of mice



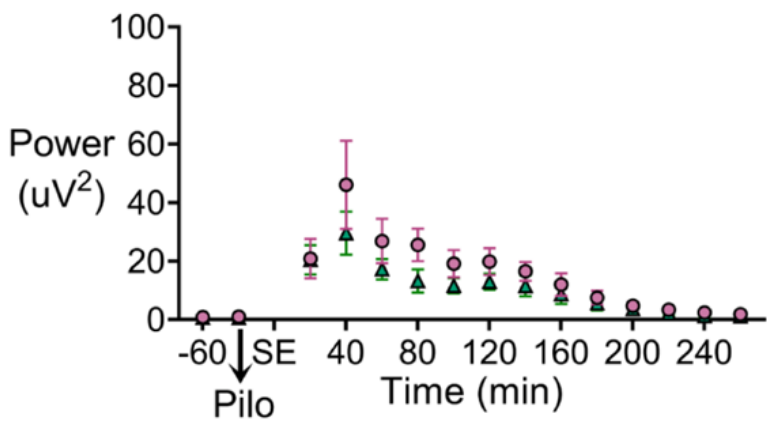
C Percentages of mice



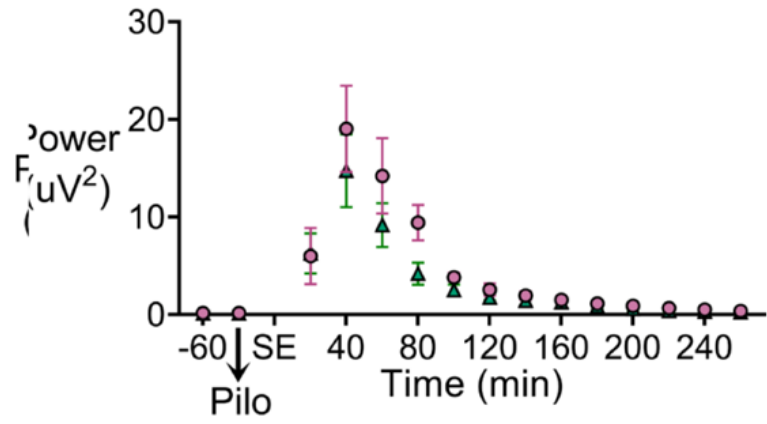
A 1-4 Hz



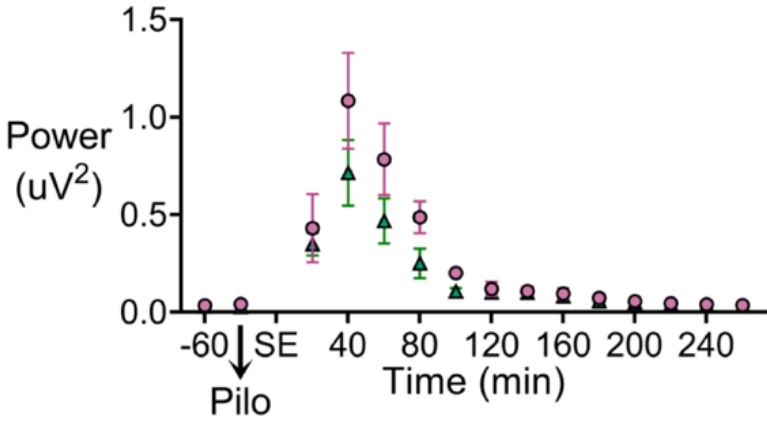
B 4-8 Hz



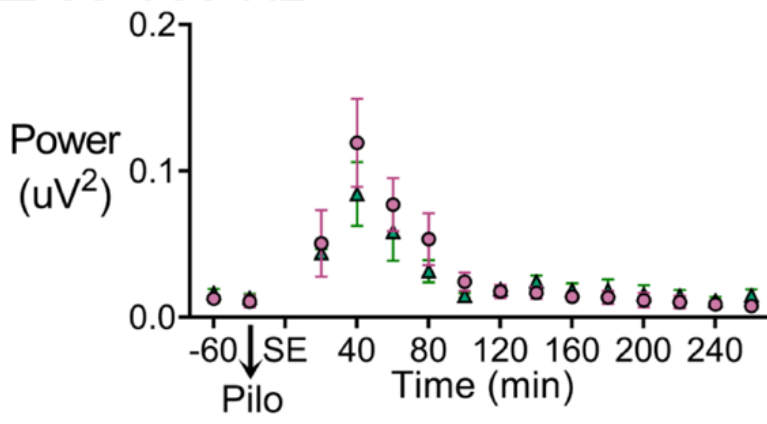
C 8-30 Hz

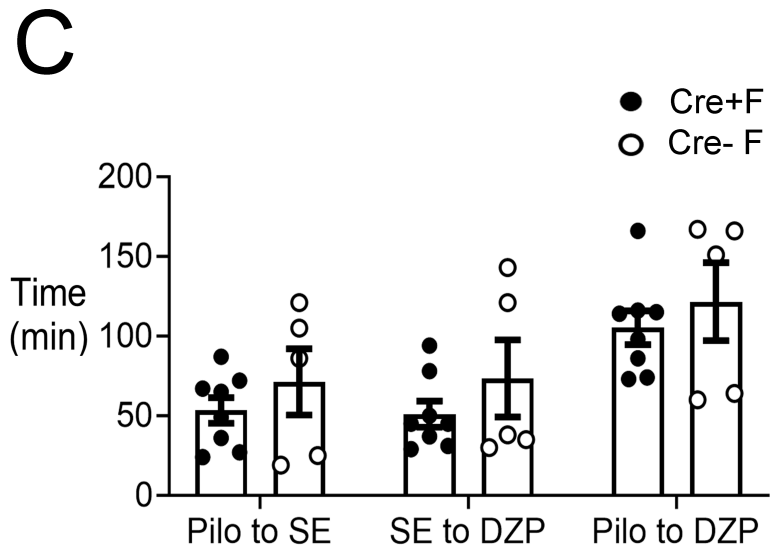
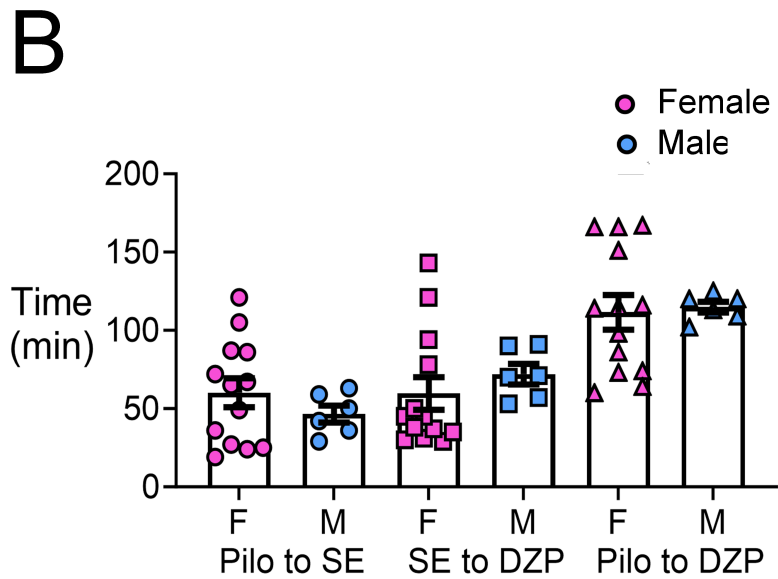
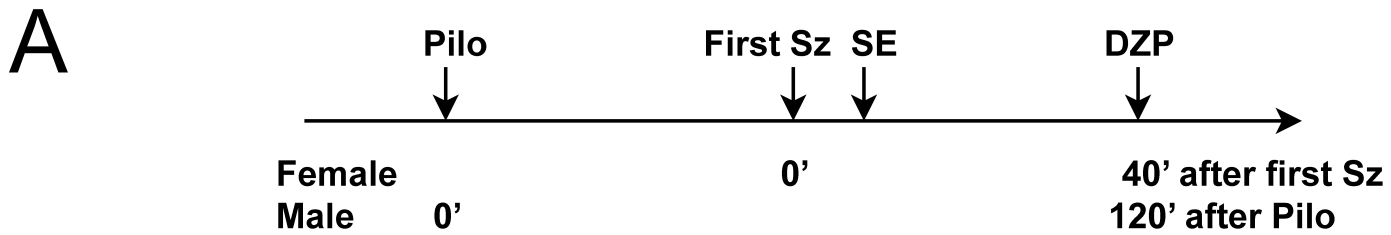


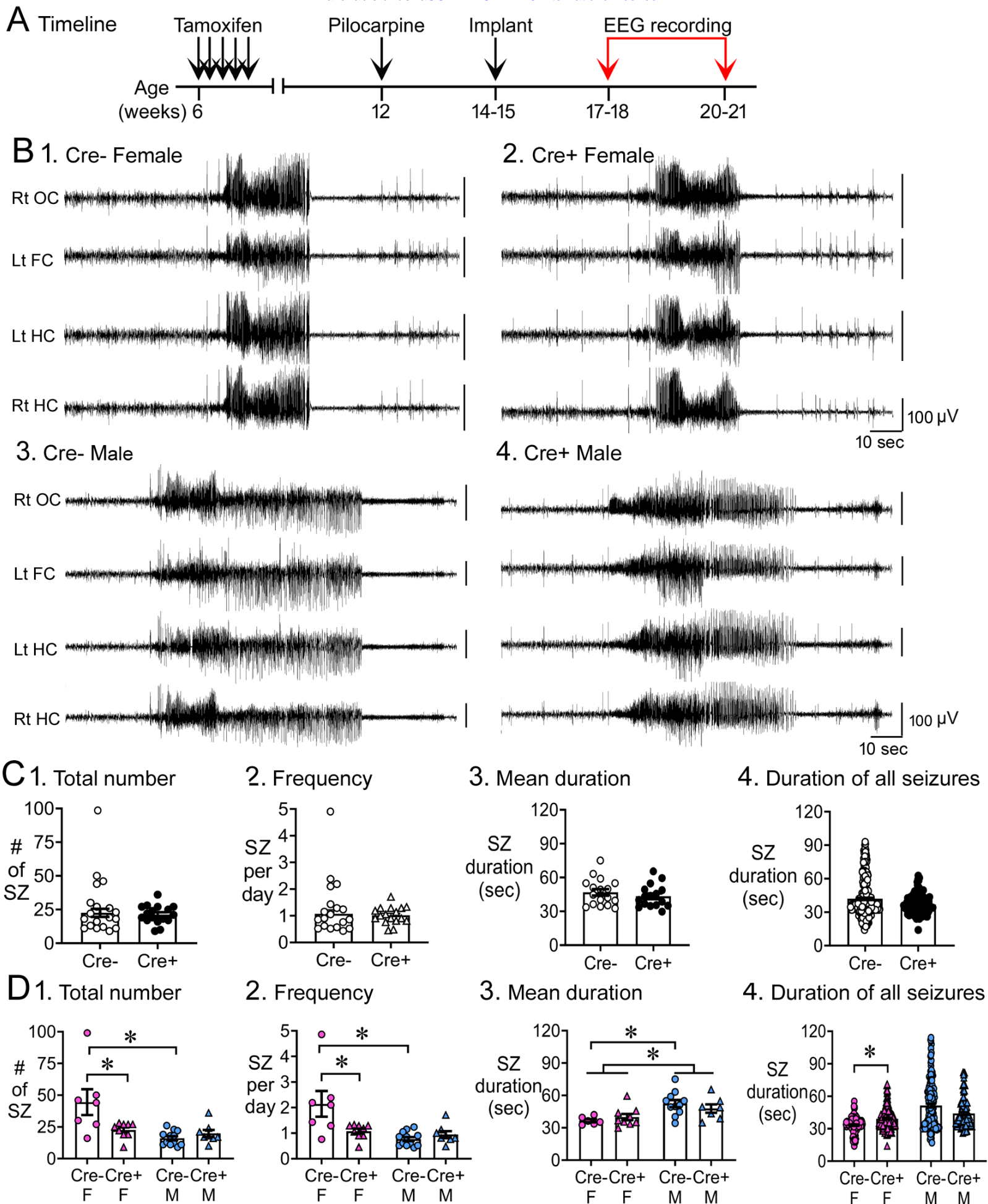
D 30-80 Hz



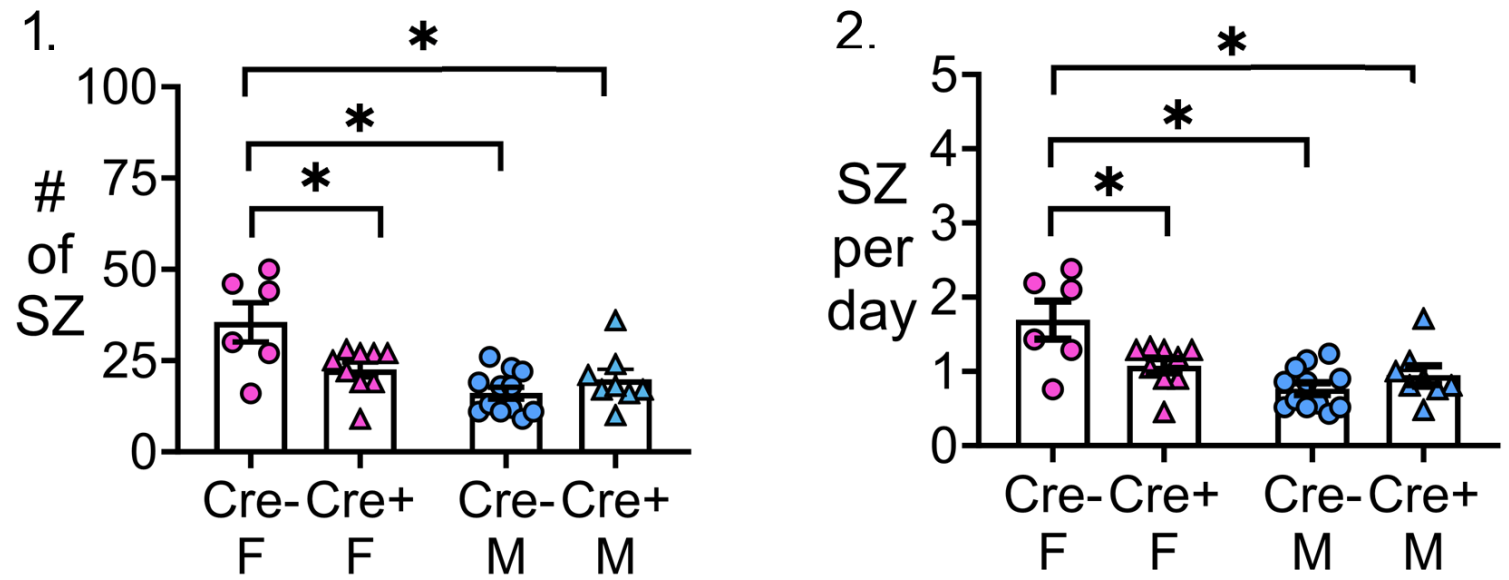
E 80-100 Hz



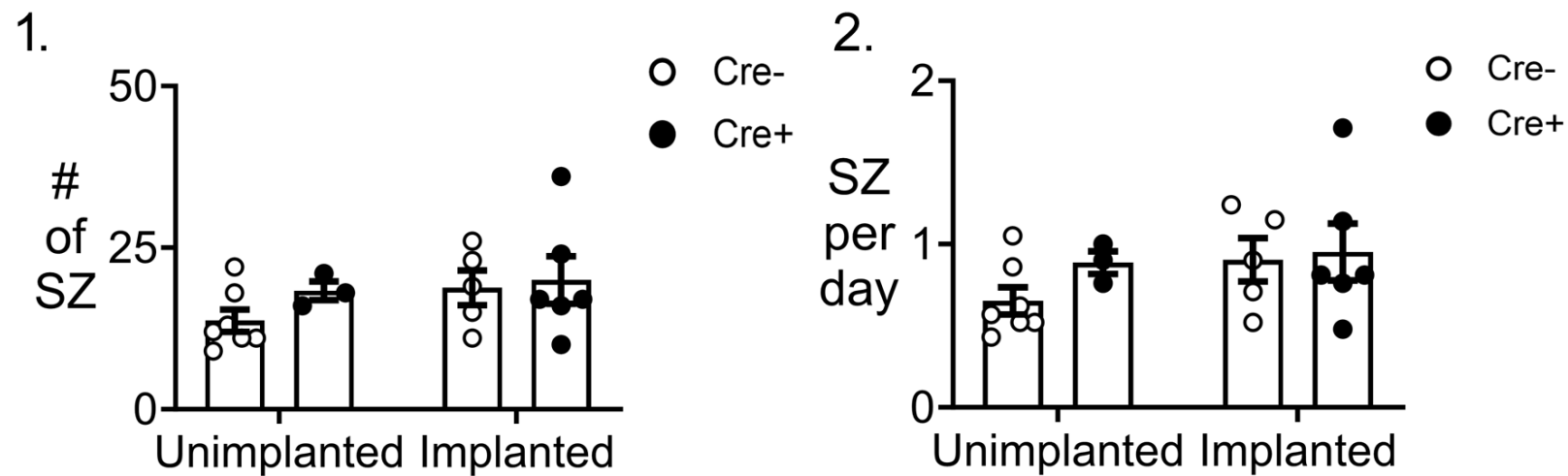




A After outlier removal

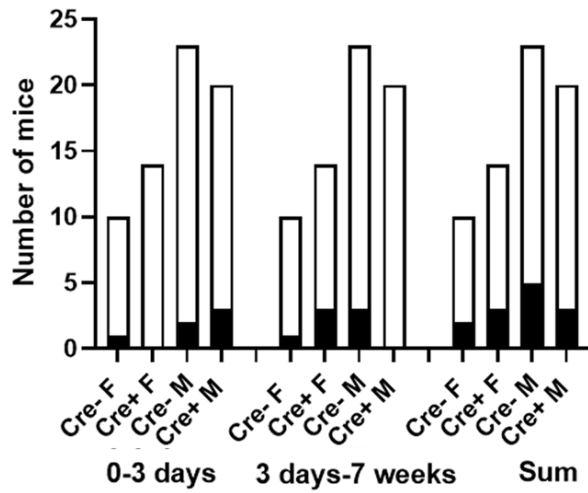


B Similar results for unimplanted and implanted mice

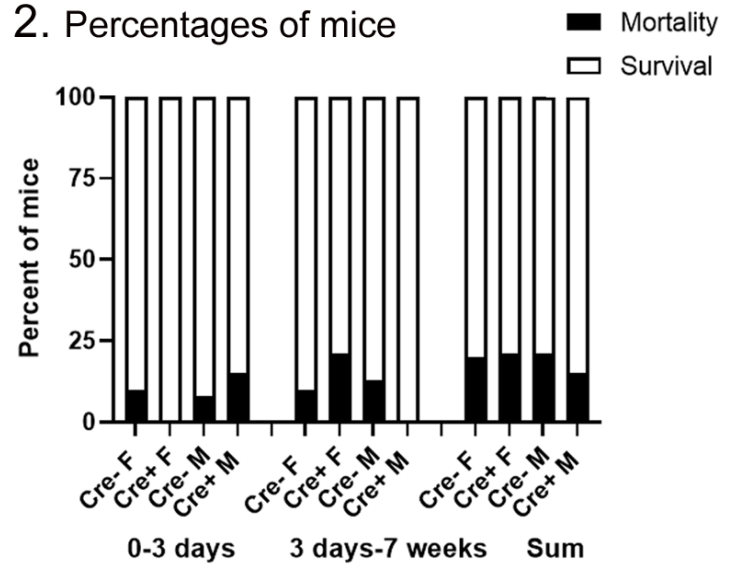




B 1. Numbers of mice



2. Percentages of mice

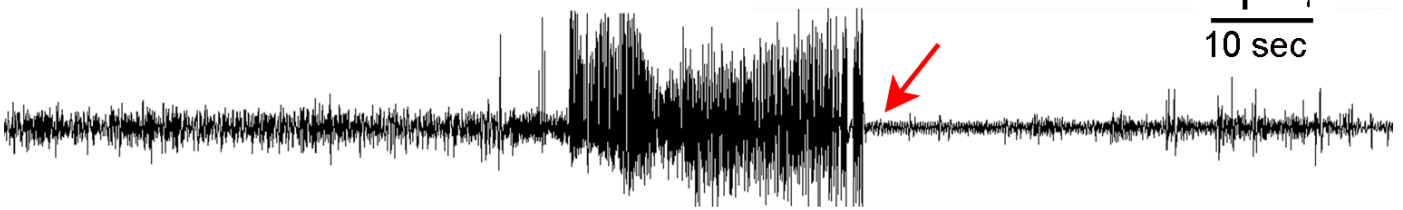


A1

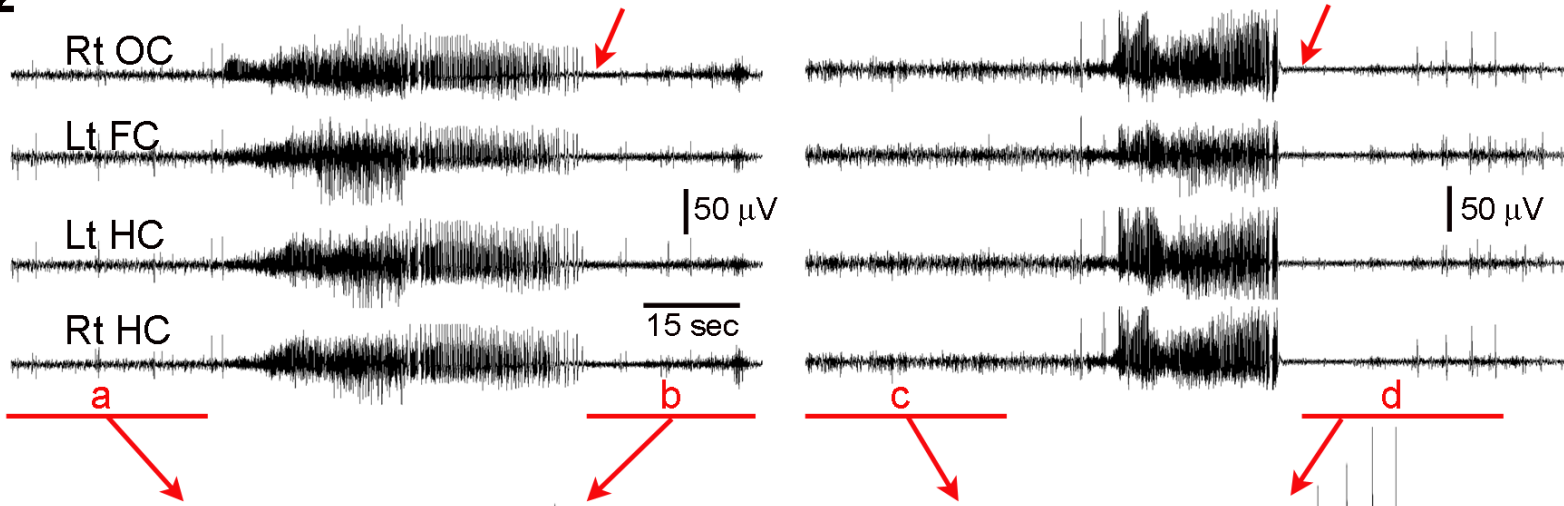
Male



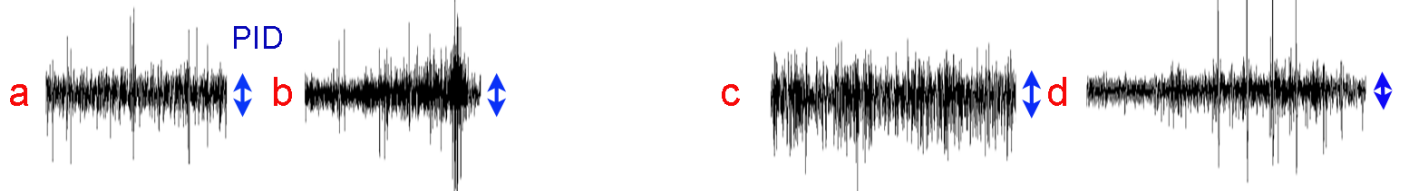
Female



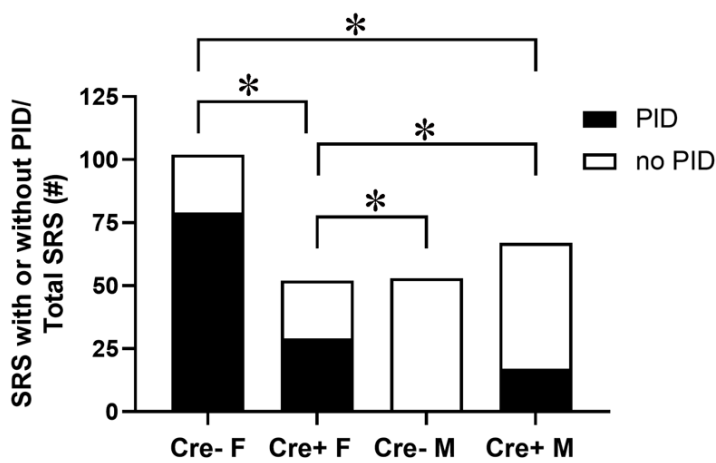
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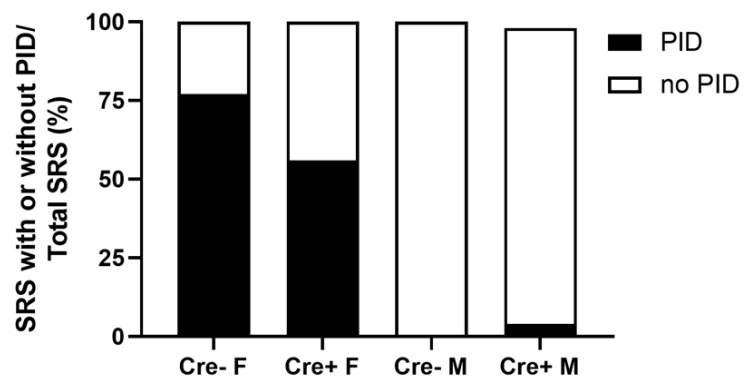
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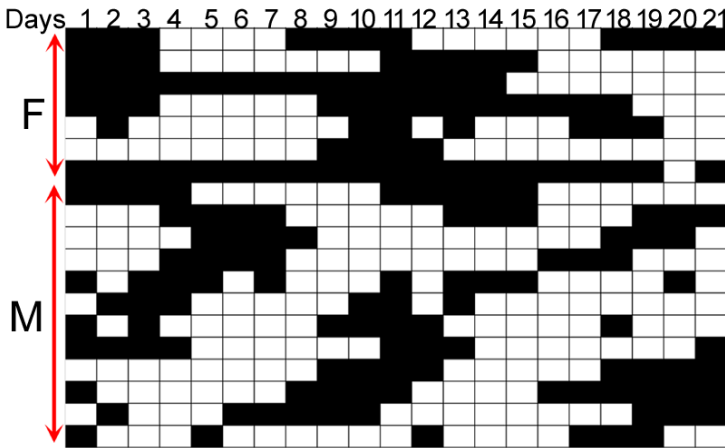
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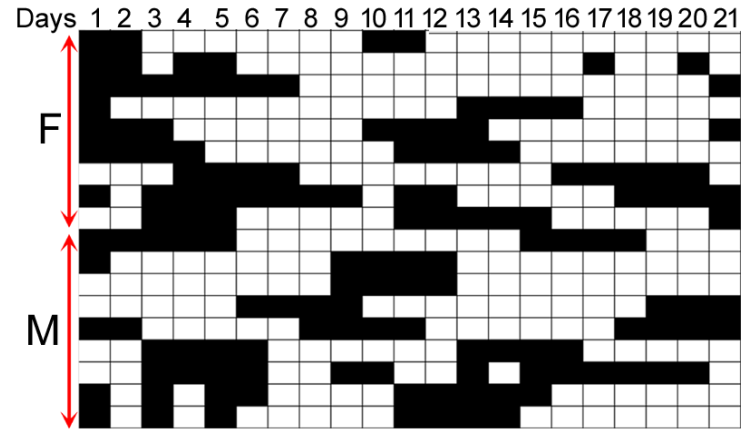
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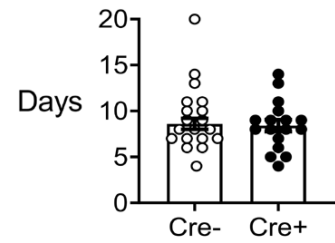
A1. Cre-



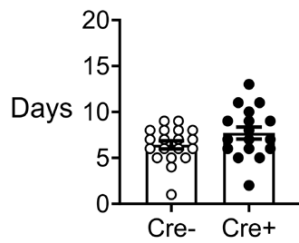
2. Cre+



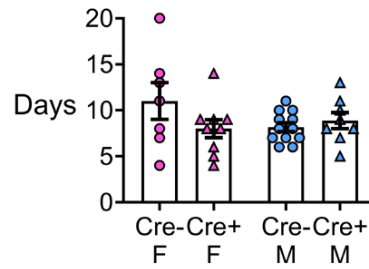
B1. Days with SZ



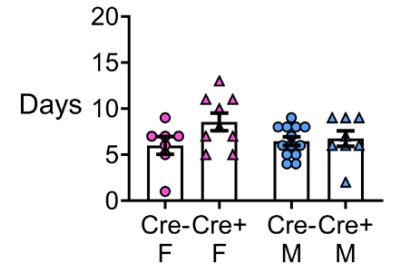
2. Max. SZ-free interval



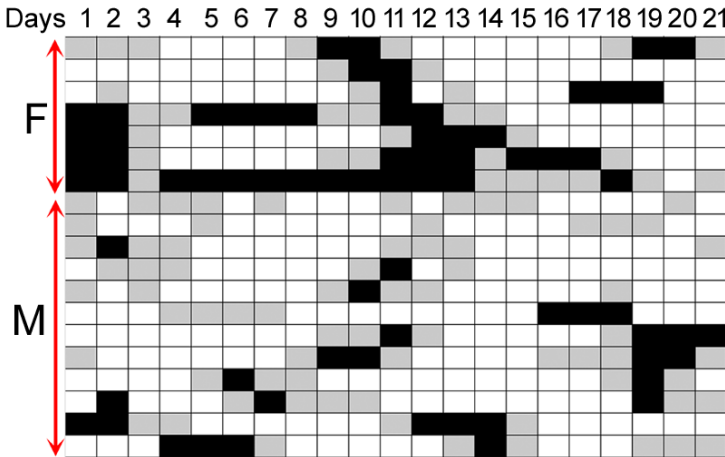
3. Days with SZ



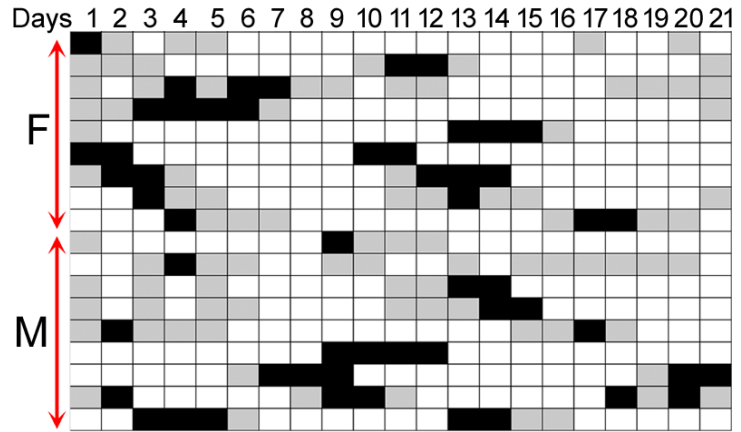
4. Max. SZ-free interval



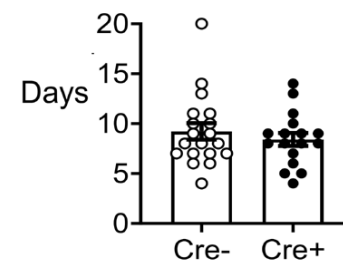
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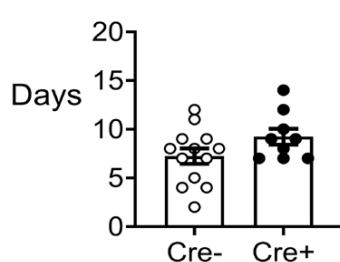
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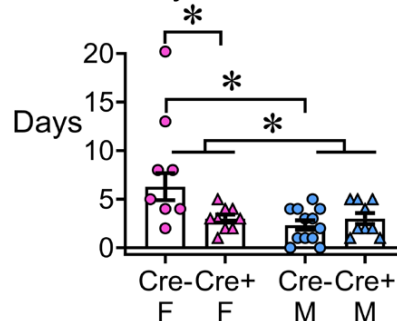
D1. Days with ≥3 SZ



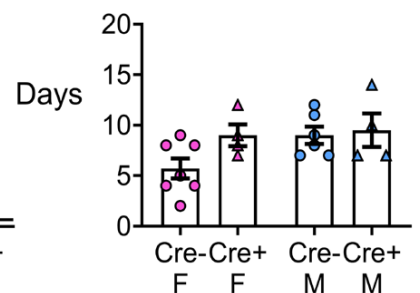
2. Max. inter-cluster interval



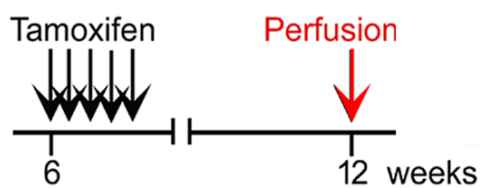
3. Days with ≥3 SZ



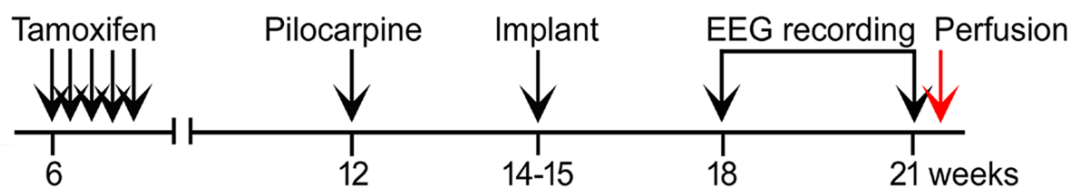
4. Max. inter-cluster interval



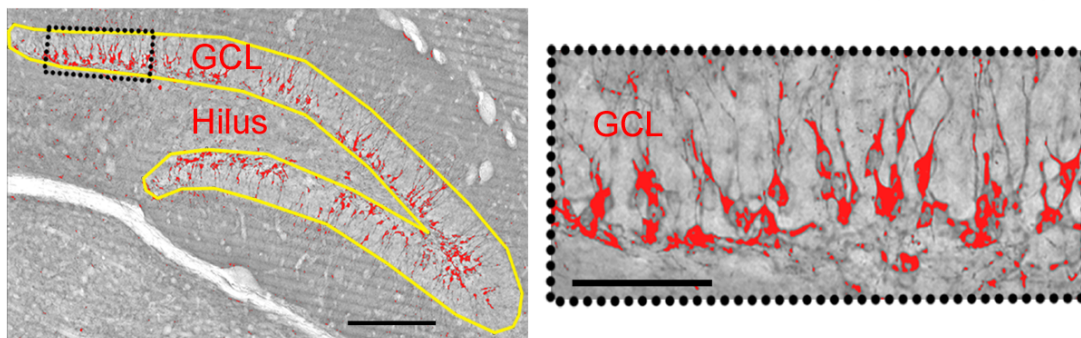
A DCX before SE



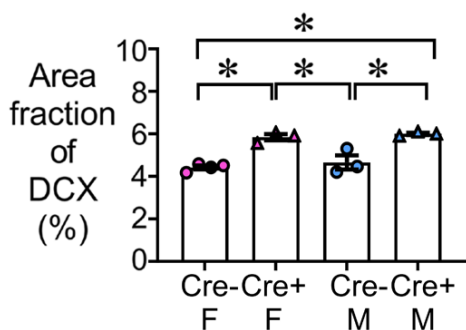
B DCX 2 months after SE



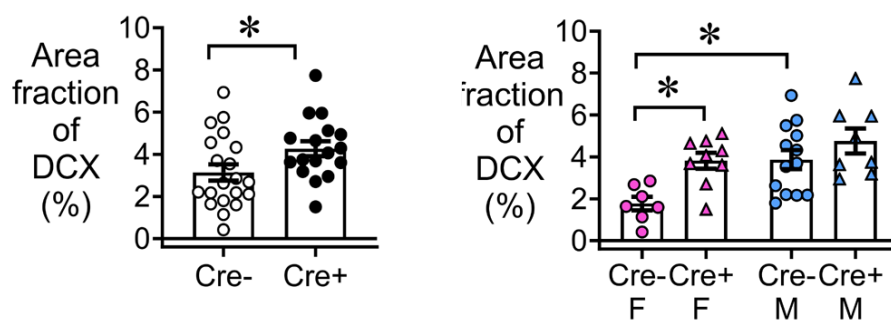
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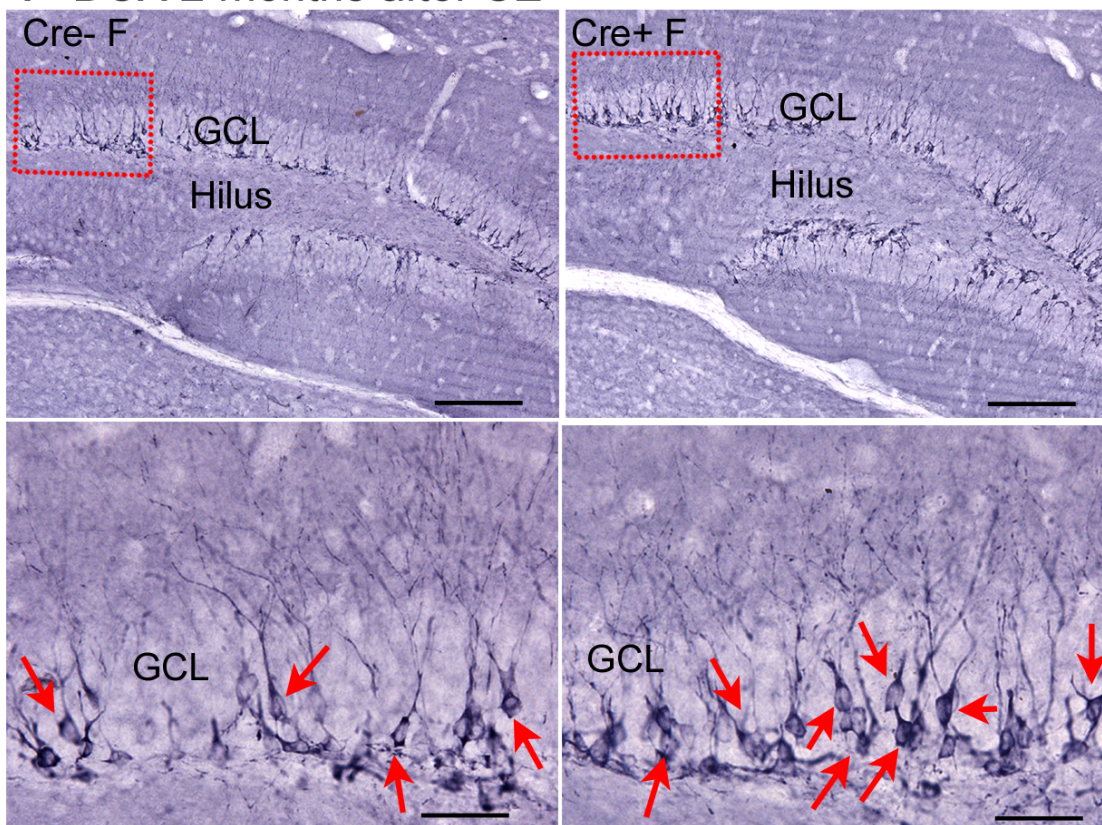
D DCX before SE



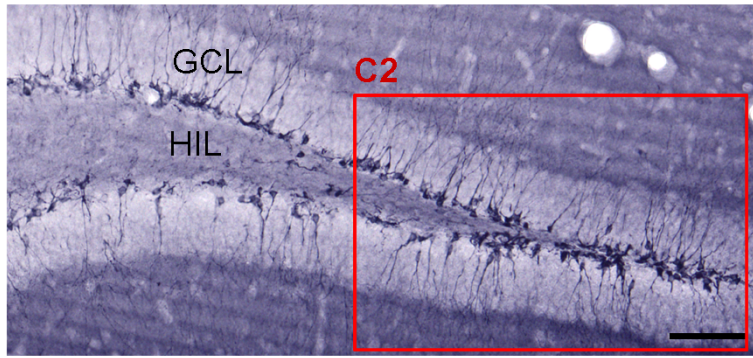
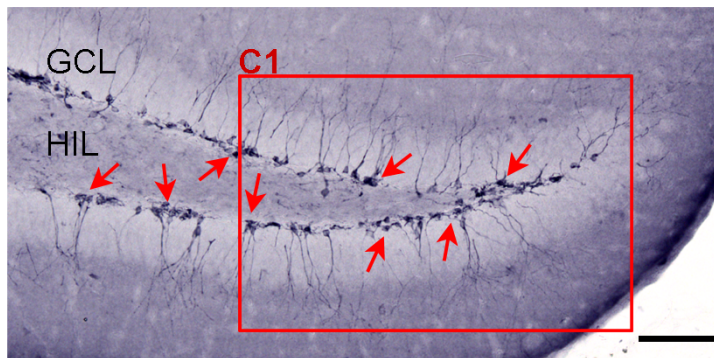
E DCX 2 months after SE



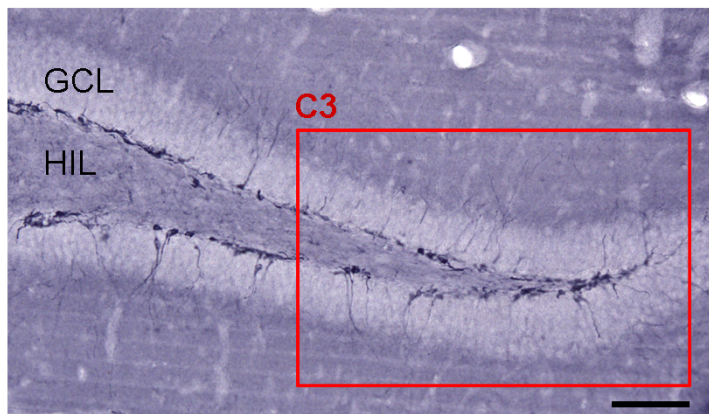
F DCX 2 months after SE



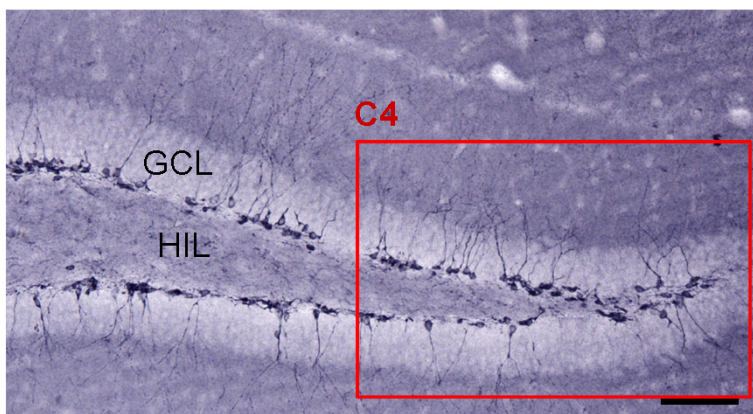
A 1. Cre- F



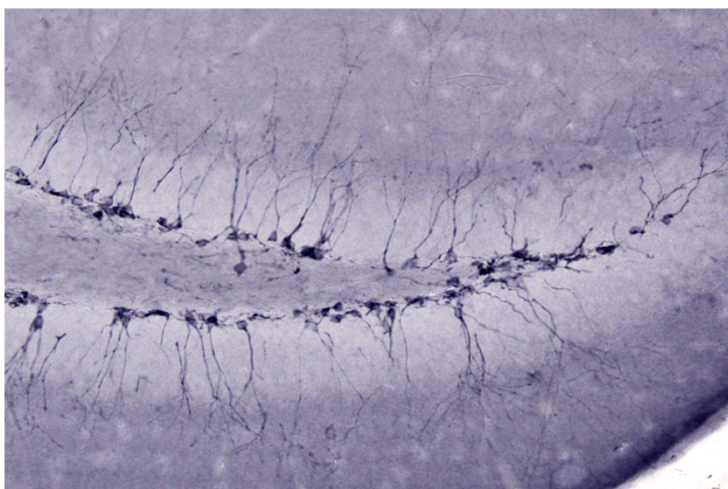
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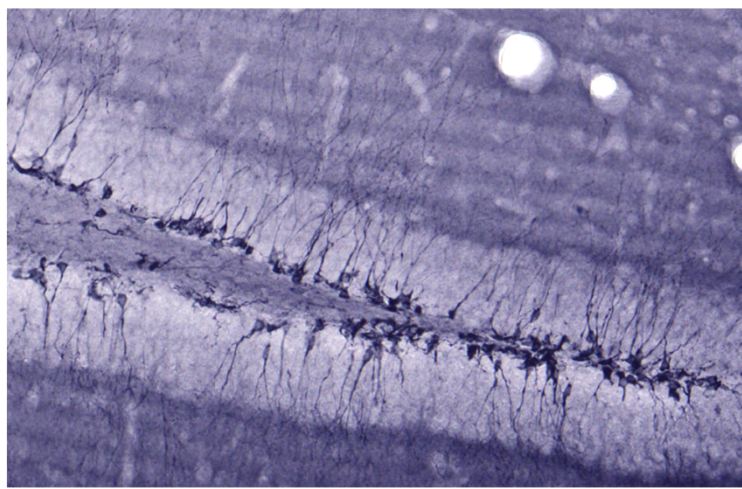
2. Cre+ M



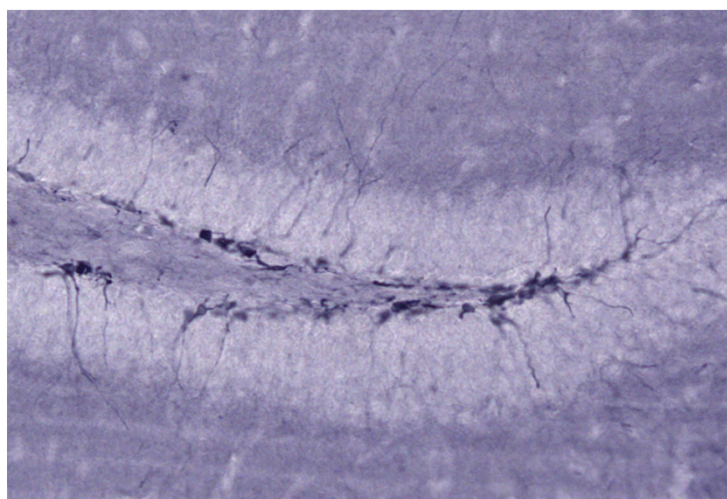
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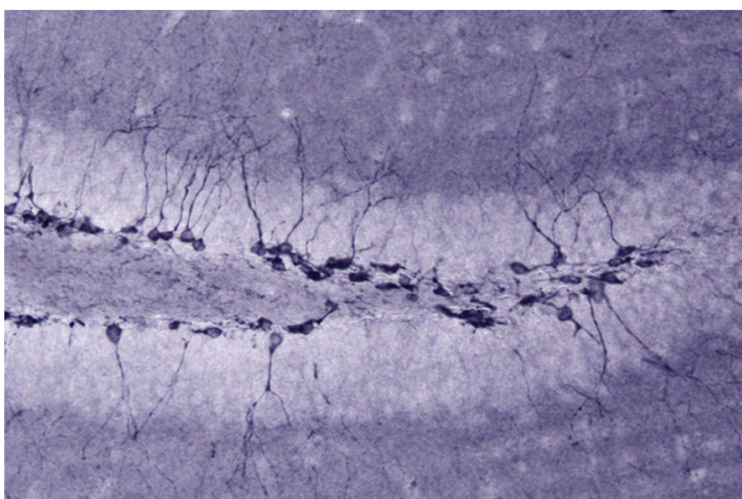
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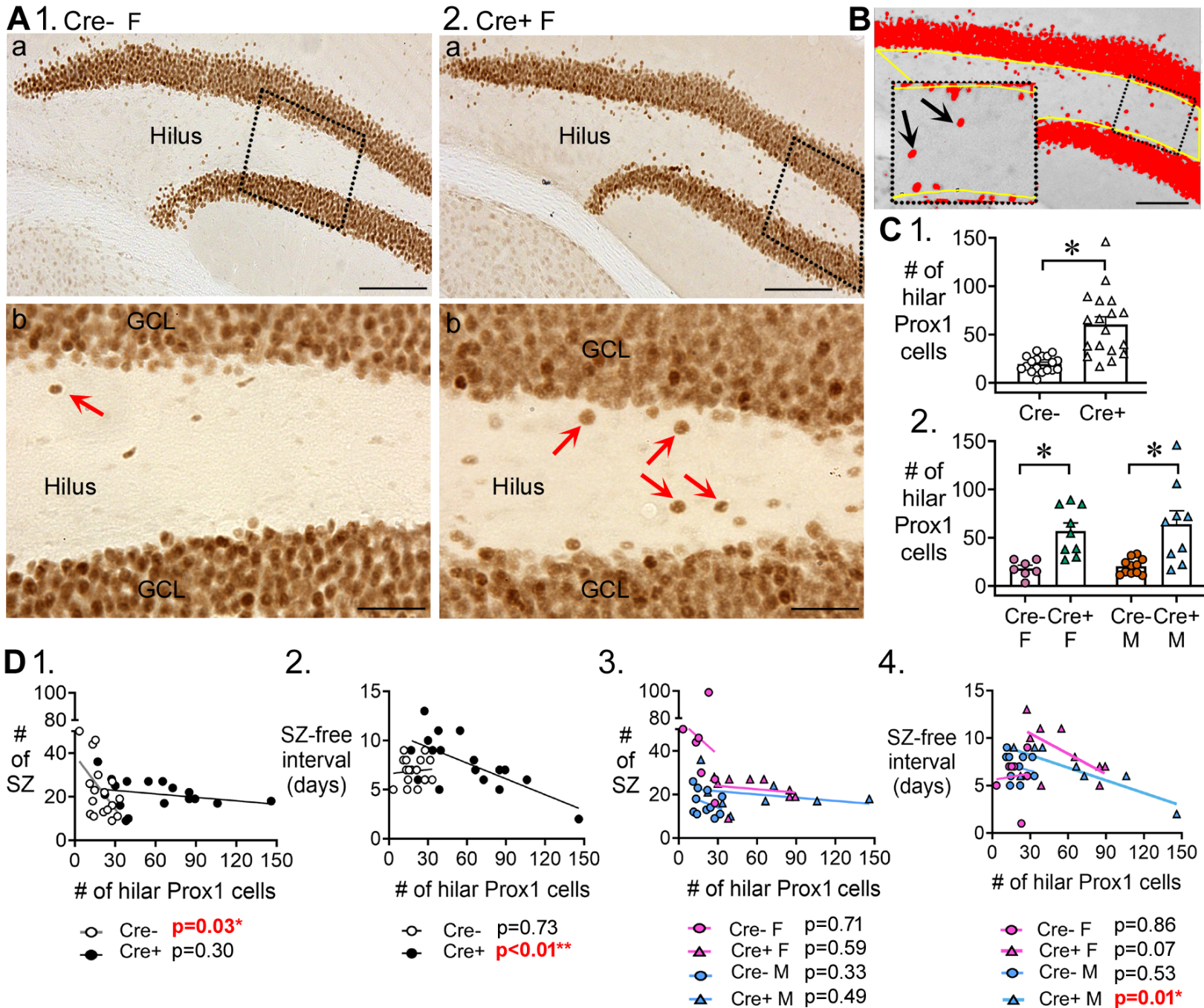


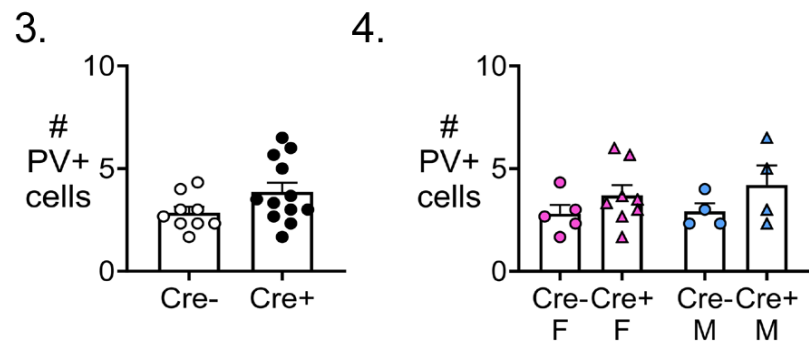
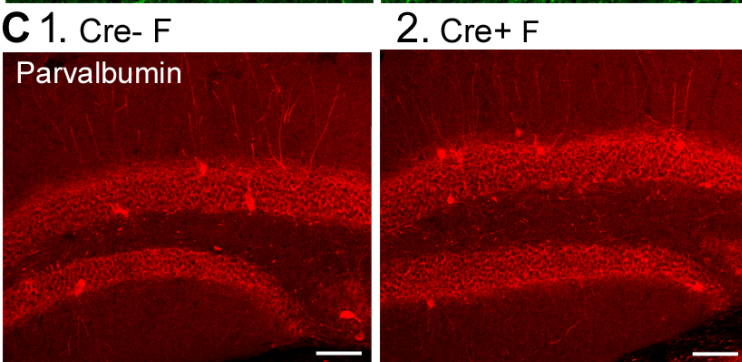
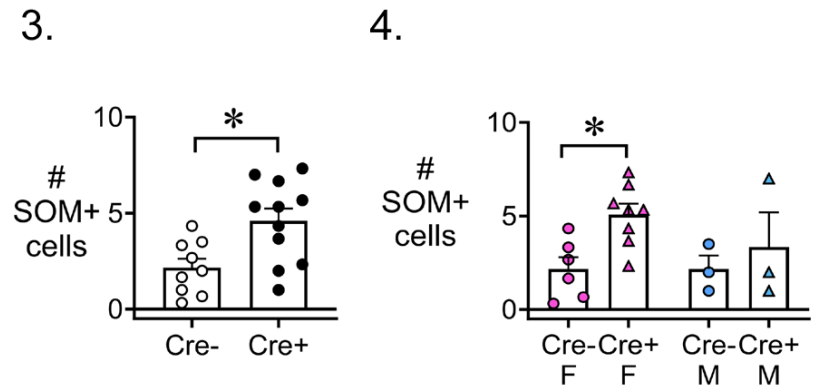
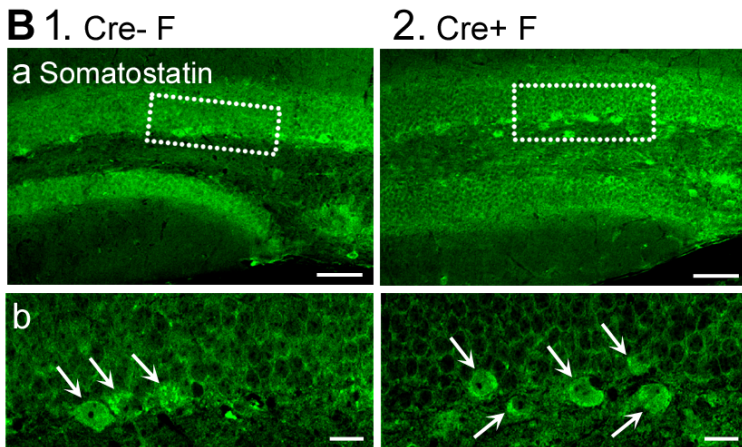
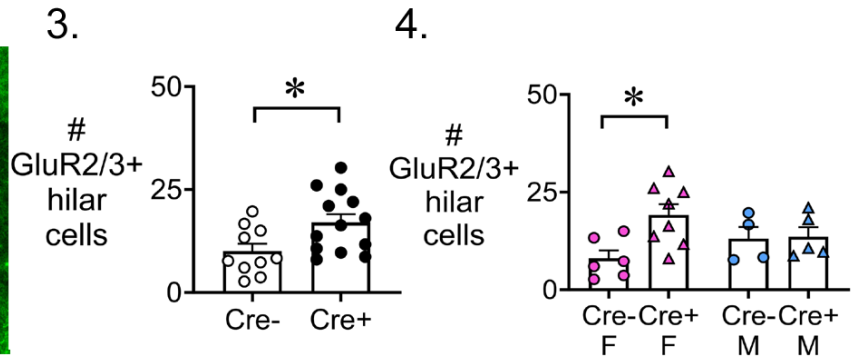
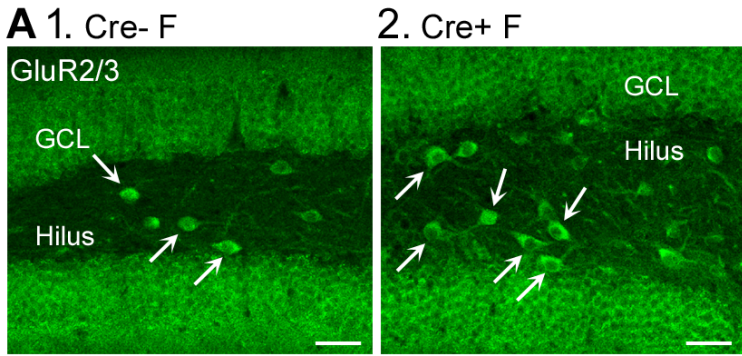
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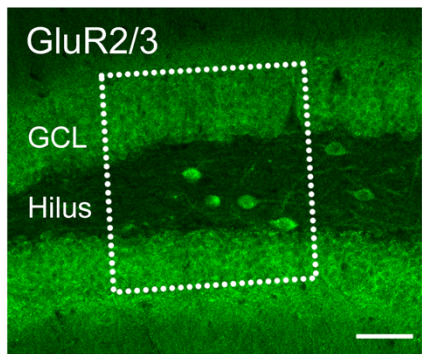
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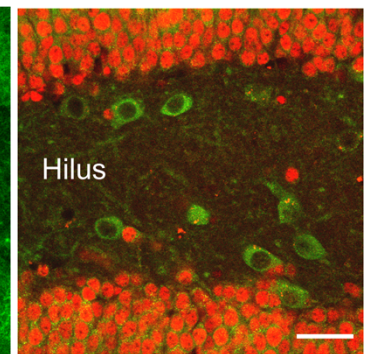
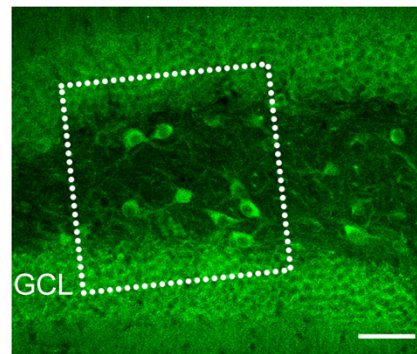
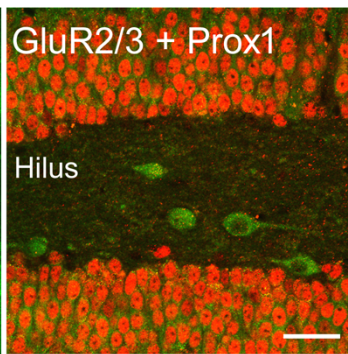




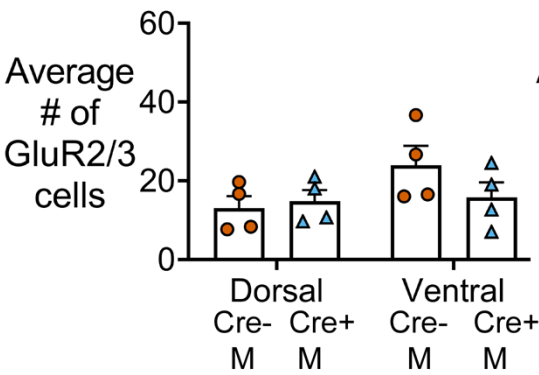
A 1. Cre- F



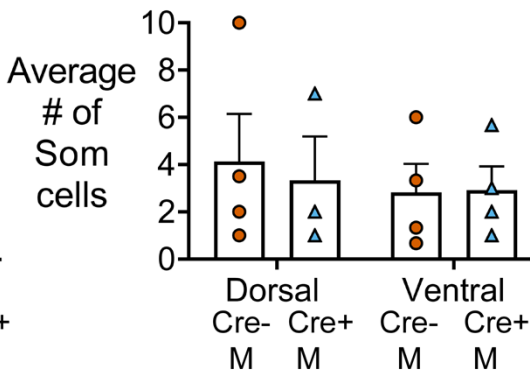
2. Cre+ F



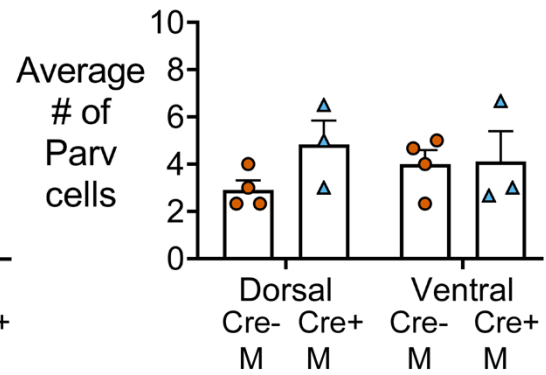
B 1. GluR2/3 cells



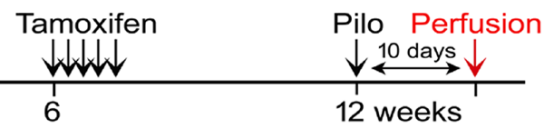
2. Somatostatin



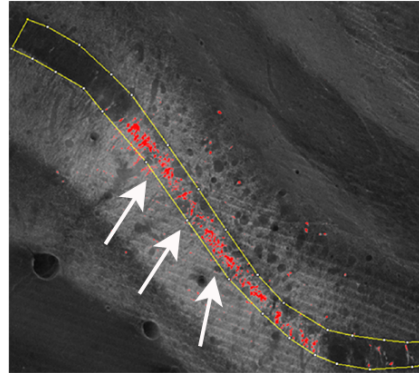
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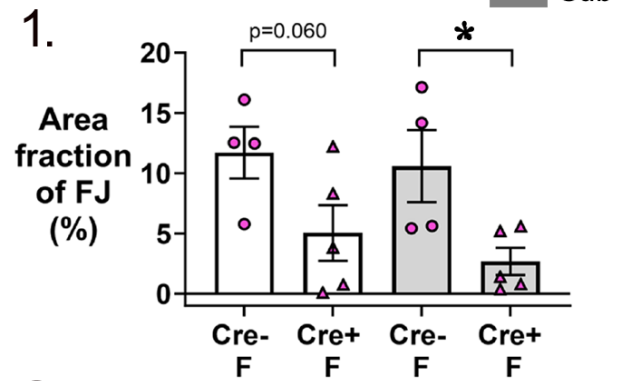
A Timeline



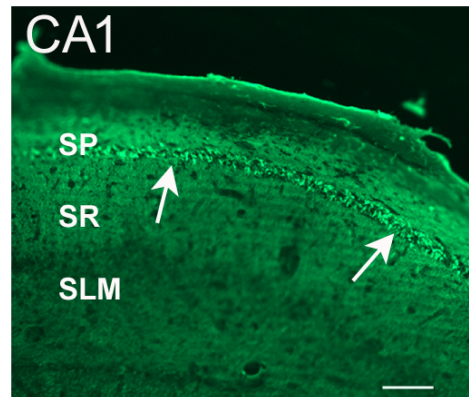
B Quantification



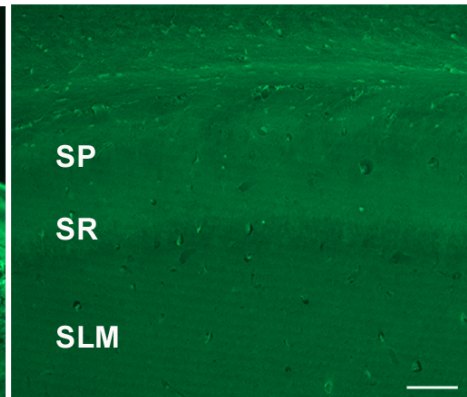
D



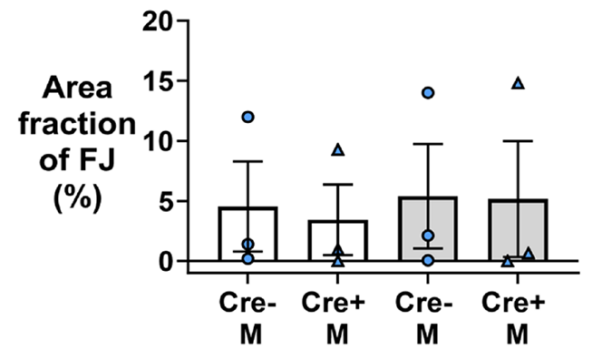
C 1. Cre- F



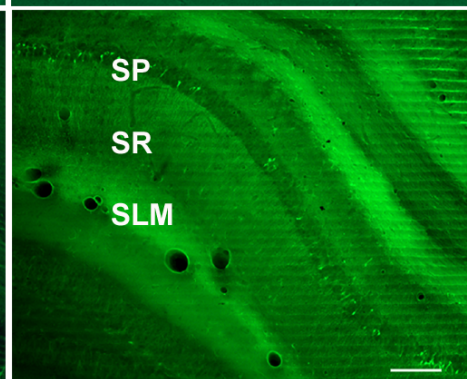
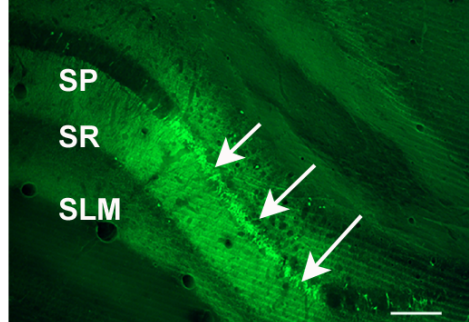
2. Cre+ F



2.



Subiculum



3.

