

# Perineuronal nets support astrocytic ion and glutamate homeostasis at tripartite synapses

**Bhanu Tewari** (✉ [bptewari@gmail.com](mailto:bptewari@gmail.com))

University of Virginia <https://orcid.org/0000-0001-8803-5950>

**AnnaLin Woo** (✉ [rsx2bn@virginia.edu](mailto:rsx2bn@virginia.edu))

University of Virginia

**Courtney Prim** (✉ [cep2sy@virginia.edu](mailto:cep2sy@virginia.edu))

University of Virginia

**Lata Chaunsali** (✉ [hbe5rr@virginia.edu](mailto:hbe5rr@virginia.edu))

University of Virginia

**Ian Kimbrough** (✉ [pku4qv@virginia.edu](mailto:pku4qv@virginia.edu))

University of Virginia

**Kaliroi Engel** (✉ [ekali6@vt.edu](mailto:ekali6@vt.edu))

School of Neuroscience, Virginia Tech

**Jack Browning** (✉ [jackb7@vt.edu](mailto:jackb7@vt.edu))

School of Neuroscience, Virginia Tech <https://orcid.org/0000-0002-3102-0667>

**Susan Campbell** (✉ [susanc08@vt.edu](mailto:susanc08@vt.edu))

Department of Animal Sciences, Virginia Tech

**Harald Sontheimer** (✉ [sontheimer@virginia.edu](mailto:sontheimer@virginia.edu))

University of Virginia <https://orcid.org/0000-0002-5843-9871>

---

## Article

### Keywords:

DOI: <https://doi.org/>

**License:**  This work is licensed under a Creative Commons Attribution 4.0 International License.

[Read Full License](#)

**Additional Declarations:** There is **NO** Competing Interest.

---

1 **Title:**

2

3 Perineuronal nets support astrocytic ion and glutamate homeostasis at tripartite synapses

4

5 **Authors:**

6

7 Bhanu P. Tewari 1, AnnaLin M. Woo\* 1, Courtney E. Prim\* 1, Lata Chaunsali 1, Ian F.  
8 Kimbrough 1, Kaliroi Engel 2, Jack L. Browning 2, Susan L. Campbell 3, Harald  
9 Sontheimer 1

10

11 **Affiliations:**

12

13 1 Department of Neuroscience, University of Virginia School of Medicine, Charlottesville,  
14 VA, USA

15 2 School of Neuroscience, Virginia Tech, Blacksburg, VA, USA

16 3 Department of Animal Sciences, Virginia Tech, Blacksburg VA, USA

17

18 **Author's contribution:**

19

20 BPT: conceived idea, design, execution, analysis, and interpretation of electrophysiology,  
21 IHC, and imaging experiments, manuscript writing, editing, and communication.

22 CP, AMW, and LC: animal surgery, IHC, and confocal imaging and analysis.

23 IFK: image analysis and 3D reconstruction.

24 KE and JLB: IHC and analysis assistance.

25 SLC: electrophysiology assistance and data analysis.

26 HWS: Conceived idea, project supervision, data interpretation, manuscript writing and  
27 editing, funding acquisition.

28 \* These authors contributed equally.

29 **Acknowledgment:**

30

31 This work was supported by the National Institutes of Health grants R01AG065836,  
32 R01NS036692, and R01NS123069 awarded to HS.

33 We acknowledge Jenny Raines Ph.D. for her assistance with animal handling and  
34 genotyping.

35

36 **Conflict of Interest:**

37

38 Authors declare no conflict of interest.

39

40 **Correspondence:**

41

42 Harald Sontheimer, sontheimer@virginia.edu

43

44

45 **Abstract:**

46 Perineuronal nets (PNNs) are dense, negatively charged extracellular matrices that cover  
47 the cell body of fast-spiking inhibitory neurons. Synapses can be embedded and  
48 stabilized by PNNs believed to prevent synaptic plasticity. We find that in cortical fast-  
49 spiking interneurons synaptic terminals localize to perforations in the PNNs, 95% of which  
50 contain either excitatory or inhibitory synapses or both. The majority of terminals also  
51 colocalize with astrocytic processes expressing Kir4.1 as well as glutamate (Glu) and  
52 GABA transporters, hence can be considered tripartite synapses. In the adult brain,  
53 degradation of PNNs does not alter axonal terminals but causes expansion of astrocytic  
54 coverage of the neuronal somata. However, loss of PNNs impairs astrocytic transmitter  
55 and K<sup>+</sup> uptake and causes spillage of synaptic Glu into the extrasynaptic space. This data  
56 suggests a hitherto unrecognized role of PNNs, to synergize with astrocytes to contain  
57 synaptically released signals.

58

59

60

61

62

63

64

65

66

67

68

69

70

71

72

73

74

75

76

77 **Introduction:**

78 With the discovery that up to 99% of excitatory synapses are associated with astrocytic  
79 processes[1], the classical view of the synapse as being formed only by two neurons has  
80 changed dramatically. Often called the “tripartite synapse”, this arrangement allows for  
81 astrocytes to sense synaptic activity and potentially modulate activity via the release of  
82 gliotransmitters[2]. Importantly, at the tripartite synapse astrocytic processes are at the  
83 perfect place to remove neurotransmitters and ions released in conjunction with synaptic  
84 activity. Astrocytic ensheathment of glutamatergic synapses is particularly important to  
85 ensure that glutamate (Glu) does not spill out of synapses and activate extra-synaptic  
86 receptors[3] as this can cause excitotoxicity[4].

87 Neurons and astrocytes are also embedded in an extracellular matrix (ECM) made up of  
88 proteoglycans, glycoproteins, and polysaccharides. Neurons and astrocytes each  
89 synthesize defined ECM constituents including hyaluronan (HA), chondroitin sulfate  
90 proteoglycans (CSPGs) such as aggrecan, brevican, versican and neurocan, Tenascin R  
91 (Tn-R), and link proteins such as Crtl1 and Bral2[5, 6]. Specific ECM constituents have  
92 been shown to directly interact with synaptic receptors and ion channels [7-9] thereby  
93 affecting synaptic vesicle release[10], dendritic spine morphology [11-14], and the  
94 structural integrity of synapses[13, 15]. Moreover, due to the negative charges associated  
95 with sulfate groups on the CSPGs, the ECM has been implicated in altering diffusion of  
96 ions in the extracellular space [16]as well as binding water molecules.

97 On some cortical inhibitory neurons, particularly those expressing the Ca<sup>2+</sup> binding protein  
98 parvalbumin (PV), the ECM forms a highly condensed corset-like structure known as  
99 perineuronal nets (PNNs). Easily recognized by the binding of wisteria floribunda  
100 agglutinin (WFA) [17], PNNs encapsulate the cell soma, dendrites, and axon initial  
101 segment. PNNs have been shown to stabilize synapses and restrict synaptic plasticity,  
102 particularly in pathways that show activity-dependent plasticity during development such  
103 as the visual system[18]. However, whether and how PNNs interact structurally and  
104 functionally with astrocytes at tripartite synapses is unknown.

105 In the present study, we show that synapses onto PNN-expressing fast-spiking neurons  
106 (FSNs) exist in small perforations or holes within the PNN and these contain excitatory  
107 and inhibitory synapses in conjunction with astrocytic processes, hence can be  
108 considered tripartite synapses. Upon PNNs disruption, the synapses retain their place on  
109 the neuron but the astrocytes expand their coverage to vacated areas on the cell body.  
110 Importantly, we show that PNN disruption impedes astrocytic uptake of synaptically  
111 released Glu and K<sup>+</sup> and causes the spillage of glutamate into the perisynaptic space.  
112 This suggests a hitherto unrecognized function of the PNNs namely to create a barrier  
113 that limits diffusion of Glu and K<sup>+</sup> from the synaptic cleft so as to synergize with astrocytes  
114 for effective reuptake of neuronally released ions and transmitters. Hence PNNs are an  
115 important structural and functional contributor to the tripartite synapse.

116



117 **Results:**

118 **1. Astrocyte-PNN spatial interface:**

119 Tripartite synapses on excitatory neurons are typically on spines that are almost  
120 completely ensheathed by astrocytic processes also called leaflets[19]. The near  
121 complete coverage of the synapse by astrocytes facilitates effective clearance of  
122 synaptically released ions and neurotransmitters. On inhibitory neurons, the PNN forms  
123 a coat around the cell body, some of its dendrites, and the axon initial segment. Small  
124 holes or perforations of the PNN provide access for synaptic terminals. Hence our first  
125 question was whether these axosomatic synapses on inhibitory neurons are tripartite  
126 synapses, i.e., contain astrocytic processes associated with the synapse.

127 To visualize astrocytic processes in PNN holes, we used FVB-N//Swiss Webster-Aldh111-  
128 eGFP mice expressing eGFP in astrocytes [20] to image astrocytes in relationship to  
129 WFA-labeled PNNs in layers 3 - 4 of the somatosensory cortex. This is an area of the  
130 highest PNN density where nearly all astrocytes are contacting PNNs. Confocal images  
131 show a majority of astrocytic processes terminating within PNN holes however, some  
132 astrocytic processes terminated on the outer side of the PNN (white and yellow arrows  
133 **Fig. 1a**). We rarely observed astrocytic processes interspaced between the PNN and  
134 neuronal cell body. To quantify these images, we generated intensity profiles along the  
135 PNNs (**Fig. 1a**, dotted line). These profiles show peaks of astrocytic AldheGFP (**Fig. 1a**,  
136 green line) in the holes of PNNs (with lowest WFA signal, red line) suggesting astrocytes  
137 preferentially occupy PNN holes (line graphs in **Fig. 1a**). A 3D rendering (**Fig. 1b**, images)  
138 and a Pearson correlation analysis of the PNN marker WFA with astrocytic AldheGFP  
139 and Kir4.1 (**Fig. 1b** bar graph), shows no correlation between WFA and both astrocytic  
140 markers consistent with a non-overlapping interdigitating spatial interface where  
141 astrocytic processes are exclusively found in the PNN holes.

142 Less condensed PNNs are also found in small populations of excitatory neurons, for  
143 example, CA2 pyramidal neurons. To ask whether the presence of PNNs would similarly  
144 place astrocytic processes into PNN holes on excitatory cells we repeated the above  
145 analysis on sections of hippocampus CA2. Indeed, CA2 PNN holes were also occupied  
146 by astrocytic processes similar to the cortical PNNs around PV neurons (**Fig. S1a, b**),  
147 however with much closer spatial proximity as seen by a positive Pearson's coefficient in  
148 both stratum pyramidale (**Fig. S1c**) and condensed interstitial matrix in the stratum oriens  
149 (**Fig. S1d**).

150 Since astrocytic processes are primarily confined to the PNN holes, it is reasonable to  
151 assume that these holes are sites where astrocytes perform their homeostatic and  
152 neuromodulatory functions analogous to the conventional tripartite synapses. Hence, we  
153 would expect astrocytic processes in PNN holes to express the necessary proteins  
154 associated with potassium, glutamate, and GABA uptake.

155 Using IHC followed by high magnification confocal imaging and line intensity profile  
156 analysis of the PNNs[21, 22], we assessed the expression of Kir4.1, Glt1, and GABA

157 transporters GAT1 and GAT3 in combination with the astrocytic marker AldheGFP in  
158 >900 holes in each experimental group. We observed immunoreactivity of AldheGFP,  
159 Kir4.1, and both in 63%, 71%, and 62% of PNN holes respectively (**Fig. 1c, d**). In a  
160 separate set of experiments, we observed 59%, 70%, and 56% PNN holes with  
161 aldheGFP, Glt1, and both immunoreactivities respectively (**Fig. 1e, f**). In similar  
162 proportions, astrocytic processes expressed both GABA transporters, GAT1 (**Fig. 1g, h**)  
163 and GAT3 (**Fig. 1i, j**) in the PNN holes. We also examined the expression of Aqp4, Cnx43,  
164 and Cnx30, however, these all showed low to almost undetectable immunoreactivity in  
165 the PNN holes (**Fig. S1e - j**). These data suggest that astrocytic processes in PNN holes  
166 express the necessary proteins to support the clearance of synaptically released ions and  
167 transmitters.

## 168 **2. PNN holes house tripartite synapses:**

169  
170 Since PNN holes provide access for the placement of axosomatic synapses on FSNs, we  
171 assessed whether synaptic terminals and astrocytic processes coexist in the PNN holes  
172 analogous to a classic tripartite synapse. As an extension, we wondered whether a given  
173 hole exclusively houses excitatory or inhibitory synapses and if so whether astrocytes  
174 show matching GABA or Glu transporters.

175 We examined >1000 PNN holes and analyzed the distribution of excitatory and inhibitory  
176 synaptic terminals in conjunction with the astrocytic marker AldheGFP on the surface of  
177 PNN-expressing cortical FSNs (**Fig. 2a, b**). ~80% of the PNN holes were occupied by  
178 either astrocytic processes or vGlut1 terminals or both leaving only ~20% holes  
179 unoccupied. Of all PNN holes, 71% contained excitatory synapses (vGlut1) and 53%  
180 contained astrocytic processes (AldheGFP) and 44% showed co-occupancy of excitatory  
181 synapses with astrocytic processes (**Fig. 2a, b**). Hence the presence of excitatory  
182 synapse in a PNN hole does not necessarily predict the presence of an astrocytic process  
183 in it; however, >60% of synaptic contacts showed the presence of astrocytic processes  
184 with them. PNN holes also contained vGlut2 expressing synapses from thalamocortical  
185 sensory projections in conjunction with astrocytic processes (**Fig. S2a, b**).

186 The inhibitory synaptic terminals in the PNN holes showed a lower astrocytic occupancy  
187 (58%), and only 33% of all PNN holes showed both astrocytic and synaptic components  
188 (**Fig. 2c, d**). However, similar to the vGlut1 terminals, ~58% of vGAT-occupied holes  
189 showed astrocytic contacts and overall ~80% of total PNN holes were occupied by either  
190 astrocytic processes or vGAT terminals or both (**Fig. 2d**).

191 We next asked whether astrocytic processes that colocalize with synapses in the PNN  
192 holes are equipped with transporters and Kir4.1 channels to clear synaptically released  
193 Glu, GABA, and K<sup>+</sup>.

194 Analysis of >1000 PNN holes showed that ~47% of all holes contained vGlut1 terminals  
195 as well as Glt1-expressing astrocytic processes (**Fig. 2e, f**) suggesting that ~ 75% of all  
196 excitatory terminals are accompanied by astrocytic processes equipped to uptake  
197 synaptically released Glu. On the other hand, ~35% of all holes contained both vGAT

2198 terminals and GAT3-expressing astrocytic processes thereby suggesting that ~67% of  
2199 total inhibitory terminals in the PNN holes are equipped with astrocytic processes capable  
2200 of GABA uptake (**Fig. 2g, h**). Once again ~80% of holes were occupied by one or more  
2201 elements from astrocytes or synapses, or both (**Fig. 2f, h**) and 20% were vacant.

2202 Interestingly vGlut1 and vGAT terminals were co-expressed in ~37% of all holes and 64%  
2203 contained astrocytic processes (**Fig. 2i, j**). Only 33% and 19% of all holes exclusively  
2204 contained vGlut1 and vGAT terminals respectively. We also observed co-expression of  
2205 vGAT with vGlut2 expressing glutamatergic synapses from thalamocortical sensory  
2206 projections in the PNN holes (**Fig. S2c, d**). Importantly, combining markers of  
2207 glutamatergic and GABAergic synapses with astrocytes increased the overall occupancy  
2208 of PNN holes from ~80% (**Fig. 2b, d, f, h**) to ~95% (**Fig. 2j**) suggesting that nearly all  
2209 PNN holes are occupied with a mixture of synapses and astrocytic processes.

2210 Taken together these results explicitly show that holes in the PNN contain both excitatory  
2211 and inhibitory synapses and the majority of them are accompanied by astrocytic  
2212 processes expressing both Glu and GABA transporters thereby suggesting the PNN  
2213 holes houses a structural and functional analogue of tripartite synapses. Also, a single  
2214 PNN hole can contain either excitatory or inhibitory or both types of synaptic terminals  
2215 thereby ruling out the possibility of PNN holes being exclusively meant to contain a  
2216 specific type of synapse or astrocytic process. Finally, our data suggest that nearly all  
2217 PNN holes are filled with a synapse and astrocytic processes rarely leaving holes  
2218 unoccupied.

### 2219 **3. Concurrent maturation of astrocytes with PNNs constrains the astrocytic** 2220 **coverage on PV neurons:**

2221  
2222 PNN deposition in the developing brain coincides with the closure of the critical period of  
2223 plasticity after which synapses are “locked” for future modifications provided PNNs are  
2224 unaltered [17, 18, 23]. In parallel with the PNN condensation, the morphological  
2225 maturation of astrocytes also occurs (**Fig. 3a**) [24, 25] during which astrocytic processes  
2226 and synaptic contacts can both be traced to the newly formed PNN holes as evidenced  
2227 in the developing brain (**Fig. 3b**). Based on the facts of concurrent development (**Fig. 3a**)  
2228 and PNN holes containing both synapses and astrocytic processes (**Fig. 2**), we  
2229 hypothesized that PNN accumulation not only locks the synapses but also stabilizes  
2230 astrocytic processes at PNN holes thereby limiting the astrocytic coverage on PNN-  
2231 expressing neurons only to those patches of membrane accessible through the PNN  
2232 perforations.

2233 To test this hypothesis, we compared the pericellular astrocytic coverage of PNN-  
2234 expressing and non-expressing neurons identified by NeuN, AldheGFP, and WFA  
2235 immunostaining (**Figs. 3c, S3**). Within the 0.8 $\mu$ m perimeter of the cell soma, PNN-  
2236 expressing (NeuN+/WFA+) neurons showed a significantly lower astrocytic coverage  
2237 than PNN-lacking (NeuN+/WFA-) neurons (**Fig. 3c, d - top bar graph**). To confirm that  
2238 the lower pericellular astrocytic coverage around PV neurons is attributed to the PNN, we

239 compared the astrocytic coverage of PNN expressing PV neurons (PV+/PNN+) with a  
240 rare cortical population of PNN lacking PV neurons (PNN-/PV+). Again, we observed a  
241 significantly higher astrocytic coverage around PNN lacking PV neurons (PV+/PNN-) than  
242 PNN expressing PV neurons (PV+/PNN-) (**Fig. 3c** right panel, **d** bottom bar graph). These  
243 data confirm that pericellular astrocytic coverage negatively correlates with the presence  
244 of the PNN most likely by restricting the access of astrocytic processes to the cell  
245 membrane to the holes within the PNN.

246 As mentioned above, in the hippocampus a small population of excitatory cells in CA2  
247 express an atypical and less condensed version of PNNs[26]. Astrocytic coverage of  
248 these CA2 neurons, visualized by using astrocytic markers AldheGFP and Kir4.1, did not  
249 differ from their CA1 and CA3 counterparts lacking PNNs (**Fig. 3e - h**) suggesting that  
250 only the “typical” condensed PNNs on cortical PV neurons leads to restricted pericellular  
251 astrocytic coverage.

#### 252 **4. PNN disruption increases astrocytic coverage of neuronal somata:**

253  
254 The most straight forward interpretation of the above findings is that the highly condensed  
255 PNNs leave only a few defined sites, namely the PNN holes, for astrocytes to interact  
256 with the soma, and these holes are most likely determined by the placement of synapses  
257 during development. However, PNNs are known as dynamic structures that undergo  
258 constant homeostatic remodeling [27], which may be pivotal for allowing experience-  
259 dependent synaptic plasticity in the adult brain, and experimental degradation of PNNs  
260 has also been shown to cause synaptic plasticity[23, 28]. Hence it stands to reason that  
261 this would concomitantly change astrocytic coverage and placement of their processes.

262 To examine such plasticity in astrocytic coverage and synaptic contacts, we degraded  
263 cortical PNNs *in-vivo* by intracranially injecting Aldhe111eGFP mice with Chondroitinase  
264 ABC and compared the pericellular astrocytic coverage as well as synaptic contacts of  
265 disrupted PNNs 6-day post-injection with intact PNNs from sham (**Fig. 4a**). At 6-day post-  
266 ChABC injection we observed a widespread decrease in WFA intensity (**Fig. 4b**) as well  
267 as increased perforations (**Fig. 4c**) thereby making PNN less dense and more porous  
268 with an increase in extracellular space in the pericellular region.

269 PNN depletion did not change the total space occupied by astrocytes as determined by  
270 Aldhe111eGFAP in a given field of view (**Fig. 4d** top whole field images, **e** top bar  
271 diagram), however, we observed a significant increase in the cell surface-associated  
272 pericellular astrocytic coverage on PNN-expressing PV neurons (PV+PNN+) (**Fig. 4d**  
273 single cell binary images, **e** middle bar diagram) suggesting a localized change confined  
274 to the pericellular area previously occupied by the PNN. Excitatory neurons (PV<sup>-</sup> PNN<sup>-</sup>)  
275 did not show any change in the pericellular coverage in the ChABC-treated condition (**Fig.**  
276 **4d** bottom single cell binary images, **e** bottom bar diagram) suggesting that the increased  
277 astrocytic coverage is indeed associated with the depletion of the PNN but not diffuse  
278 CSPG. The increased astrocytic coverage also changed the non-overlapping spatial

279 arrangement of the astrocytic processes with the remnant of the PNNs as observed in the  
280 positive spatial correlation between astrocyte and WFA (**Fig. 4f**).

281 We next examined whether the increased pericellular astrocytic coverage following  
282 ChABC treatment also increases the number of astrocytic contacts in newly created holes  
283 in a now highly perforated PNN (**Fig. 4c**). Indeed, using a combination of astrocytic  
284 markers Glt1 and AldheGFP, we observed a significantly higher number of PNN holes  
285 that were now occupied with astrocytic processes (**Fig. S4a - c**). In control mice, we  
286 observed ~60-70% of the PNN holes being occupied by at least one of the astrocytic  
287 markers (**Figs. 1, S4b, c**), whereas, in the ChABC-treated group, this number increased  
288 to 80-90% (**Fig. S4b, c**).

289 PNNs are known to stabilize synapses and depletion of PNNs is known to alter  
290 axosomatic synaptic contacts in several brain areas[28, 29]. Since astrocytes seem to be  
291 an integral component of the axosomatic synapses in the PNN holes, we next asked  
292 whether the observed changes in pericellular astrocytic coverage resulting from PNN  
293 depletion also alter or destabilize the axosomatic synaptic contacts. To this end, we first  
294 compared the total number of vGlut1 and vGAT expressing synaptic terminals;  
295 surprisingly ChABC treatment neither changed the total vGlut1 (**Fig. 4g**) nor the total  
296 vGAT terminals (**Fig. 4h**). Next, we restricted our analysis to the pericellular area in which  
297 astrocytic coverage was increased; here too no significant change in the numerical  
298 densities of pericellular vGlut1 (**Fig. 4i, j**) and vGAT (**Fig. 4k, l**) contacts were found upon  
299 ChABC treatment in either PNN expressing or non-expressing neurons. We also  
300 assessed the pericellular vGlut1 (**Fig. 4i, j**) or vGAT (**Fig. 4k, l**) contacts closely  
301 associated with the astrocytic processes however no significant changes were observed  
302 in these groups too.

303 Finally, we analyzed whether following ChABC treatment PNN holes show any changes  
304 in their occupancy of the synaptic terminals. Using the line intensity profile method, we  
305 assessed the occupancy of PNN holes in the ChABC-treated group. Despite a  
306 significantly higher occupancy of PNN holes by astrocytic processes (**Fig. S4b, c**), no  
307 significant changes were observed either in the occupancy of vGlut1+ terminals (**Fig.**  
308 **S4d, e**) or vGlut1+ terminals with astrocytic contacts (**Fig. S4f**) within PNN perforations.  
309 Similarly, no significant differences were found in total vGAT+ terminals (**Fig. S4g, h**) as  
310 well as in vGAT+ terminals in close association with astrocytes (**Fig. S4i**) within PNN  
311 perforations.

312 Taken together these studies suggest that axosomatic synapses embedded in PNN holes  
313 are highly stable and resistant to degradation of the PNNs in the adult somatosensory  
314 cortex; however astrocytic processes in the same compartment are highly plastic and  
315 undergo structural changes independent of synaptic terminals. It, therefore, appears as if  
316 astrocytes tend to occupy as much of the free neuronal surface as is accessible.

317 **5. Permanent PNN depletion induces astrocytic plasticity without altering synaptic**  
318 **stability:**

319 Since temporary degradation of PNNs using ChABC changed astrocytic coverage by  
320 occupying newly created perforations without changing the overall pericellular numerical  
321 abundance of presynaptic terminals or synaptic terminals with astrocytic contacts on PV  
322 neurons, we sought to investigate whether a permanent genetic deletion of PNNs  
323 destabilizes the axosomatic synapses in conjunction with changing the pericellular  
324 astrocytic coverage.

325  
326 We disrupted PNNs permanently by intracranially injecting a viral vector carrying Cre  
327 recombinase with eGFP reporter AAV9.hSyn.HI.eGFP.WPRE.SV40 (AAV9.Cre) in the  
328 prefrontal cortex of adult *Acan fl/fl* mice (**Fig. 5a**) as described previously[27]. Consistent  
329 with the previous studies[27], PNNs show partial degradation at 4 weeks (**Fig. 5b**)  
330 followed by a complete elimination after 8 weeks (**Fig. 5c**) of AAV9Cre injection in all  
331 transduced PV neurons. Since *Acan fl/fl* mice lack genetic labelling of astrocytes with  
332 eGFP (AldheGFP), we used 3 different astrocytic markers S100b, Glt1, and Kir4.1 in  
333 conjunction with PV and WFA to quantify the pericellular coverage of astrocytes upon  
334 permanent PNN depletion. We compared the pericellular astrocytic coverage and  
335 numerical density of synapses of AAV9Cre-transduced PV neurons showing PNN KO  
336 with non-transduced PV neurons showing intact PNN (**Fig. 5d, f, h**) present in the close  
337 vicinity.

338  
339 With all three markers including S100 $\beta$  (**Fig. 5d, e**), Glt1 (**Fig. 5f, g**), and Kir4.1 (**Fig. 5h,**  
340 **i**), the pericellular astrocytic coverage showed a consistent increase around the PV  
341 neurons with PNN deletion compared to their control counterparts with intact PNNs  
342 suggesting that once changed, astrocytes maintain the coverage in the absence of the  
343 PNNs. However, to our surprise, we did not observe any changes in either the pericellular  
344 density of all vGlut1 terminals (**Fig. 5j, k**) or vGlut1 terminals associated with s100b-  
345 expressing astrocytic processes (**Fig. 5l**). Similarly pericellular density of all vGAT-  
346 labelled inhibitory synaptic terminals (**Fig. 5m, n**) or vGAT terminals associated with  
347 astrocytic processes (**Fig. 5o**) remained unaltered on PNN elimination around PV  
348 neurons.

349  
350 These data suggest that astrocytic processes undergo localized structural changes upon  
351 PNN elimination leading to an increase in the astrocytic coverage around the PV neurons.  
352 A temporary degradation of PNNs is sufficient to induce this astrocyte plasticity.  
353 Permanent and selective PNN elimination induce similar astrocytic structural plasticity  
354 around PV neurons however in both cases pericellular abundance of excitatory and  
355 inhibitory axosomatic synapses with and without astrocytic contacts remained unaltered.

## 356 **6. PNN facilitates astrocytic uptake of synaptically released Glu:**

357 Since astrocytic processes tightly associate with the axosomatic tripartite synapses  
358 embedded in the PNNs, it is possible that PNNs may also participate functionally in  
359 neurotransmitter and ion homeostatic functions of astrocytes. Specifically, we  
360 hypothesize that the negatively charged CSPGs associated with PNNs may

361 electrostatically interact and aid to contain synaptic activity-released charged ions and  
362 neurotransmitters, particularly Glu and K<sup>+</sup>. This may be of particular importance in fast-  
363 spiking PNN expressing PV interneurons.

364 If true, we predict that disruption of PNN assembly should reduce astrocytic Glu uptake  
365 upon synaptic activity. Since this hypothesis solely relies on the physical assembly of the  
366 PNN, an acute depletion of PNN should be disruptive to astrocytic Glu uptake without any  
367 long-term changes in the expression of Glu transporter expression in astrocytic  
368 processes.

369 Incubation of brain slices with ChABC completely dissolved PNNs (**Fig. S5a**) without  
370 altering biophysical properties of patch-clamped astrocytes compared to undigested  
371 control slices each showing characteristic negative and stable resting membrane potential  
372 (**Fig. S5b, c**), and low membrane capacitance (**Fig. S5d**) as well as input resistance (**Fig.**  
373 **S5e**). Also, input-output curves were similar in control and ChABC-treated slices (**Fig.**  
374 **S5f, g**).

375 To test the hypothesis that PNNs affect the clearance of synaptically released Glu, we  
376 stimulated L5-6 axonal fibers and recorded from astrocytes in L3-4, wherein a high  
377 density of PNNs makes it possible for each astrocyte to contact multiple PNNs (**Fig. 6a**).  
378 The distance between the stimulator and patch pipette was kept identical for all  
379 experiments. Synaptically-evoked glutamate currents were recorded in the presence of  
380 Bicuculline, BaCl<sub>2</sub>, AP-5, and CNQX as described previously[30-32]. Astrocytic Glu  
381 transporters current was blocked with a cocktail of TBOA and DHK, with the small fraction  
382 of remaining current being completely blocked by 0.5μm tetrodotoxin confirming that the  
383 recorded current was indeed the synaptically evoked glutamate uptake current (**Fig. S5h**).

384 In the first set of experiments, we determined the minimum amount of current (threshold  
385 stimulation) required to generate a reliably detectable threshold response. In ChABC-  
386 treated slices, astrocytic responses to synaptically released glutamate required  
387 significantly higher current injections than controls (**Fig. 6b**) and still generated  
388 significantly smaller threshold responses (**Fig. 6c**) compared to controls.

389 In the next set of experiments, we generated input-output curves of astrocytic glutamate  
390 uptake currents. After assessing multiple ranges of stimuli, we observed a near linear  
391 range with 10-200μA stimulation with a 20μA increment suitable for the input-output  
392 curve. ChABC-treated slices showed a significant decrease in the peak glutamate uptake  
393 (**Fig. 6d - f**) as well as reduced total charge transfers (**Fig. 6g**) suggesting that the  
394 presence of intact PNNs yields an increase in astrocytic glutamate uptake in response to  
395 synaptic activation.

396 To ensure that the reduced Glu uptake is not due to a non-specific loss of astrocytic Glt-  
397 1 transporter expression or function, we repeated these experiments supplying  
398 exogenous glutamate pulses from a set distance to astrocytes while blocking synaptic  
399 transmissions and other nonspecific currents as described previously [33, 34] (**Fig. 6h**).  
400 Under these conditions, astrocytes showed no significant change in the glutamate uptake

401 current (**Fig. 6i-k**) or uptake kinetic (**Fig. 6l, m**) after ChABC digestion. Furthermore,  
402 immunohistochemical analysis of Glut-1 expression as well as AldheGFP membrane area  
403 remained unaltered after ChABC digestion (**Fig. S6a-e**) suggesting that altered glutamate  
404 uptake by astrocytes was not attributed to a change in the glutamate transporter  
405 expression but more likely to a spillage of Glu out of the synapse and out of the reach of  
406 the patch-clamped astrocyte.

#### 407 **7. PNN facilitates astrocytic uptake of depolarization released K<sup>+</sup>:**

408 The above data suggest that PNNs harbor synapses and astrocytic processes  
409 act as a container to prevent synaptically released glutamate to spill from the synaptic  
410 cleft. Fast-spiking interneurons also release copious amounts of K<sup>+</sup>. Hence, we  
411 questioned whether PNN may also aid in containing K<sup>+</sup> diffusion into the perisynaptic  
412 space.

413 To this end, we recorded L5-6 induced depolarization K<sup>+</sup> currents in L3-4 astrocytes in  
414 the presence of a cocktail blocking postsynaptic and astrocytic glutamate currents  
415 comparing control and ChABC-treated slices (**Fig. 7a**). As was the case with Glu uptake,  
416 astrocytic K<sup>+</sup> uptake was also significantly attenuated on PNN disruption. Although  
417 astrocytes required similar magnitudes of threshold stimuli (**Fig. 7b**) to evoke a detectable  
418 K<sup>+</sup> current, the threshold response was significantly lower in the ChABC-treated slices  
419 (**Fig. 7c**). Complementing the threshold response current, the input-output curve of the  
420 synaptically evoked K<sup>+</sup> currents showed a significantly lower K<sup>+</sup> uptake current (**Fig. 7d,**  
421 **e, f**) resulting in a reduced total charge transfer (**Fig. 7d, e, g**). To eliminate the possibility  
422 of altered expression of Kir4.1 contributing to the lower K<sup>+</sup> uptake by astrocytes, we  
423 performed immunohistochemical analysis of Kir4.1 in recorded slices. ChABC treatment  
424 eliminated PNNs as seen in significantly lower WFA reactivity (**Fig. 7h, i**), however Kir4.1  
425 (**Fig. 7h, j**) AldheGFP (**Fig. 7h, k**) expressions were unaltered in ChABC-treated slices  
426 suggesting that altered K<sup>+</sup> currents could not be attributed to a change in Kir4.1  
427 expression in astrocytes.

428 Taken together this data suggests that intact PNNs ensure the effective uptake of K<sup>+</sup> and  
429 Glu into astrocytes, and PNN disruption causes reduced uptake and accumulation of  
430 extrasynaptic Glu most likely as synaptically released molecules can now diffuse into the  
431 extrasynaptic space.

432

433

434

435

436

437



438 **Discussion:**

439 PNNs have fascinated neuroscientists since their description by Golgi over a century ago.  
440 Their role in stabilizing synapses has been well documented in the visual system where  
441 PNN disruption can re-establish synaptic plasticity. Those studies, however, were  
442 unaware that most excitatory and many inhibitory synapses are ensheathed by astrocytic  
443 processes or leaflets and together form the tripartite synapse. Here astrocytes are well  
444 positioned to support synaptic function through the effective clearance of neuronally  
445 released transmitters and ions. The major objective of our study was to examine the  
446 hypothesis that PNNs may be an important structural and functional component at the  
447 tripartite synapse.

448 To do so we elected to focus our work on cortical fast-spiking PV+ interneurons, as 90%  
449 of them express PNNs. Our studies in adult animals shed light on a number of structural  
450 and functional roles of PNNs that have not been recognized before. Notably, to the best  
451 of our knowledge, ours is the first study to describe the functional cooperation of PNNs  
452 and astrocytes in the clearance of neuronally released Glu and K<sup>+</sup> at tripartite synapses.  
453 Firstly, we show that about 90% of PNN holes contain either excitatory synapses,  
454 inhibitory synapses, or both. Secondly, about 70% of these synapses are tripartite, hence  
455 contain astrocytic processes or leaflets. In all instances, these express Kir4.1 the  
456 astrocytic channel tasked with K<sup>+</sup> uptake as well as Glt-1, the major excitatory amino acid  
457 transporter in astrocytes irrespective of whether they are associated with an excitatory or  
458 inhibitory synapse. Interestingly they do not harbor aquaporins, the water channels  
459 abundantly expressed in conjunction with Kir4.1 channels on astrocytic endfeet touching  
460 blood vessels. Thirdly we used transient enzymatic or permanent genetic disruption of  
461 the PNNs to ascertain whether the astrocytic investment in tripartite synapses on  
462 inhibitory neurons is stable or plastic, and to ask whether the synapse itself is static or  
463 shows plasticity when PNNs are lost. To our surprise, we found that in the adult cortex,  
464 synapses on inhibitory neurons are static and do not change with PNN removal. However,  
465 astrocytic processes are plastic and upon PNN removal astrocytes expand their territory  
466 on the cell soma. It almost appears as if astrocytes take possession of any membrane  
467 that was previously occupied by PNNs. Interestingly, this picture appears different from  
468 the developing cortex where PNN degradation has been shown to reestablish synaptic  
469 plasticity in the visual system[35].

470 While the structural interactions of synapses and astrocytes, and the structural astrocyte  
471 plasticity are fascinating, we also uncovered an important functional synergism between  
472 PNNs and astrocytes. Having the synapse embedded in negatively charged ECM  
473 constituents, the PNN walls off the synapse prevent lateral diffusion of ions and  
474 neurotransmitters, and contain them in the local pocket or cavity where astrocytic leaflets  
475 can effectively remove all the synaptically released transmitter as well as K<sup>+</sup> released in  
476 the process (**Fig. 8**). Not so when the PNNs are disrupted and Glu spills out of the  
477 synapses as evident from increased [Glu] in the perisynaptic space. As PNNs are known  
478 to be degraded by proteolytic enzymes including MMPs and ADAMTSs [6, 36, 37], it is  
479 easy to envision how brain inflammation in the context of injury or disease will break down

480 PNNs and hence allow spillage of  $K^+$  and Glu into the perisynaptic space. Increases in  $K^+$   
481 and Glu are known contributors to epilepsy, and at least in glioma-associated epilepsy, it  
482 has been demonstrated that inhibition of PNN proteolysis suppresses seizures[22].  
483 Furthermore, it is well documented that activation of extrasynaptic NMDA-R by spillage  
484 of Glu can induce neuronal death via activation of the p38MAPK pathway[4]. Hence, we  
485 propose that the containment of axosomatic synapses on PV interneurons is a protective  
486 strategy. This may be particularly important for this group of neurons, which are among  
487 the fastest-firing neurons in the brain. Reported to be capable of firing 100s of action  
488 potentials per second, the release of  $K^+$  by these neurons will far exceed that of most  
489 normal spiking neurons in the brain. The negative charges on the walls of the PNN holes  
490 may temporarily bind the positively charged  $K^+$  ions thereby neutralizing them temporarily  
491 allowing astrocytes additional time to buffer  $K^+$  during following bursts of neuronal activity  
492 without risking depolarization of the membrane. This is not far-fetched in light of the  
493 previously reported charge density of PNNs estimated at over 100 mM[38]. Hence PNNs  
494 may be a structure that evolved specifically to endow fast-spiking neurons to handle the  
495 shift in ions and charged neurotransmitters that are associated with rapid firing. We  
496 recently showed that PNNs around cortical PV neurons also change the dielectric  
497 properties of the cell membrane such as to reduce the effective membrane capacitance.  
498 This facilitates rapid burst firing of the neurons and we show in a glioma mouse model  
499 that enzymatic degradation of PNNs, by MMPs released from the tumor degrades PNNs,  
500 increases the membrane capacitance, and impairs fast burst firing[22].

501 The finding that PNN degradation impairs Glu uptake and  $K^+$  buffering by astrocytes at  
502 tripartite synapses may have general applicability relevant to numerous acute and chronic  
503 neurological conditions. This may obviously contribute to the generation of seizures which  
504 are known to cause the release of matrix metalloproteinases (MMPs) [39, 40]. Moreover,  
505 changes in PV interneuron firing have been reported in the PFC in the context of  
506 Schizophrenia [41], and in the substantia nigra the release of dopamine comes from fast-  
507 spiking interneurons that may be negatively affected in Parkinson disease [42, 43]. It is  
508 possible that in these diseases and others, loss of PNN integrity may also affect the  
509 astrocytes' ability to support effective clearance of neuronally released  $K^+$  and Glu. The  
510 idea that synaptic function and not just structure critically depends on functional  
511 contribution of astrocytes working synergistically with the PNN, dubbed the "tetrapartite  
512 synapse" [44, 45], is an exciting concept that clearly warrants further study.

513

514

515

516

517

518

519 **Materials and methods:**

520 **Mice:**

521 All animal procedures were approved and performed following the ethical guidelines set  
522 by the University of Virginia Institutional Animal Care and Use Committee (IACUC) ACUC.  
523 Mice were housed in groups of five in a facility in 12 h light/dark cycles and had access  
524 to food and water *ad libitum*. AldheGFP mice expressing enhanced green fluorescence  
525 protein (eGFP) under astrocyte-specific promotor AldheGFP were generated as  
526 described previously [33, 34] and housed and bred according to ACUC guidelines. We  
527 received C57BL/6N-Acan<sup>tm1a(EUCOMM)Hmgu</sup>/H mice (common name  
528 HEPD0602\_5\_G11) from EUCOMM (UK Research & Innovation, Mary Lyon Center,  
529 Harwell Campus, strain ID # EM:10224). The heterozygous mice were bred together to  
530 generate Acan fl/fl mice and confirmed by genotyping before being used for any  
531 experimentation.

532 **Intracranial surgeries and injections:**

533 *ChABC injection*

534  
535 Chondroitinase ABC from *Proteus vulgaris* (cat # C3667-10UN, Sigma Aldrich) was  
536 dissolved in sterilized PBS (50mU/μl); subsequently, 2 μl solution was injected unilaterally  
537 at an infusion rate of 200nl/minute. Mice were anesthetized with 2–5% isoflurane and  
538 fixed to a stereotaxic apparatus (Leica Angleone stereotaxic model 39464710) followed  
539 by a midline scalp incision and a 0.5 mm burr hole 2.0mm lateral and 1.0mm ventral to  
540 bregma and infused ChABC at ~2.0mm deep from the cortical surface using a 10μl  
541 syringe (World Precision Instruments #SGE010RNS). Sham control mice were injected  
542 with sterile PBS with an identical procedure. Mice were dosed with Buprenorphine /  
543 Rimadyl and allowed to recover on a heating pad until mobile and were monitored daily  
544 for up to 5 days from surgery. Body weight was measured for 3 consecutive days after  
545 surgery and all mice were perfused on 6<sup>th</sup>-day post-injection.

546 *AAV injection and Acan knockout:*

547  
548 To achieve PNN knockout in adult mice brains, we injected pENN.AAV.hSyn.HI.eGFP-  
549 Cre.WPRE.SV40 (Addgene #105540-AAV9) in 7-8 weeks old Acan fl/fl mice. In brief  
550 AAV9 (2.7x 10<sup>13</sup> vg/mL) was diluted in sterile PBS to achieve 1x 10<sup>13</sup> vg/mL concentration  
551 and 1.5μl was injected in each hemisphere (from bregma: 0.5mm posterior, 2.0mm  
552 lateral, 1.0mm ventral) with 200nl/minute infusion rate as described above. Mice were  
553 transcardially perfused after 8-10 weeks of AAV9 injections to perform IHC.

554 **Acute slice electrophysiology:**

555  
556 Whole-cell patch-clamp recordings were obtained from astrocytes *in situ* acute brain  
557 slices as described previously[30, 31, 34]. In brief, mice underwent cervical dislocation  
558 followed by a quick decapitation and dissection to remove brains and were kept in an ice-  
559 cold ACSF (135mM NMDG, 1.5mM KCl, 1.5mM KH<sub>2</sub>PO<sub>4</sub>, 23mM choline bicarbonate,

560 25mM D glucose, 0.5mM CaCl<sub>2</sub>, 3.5mM MgSO<sub>4</sub>; pH 7.35, 310 ± 5 mOsm) (All from  
561 Sigma-Aldrich) saturated with carbogen (95% O<sub>2</sub> + 5% CO<sub>2</sub>). We prepared 300 µm thick  
562 coronal slices using Leica VT 1000P or 1200S tissue slicers. Slices were transferred into  
563 a custom-made recovery chamber filled with ACSF (125mM NaCl, 3mM KCl, 1.25mM  
564 NaH<sub>2</sub>PO<sub>4</sub>, 25mM NaHCO<sub>3</sub>, 2mM CaCl<sub>2</sub>, 1.3mM MgSO<sub>4</sub>, 25mM glucose, pH 7.35, 310  
565 ± 5 mOsm) constantly bubbled with carbogen (95% CO<sub>2</sub> + 5% O<sub>2</sub>) to recover at 32 °C  
566 for 1 hr. Subsequently, slices were transferred to room temperature conditions until used  
567 for recordings. Individual slices were transferred to a recording chamber that was  
568 continuously superfused with ACSF at a flow rate of 2 ml/min. GFP-positive astrocytes in  
569 Aldh1l1-eGFP mice cortical slices were visualized using an upright microscope (Leica  
570 DMLFSA) with 5x and 40x water immersion lens and epifluorescence and infrared  
571 illuminations to identify eGFP-expressing astrocytes.

572  
573 Whole-cell voltage-clamp and current-clamp recordings were achieved using an  
574 Axopatch 200B amplifier (Molecular Devices) with an Axon Digidata 1550A interface  
575 (molecular devices). Patch pipettes of 7–10MΩ open-tip resistance were created from  
576 standard borosilicate capillaries (WPI, 4IN THINWALL GI 1.5OD/1.12ID) using HEKA PIP  
577 6 (HEKA) or PMP-102 (Warner instruments) programmable pipette pullers. We filled  
578 patch pipettes with an intracellular solution containing 134mM potassium gluconate, 1mM  
579 KCl, 10mM 4-(2-hydroxyethyl)- 1-piperazineethanesulfonic acid (HEPES), 2mM  
580 adenosine 5'-triphosphate magnesium salt (Mg-ATP), 0.2mM guanosine 5'-triphosphate  
581 sodium salt (Na-GTP) and 0.5mM ethylene glycol tetraacetic acid (EGTA) (pH 7.4, 290–  
582 295 mOsm). MM-225 micromanipulator (Sutter Instrument, Navato, CA) was used to  
583 visually guide the patch pipette to the cell. After making a tight seal of >5GΩ resistance,  
584 brief suction was applied to achieve the whole-cell mode and cells were immediately  
585 clamped at -80mV. The membrane capacitance (C<sub>m</sub>) and series resistance were not  
586 compensated unless otherwise stated. Data were acquired using Clampex 10.4 software  
587 and Axon Digidata 1550A interface (Molecular Devices), filtered at 5 kHz, digitized at 10–  
588 20 kHz, and analyzed using Clampfit 10.6 or Clampfit 11.2 software (Molecular Devices).  
589 Unless otherwise stated, throughout all the recordings carbogen-bubbled ACSF was  
590 continuously superfused (2 ml/min) and the bath temperature inside the reordering  
591 chamber was maintained at 32–33 °C using an inline feedback heating system (Cat# TC  
592 324B, Warner Instruments).

593

#### 594 **PNN degradation in ex-situ brain slices:**

595

596 Chondroitinase ABC (ChABC) from *Proteus vulgaris* (Cat# C3667, Sigma-Aldrich) was  
597 reconstituted in a 0.01% bovine serum albumin aqueous solution according to the  
598 manufacturer's instruction to make a 1 U/40 µl stock solution. Aliquots of 1U were  
599 prepared and stored at -20 °C until used. After slice recovery slices were treated with  
600 ChABC and subsequent recordings were made as previously described[21, 22]. In brief,  
601 after recovery, 2–3 cortical half slices were incubated in 3ml ChABC solution (0.5 U  
602 ChABC / ml ACSF) in an incubation chamber continuously supplied with carbogen at 33

603 °C for 45 min. Next, the slices were rinsed twice and incubated in ACSF until used for  
604 electrophysiological recordings. These parameters of PNNs digestion by ChABC  
605 (enzyme concentration—0.5 U/ml, incubation time - 45 min, incubation temperature - 33  
606 °C) reliably degraded PNNs (**Fig. S5a**) as described previously[21, 22]. For controls,  
607 previously separated contralateral halves of the ChABC-treated slices were incubated in  
608 3ml of ACSF without ChABC, and subsequently, both ChABC-treated and non-treated  
609 slices were kept in ACSF together until used for the recordings.

610

### 611 **Measurement of intrinsic biophysical properties of astrocytes:**

612

613 The resting membrane potential ( $V_m$ ) was measured by setting  $I = 0$  mode immediately  
614 after achieving the whole-cell configuration. Membrane capacitance ( $C_m$ ) was measured  
615 directly from the amplifier by adjusting capacitance and monitoring the capacitive  
616 transients as described previously[30]. To calculate the input resistance ( $R_{in}$ ) of  
617 astrocytes, we calculated the steady-stated membrane voltage deflection ( $\Delta V$ ) on  
618 injecting 15 hyperpolarization current steps ( $-100$  pA each for 1000 ms). The ratio ( $\Delta V/I$ )  
619 of steady-state change in the membrane voltage ( $\Delta V$ ) and the corresponding injected  
620 current ( $I$ ) was computed as  $R_{in}$ . The I-V curve of astrocytes was computed in both the  
621 current clamp (31 steps,  $-100$ pA to  $+500$ pA, step size 20pA, step duration 1100ms) and  
622 voltage clamp (25 sweeps,  $-180$ mV to  $+60$ mV, step size 10mV) modes (**Fig. S5f - i**)  
623 followed by plotting the voltage/current responses. Astrocytes with nonlinear IV  
624 responses were not continued for recordings and analysis.

625

### 626 **Measurement of astrocytic currents:**

627

#### 628 Synaptically evoked Glutamate uptake current:

629

630 We recorded synaptically evoked currents from cortical astrocytes according to the  
631 previously published studies with some modifications[31, 47]. In brief, we placed a  
632 concentric bipolar electrode (FHC, # CBABD75) in L5-6 of the prefrontal cortical slices  
633 and patched astrocytes in L 3-4 (**Fig. 6a**). The stimulation protocol consists of initial  $10\mu$ A  
634 and  $20\mu$ A pulses followed by a  $20\mu$ A increment in each subsequent pulse capping at  
635  $200\mu$ A (pulse duration 200 $\mu$ s). All recordings were performed in presence of  $20\mu$ M  
636 bicuculline,  $50\mu$ M D-2-amino-5-phosphonovalerate (D-AP-5),  $20\mu$ M 6-cyano-7-  
637 nitroquinoxaline-2,3-dione (CNQX), and  $100\mu$ M BaCl<sub>2</sub>. In the initial few recordings, we  
638 confirm that the recorded current is glutamate by observing a near-complete blockade of  
639 evoked current upon  $100\mu$ M TBOA and  $300\mu$ M DHK application (**Fig. S5h**). The  
640 remaining current was completely abolished by superfusing  $0.5\mu$ M TTX, confirming it as  
641 a neuronal-evoked glutamate current (**Fig. S5h**). Each stimulation pulse was repeated 5  
642 times (sweeps) and a minimum of two sweeps were averaged to compute the peak  
643 current and charge transfer after excluding the sweeps with baseline fluctuation or noise.

644

#### 645 Depolarization evoked potassium uptake current:

646  
647 To record depolarization evoked astrocytic potassium uptake current, we positioned the  
648 stimulator and patch pipette as described above and incubated slices in a mixture of 20 $\mu$ M  
649 Bicuculin, 50  $\mu$ M D-AP-5, 20  $\mu$ M CNQX, 100  $\mu$ M TBOA, and 300  $\mu$ M DKH. The stimulation  
650 protocol consisted of initial 0.1mA and 0.2mA pulses followed by 0.5mA and 5 subsequent  
651 pulses with 0.5mA increment capping at 3mA (pulse duration 200 $\mu$ s). Each stimulation  
652 pulse was repeated 3 times (sweeps) and a minimum of two sweeps were averaged to  
653 compute the peak current and charge transfer after excluding the sweeps with baseline  
654 fluctuation or noise.

655

#### 656 *Astrocytic uptake of exogenously applied glutamate:*

657

658 To measure the glutamate uptake capacity of astrocytes we adopted the exogenous  
659 glutamate puffing method as described previously with minor modifications[30, 34]. In  
660 brief, we constantly perfused slices with ACSF containing 500 nM TTX, 20  $\mu$ M bicuculline,  
661 100  $\mu$ M CdCl<sub>2</sub>, 50  $\mu$ M D-AP5, and 50  $\mu$ M CNQX and 100  $\mu$ M BaCl<sub>2</sub>. After patching an  
662 astrocyte, a 500 msec puff (2PSI pressure using a Pico-liter Injector PLI-10 from Warner  
663 Instruments) of 200  $\mu$ M glutamate solution (120mM NaCl, 3.5mM KCl, 25mM HEPES,  
664 10mM glucose, 200  $\mu$ M Glutamate) was applied from a distance of ~100  $\mu$ m by a 5-8m $\Omega$   
665 open tip resistance glass pipette. In several random recordings, we applied a mixture of  
666 100  $\mu$ M TBOA and 300  $\mu$ M DHK to confirm that the recorded current was glutamate (**Fig.**  
667 **6i**). We recorded 5 sweeps and averaged a minimum of 3 sweeps to generate a result  
668 sweep that was utilized to compute the data. The sweeps with fluctuating baseline and  
669 noise were excluded from the analysis. The averaged trace of uptake current was  
670 analyzed using Clampfit 10.6 or Clampfit 11.2 program to generate the below-described  
671 measurements. The peak current was calculated by subtracting the baseline from the  
672 peak response. The total charge transfer was computed by calculating the total areas  
673 under the curve of glutamate uptake current response. Decay time and decay slope were  
674 calculated from the decaying phase (100 % to 37% of the peak) of the uptake current.

#### 675 **Immunohistochemistry (IHC) and confocal imaging:**

676

##### 677 *IHC:*

678

679 Mice were injected with a mixture of ketamine and xylazine (100 mg/kg and 10 mg/kg,  
680 respectively) and subsequently perfused transcardially with PBS followed by 4% PFA.  
681 We dissected out the brains and stored them overnight in 4% PFA at 4 °C followed by  
682 storing them in PBS at 4 °C until sectioning was done. We cut 50- $\mu$ m-thick floating  
683 sections using 5100MZ vibratome from Campden instruments or Pelco EasiSlicer from  
684 Ted Pella. The sections were either used for IHC immediately or stored at -20 °C in a  
685 custom-made storage medium (10% (v/v) 0.2mM phosphate buffer, 30% (v/v) glycerol,  
686 30% (v/v) ethylenglycol in deionized water, pH 7.2–7.4) for future uses. To minimize  
687 procedure-associated variations, we stained duplicate sections from 5 to 7 mice of each  
688 experimental group in a single batch. In brief, sections were retrieved from -20 °C

689 storage, rinsed thrice with PBS, and permeabilized and blocked by incubating them in  
690 blocking buffer (0.5% Triton X-100 and 10% goat serum in PBS) for 2 hr at RT in a 24-  
691 well plate. Sections were incubated for 18-24hrs at 4 °C with appropriate primary  
692 antibodies or biotinylated WFA (Cat# B-1355, Vector Laboratories) in diluted blocking  
693 buffer (1:3 of blocking buffer and PBS). Following this, we incubated sections with  
694 appropriate secondary antibodies and Alexa Fluor® 555-conjugated streptavidin (Cat#  
695 S32355, ThermoFisher Scientific, 1:500) in diluted blocking buffer overnight at 4 °C in  
696 dark. Further, the sections were rinsed with PBS and were mounted on the glass slides  
697 (Fisherfinest 25 × 25 × 1, Cat# 12-544-2) covered with cover glass, and the edges of the  
698 slides were sealed with nail polish. The primary and secondary antibodies used are  
699 enlisted in Supplementary Table 1.

700

#### 701 Confocal imaging:

702

703 Representative images and data in Figs. 1a-f, S1, 2a-e, S2, 3, S3, 4, S4, and S5 e-h,  
704 were acquired using Nikon A1 confocal microscope, and quantification was performed by  
705 associated NIS-Elements AR analysis program. Images and data in the remaining figures  
706 were acquired using Olympus FV 3000 confocal microscope and images were analyzed  
707 using the ImageJ program. We utilized several different objective lenses including 10x  
708 (air), 20x(air), 40x(oil), 60x (oil), or 100x (oil) with a range of optical zoom based on the  
709 experimental requirement. Images were acquired as 12 bits, and acquisition settings were  
710 minimally adjusted to accommodate a few unsaturated and saturated pixels.

711

#### 712 **Quantification of IHC images:**

713

#### 714 PNN disruption analysis

715

716 The spread of PNN disruption by ChABC injection in mice brains was quantified from  
717 whole coronal section images (**Fig. 4b**). We drew uniform-sized ROIs (0.4x0.4 mm<sup>2</sup>)  
718 adjacent to each other starting from the ChABC incision site towards the lateral side of  
719 the coronal plane. The mean fluorescence intensity was computed. All analyzed  
720 images/ROIs at similar distances from the incision site were tabulated to plot the mean  
721 and SD of the fluorescence intensity. To assess the PNN disruption on PV cells after  
722 AAV-mediated Acan KO, we selected a 0.8µm perimeter area of cell soma and binarized  
723 the WFA signal using an automated thresholding method (OTSU) in ImageJ. WFA area  
724 from individual cells from different groups was utilized for plotting the graph (**Fig. 5c**).

725

#### 726 PNN holes analysis:

727

728 We assessed the PNN holes for the presence of astrocytic processes (**Fig. 1**) and  
729 synaptic contacts (**Fig. 2**) and their fate after ChABC treatment (**Fig. S4**) using the PNN  
730 line intensity profile method with slight modifications as described previously[21, 22]. In  
731 brief, we acquired high magnification (200x or higher) images of individual PNNs at their

732 maximum perimeter plane (**Fig. 1a** main image). Subsequently, we drew a polyline on  
733 PNN (WFA) along the entire periphery of the cell and generated an intensity profile  
734 consisting of high-intensity peaks (PNN-CSPGs) and low-intensity drops (PNN holes).  
735 We set a threshold of WFA intensity (ranging from 50-66% of the peak WFA intensity)  
736 that covered the maximum number of drops as PNN holes. The WFA intensity drops  
737 under the threshold (**Fig. 1a**, blue area in top graph) were considered holes. The presence  
738 of a specific fluorophore peak in the PNN hole was determined by the presence of a  
739 clearly distinguishable peak within the two consecutive peaks of WFA (**Fig. 1a** red areas  
740 in bottom graph). Subsequently, we provided unique identifying marks to each PNN hole  
741 and computed the presence or absence of astrocytic/synaptic components within it. To  
742 quantify the degree of perforations in PNNs after ChABC disruption, we counted the  
743 number of PNN holes as described above and normalized it to the cell perimeter. This  
744 method was utilized for analyzing PNN holes in Figs. 1, 2, S1, S2, and S4.

745

746 Quantification of PNN disruption and astrocytic Kir4.1 and Glt1 expressions in fixed ex-  
747 vivo acute brain slices:

748

749 To assess whether ChABC-mediated PNN digestion in acute brain slices alters the  
750 expressions of Kir4.1 and Glt1 to influence the astrocytic potassium and glutamate  
751 uptake, we fixed acute brain slices after electrophysiological recordings and performed  
752 immunostaining using specific antibodies and WFA. 40x magnification images were  
753 acquired using Olympus FV 3000 and analyzed using ImageJ. The signal of the individual  
754 channel (Kir4.1/Glt1, WFA, and AldheGFP) was binarized using an in-built thresholding  
755 function OTSU, and the resulting total area was tabulated to plot the graphs (**Figs. 7h -**  
756 **k, S6**).

757

758 Quantification of astrocytic coverage and synaptic puncta analysis:

759

760 To quantify the pericellular astrocytic coverage of PV / excitatory neurons, we acquired  
761 high-magnification images using either Nikon A1 (40x5 optical zoom oil immersion lens)  
762 or Olympus Fluoview FV 3000 (100x3 oil immersion objective lens) with a 0.2 $\mu$ m optical  
763 plane thickness (**Fig. S3**). After image acquisition, we generated a binary representation  
764 of the cell soma using inbuilt functions in ImageJ and Nikon elements programs. We  
765 defined a 0.8 $\mu$ m wide perimeter from the cell surface as a pericellular area (based on the  
766 measurement of the thickness of the PNN). Subsequently, we binarized the individual  
767 channels (AldheGFP / s100b / Glt1 / Kir4.1) using inbuilt auto thresholding functions in  
768 Nikon elements or ImageJ (OTSU). Using Boolean operations, we computed the binary  
769 areas of different astrocytic markers confined to the cell perimeter defined above (**Fig.**  
770 **S3**). We normalized the pericellular area with the perimeter of the same cell before pooling  
771 images from a section/mouse (**Fig. 4**). In **Figs. 3d and 5** controls and experimental cells  
772 were from the same section, therefore we used individual cells data for plotting graphs  
773 and statistical tests.



774 We added one more step of *find maxima* in ImageJ or an analogous function in Nikon AR  
775 analysis programs to quantify the overall and pericellular numerical densities of vGlut1  
776 and vGAT puncta in the above protocol (**Fig. S3**). A prominence setting of 500 (for vGlut1  
777 puncta) or 2000 (for vGAT puncta) was found appropriate to capture all puncta and was  
778 used for images. The total number of synapses in the entire image was used to plot the  
779 total vGlut1/vGAT puncta (**Fig. 4d - e**). We used Boolean operations to compute the total  
780 pericellular synapses and pericellular synapses with astrocytic processes in contact with  
781 them. The resultant absolute values were normalized to the perimeter of the individual  
782 cells and were used for data pooling or directly for plotting graphs.

783

#### 784 Volumetric analysis in Imaris:

785

786 The representative 3D reconstruction images of the PNN and pericellular astrocytic  
787 processes in Figs. 1, 2, and 4 were generated using Imaris v9.90 (Oxford Instruments).  
788 In brief, we generated volumetric masks from the PV channel that were expanded by  
789 0.8 $\mu$ m – 1.0 $\mu$ m to accommodate pericellular PNN structures. These masks were then  
790 used to create new astrocytic and synaptic data channels that excluded structures outside  
791 of the pericellular domain. The enlarged PV channel volume was created using the  
792 surface creation tool with smoothing detail enabled and a surface grain size set to  
793 0.103 $\mu$ m. Background subtraction was also enabled with the diameter of the largest  
794 sphere set to 0.388 $\mu$ m and manual thresholding set to a value of 200. Astrocytic and  
795 synaptic channel volumes were created using the surface creation tool with smoothing  
796 detail enabled and a surface grain size set to 0.103 $\mu$ m. Background subtraction was also  
797 enabled with the diameter of the largest sphere set to 0.388 $\mu$ m and manual thresholding  
798 set to a value of 200.

799 Illustrations / cartoons in Figs. 4, 5, 6, 7, and 8 were created with BioRender.com under  
800 a paid subscription.

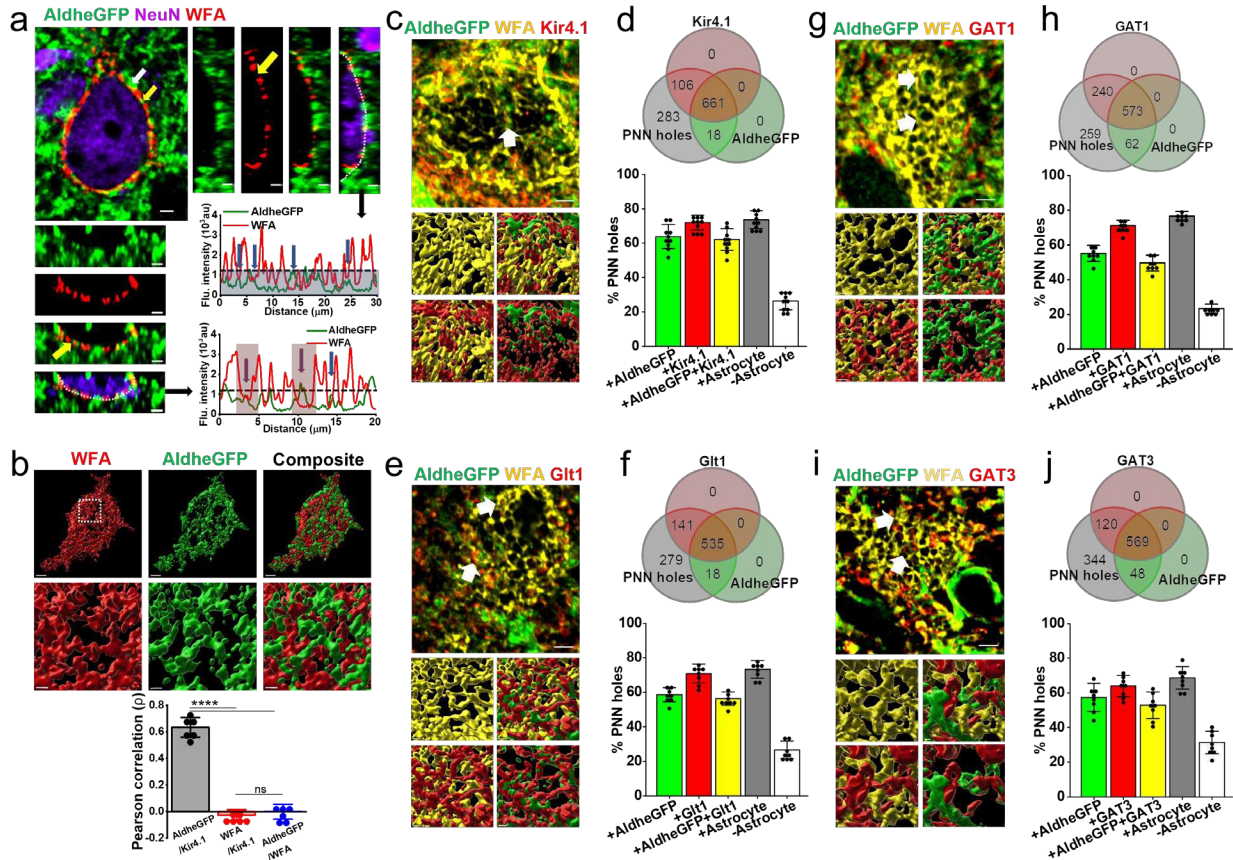
801

#### 802 **Statistical analysis**

803

804 Data in the bar diagrams are expressed as mean  $\pm$  SD unless stated otherwise in the  
805 specific figure. Individual data points are represented by dots. Figure legends contain the  
806 essential details including numerical values of mean, standard deviation, biological or  
807 technical replicates, statistical tests, and corrections. The detailed statistical analysis data  
808 including test statistics, P values, post-hoc comparisons, and corrections are summarized  
809 in Supplementary Table 2. The sample size was not predetermined. We ran appropriate  
810 normality tests and found that data distribution was sufficiently normal and variance within  
811 groups was sufficiently similar to be used for parametric tests. Therefore, experimental  
812 designs with two treatment groups were analyzed by two-tailed unpaired t-test unless  
813 stated otherwise in the figure legends. Welch's correction was applied regardless of  
814 statistically different variances. Experimental designs with more than two groups were  
815 analyzed using one-way ANOVA or two-way ANOVA followed by Tukey's post-hoc

816 multiple comparison tests. Statistically significant difference between groups were notified  
817 in graphs using asterisk(s) (\*P < 0.05, \*\*P < 0.01, \*\*\*P < 0.001, \*\*\*\*P < 0.0001). Data  
818 analysis was performed using Microsoft Excel and Origin 2021 (OriginLab).  
819



820  
 821 **Fig. 1 Astrocytic processes and homeostatic proteins (Kir4.1. And GLT-1) are largely**  
 822 **confined to PNN holes.**

823 **a** Confocal micrograph showing WFA (red)-labelled PNN surrounded by AldheGFP  
 824 (green) labelled astrocytic processes. White and yellow arrows point to astrocytic process  
 825 on outer surface and within PNN holes respectively. The side and bottom panels show  
 826 orthogonal planes of the 3D image. WFA and AldheGFP fluorescence intensities of a line  
 827 drawn on the PNNs in orthogonal view planes showing PNN holes occupied by astrocytic  
 828 processes (blue arrows). The blue area below the dotted line represents WFA threshold  
 829 intensity. Vertical red bars in between two consecutive WFA peaks represent the area of  
 830 PNN holes wherein astrocytic processes can be confined. Scale 2µm.

831 **b** 3D reconstruction of PV neuron surface area occupied by PNN (WFA - red) and  
 832 astrocytic processes (AldheGFP - green) showing a non-overlapping interdigitating  
 833 spatial interface. The marked area in the white square is magnified in the bottom panels.  
 834 Scale bar 2µm on top and 1µm on bottom panels. Error bars show a highly positive  
 835 Pearson correlation between astrocytic markers AldheGFP and Kir4.1 ( $0.63 \pm 0.07$ ) and  
 836 a significantly negative correlation between PNN marker (WFA) and astrocytic markers  
 837 Kir4.1 ( $-0.02 \pm 0.04$ ) and AldheGFP ( $-0.001 \pm 0.05$ ).  $n = 6\text{PNNs}/3\text{m}$ ; \*\*\*\* $P < 0.0001$ ; one-  
 838 way ANOVA, Tukey's post-hoc test.

839 **c** Confocal micrograph (top) showing astrocytic processes (AldheGFP - green)  
 840 expressing Kir4.1(red) in PNN (WFA - yellow) holes. Bottom panels show a 3D

841 reconstruction of PNN holes (white arrows) occupied by astrocytic processes expressing  
842 Kir4.1.

843 **d** Venn diagram showing proportional occupancy of PNN holes by Kir4.1-expressing  
844 astrocytic processes. Bar diagram showing the percent of total PNN holes occupied by  
845 AldheGFP ( $63.83 \pm 7.0$ ), Kir4.1 ( $71.9 \pm 4.22$ ), both ( $62.13 \pm 6.33$ ), occupied by any  
846 astrocytic marker ( $73.64 \pm 5.14$ ), and not occupied by any astrocytic marker ( $26.35 \pm$   
847  $5.14$ ).  $n = >40\text{PNNs}/10\text{s}/4\text{m}$ .

848 **e** Confocal micrograph (top) showing astrocytic processes (AldheGFP - green)  
849 expressing Glt1 (red) in PNN (WFA - yellow) holes. Bottom panels show 3D reconstruction  
850 of PNN holes (white arrows) occupied by astrocytic processes expressing Glt1.

851 **f** Venn diagram showing proportional occupancy of PNN holes by Glt1 expressing  
852 astrocytic processes. Bar diagram showing the percent of total PNN holes occupied by  
853 AldheGFP ( $58.68 \pm 4.02$ ), Glt1 ( $70.85 \pm 5.55$ ), both ( $56.35 \pm 3.87$ ), occupied by any  
854 astrocytic marker ( $73.37 \pm 5.11$ ), and not occupied by any astrocytic marker ( $26.62 \pm$   
855  $5.11$ ).  $n = >40\text{PNNs}/8\text{s}/4\text{m}$ .

856 **g** Confocal micrograph (top) showing astrocytic processes expressing AldheGFP (green)  
857 and GAT1 (red) in PNN (WFA - yellow) holes. Bottom panels show 3D reconstruction of  
858 PNN holes occupied by astrocytic processes expressing GAT1 (red).

859 **h** Venn diagram showing proportional occupancy of PNN holes by GAT1 expressing  
860 astrocytic processes. Bar diagram showing the percent of total PNN holes occupied by  
861 AldheGFP ( $55.11 \pm 4.59$ ), GAT1 ( $71.06 \pm 3.18$ ), both ( $49.62 \pm 4.56$ ), occupied by any  
862 astrocytic marker ( $76.64 \pm 2.57$ ), and not occupied by any astrocytic marker ( $23.35 \pm$   
863  $2.57$ ).  $n = >40\text{PNNs}/8\text{s}/4\text{m}$ .

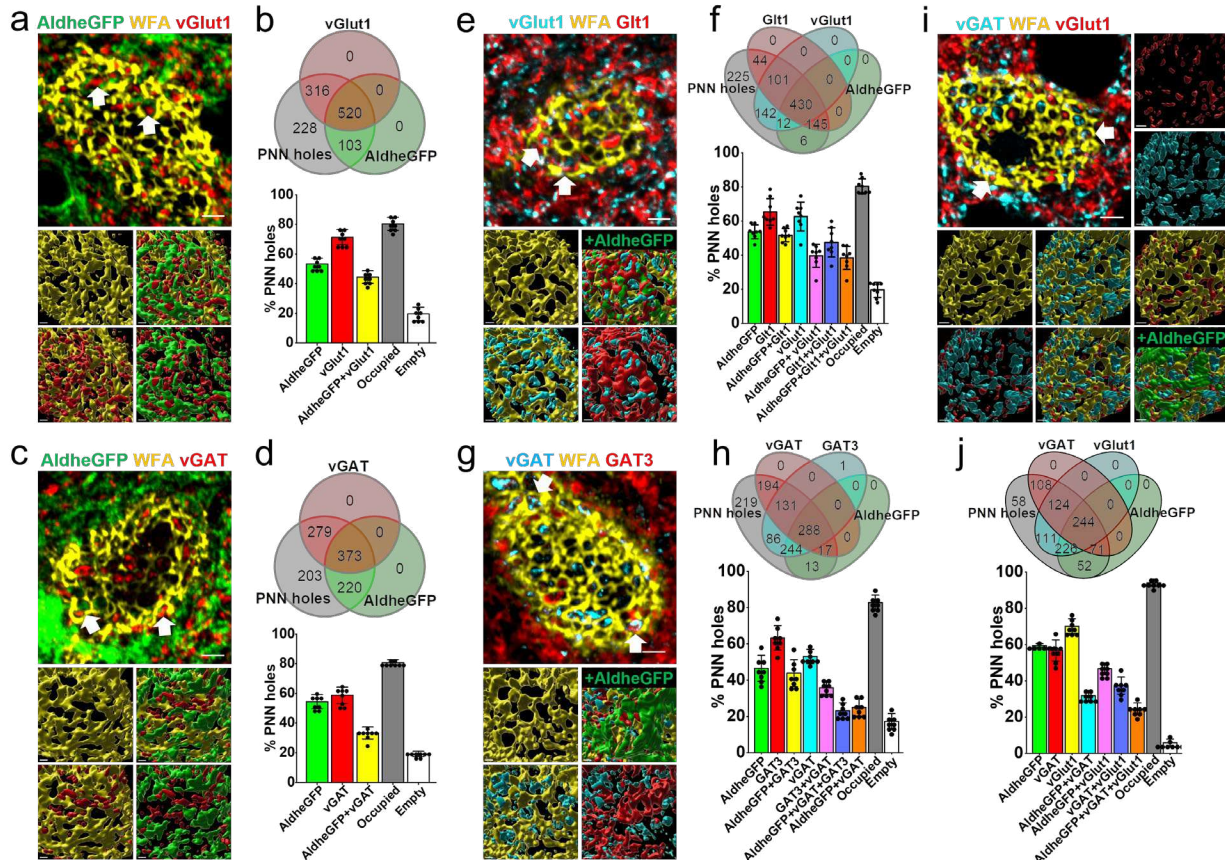
864 **i** Confocal micrograph showing astrocytic processes expressing AldheGFP (green) and  
865 GAT3 (red) in PNN (WFA - yellow) holes. Bottom panels show 3D reconstruction of PNN  
866 holes occupied by astrocytic processes expressing GAT3.

867 **j** Venn diagram showing proportional occupancy of PNN holes by GAT3 expressing  
868 astrocytic processes. Bar diagram showing the percent of total PNN holes occupied by  
869 AldheGFP ( $57.41 \pm 8.14$ ), GAT3 ( $64.0 \pm 6.18$ ), both ( $52.95 \pm 7.74$ ), occupied by any  
870 astrocytic marker ( $68.62 \pm 6.48$ ), and not occupied by any astrocytic marker ( $31.37 \pm$   
871  $6.48$ ).  $n = >40\text{PNNs}/8\text{s}/4\text{m}$ .

872 s and m represent the number of sections and mice respectively. Bar data are expressed  
873 as mean  $\pm$  standard deviation (SD). Dots in the bars represent individual data points.  
874 Scale bar  $2\mu\text{m}$  in top and  $1\mu\text{m}$  in bottom images in c, e g, and i.

875

876



877  
878 **Fig. 2 PNN holes contain excitatory and inhibitory synaptic terminals together with**  
879 **astrocytic processes that express corresponding neurotransmitter transporters.**

880 **a** Representative confocal micrographs showing PNN (WFA - yellow), astrocytic  
881 processes (AldheGFP - green), and excitatory presynaptic terminals (vGlut1 - red)  
882 expression around cortical FSNs. Arrows point to the PNN holes containing both vGlut1  
883 and AldheGFP. Bottom images represent 3D reconstruction of the PNN lattice,  
884 containing astrocytic processes and vGlut1 terminals.

885 **b** Venn diagram showing the proportional occupancy of PNN holes by vGlut1 expressing  
886 excitatory synapses and AldheGFP expressing astrocytic processes. Bar diagram  
887 showing the percent of total PNN holes occupied by AldheGFP ( $53.28 \pm 4.05$ ), vGlut1  
888 ( $71.37 \pm 3.18$ ), and both ( $44.45 \pm 4.26$ ).  $80.39 \pm 4.46$  % of total holes are occupied (by  
889 AldheGFP or vGlut1 or both) leaving  $19.60 \pm 4.4$  % of holes empty.  $n = >40$ PNNs/8s/4m.

890 **c** Representative confocal micrographs showing PNN (WFA - yellow), astrocytic  
891 processes (AldheGFP - green), and inhibitory presynaptic terminals (vGAT - red)  
892 expression around cortical FSNs. Arrows point to the PNN holes containing both vGAT  
893 and AldheGFP. Bottom images represent 3D reconstruction of the PNN lattice  
894 containing astrocytic processes and vGAT terminals.

895 **d** Venn diagram showing the proportional occupancy of PNN holes by astrocytic  
896 processes and GABAergic synapses. Bar diagram showing the percent of total PNN holes



897 occupied by AldheGFP ( $54.45 \pm 4.7$ ), vGAT ( $58.74 \pm 5.5$ ), and both ( $33.31 \pm 4.02$ ). 80.89  
898  $\pm 1.8$  % of total holes are occupied (by AldheGFP or vGAT or both) leaving  $19.10 \pm 1.8$  %  
899 holes empty.  $n = >40\text{PNNs}/8\text{s}/4\text{m}$ .

900 **e** Representative confocal micrographs showing PNN (WFA - yellow), astrocytic  
901 processes labelled with AldheGFP (green) and Glt1 (red), and excitatory presynaptic  
902 terminals (vGlut1 - cyan) expression around cortical FSNs. Arrows point to the PNN holes  
903 containing both Glt1 and vGlut1. Bottom images represent 3D reconstruction of PNN  
904 lattice containing astrocytic processes and vGAT terminals.

905 **f** Venn diagram showing the proportional occupancy of PNN holes by glutamatergic  
906 synapses and astrocytic processes with glutamate transporter expression. Bar diagram  
907 shows the percent of total PNN holes occupied by AldheGFP ( $53.73 \pm 4.08$ ), Glt1 ( $65.36$   
908  $\pm 7.83$ ), AldheGFP + Glt1 ( $51.81 \pm 3.87$ ), vGlut1 ( $62.72 \pm 8.45$ ), AldheGFP + vGlut1 ( $39.68$   
909  $\pm 6.76$ ), Glt1 + vGlut1 ( $47.56 \pm 8.55$ ), and AldheGFP + Glt1 + vGlut1 ( $38.44 \pm 6.79$ ). 80.31  
910  $\pm 4.4$  % of total holes are occupied (by astrocytic markers or synapses or both) leaving  
911  $19.68 \pm 4.4$  % holes empty.  $n = >40\text{PNNs}/8\text{s}/4\text{m}$ .

912 **g** Representative confocal micrographs showing PNN (WFA - yellow), astrocytic  
913 processes labelled with AldheGFP (green) and GAT3 (red), and inhibitory presynaptic  
914 terminals (vGAT - cyan) expression around cortical PV neuron. Arrows point to the PNN  
915 holes containing both GAT3 and vGAT. Bottom images represent 3D reconstruction of  
916 PNN showing PNN holes containing astrocytic processes with GABA transporter and  
917 GABAergic synapses.

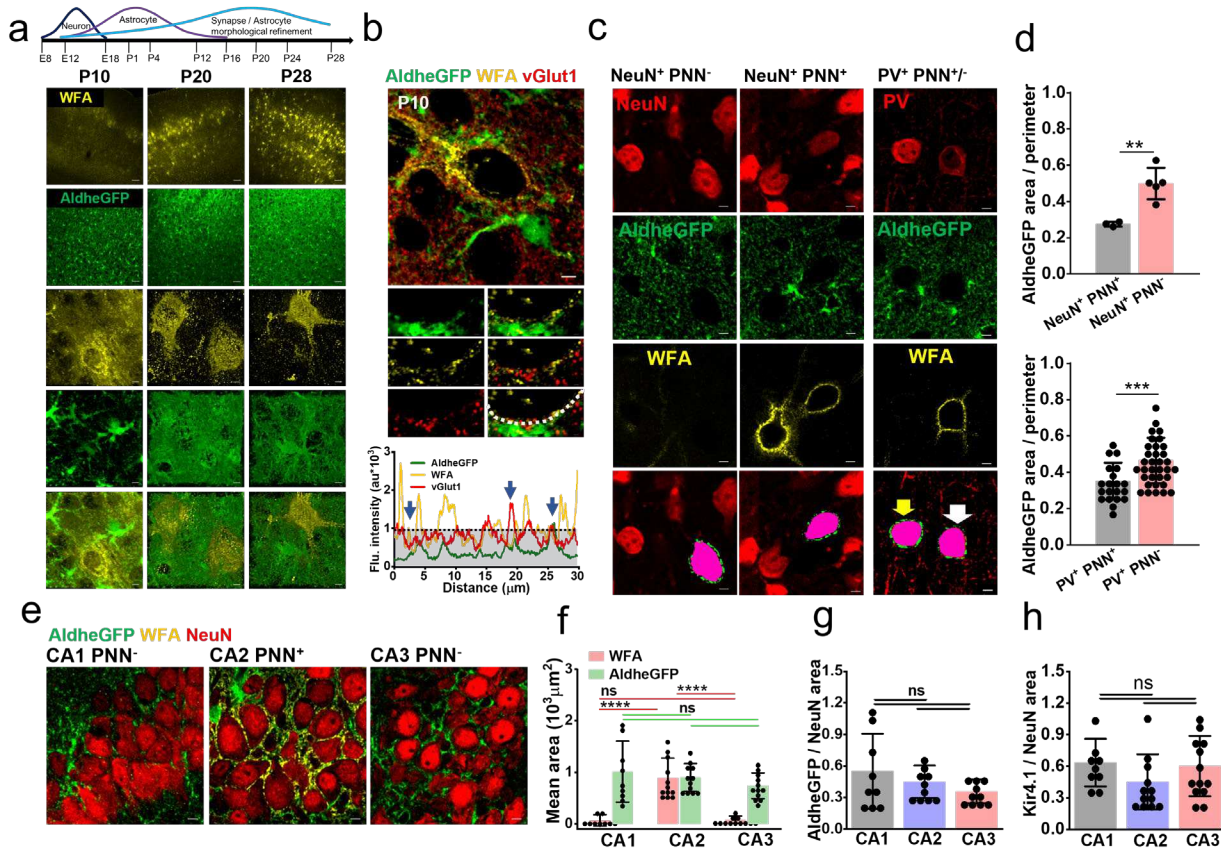
918 **h** Venn diagram showing the proportional occupancy of PNN holes by GABAergic  
919 synapses and astrocytic processes with GABA transporter expression. Bar diagram  
920 shows the percent of total PNN holes occupied by AldheGFP ( $46.51 \pm 7.17$ ), GAT3 ( $63.37$   
921  $\pm 6.81$ ), AldheGFP + GAT3 ( $43.89 \pm 7.4$ ), vGAT ( $53.11 \pm 3.99$ ), GAT3 + vGAT ( $35.89 \pm$   
922  $3.63$ ), AldheGFP + vGAT + GAT3 ( $23.12 \pm 4.45$ ), and AldheGFP + vGAT ( $25.02 \pm 4.76$ ).  
923  $82.72 \pm 4.2$  % of total holes are occupied (by astrocytic markers or synapses or both)  
924 leaving  $17.27 \pm 4.2$  % holes empty.  $n = >40\text{PNNs}/8\text{s}/4\text{m}$ .

925 **i** Representative confocal micrographs showing PNN (WFA - yellow), astrocytic  
926 processes (AldheGFP - green), and inhibitory (vGAT - cyan) and excitatory (vGlut1 - red)  
927 synaptic terminals around cortical FSNs. Arrows point to the PNN holes containing both  
928 vGlut1 and vGAT. Bottom images represent 3D reconstruction of PNN lattice showing  
929 PNN holes containing astrocytic processes (AldheGFP-Green) as well as GABAergic  
930 (vGAT - cyan) and glutamatergic (vGlut1 - red) synaptic terminals.

931 **j** Venn diagram showing the proportional occupancy of PNN holes by glutamatergic and  
932 GABAergic synapses with astrocytic processes. Bar diagram (bottom) shows percent of  
933 total PNN holes occupied by AldheGFP ( $59.41 \pm 1.11$ ), vGAT ( $56.74 \pm 5.83$ ), vGlut1 ( $70.15$   
934  $\pm 4.08$ ), AldheGFP + vGAT ( $31.85 \pm 2.73$ ), AldheGFP + vGlut1 ( $46.60 \pm 3.26$ ), vGAT +  
935 vGlut1 ( $37.35 \pm 4.71$ ), AldheGFP + vGAT + vGlut1 ( $24.60 \pm 3.19$ ).  $94.05 \pm 1.9$  % of total  
936 holes are occupied (by astrocytic markers or synapses or both) leaving  $5.94 \pm 1.96$  %  
937 holes empty.  $n = >40\text{PNNs}/8\text{s}/4\text{m}$ .

938 s and m represent the number of sections and mice. Bar data are expressed as mean  $\pm$   
939 SD. Scale bar 2 $\mu$ m in the top and 1 $\mu$ m in bottom images in a, c, e, g, and i.

940



941 **Fig. 3 Concurrent maturation of PNN and astrocytes leads to a lower pericellular**  
 942 **astrocytic coverage of cortical PV neurons.**  
 943

944 **a** Developmental formation and maturation of neurons, astrocytes, and synapses (top)  
 945 (adapted and modified from [46]). Confocal micrographs of WFA (yellow) and AldheGFP  
 946 (green) immunofluorescence in mouse cerebral cortex in postnatal days 10 (P10), 20  
 947 (P20) and 28 (P28). High magnification 3D volume images of PNNs (3<sup>rd</sup> row from top)  
 948 and astrocytes (4<sup>th</sup> row from top) in different postnatal ages showing concurrent  
 949 maturation. Scale bar 50 $\mu$ m in top two rows, 2 $\mu$ m in bottom 3 rows.

950 **b** Top, confocal micrograph of developing PNN (WFA – yellow) showing astrocytic  
 951 processes (AldheGFP – green) and excitatory synaptic terminals (vGlut1 – red). The  
 952 rectangular area (white) is magnified in the bottom panel images. line intensity profile of  
 953 PNN (white dotted line in bottom right image) shows peaks of vGlut1 and AldheGFP in  
 954 the PNN holes (marked by arrows). Gray area represents the WFA threshold. Scale 5 $\mu$ m  
 955 (top main image) and 2 $\mu$ m (bottom panel images).

956 **c** Representative confocal micrographs showing NeuN (red), AldheGFP (green), and  
 957 WFA (yellow) fluorescence in cortical neurons without (left panel) and with PNN (middle  
 958 panel). Bottom images show the binarized pericellular astrocytic coverage area (green)  
 959 of the cell (pink) of interest. Right panel shows confocal micrographs of PV (red),  
 960 AldheGFP (green), and WFA (yellow) in the PV neurons with and without PNN. Bottom



961 images show binarized pericellular astrocytic coverage areas (green) in PV+ neurons with  
962 (white arrow) and without (yellow arrow) a PNN. Scale 5 $\mu$ m.

963 **d** Top graph showing a significantly lower pericellular AldheGFP area (normalized to cell  
964 perimeter) in NeuN+ PNN+ neurons ( $0.27 \pm 0.01$ ) compared to the NeuN+ PNN- neurons  
965 ( $0.49 \pm 0.08$ ). n = 22c/3m (NeuN+ PNN+); n = 28c/5m (NeuN+ PNN-). Bottom graph shows  
966 a significantly lower pericellular AldheGFP area (normalized to cell perimeter) of PV+  
967 PNN+ neurons ( $0.35 \pm 0.1$ ) compared to the PV+ PNN- neurons ( $0.46 \pm 0.12$ ). n = 20c/4m  
968 (PV+ PNN+); n = 34c/7m (PV+ PNN-).

969 **e** Representative confocal micrographs showing NeuN (red), AldheGFP (green), and  
970 WFA (yellow) fluorescence in hippocampal CA1 (left), CA2 (middle), and CA3 (right)  
971 *stratum pyramidale*. Scale 5 $\mu$ m.

972 **f** Bar graph representing the mean total coverage area of AldheGFP (green) and WFA  
973 (red) in CA1, CA2, and CA3 areas. Due to PNNs, WFA covered area in CA2 is  
974 significantly higher ( $893.23 \pm 383.56$ ) compared to CA1 ( $65.16 \pm 112.02$ ) and CA3 ( $82.03$   
975  $\pm 71.57$ ); however, AldheGFP covered area in CA2 ( $900.02 \pm 268.61$ ) remains statistically  
976 indifferent from CA1 ( $1014.81 \pm 594.63$ ) and CA3 ( $740.86 \pm 251.12$ ). n = 9s/3m in CA1  
977 group; 12s/4m in CA2 and CA3 groups. Red and green lines show comparisons between  
978 red (WFA) bars and green bars (AldheGFP) respectively.

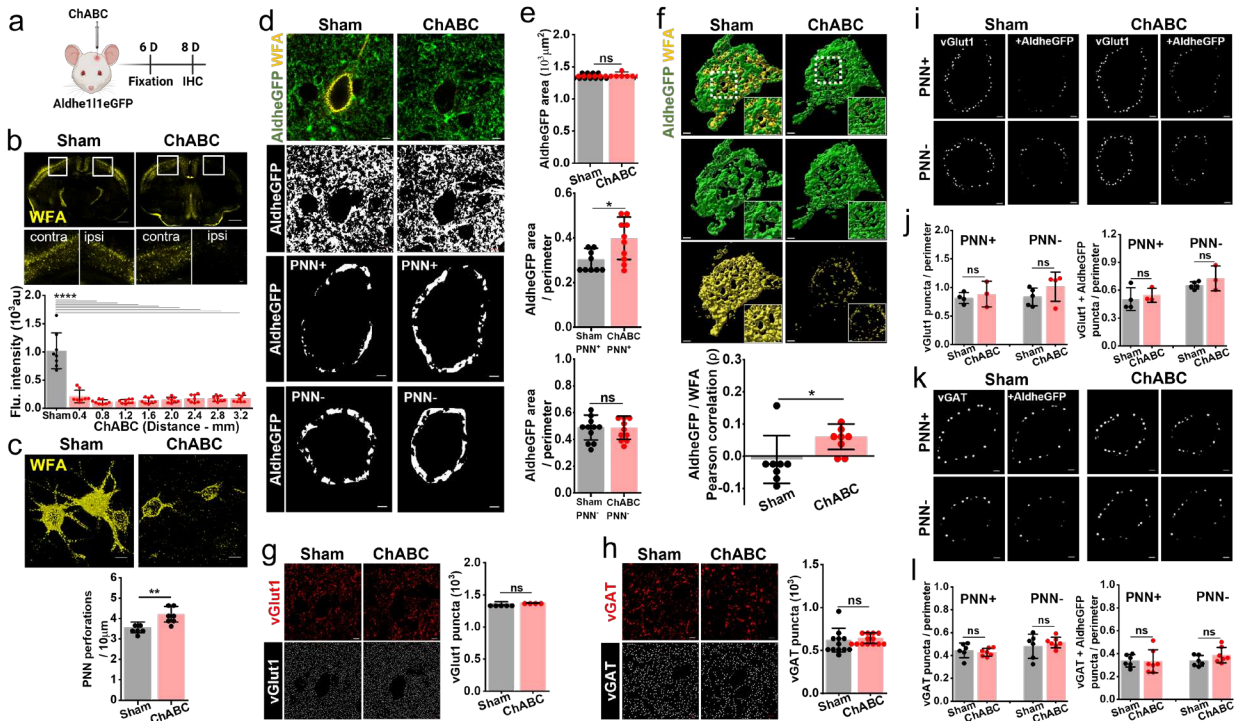
979 **g** Graphs show non-significant total astrocytic area (AldheGFP) normalized to neuronal  
980 area (NeuN) in CA1 ( $0.55 \pm 0.35$ ) CA2 ( $0.45 \pm 0.15$ ) and CA3 ( $0.35 \pm 0.11$ ) regions. n =  
981 9s/3m in CA1; 10s/4m in CA2 and CA3.

982 **h** Graphs showing statistically non-significant total astrocytic area (Kir4.1) normalized to  
983 neuronal area (NeuN) in CA1 ( $0.63 \pm 0.22$ ) CA2 ( $0.45 \pm 0.26$ ) and CA3 ( $0.60 \pm 0.28$ )  
984 regions. n = 9s / 3m in CA1; 12s / 4m in CA2 and 13s/4m CA3.

985 c, s, and m indicate the number of cells, sections, and mice respectively. Bar data are  
986 expressed as mean $\pm$ SD; dots on the bars represent the individual data points. \*\*\*\*P <  
987 0.0001, \*\*\*P < 0.001, \*\*P < 0.01, \*P < 0.05, ns = P > 0.05. unpaired two-tailed t-test with  
988 welch correction in d; One-way ANOVA, Tukey's post-hoc test in f, g, and h.

989

990



991 **Fig. 4 PNN disruption increases pericellular astrocytic coverage without altering**  
 992 **synaptic contacts.**  
 993

994 **a** Schematics of intracranial ChABC injection and subsequent experiments.

995 **b** Confocal micrographs showing immunofluorescence of WFA (yellow) in sham and  
 996 ChABC injected mouse brains. Marked rectangular areas in ipsilateral (right) and  
 997 contralateral (left) hemispheres are magnified in bottom panels. Scale bars, 1mm in top  
 998 and 100µm in bottom images. Bar diagram of fluorescence intensity of WFA in sham and  
 999 ChABC-injected groups showing widespread PNN disruption throughout the cerebral  
 1000 cortex. (Sham: 1019.07 ± 318.16; ChABC: 0.4mm 209.38 ± 111.13, 0.8mm 112.95 ±  
 1001 37.09, 1.2mm 121.11 ± 35.20, 1.6mm 137.94 ± 52.69, 2.0mm 147.37 ± 49.09, 2.4mm  
 1002 170.54 ± 65.98, 2.8mm 168.53 ± 48.71, 3.2mm 168.05 ± 61.36), n = 8s/4m in sham,  
 1003 8s/8m in ChABC.

1004 **c** 3D volume images of PNNs (WFA - yellow) from sham with intact and dense PNNs  
 1005 compared to the ChABC injected mice showing disrupted PNNs with granulated and  
 1006 fragmented WFA labelling. Scale 10µm. Bar diagrams showing a significant increase in  
 1007 PNN perforation after ChABC-mediated PNN degradation (Sham 3.57 ± 0.25, n = 6s/4m;  
 1008 ChABC 4.19 ± 0.38, n = 8s/3m).

1009 **d** Confocal micrographs showing AldheGFP (green) and WFA (yellow) fluorescence (top)  
 1010 and AldheGFP binarized signal of total area (2<sup>nd</sup> row from top), the pericellular area  
 1011 around PNN-expressing (3<sup>rd</sup> row from top) and non-expressing (bottom) neurons in sham  
 1012 (left panels) and ChABC-injected (right panels) groups.

1013 **e** Bar diagram (top) of total AldheGFP area in a field of view of PNNs was unchanged in  
1014 ChABC treated group (sham  $1380.97 \pm 23.04$ ,  $n = 21i$ , 3m; ChABC  $1393.50 \pm 24.84$ ,  $n =$   
1015  $20i$ , 3m). Bar diagram (middle) of normalized pericellular AldheGFP area increased  
1016 significantly in PNN+ neurons in ChABC treated group (sham  $0.30 \pm 0.05$ ,  $n = 48c/9s/7m$ ;  
1017 ChABC  $0.39 \pm 0.09$ ,  $n = 38c/10s/7m$ ) however remained unaltered in PNN- neurons in  
1018 bottom bar diagram (sham  $0.48 \pm 0.09$ ,  $n = 53c/11s/9m$ ; ChABC  $0.48 \pm 0.86$ ,  $n =$   
1019  $38c/10s/8m$ ).

1020 **f** 3D reconstruction of PNN (yellow) and pericellular astrocytic coverage (green) showing  
1021 increased pericellular coverage and disintegrated PNN. Inset images represent the  
1022 magnified areas marked by white squares. Bar data (bottom) show an altered spatial  
1023 correlation between AldheGFP and WFA in ChABC treated condition (sham  $-0.01 \pm 0.07$ ;  
1024 ChABC  $0.06 \pm 0.03$ ,  $n = 8c/3s/3m$  in each group).

1025 **g** Representative confocal images of vGlut1 fluorescence (top-red) and binary form of  
1026 vGlut1 puncta (bottom-white) showing the unaltered numerical density of excitatory  
1027 presynaptic puncta in ChABC treated condition ( $1368.63 \pm 24.14$ ,  $n = 55c/5s/5m$ )  
1028 compared to sham ( $1374 \pm 8.61$ ,  $n = 21c/4s/4m$ ).

1029 **h** Representative confocal images of vGAT fluorescence (top-red) and binary form of  
1030 vGAT puncta (bottom-white) showing the unaltered numerical density of inhibitory  
1031 presynaptic puncta on PNN disruption in ChABC treated condition ( $643.95 \pm 62.87$ ,  $n =$   
1032  $51c/14s/7m$ ) compared to sham ( $620.75 \pm 138.95$ ,  $n = 59c/13s/5m$ ).

1033 **i** Representative binary images from sham (left panels) and ChABC treated (right panels)  
1034 groups showing pericellular vGlut1 puncta (left) and vGlut1 puncta with astrocytic  
1035 contacts (right) in PNN expressing (top row) and PNN non-expressing (bottom row)  
1036 cortical neurons.

1037 **j** Bar graphs showing total pericellular vGlut1 puncta (left) and vGlut1 puncta with  
1038 AldheGFP contacts (right) in sham and ChABC-treated groups. The numerical density of  
1039 vGlut1 puncta remained unchanged in PNN-expressing neurons (sham  $0.81 \pm 0.09$ ,  $n =$   
1040  $22c/4s/4m$ ; ChABC  $0.88 \pm 0.02$ ,  $n = 12c/3s/3m$ ) as well as in PNN non-expressing  
1041 neurons (sham  $0.83 \pm 0.15$ ,  $n = 27c/5s/5m$ ; ChABC  $1.01 \pm 0.22$ ,  $n = 11c/4s/3m$ ). Similarly,  
1042 pericellular numerical density of vGlut1 puncta with astrocytic contacts(+AldheGFP)  
1043 remained unchanged in PNN expressing neurons (sham  $0.50 \pm 0.12$ ,  $n = 22c/4s/4m$ ;  
1044 ChABC  $0.54 \pm 0.07$ ,  $n = 12c/3s/3m$ ) as well as in PNN non expressing neurons (sham  
1045  $0.65 \pm 0.03$ ,  $n = 27c/5s/5m$ ; ChABC  $0.72 \pm 0.13$ ,  $n = 11c/3s/3m$ ).

1046 **k** Representative binary images from sham (left panels) and ChABC treated (right panels)  
1047 groups showing total pericellular vGAT puncta (left) and vGAT puncta with astrocytic  
1048 contacts (right) in PNN expressing (top row) and PNN non-expressing (bottom row)  
1049 cortical neurons.

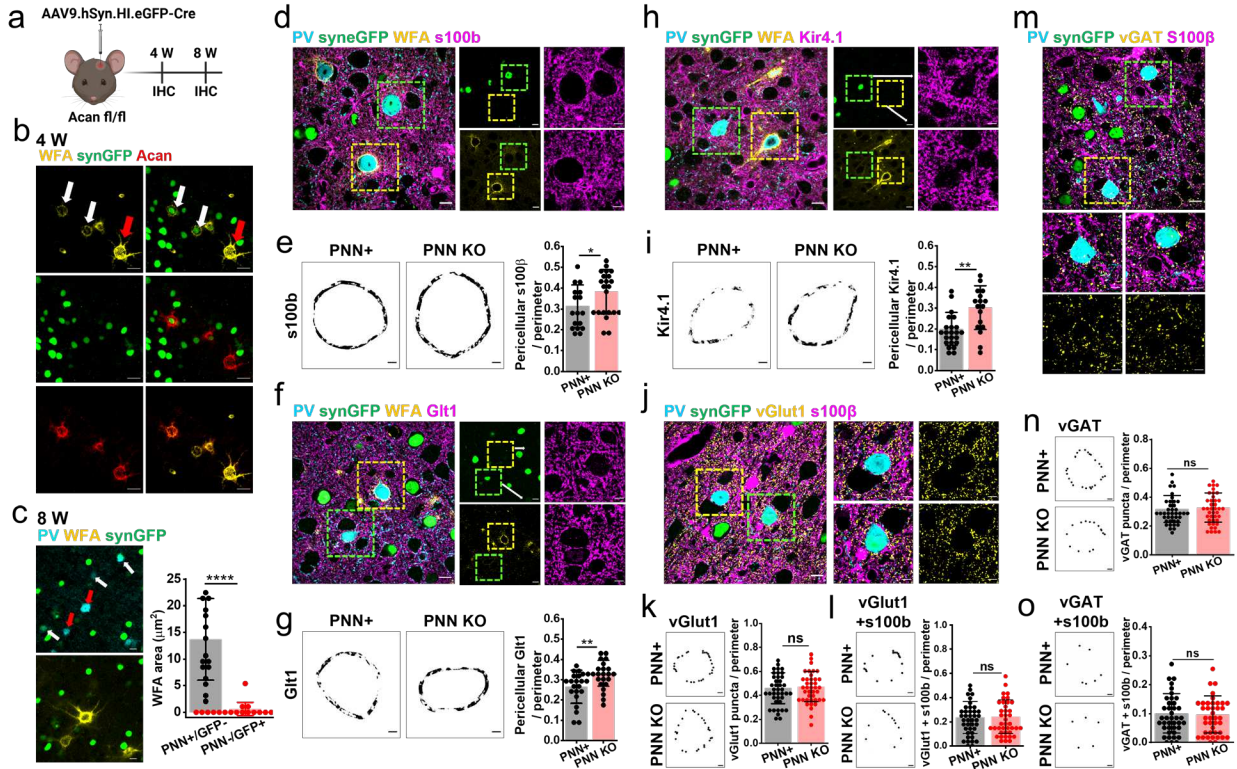
1050 **l** Bar graphs showing total pericellular vGAT puncta (left) and vGAT puncta with  
1051 AldheGFP contacts (right) in sham and ChABC-treated groups. The numerical density of  
1052 vGAT puncta remained unchanged in PNN-expressing neurons (sham  $0.44 \pm 0.6$ ,  $n =$

1053 26c/6s/5m; ChABC  $0.42 \pm 0.03$ ,  $n = 24c/7s/6m$ ) as well as in PNN non-expressing  
1054 neurons (sham  $0.47 \pm 0.10$ ,  $n = 29c/6s/5m$ ; ChABC  $0.51 \pm 0.04$ ,  $n = 27c/7s/6m$ ). Similarly,  
1055 pericellular numerical density of vGAT puncta with astrocytic contacts(+AldheGFP)  
1056 remained unchanged in PNN expressing neurons (sham  $0.33 \pm 0.05$ ,  $n = 26c/6s/5m$ ;  
1057 ChABC  $0.33 \pm 0.10$ ,  $n = 24c/7s/6m$ ) as well as in PNN non expressing neurons (sham  
1058  $0.33 \pm 0.04$ ,  $n = 29c/6s/5m$ ; ChABC  $0.38 \pm 0.06$ ,  $n = 27c/7s/6m$ ).

1059 c, s, and m indicate the number of cells, sections, and mice respectively. Bar graph data  
1060 are expressed as mean $\pm$ SD; dots on the bars represent the individual data points . \*P <  
1061 0.05, ns = P > 0.05. unpaired two-tailed t-test with welch correction in c-l, One-way  
1062 ANOVA, Tukey's post-hoc test in b. Scale bar 5 $\mu$ m in field images in d, g, h; and 2 $\mu$ m in  
1063 magnified images in d, f, i, and k.

1064

1065



1066  
1067  
1068

**Fig. 5 AAV-mediated permanent PNN disruption increases pericellular astrocytic coverage without altering synaptic contacts.**

1069 **a** Schematics of AAV9.hSyn.HI.eGFP-Cre mediated PNN knockout in adult Acan fl/fl  
1070 mice.

1071 **b - c** Confocal micrographs of PV (cyan), WFA (yellow), synGFP (green), and aggrecan  
1072 (red) immunofluorescence from the cortical section of acan fl/fl mice showing (b) partial  
1073 PNN disruption after 4 weeks and (c) complete disruption after 8 weeks of  
1074 AAV9.hSyn.HI.eGFP-Cre injection. AAV-infected and non-infected cells are marked by  
1075 white and red arrows respectively. Bar graph in c showing a negligible WFA area in  
1076 AAV9.hSyn.HI.eGFP-Cre infected (PNN-/GFP+) cells compared to the non-infected cells  
1077 (PNN+/GFP-). Control  $13.66 \pm 7.65$ ,  $n = 17$  i/3m; ChABC  $0.61 \pm 1.21$ ,  $n = 21$ i/3m. Scale  
1078  $20\mu\text{m}$  in b and  $10\mu\text{m}$  in c.

1079 **d** Confocal micrograph of PV (cyan), SynCreGFP (green), PNN (WFA - yellow), and  
1080 astrocyte (s100b - magenta) fluorescence in Acan fl/fl injected with SynCreGFP prefrontal  
1081 cortical section. Transduced PV neurons (green square) show PNN elimination compared  
1082 to the non-transduced PV neurons with intact PNNs (yellow square). The left panels show  
1083 magnified areas in the squares.

1084 **e** Binary images of pericellular s100b area from PV neuron with intact PNN (PNN+, left)  
1085 and AAV-mediated PNN KO (right). Bar graph (right) shows increased pericellular s100b  
1086 coverage on PV neurons with PNN KO (PNN+  $0.31 \pm 0.10$ ,  $n = 17$ c/3m; PNN KO  $0.38 \pm$   
1087  $0.10$ ,  $n = 22$ c/3m).

1088 **f** Confocal micrographs of PV (cyan), SynCreGFP (green), PNN (WFA - yellow), and  
1089 astrocyte (Glt1 - magenta) fluorescence in Acan fl/fl injected with SynCreGFP prefrontal  
1090 cortical section. Transduced PV neurons (green square) show PNN elimination compared  
1091 to the non-transduced PV neurons with intact PNNs (yellow square). Right panels show  
1092 magnified areas in the squares.

1093 **g** Binary images of pericellular Glt1 area from PV neuron with intact PNN (PNN+, left)  
1094 and AAV-mediated PNN KO (right). Bar diagram (right) shows increased pericellular Glt1  
1095 coverage on PV neurons with PNN KO (PNN+  $0.26 \pm 0.08$ , n = 20c/3m; PNN KO  $0.33 \pm$   
1096  $0.06$ , n = 26c/3m).

1097 **h** Confocal micrographs of PV (cyan), SynCreGFP (green), PNN (WFA - yellow), and  
1098 astrocyte (Kir4.1 - magenta) fluorescence in Acan fl/fl injected with SynCreGFP prefrontal  
1099 cortical section. Transduced PV neurons (green square) show PNN elimination compared  
1100 to the non-transduced PV neurons with intact PNNs (yellow square). Right panels show  
1101 magnified areas in the squares.

1102 **i** Binary images of pericellular Kir4.1 area from PV neuron with intact PNN (PNN+, left)  
1103 and AAV-mediated PNN KO (right). Bar diagram (right) shows increased pericellular  
1104 Kir4.1 coverage on PV neurons with PNN KO (PNN+  $0.19 \pm 0.08$ , n = 23c/3m; PNN KO  
1105  $0.30 \pm 0.10$ , n = 17c/3m).

1106 **j** Confocal micrographs of PV (cyan), SynCreGFP (green), glutamatergic synapses  
1107 (vGlut1 - yellow), and astrocyte (s100b - magenta) fluorescence in Acan fl/fl injected with  
1108 SynCreGFP prefrontal cortical section. Yellow and green squares mark the magnified  
1109 areas in top and bottom panels showing non-transduced and transduced PV neurons  
1110 respectively.

1111 **k** Binary images of pericellular vGlut1 puncta on PV neuron with intact PNN (PNN+, top)  
1112 and AAV-mediated PNN KO (bottom). Bar diagram (right) shows unaltered pericellular  
1113 vGlut1 puncta on PV neurons with PNN KO (PNN+  $0.46 \pm 0.12$ , n = 40c/4m; PNN KO  
1114  $0.47 \pm 0.12$ , n = 40c/4m).

1115 **l** Binary images of pericellular vGlut1 puncta with s100b processes from PV neuron with  
1116 intact PNN (PNN+, top) and AAV-mediated PNN KO (bottom). Bar diagram (right) shows  
1117 unaltered pericellular vGlut1 puncta with s100b processes on PV neurons with PNN KO  
1118 (PNN+  $0.23 \pm 0.13$ , n = 40c/4m; PNN KO  $0.24 \pm 0.13$ , n = 40c/4m).

1119 **m** Confocal micrographs of PV (cyan), SynCreGFP (green), GABAergic synapses (vGAT  
1120 - yellow), and astrocyte (s100b - magenta) fluorescence in Acan fl/fl injected with  
1121 SynCreGFP prefrontal cortical section. Yellow and green squares mark the magnified  
1122 areas in the bottom panels showing non-transduced and transduced PV neurons  
1123 respectively.

1124 **n** Binary images of pericellular vGAT puncta on PV neuron with intact PNN (PNN+, top)  
1125 and AAV-mediated PNN KO (bottom). Bar diagram (right) shows unaltered pericellular

1126 vGAT puncta on PV neurons with PNN KO (PNN+  $0.31 \pm 0.09$ , n = 40c/4m; PNN KO  $0.32$   
1127  $\pm 0.10$ , n = 40c/4m).

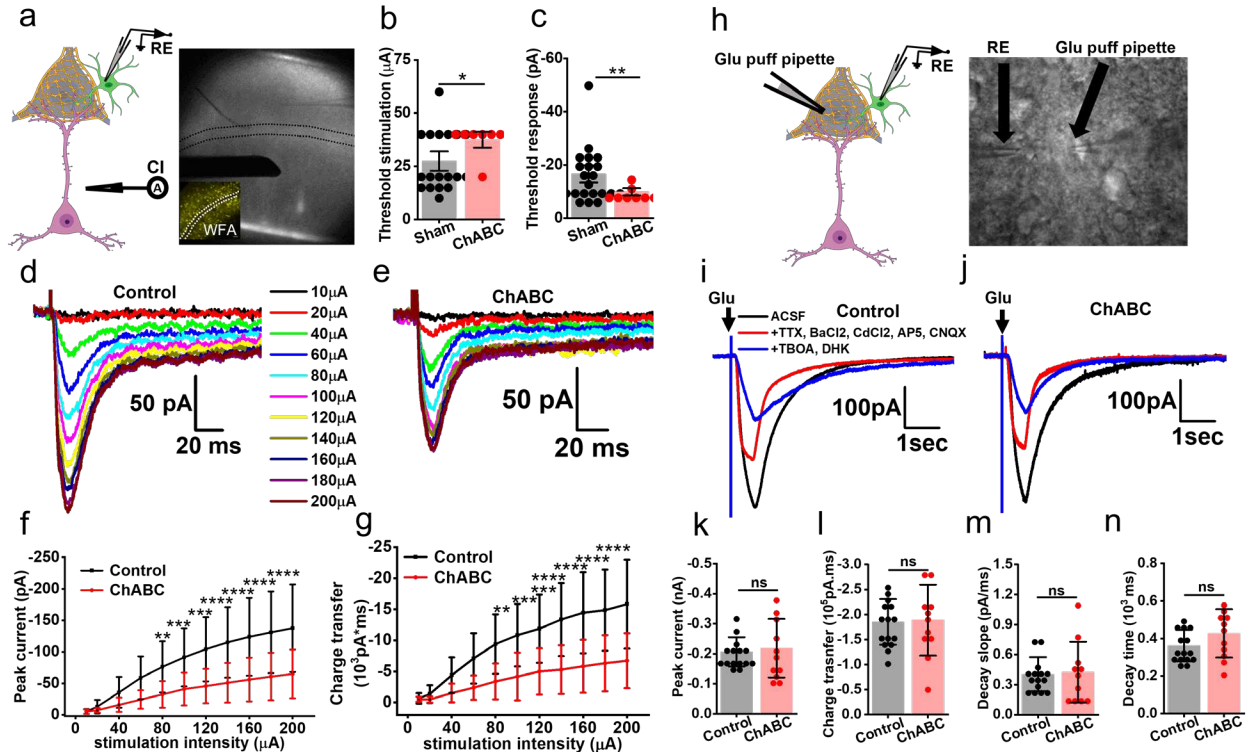
1128 ○ Binary images of pericellular vGAT puncta with s100b processes from PV neuron with  
1129 intact PNN (PNN+, top) and AAV-mediated PNN KO (bottom). Bar diagram (right) shows  
1130 unaltered pericellular vGAT puncta with s100b processes on PV neurons with PNN KO  
1131 (PNN+  $0.10 \pm 0.06$ , n = 40c/4m; PNN KO  $0.09 \pm 0.06$ , n = 40c/4m).

1132 c, i, and m indicate the number of cells, images, and mice respectively. \*\*\*\*P < 0.0001,  
1133 \*\*P < 0.01, \*P < 0.05, ns = P > 0.05 unpaired two-tailed t-test with Welch correction. Bar  
1134 data are expressed as mean $\pm$ SD; dots on the bars represent the individual data points.  
1135 Scale bar d - o: 10 $\mu$ m large images, 5 $\mu$ m magnified images, 2 $\mu$ m binary images.

1136

1137





1138  
1139

**Fig. 6 Acute PNN depletion disrupts glutamate uptake by astrocytes.**

1140 **a** Schematics showing stimulation of glutamatergic neurons by a current injector (CI)  
1141 causing glutamate release by synapses and subsequent uptake by astrocytic processes  
1142 in the PNN holes. Brightfield image showing placement of stimulator (CI) and recording  
1143 electrode (RE) in the cortical slice. The Inset image shows WFA labelling in a cortical  
1144 slice.

1145 **b - c** Bar diagrams showing significantly higher threshold stimulation (**b**) (control  $27.5 \pm$   
1146  $13.52$ ,  $n = 20c/10m$ ; ChABC  $37.5 \pm 7.07$ , units  $\mu A$ ,  $n = 8c/5m$ ) and lower glutamate uptake  
1147 threshold response (**c**) (control  $-16.6 \pm 10.07$ ,  $n = 22c/10m$ ; ChABC  $-9.9 \pm 2.5$ ,  $n = 8c/5m$ ,  
1148 units pA) in ChABC treated slices than control slices.

1149 **d - e** Representative glutamate uptake current traces recorded from cortical astrocytes in  
1150 response to a series of incrementing field stimulate in control (**d**), and in ChABC treated  
1151 (**e**) acute slices.

1152 **f - g** Input-output curves recorded from cortical astrocytes showing significantly lower  
1153 peak glutamate uptake currents (**e**), as well as a lower charge transfer (**f**) in ChABC-  
1154 treated slices compared to control.  $n = 24c/10m$ ; ChABC  $16c/6m$  both **f** and **g**.

1155 **h** Schematics of recording method (left) and bright-field image (right) of an acute slice  
1156 showing the recording electrode (RE) and glutamate puff electrode.

1157 **i - j** Representative voltage clamp traces of astrocytic currents on puffing  $200\mu M$   
1158 glutamate in presence of various blockers to isolate glutamate uptake current in control  
1159 (**i**) and ChABC-treated (**j**) acute cortical slices.

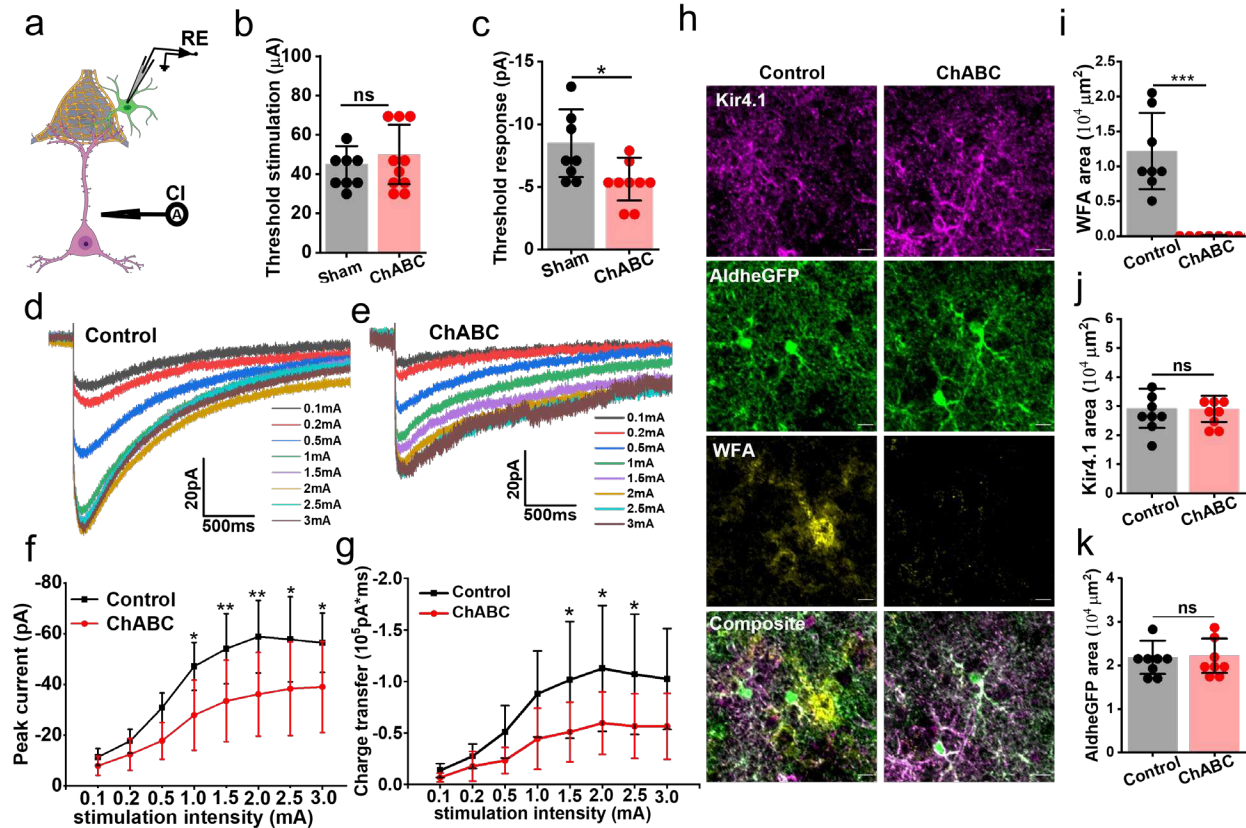


1160 **k - n** Bar diagrams showing unchanged (**k**) peak uptake current (control  $-206.51 \pm 48.59$ ;  
1161 ChABC  $-218.77 \pm 97.48$ ), (**l**) total charge transfer (control  $-154042.79 \pm 45650.84$ ; ChABC  
1162  $-188759.21 \pm 70795.66$ ), (**m**) decay slope (control  $0.405 \pm 0.17$ ; ChABC  $0.425 \pm 0.30$ ),  
1163 and (**n**) current decay time (control  $362.80 \pm 83.68$ ; ChABC  $427.35 \pm 128.51$ ) of astrocytes  
1164 in ChABC treated slices compared to control slices.  $n = 15c/8m$  in the control and  $11c/5m$   
1165 in ChBAC treated group. Units, pA (c), pA.ms (d), pA/ms (e), and ms (e).

1166 c and m indicate the number of cells and mice respectively. \*\*\*\*P < 0.0001, \*\*\*P < 0.001,  
1167 \*\*P < 0.01, \*P < 0.05, ns = P > 0.05. unpaired two-tailed student t-test (equal variance  
1168 not assumed) in b, c, j-m; two-way ANOVA, Tukey's post-hoc test in f and g. Bar data are  
1169 expressed as mean $\pm$ SD; dots on the bars represent the individual data points.

1170

1171



1172  
1173

**Fig. 7 Acute PNN digestion disrupts potassium uptake by astrocytes.**

1174 **a** Schematics showing stimulation of glutamatergic neuron by a current injector (CI)  
1175 causing neuronal depolarization and release of extracellular potassium and subsequent  
1176 uptake by astrocytic processes.

1177 **b-c** Bar diagrams showing **(b)** threshold stimulation (control:  $45 \pm 9.25\mu\text{A}$ ,  $n = 8\text{c}/3\text{m}$ ;  
1178 ChABC:  $47.77 \pm 14.16 \mu\text{A}$ ,  $10\text{c}/5\text{m}$ ) and **(c)** significantly lower threshold  $\text{K}^+$  current  
1179 response (control:  $-8.62 \pm 2.69\text{pA}$ ,  $n = 8\text{c}/3\text{m}$ ; ChABC:  $-5.627 \pm 1.70\text{pA}$ ,  $n = 9\text{c}/5\text{m}$ ) in  
1180 ChABC treated slices than control slices.

1181 **d - e** Representative  $\text{K}^+$  uptake current traces recorded from cortical astrocytes in  
1182 response to a series of incrementing field stimuli in control **(d)** and ChABC-treated **(e)**  
1183 acute slices.

1184 **f - g** Input-output curves recorded from cortical astrocytes showing **(f)** significantly lower  
1185 peak  $\text{K}^+$  uptake currents, as well as **(g)** a lower charge transfer in ChABC-treated slices  
1186 compared to control.  $n = \text{control } 11\text{c}/7\text{m}$ ; ChABC  $15\text{c}/6\text{m}$  in **f**; and  $n = \text{control } 11\text{c}/7\text{m}$ ;  
1187 ChABC  $13\text{c}/6\text{m}$  in **g**.  $***P < 0.001$ ,  $**P < 0.01$ ,  $*P < 0.05$ ,  $\text{ns} = P > 0.05$ . Two-way ANOVA,  
1188 Tukey's post-hoc test.

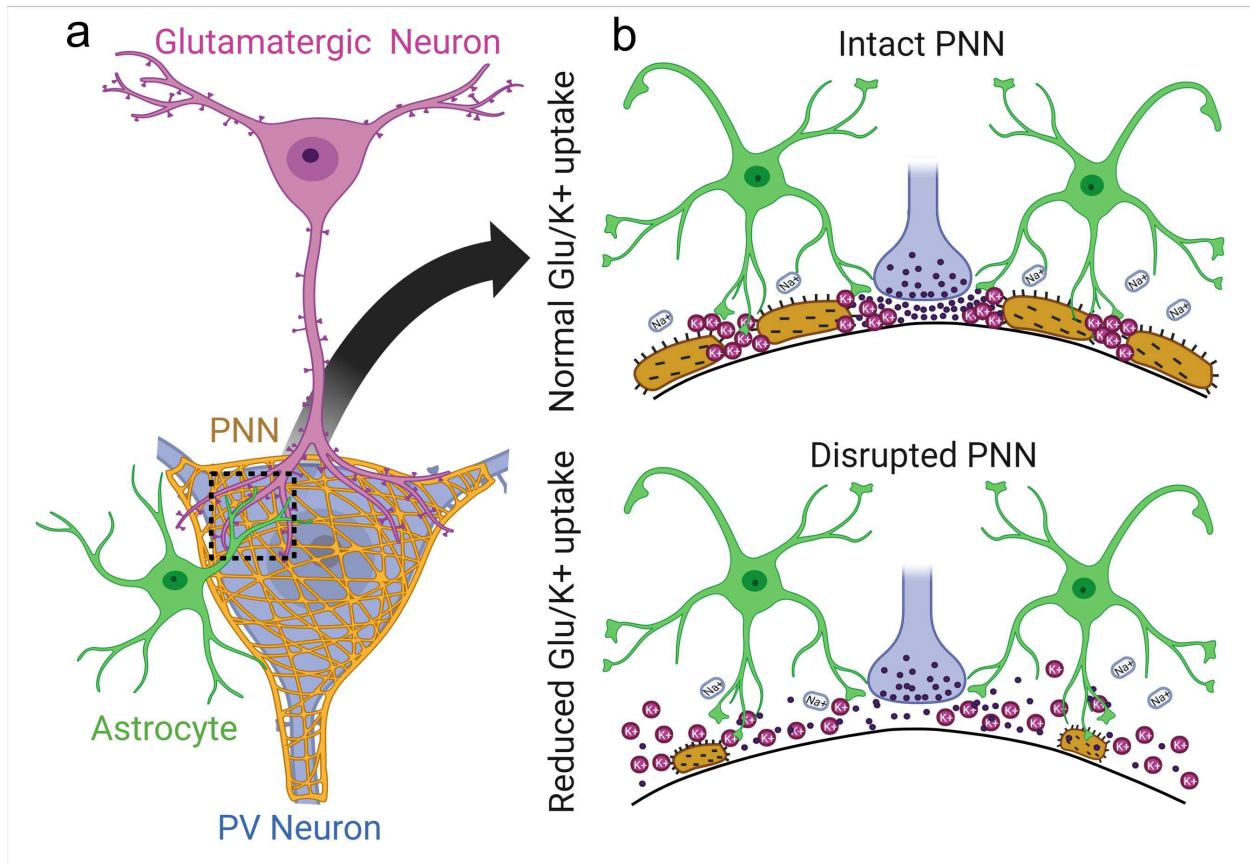
1189 **h** Confocal micrographs of Kir4.1 (magenta), aldheGFP (green), and WFA (yellow) from  
1190 fixed acute slices from control and after ChABC treatment. Scale bar  $10\mu\text{m}$ .

1191 **i - k** Bar diagrams of immunofluorescence area of WFA (**i**) (control  $12175.74 \pm 5481.42$ ,  
1192 ChABC  $84.24 \pm 40.79 \mu\text{m}^2$ ), Kir4.1 (**j**) (control  $29277.79 \pm 6735.50$ , ChABC  $29123.26 \pm$   
1193  $4496.46 \mu\text{m}^2$ ), and AldheGFP (**k**) (control  $21833.27 \pm 3826.81$ , ChABC  $22221.49 \pm$   
1194  $3948.68 \mu\text{m}^2$ ) showing PNN disruption without any changes in astrocytic Kir4.1  
1195 expression. N = 8s/4m in each group.

1196 c, s, and m indicate the number of cells, slices, and mice respectively. \*\*\*\*P < 0.0001, \*\*\*P  
1197 < 0.001, \*\*P < 0.01, \*P < 0.05, ns = P > 0.05. unpaired two-tailed student t-test with Welch  
1198 correction in b, c, l, j, and k. Two-way ANOVA with Tukey's post-hoc test in f and g. Bar  
1199 data are expressed as mean $\pm$ SD; dots on the bars represent the individual data points.

1200

1201



1202  
1203  
1204

**Figure 8: PNN facilitates astrocytic uptake of synaptic activity released glutamate and potassium.**

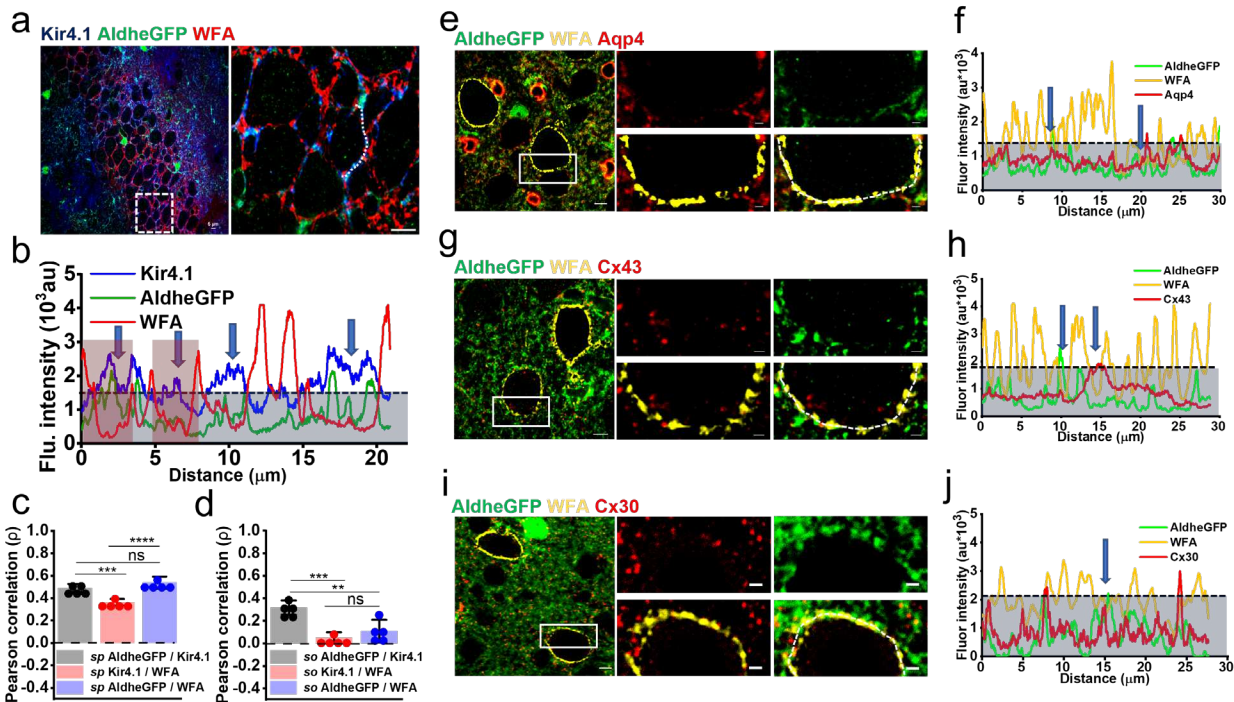
1205 **a** PNN-enclosed PV neurons receive glutamatergic inputs in the PNN holes wherein  
1206 astrocytic processes are also confined (magnified area) and make this space analogous  
1207 to the tripartite synapse.

1208 **b** Under physiological conditions, when PNN is intact (top), glutamate released during  
1209 synaptic activity and extracellular potassium ions released upon neuronal repolarization  
1210 are directed towards astrocytic processes by the PNN to aid efficient clearance of  
1211 extracellular glutamate and potassium. When PNNs are disrupted (bottom), glutamate  
1212 and potassium ions escape from and diffused laterally into the extracellular space.

1213

1214 **Supplementary figures**

1215



1216  
1217

**Fig. S1 In CA2 of hippocampus, PNNs holes contain astrocytic processes.**

1218 **a** Representative confocal micrographs showing hippocampal CA2 PNNs (WFA - red)  
1219 and astrocytes labelled with AldheGFP (green) and Kir4.1 (blue). The right image shows  
1220 the magnified area of the white square in left large image. Scale bar 5μm.

1221 **b** Line intensity profiles of the white dotted line drawn in Fig. **a** (right), showing  
1222 fluorescence intensity of PNN and astrocytic markers. Blue arrows point to the PNN holes  
1223 occupied by astrocytic processes expressing Kir4.1. Red bars in between two  
1224 consecutive WFA peaks represent the area of PNN holes wherein astrocytic processes  
1225 can be confined.

1226 **c-d** Bar diagrams showing Pearson correlation of spatial overlap between astrocytic  
1227 markers AldheGFP and Kir4.1 with each other and with PNN marker WFA in **c** stratum  
1228 pyramidale (Kir4.1-AldheGFP  $0.49 \pm 0.03$ , Kir4.1-WFA  $0.36 \pm 0.02$ , AldheGFP-WFA  $0.53$   
1229  $\pm 0.05$ ) and **d** stratum radiatum of CA2 (Kir4.1-AldheGFP  $0.31 \pm 0.06$ , Kir4.1-WFA  $0.05 \pm$   
1230  $0.04$ , AldheGFP-WFA  $0.10 \pm 0.10$ ).  $n = 5$  sections/5 mice, \*\*\*\* $P < 0.0001$ , \*\*\* $P < 0.001$ ,  
1231 \*\* $P < 0.01$ , \* $P < 0.05$ , ns =  $P > 0.05$ . One-way ANOVA, Tukey's post-hoc test. Bar data  
1232 are expressed as mean  $\pm$  SD; dots on the bars represent the individual data points.

1233 **e** Representative confocal micrographs showing expression of aquaporin 4 (red) in  
1234 astrocytic processes (AldheGFP – green) in holes of cortical PNNs (WFA - yellow). The  
1235 right side panels show the magnified area marked by a white rectangle in the left image.

1236 **f** Intensity profiles of the white dotted line drawn on the PNN (bottom right panel)  
1237 represented PNN holes (marked by blue arrows) occupied with astrocytic processes  
1238 (AldheGFP-green) expressing Aqp4 (red).

1239 **g** Representative confocal micrographs showing expression of connexin 43 (red) in  
1240 astrocytic processes (AldheGFP – green) in holes of cortical PNNs (WFA - yellow). The  
1241 right side panels show the magnified area marked by the white rectangle in the left image.

1242 **h** Intensity profiles of the white dotted line drawn on the PNN (bottom right panel)  
1243 represented PNN holes (marked by blue arrows) occupied with astrocytic processes  
1244 (AldheGFP-green) expressing connexin 43 (red).

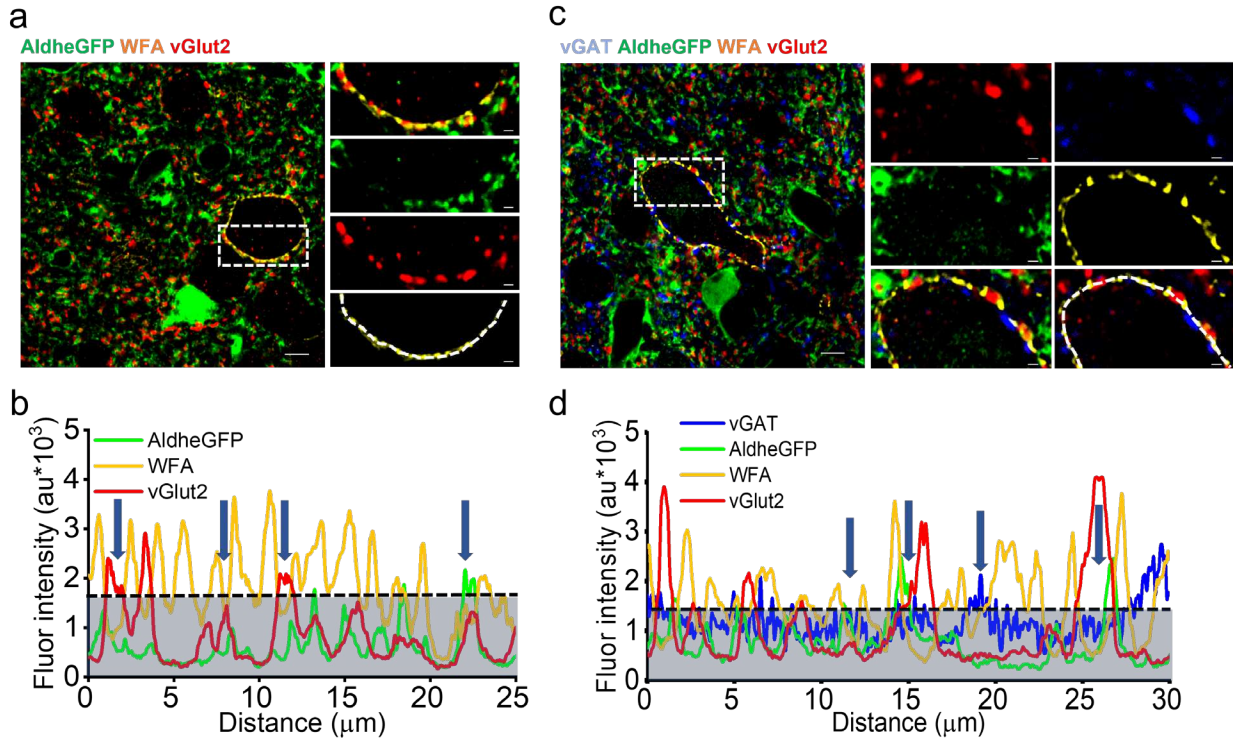
1245 **i** Representative confocal micrographs showing expression of connexin 30 (red) in  
1246 astrocytic processes (AldheGFP – green) in holes of cortical PNNs (WFA - yellow). The  
1247 right side panels show a magnified area marked by the white rectangle in left image.

1248 **j** Intensity profiles of the white dotted line drawn on the PNN (bottom right panel)  
1249 represented PNN holes (marked by blue arrows) occupied with astrocytic processes  
1250 (AldheGFP-green) expressing connexin 30 (red).

1251 Scale bars 5 $\mu$ m in large images, 1 $\mu$ m in magnified images in e-i. Blue area under the  
1252 dotted lines in line profiles in b, f, h, and j represents the WFA threshold.

1253





1254  
1255  
1256

**Fig. S2 PNN holes contain astrocytic processes and thalamocortical synaptic contacts.**

1257 **a** Representative confocal micrographs showing expression of vGlut2 (red) expressing  
1258 synapses as well as astrocytic processes (AldheGFP – green) in holes of cortical PNNs  
1259 (WFA - yellow). Right side panels show a magnified area marked by the white rectangle  
1260 in left image.

1261 **b** Intensity profiles of the white dotted line drawn on the PNN (bottom right panel)  
1262 represented PNN holes (marked by blue arrows) occupied with astrocytic processes  
1263 (AldheGFP-green) and vGlut2 expressing synapses (red).

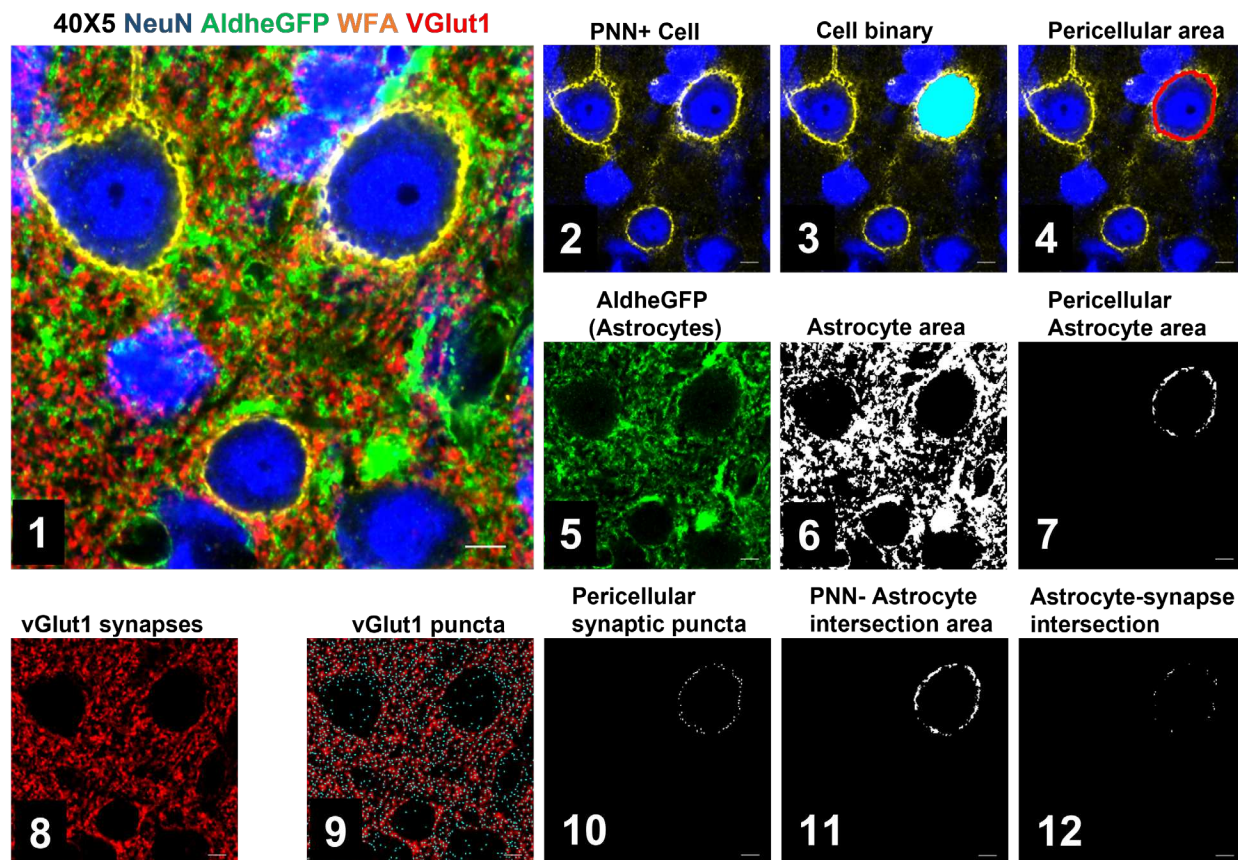
1264 **c** Representative confocal micrographs showing expression of glutamatergic (vGlut2 -  
1265 red) and GABAergic (vGAT - blue) synapses as well as astrocytic processes (AldheGFP  
1266 – green) in holes of cortical PNNs (WFA - yellow). Right side panels show a magnified  
1267 area marked by the white rectangle in left image.

1268 **d** Intensity profiles of the white dotted line drawn on the PNN (bottom right panel)  
1269 represented PNN holes (marked by blue arrows) occupied with astrocytic processes  
1270 (AldheGFP-green) and glutamatergic (vGlut2 - red) and GABAergic (vGAT - blue)  
1271 synapses.

1272 Scale bar 5 $\mu$ m in large images, 1 $\mu$ m in magnified images in both a and c. Blue area under  
1273 the dotted lines in line profiles in b and d represents the WFA threshold.

1274



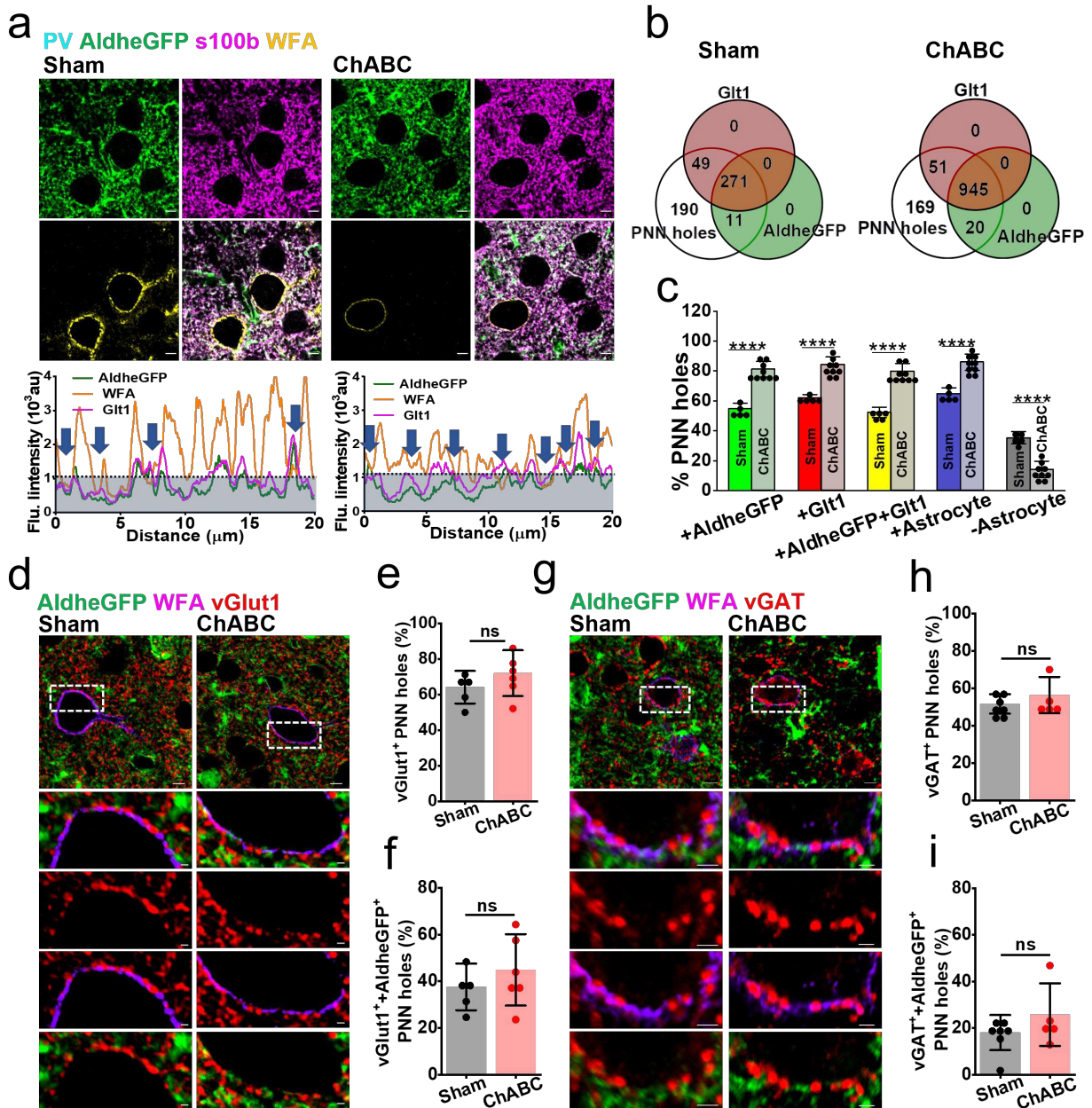


1275  
1276

**Fig. S3 Pericellular astrocytic coverage and synaptic contacts analysis method.**

1277 Multichannel confocal image (1), showing immunofluorescence labeling of neurons  
 1278 (NeuN), astrocytes (AldheGFP), PNNs (WFA), and synaptic terminals (vGlut1). The NeuN  
 1279 signal of the soma of PNN-expressing neurons (2), is binarized (3), and the pericellular  
 1280 0.8 - 1 $\mu$ m area (4) is defined. AldheGFP signal (5), is binarized using automated OTSU  
 1281 function (6), and pericellular AldheGFP area (7), is extracted by intersecting (4) and (6)  
 1282 binary images. Synaptic marker vGlut1 (8), image is processed with an automated peak  
 1283 detection function to detect vGlut1 puncta (8). Intersecting (9) with (4) generates  
 1284 pericellular synaptic puncta (10). Intersecting the PNN signal with (7) generates a PNN-  
 1285 astrocyte intersection area (11), and intersecting (7) with (10) generates vGlut1 puncta in  
 1286 contact with the pericellular astrocytic area (12). Scale bar: 5 $\mu$ m.

1287  
1288



1289  
1290  
1291

**Fig. S4 Astrocytic processes occupy newly formed PNN holes after ChABC treatment.**

1292 **a** Confocal micrographs showing immunofluorescence of astrocytes (AldheGFP – green),  
1293 astrocytic glutamate transporter Glt1 (magenta), and PNNs (WFA - yellow), from sham  
1294 and ChABC-injected mouse cerebral cortex. Scale bar 5 $\mu$ m. Line intensity profiles of a  
1295 typical PNN from sham (left) and ChABC treated (right) conditions showing low WFA  
1296 intensity and high occupancy of PNN perforations with astrocytic processes. Blue area  
1297 under the dotted line represents the threshold WFA intensity.

1298 **b** Venn diagrams showing the proportional occupancy of PNN holes by astrocytic  
1299 processes (AldheGFP + Glt1) in sham (left) and ChABC-treated (right) conditions.

1300 **c** Bar diagram showing the percent of total PNN holes in sham and ChABC treated groups  
1301 occupied by AldheGFP (Control  $54.76 \pm 3.75$ , ChABC  $81.05 \pm 5.24$ ), Glt1 (Control  $62.09$   
1302  $\pm 2.16$ , ChABC  $84.15 \pm 5.26$ ) and both (Control  $52.32 \pm 3.50$ , ChABC  $79.64 \pm 5.38$ ), any  
1303 astrocytic marker positive (Control  $64.67 \pm 4.09$ , ChABC  $85.80 \pm 5.51$ ) and any astrocytic  
1304 marker negative (Control  $35.32 \pm 4.09$ , ChABC  $14.19 \pm 5.51$ ) holes.  $n \geq 40$  PNNs/8s/4m  
1305 in each group.

1306 **d** Confocal micrographs showing immunofluorescence of astrocytes (AldheGFP – green),  
1307 excitatory terminals vGlut1 (red), and PNNs (WFA - magenta), from sham and ChABC-  
1308 injected mouse brains. Magnified images of different combinations showing synaptic  
1309 contacts in PNN holes in sham and ChABC-injected mouse brains. Scale bar  $5\mu\text{m}$  in the  
1310 large image,  $1\mu\text{m}$  in magnified images.

1311 **e** Pericellular density of vGlut1 terminal in the PNN holes in the ChABC-treated group  
1312 remained unaltered compared to sham (Control  $64.10 \pm 9.14$ ,  $n = 22$ PNNs/5m, ChABC  
1313  $71.98 \pm 12.89$ , 35PNNs/5m).

1314 **f** Pericellular density of vGlut1 terminal with astrocytic contacts in the PNN holes in  
1315 ChABC treated group remained unaltered compared to sham (Control  $37.58 \pm 9.97$ ,  $n =$   
1316  $22$ PNNs/5m ChABC  $44.87 \pm 15.29$ , 35PNNs/5m).

1317 **g** Confocal micrographs showing immunofluorescence of astrocytes (AldheGFP – green),  
1318 inhibitory terminals vGAT (red), and PNNs (WFA – magenta), from sham and ChABC-  
1319 injected mouse brains. Magnified images of different combinations showing synaptic  
1320 contacts in PNN holes in sham and ChABC-injected mouse brains. Scale bar  $5\mu\text{m}$  in the  
1321 large image,  $1\mu\text{m}$  in magnified images.

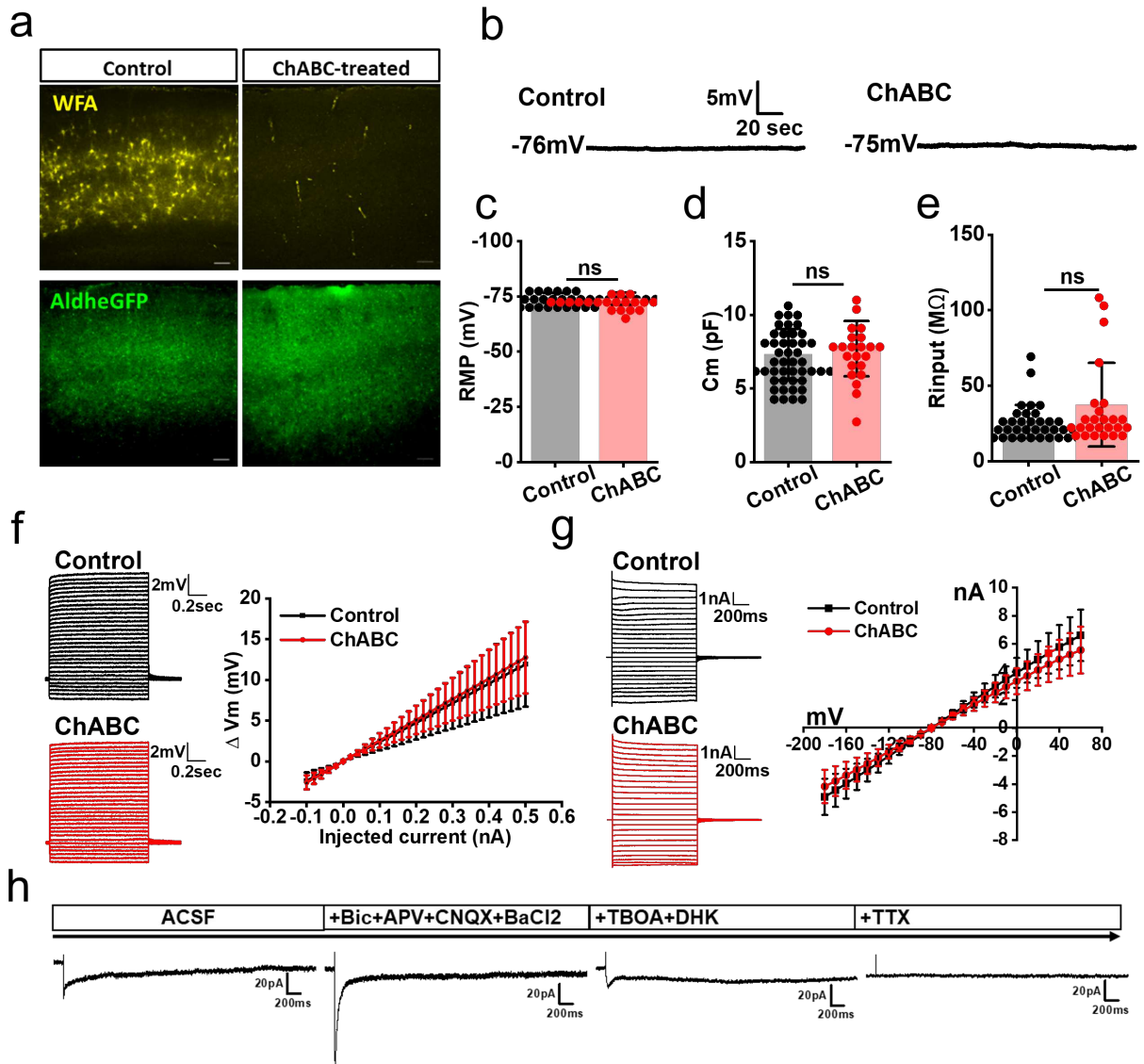
1322 **h** Pericellular density of vGAT terminal in the PNN holes in ChABC treated group  
1323 remained unaltered compared to sham (Control  $51.62 \pm 5.20$ ,  $n = 40$ PNNs/6m; ChABC  
1324  $56.36 \pm 9.69$ , 26PNNs/5m).

1325 **i** Pericellular density of vGAT terminal with astrocytic contacts in the PNN holes in ChABC  
1326 treated group remained unaltered compared to sham (Control  $18.11 \pm 7.53$ ,  $n =$   
1327  $22$ PNNs/5m ChABC  $25.81 \pm 13.43$ , 35PNNs/5m).

1328 s and m indicate the number of sections and mice respectively. \*\*\*\*P < 0.0001, \*\*\*P <  
1329 0.001, \*\*P < 0.01, \*P < 0.05, ns = P > 0.05. One-way ANOVA, Tukey's post-hoc test in d;  
1330 Unpaired two-tailed t-test with Welch's correction in f, g, i, and j. Bar data are expressed  
1331 as mean $\pm$ SD; dots represent individual data points.

1332

1333



1334  
1335  
1336

**Figure S5. Biophysical properties of astrocytes remain unchanged on PNN disruption with ChABC.**

1337 **a** Confocal images of WFA and AldheGFP fluorescence in control and ChABC-treated  
1338 acute slices fixed and stained after electrophysiological recordings. Scale 100μm.

1339 **b** Representative current-clamp traces of astrocytic resting membrane potential from  
1340 control and ChABC-treated slices.

1341 **c - e** Bar diagrams showing unchanged (**c**) resting membrane potential (control  $-74.47 \pm$   
1342  $2.24\text{mV}$ ,  $n = 49\text{c}/18\text{m}$ ; ChABC  $-74.04 \pm 2.7\text{mV}$ ,  $24\text{c}/6\text{m}$ ), (**d**) membrane capacitance  
1343 (control  $7.29 \pm 1.3\text{pF}$ ,  $n = 46\text{c}/17\text{m}$ , ChABC  $7.70 \pm 1.8$ ,  $n = 22\text{c}/7\text{m}$ ), and (**e**) input  
1344 resistance (control  $-26.29 \pm 37.46\text{m}\Omega$ ,  $n = 41\text{c}/18\text{m}$ ; ChABC  $-37.46 \pm 27.69 \text{m}\Omega$ ,  $25\text{c}/7\text{m}$ )  
1345 of astrocytes in ChABC treated slices compared to control slices. ns signifies  $P > 0.05$ ,  
1346 Unpaired two-tailed student's t-test with Welch correction in **c - e**.



1347 **f** Representative current clamp traces and IV plot showing the current-voltage relationship  
1348 of astrocytes in control (n = 49c/18m) and ChABC treated (n = 28c/9m) slices.

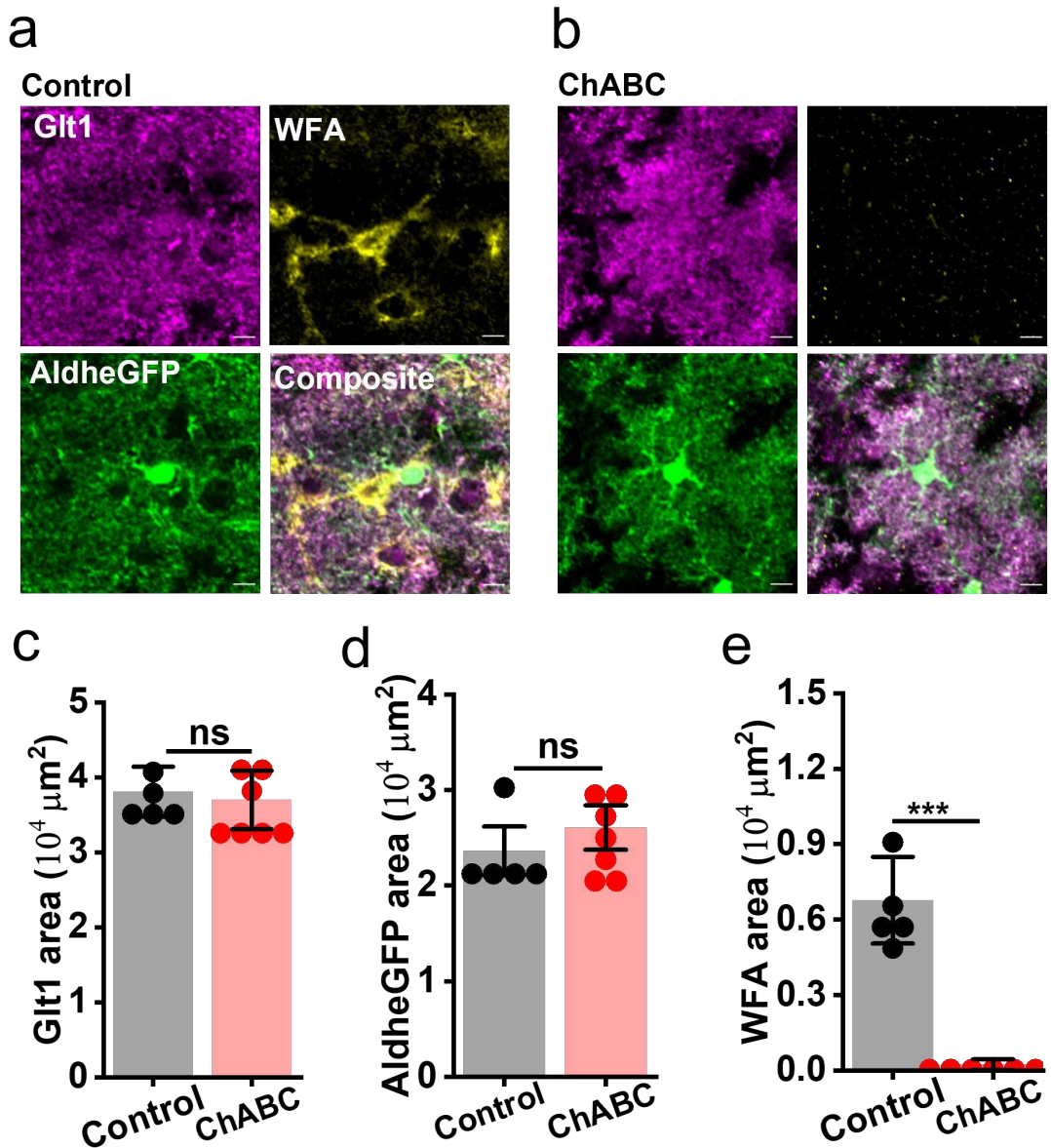
1349 **g** Representative voltage clamp traces, and IV plot showing the current-voltage  
1350 relationship of astrocytes in control (n = 59c/18m) and ChABC treated (n = 38c/8m) slices.

1351 **h** Representative voltage clamp traces of synaptically evoked currents in astrocytes in  
1352 presence of different blockers to isolate glutamate current.

1353 c and m represent the number of cells and mice respectively.

1354

1355



1356

1357 **Figure S6. Unaltered glutamate transporter expression on PNN depletion in acute**  
 1358 **brain slices.**

1359 **a - b** Confocal micrographs of Glt1 (magenta), aldheGFP (green), and WFA (yellow)  
 1360 fluorescence from fixed acute slices from control (**a**) and after ChABC treatment (**b**). Scale  
 1361 bar 10 $\mu\text{m}$ .

1362 **c-e** Bar diagrams of immunofluorescence area of, (**c**) Glt1 (control  $38077.65 \pm 3342.90$ ;  
 1363 ChABC  $36986.21 \pm 3914.77$ ), (**d**) AldheGFP (control  $23635.67 \pm 3772.71$ , ChABC  
 1364  $26070.47 \pm 4100.87$ ), and (**e**) WFA (control  $6769.05 \pm 1723.79$ , ChABC  $226.70 \pm 214.73$ )  
 1365 showing PNN disruption without any changes in astrocytic Glt1 expression. Control n =  
 1366 5s/3m; ChABC 7s/3m in c-e. Units,  $\mu\text{m}^2$  in c-e.

1367 s and m represent the number of slices and mice respectively. \*\*\*\*P < 0.0001, \*\*\*P <  
1368 0.001, \*\*P < 0.01, \*P < 0.05, ns = P > 0.05. unpaired two-tailed student t-test (equal  
1369 variance not assumed). Bar data are expressed as mean±SD; dots on the bars represent  
1370 the individual data points.

1371

1372

## 1373 **References:**

- 1374 1. Watanabe, K., et al., *Three-dimensional organization of the perivascular glial limiting membrane*  
1375 *and its relationship with the vasculature: a scanning electron microscope study.* Okajimas Folia  
1376 Anat.Jpn., 2010. **87**(3): p. 109-121.
- 1377 2. Araque, A., et al., *Tripartite synapses: glia, the unacknowledged partner.* Trends in neurosciences,  
1378 1999. **22**(5): p. 208-215.
- 1379 3. Asztely, F., G. Erdemli, and D.M. Kullmann, *Extrasynaptic glutamate spillover in the hippocampus:*  
1380 *dependence on temperature and the role of active glutamate uptake.* Neuron, 1997. **18**(2): p. 281-  
1381 293.
- 1382 4. Parsons, M.P. and L.A. Raymond, *Extrasynaptic NMDA receptor involvement in central nervous*  
1383 *system disorders.* Neuron, 2014. **82**(2): p. 279-293.
- 1384 5. Chaunsali, L., B.P. Tewari, and H. Sontheimer, *Perineuronal Net Dynamics in the Pathophysiology*  
1385 *of Epilepsy.* Epilepsy Currents, 2021. **21**(4): p. 273-281.
- 1386 6. Tewari, B.P., et al., *A glial perspective on the extracellular matrix and perineuronal net remodeling*  
1387 *in the central nervous system.* Frontiers in Cellular Neuroscience, 2022. **16**: p. 1022754.
- 1388 7. Frischknecht, R., et al., *Brain extracellular matrix affects AMPA receptor lateral mobility and short-*  
1389 *term synaptic plasticity.* Nature neuroscience, 2009. **12**(7): p. 897-904.
- 1390 8. Kochlamazashvili, G., et al., *The extracellular matrix molecule hyaluronic acid regulates*  
1391 *hippocampal synaptic plasticity by modulating postsynaptic L-type Ca<sup>2+</sup> channels.* Neuron, 2010.  
1392 **67**(1): p. 116-128.
- 1393 9. Favuzzi, E., et al., *Activity-dependent gating of parvalbumin interneuron function by the*  
1394 *perineuronal net protein brevican.* Neuron, 2017. **95**(3): p. 639-655. e10.
- 1395 10. Bal, M., et al., *Reelin mobilizes a VAMP7-dependent synaptic vesicle pool and selectively augments*  
1396 *spontaneous neurotransmission.* Neuron, 2013. **80**(4): p. 934-46.
- 1397 11. Wang, X.-b., et al., *Extracellular proteolysis by matrix metalloproteinase-9 drives dendritic spine*  
1398 *enlargement and long-term potentiation coordinately.* Proceedings of the National Academy of  
1399 Sciences, 2008. **105**(49): p. 19520-19525.
- 1400 12. Orlando, C., et al., *Perisynaptic chondroitin sulfate proteoglycans restrict structural plasticity in an*  
1401 *integrin-dependent manner.* Journal of Neuroscience, 2012. **32**(50): p. 18009-18017.
- 1402 13. Roszkowska, M., et al., *CD44: a novel synaptic cell adhesion molecule regulating structural and*  
1403 *functional plasticity of dendritic spines.* Mol Biol Cell, 2016. **27**(25): p. 4055-4066.
- 1404 14. de Vivo, L., et al., *Extracellular matrix inhibits structural and functional plasticity of dendritic spines*  
1405 *in the adult visual cortex.* Nature Communications, 2013. **4**(1): p. 1484.
- 1406 15. Sonntag, M., et al., *Synaptic coupling of inner ear sensory cells is controlled by brevican-based*  
1407 *extracellular matrix baskets resembling perineuronal nets.* BMC Biology, 2018. **16**(1): p. 99.
- 1408 16. Brückner, G., et al., *Extracellular matrix organization in various regions of rat brain grey matter.*  
1409 *Journal of neurocytology,* 1996. **25**(1): p. 333-346.
- 1410 17. Fawcett, J.W., T. Oohashi, and T. Pizzorusso, *The roles of perineuronal nets and the perinodal*  
1411 *extracellular matrix in neuronal function.* Nature Reviews Neuroscience, 2019. **20**(8): p. 451-465.

- 1412 18. Pizzorusso, T., et al., *Reactivation of ocular dominance plasticity in the adult visual cortex*. Science, 2002. **298**(5596): p. 1248-1251.
- 1413
- 1414 19. Torres-Ceja, B. and M.L. Olsen, *A closer look at astrocyte morphology: Development, heterogeneity, and plasticity at astrocyte leaflets*. Curr Opin Neurobiol, 2022. **74**: p. 102550.
- 1415
- 1416 20. Cahoy, J.D., et al., *A transcriptome database for astrocytes, neurons, and oligodendrocytes: a new resource for understanding brain development and function*. J Neurosci, 2008. **28**(1): p. 264-78.
- 1417
- 1418 21. Tewari, B.P. and H. Sontheimer, *Protocol to Quantitatively Assess the Structural Integrity of Perineuronal Nets ex vivo*. Bio-protocol, 2019. **9**(10): p. e3234-e3234.
- 1419
- 1420 22. Tewari, B.P., et al., *Perineuronal nets decrease membrane capacitance of peritumoral fast spiking interneurons in a model of epilepsy*. Nature Communications, 2018. **9**(1): p. 4724.
- 1421
- 1422 23. Carceller, H., et al., *Perineuronal Nets: Subtle Structures with Large Implications*. Neuroscientist, 2022: p. 10738584221106346.
- 1423
- 1424 24. Yang, Y., H. Higashimori, and L. Morel, *Developmental maturation of astrocytes and pathogenesis of neurodevelopmental disorders*. Journal of Neurodevelopmental Disorders, 2013. **5**(1): p. 22.
- 1425
- 1426 25. Bruckner, G., et al., *Postnatal development of perineuronal nets in wild-type mice and in a mutant deficient in tenascin-R*. J Comp Neurol, 2000. **428**(4): p. 616-29.
- 1427
- 1428 26. Carstens, K.E., et al., *Perineuronal nets suppress plasticity of excitatory synapses on CA2 pyramidal neurons*. Journal of Neuroscience, 2016. **36**(23): p. 6312-6320.
- 1429
- 1430 27. Rowlands, D., et al., *Aggrecan directs extracellular matrix-mediated neuronal plasticity*. Journal of Neuroscience, 2018. **38**(47): p. 10102-10113.
- 1431
- 1432 28. Carceller, H., et al., *Perineuronal Nets Regulate the Inhibitory Perisomatic Input onto Parvalbumin Interneurons and  $\gamma$  Activity in the Prefrontal Cortex*. Journal of Neuroscience, 2020. **40**(26): p. 5008-5018.
- 1433
- 1434
- 1435 29. Christensen, A.C., et al., *Perineuronal nets stabilize the grid cell network*. Nature Communications, 2021. **12**(1): p. 253.
- 1436
- 1437 30. Somaiya, R.D., et al., *Development of astrocyte morphology and function in mouse visual thalamus*. Journal of Comparative Neurology, 2022. **530**(7): p. 945-962.
- 1438
- 1439 31. Campbell, S.L., J.J. Hablitz, and M.L. Olsen, *Functional changes in glutamate transporters and astrocyte biophysical properties in a rodent model of focal cortical dysplasia*. Frontiers in Cellular Neuroscience, 2014. **8**(425).
- 1440
- 1441
- 1442 32. Bergles, D.E. and C.E. Jahr, *Synaptic Activation of Glutamate Transporters in Hippocampal Astrocytes*. Neuron, 1997. **19**(6): p. 1297-1308.
- 1443
- 1444 33. Campbell, S.C., et al., *Potassium and glutamate transport is impaired in scar-forming tumor-associated astrocytes*. Neurochemistry International, 2020. **133**: p. 104628.
- 1445
- 1446 34. Robel, S., et al., *Reactive astrogliosis causes the development of spontaneous seizures*. Journal of Neuroscience, 2015. **35**(8): p. 3330-3345.
- 1447
- 1448 35. Sorg, B.A., et al., *Casting a Wide Net: Role of Perineuronal Nets in Neural Plasticity*. J Neurosci, 2016. **36**(45): p. 11459-11468.
- 1449
- 1450 36. Yong, V.W., et al., *Metalloproteinases in biology and pathology of the nervous system*. Nature Reviews Neuroscience, 2001. **2**(7): p. 502.
- 1451
- 1452 37. Gottschall, P.E. and M.D. Howell, *ADAMTS expression and function in central nervous system injury and disorders*. Matrix Biology, 2015. **44**: p. 70-76.
- 1453
- 1454 38. Morawski, M., et al., *Ion exchanger in the brain: quantitative analysis of perineuronally fixed anionic binding sites suggests diffusion barriers with ion sorting properties*. Scientific reports, 2015. **5**.
- 1455
- 1456
- 1457 39. Dubey, D., et al., *Increased metalloproteinase activity in the hippocampus following status epilepticus*. Epilepsy Research, 2017. **132**: p. 50-58.
- 1458



- 1459 40. McRae, P.A. and B.E. Porter, *The perineuronal net component of the extracellular matrix in*  
1460 *plasticity and epilepsy*. *Neurochemistry international*, 2012. **61**(7): p. 963-972.
- 1461 41. Gonzalez-Burgos, G., K.N. Fish, and D.A. Lewis, *GABA Neuron Alterations, Cortical Circuit*  
1462 *Dysfunction and Cognitive Deficits in Schizophrenia*. *Neural Plasticity*, 2011. **2011**: p. 723184.
- 1463 42. Brückner, G., M. Morawski, and T. Arendt, *Aggrecan-based extracellular matrix is an integral part*  
1464 *of the human basal ganglia circuit*. *Neuroscience*, 2008. **151**(2): p. 489-504.
- 1465 43. Gittis, Aryn H., et al., *Rapid Target-Specific Remodeling of Fast-Spiking Inhibitory Circuits after Loss*  
1466 *of Dopamine*. *Neuron*, 2011. **71**(5): p. 858-868.
- 1467 44. Dityatev, A., C.I. Seidenbecher, and M. Schachner, *Compartmentalization from the outside: the*  
1468 *extracellular matrix and functional microdomains in the brain*. *Trends Neurosci*, 2010. **33**(11): p.  
1469 503-12.
- 1470 45. Dityatev, A. and D.A. Rusakov, *Molecular signals of plasticity at the tetrapartite synapse*. *Current*  
1471 *Opinion in Neurobiology*, 2011. **21**(2): p. 353-359.
- 1472 46. Freeman, M.R., *Specification and morphogenesis of astrocytes*. *Science*, 2010. **330**(6005): p. 774-  
1473 778.
- 1474 47. Hanson, E., N.C. Danbolt, and C.G. Dulla, *Astrocyte membrane properties are altered in a rat model*  
1475 *of developmental cortical malformation but single-cell astrocytic glutamate uptake is robust*.  
1476 *Neurobiol Dis*, 2016. **89**: p. 157-68.

1477

# Figures

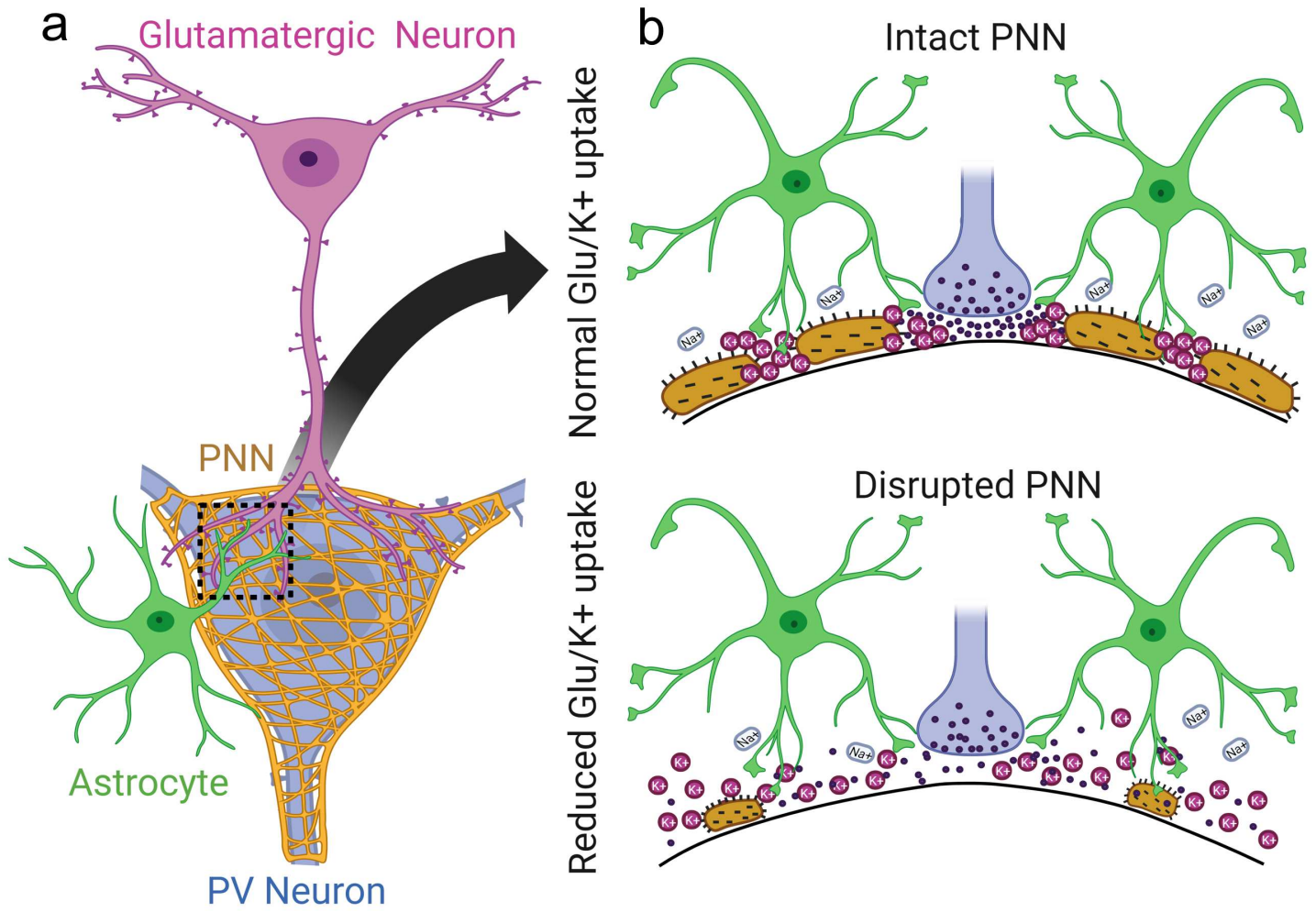


Figure 1

Figure 8

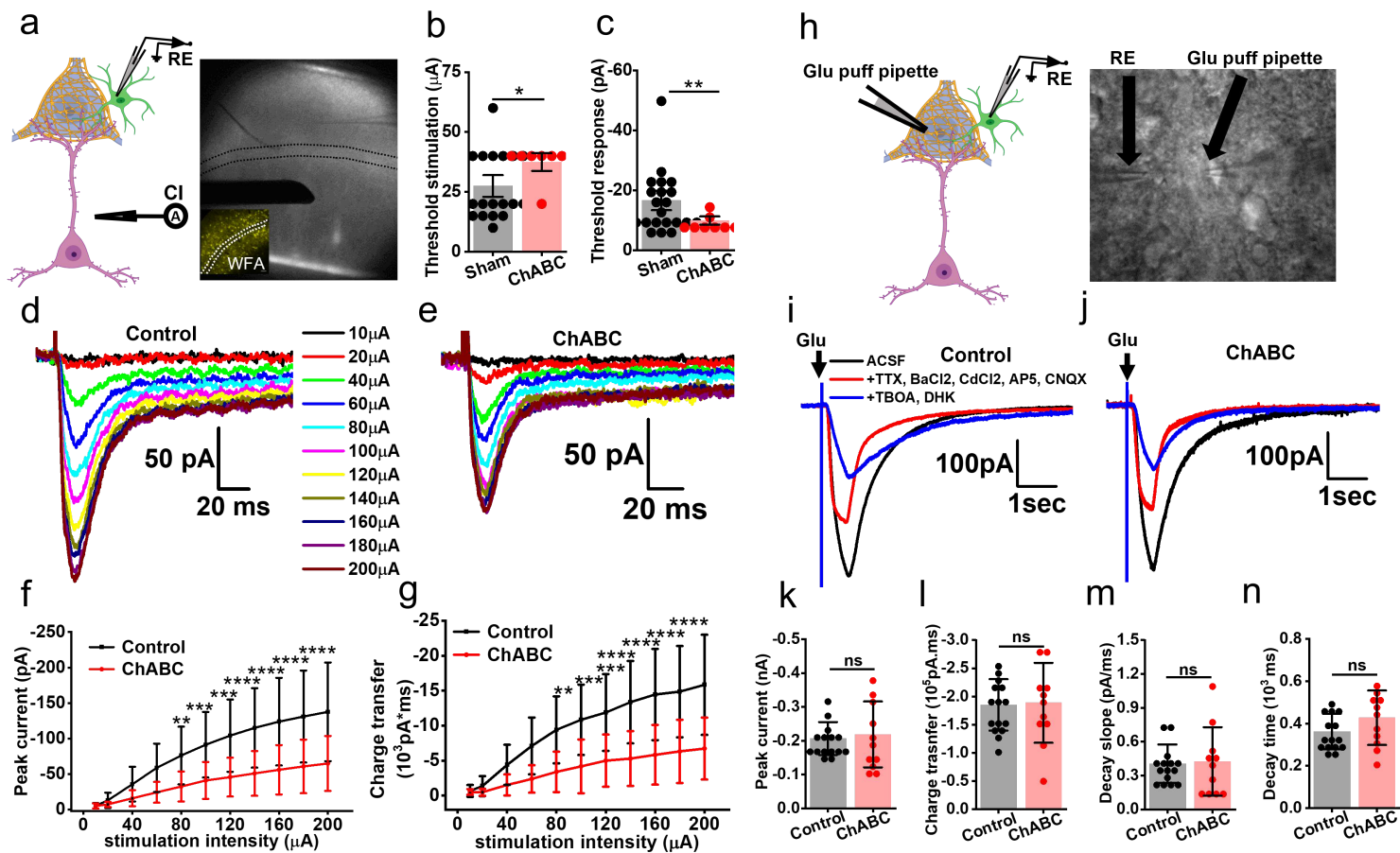


Figure 2

Figure 6

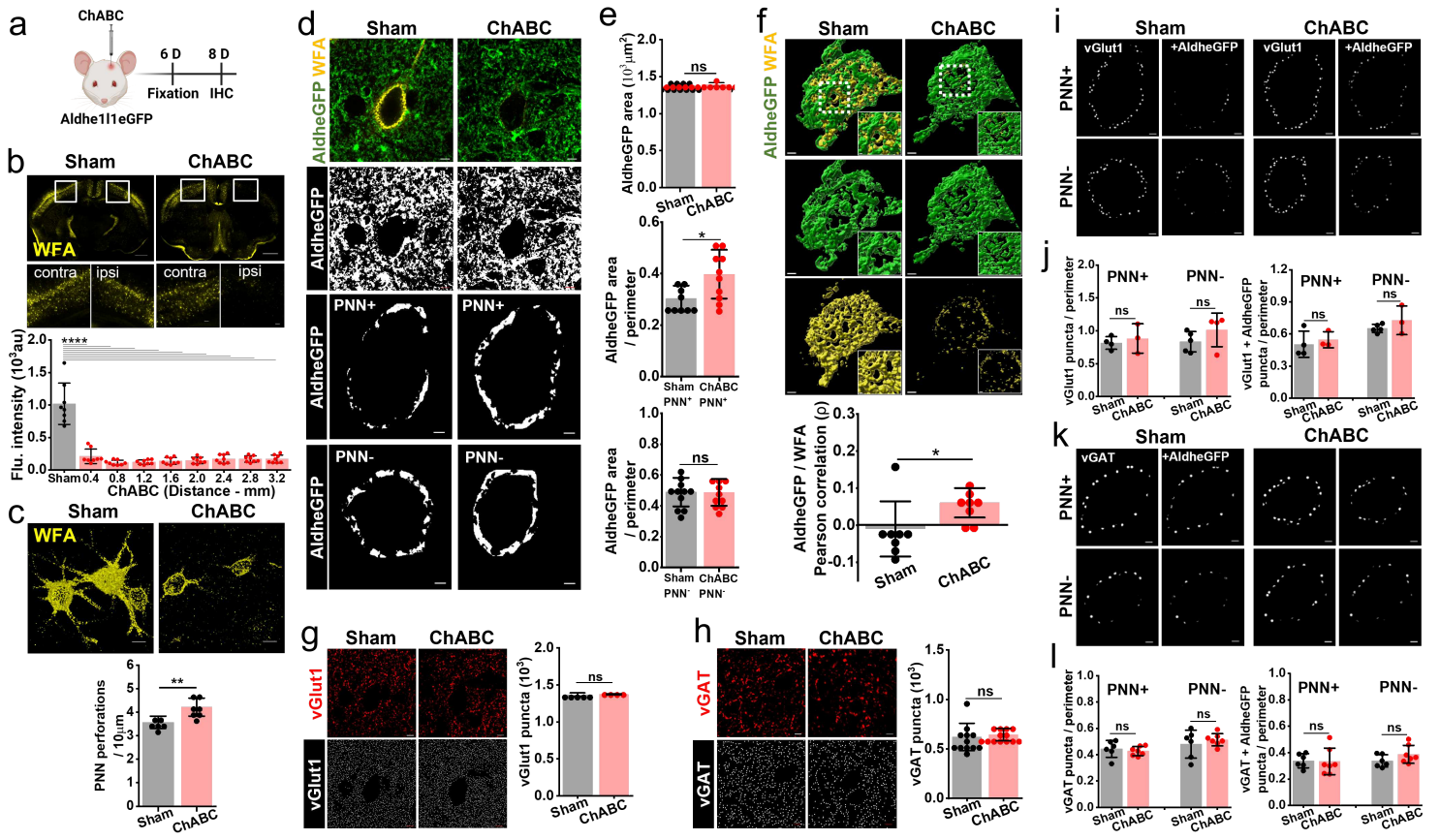


Figure 3

Figure 4

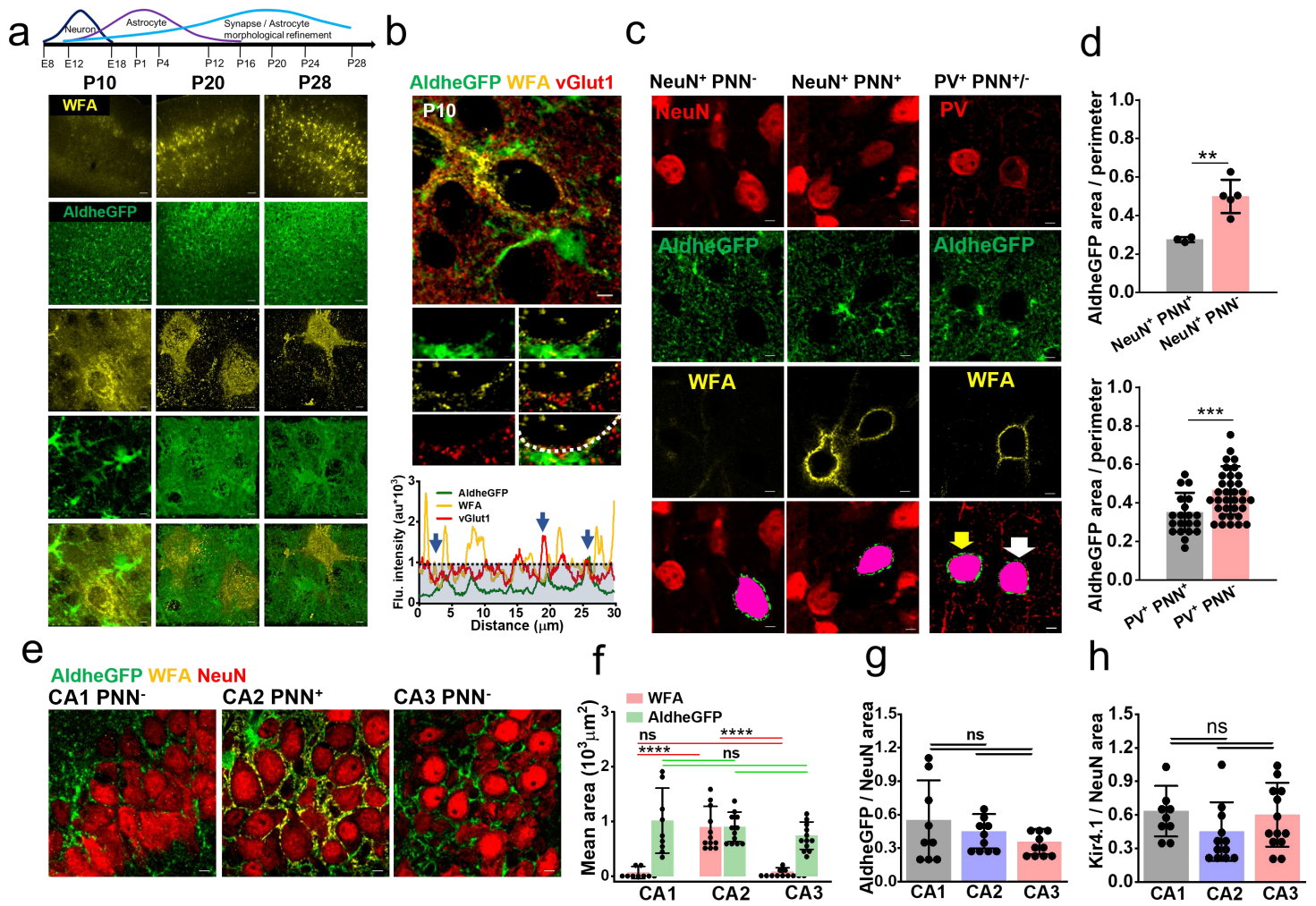


Figure 4

Figure 3



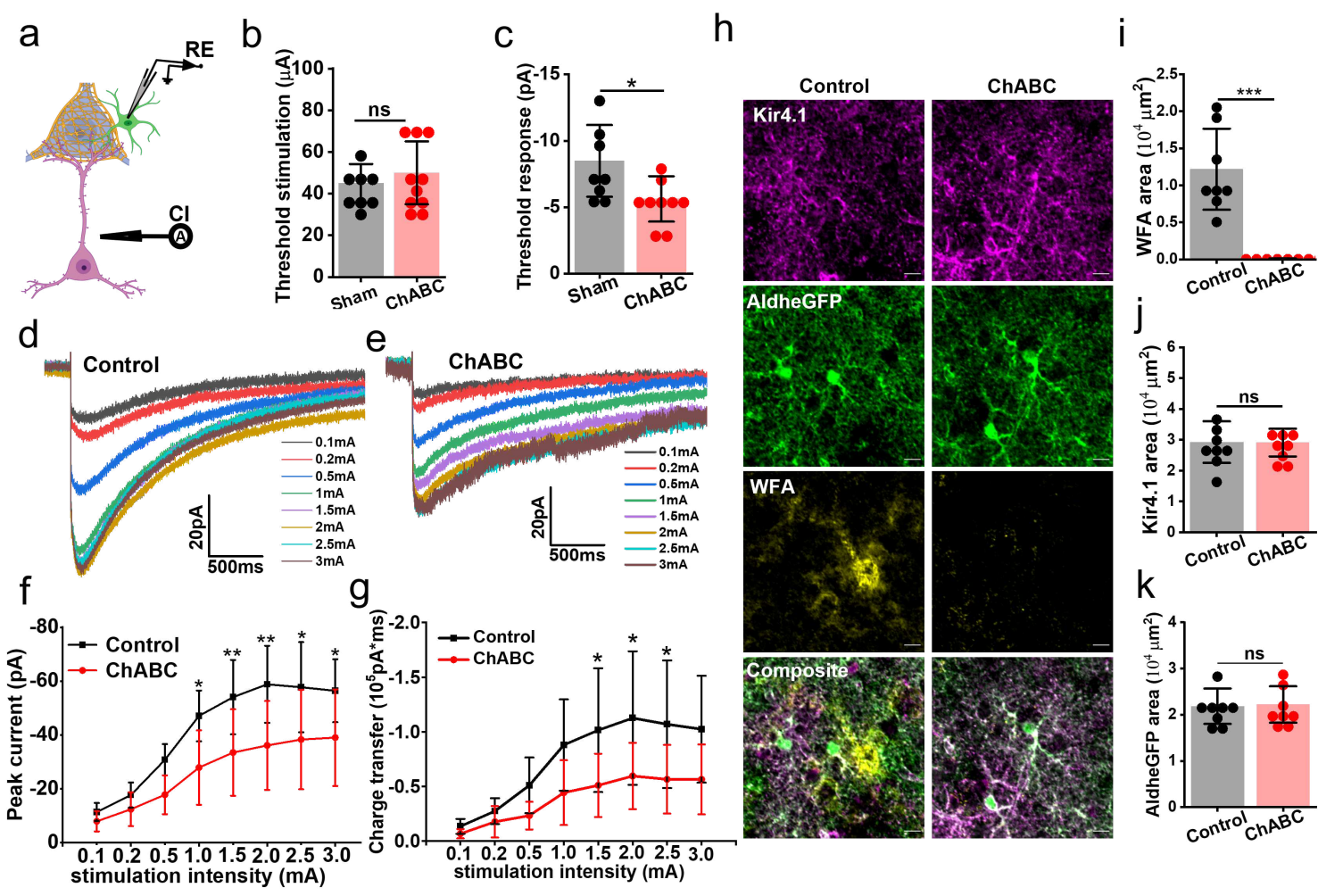


Figure 5

Figure 7

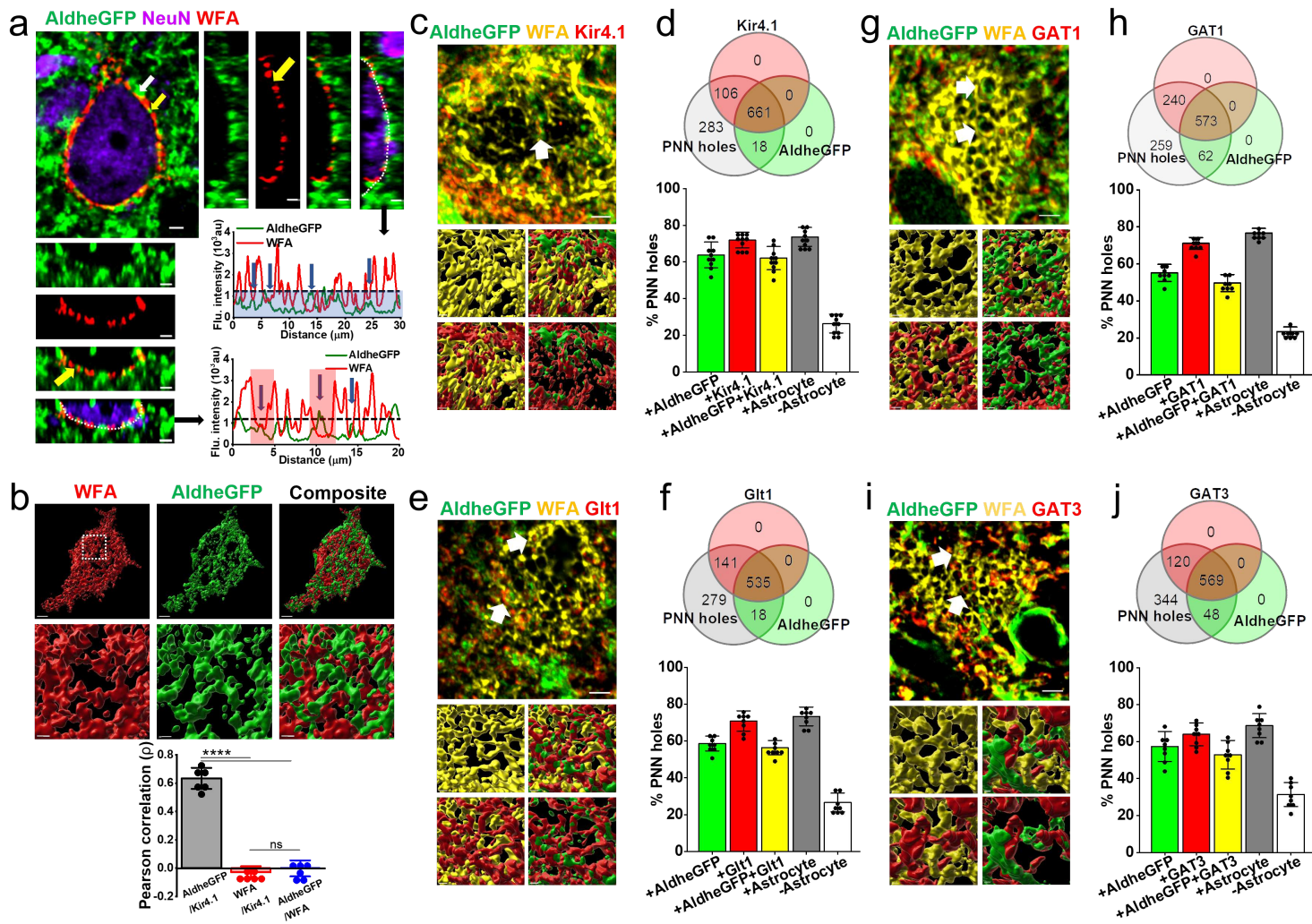


Figure 6

Figure 1

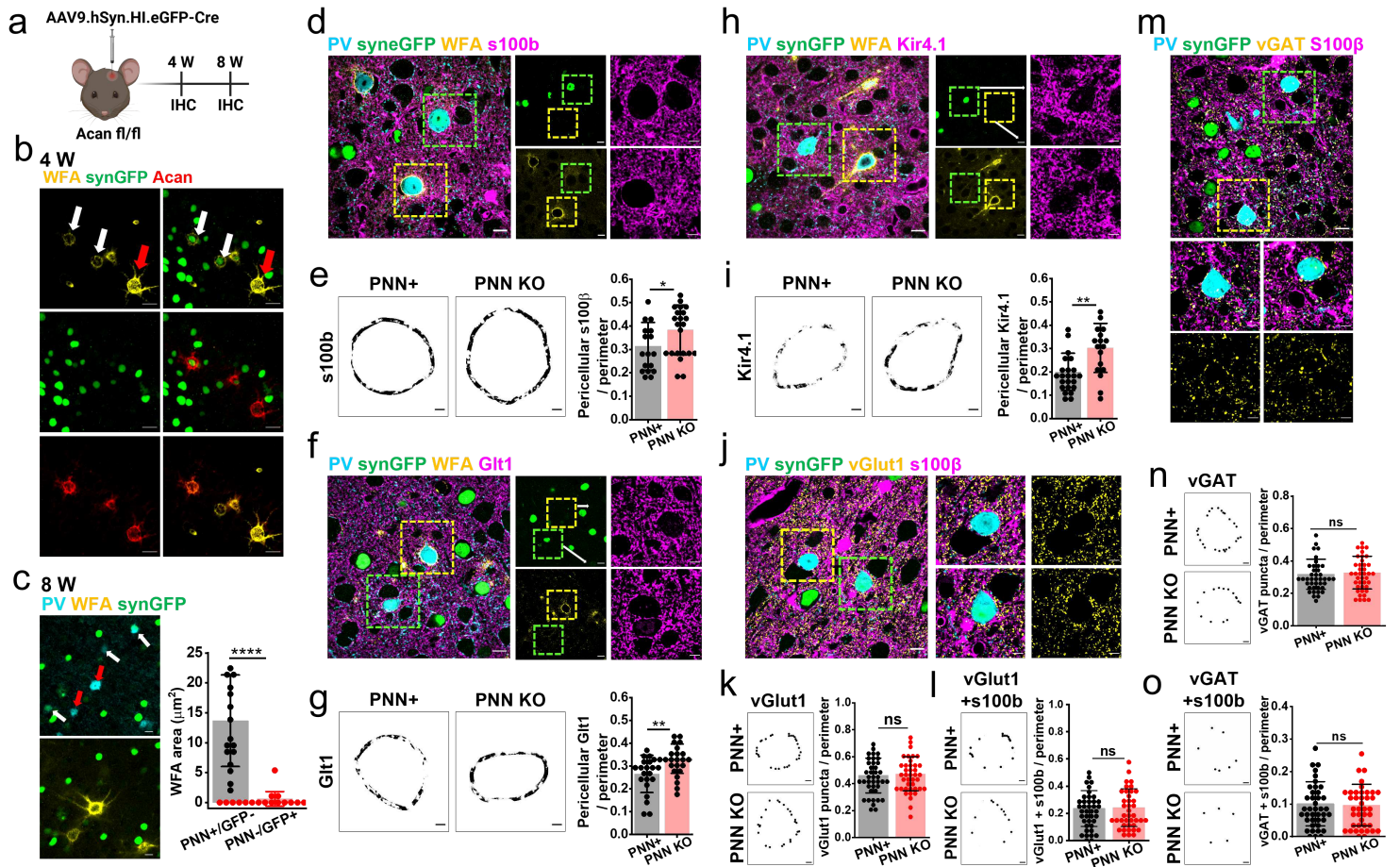


Figure 7

Figure 5



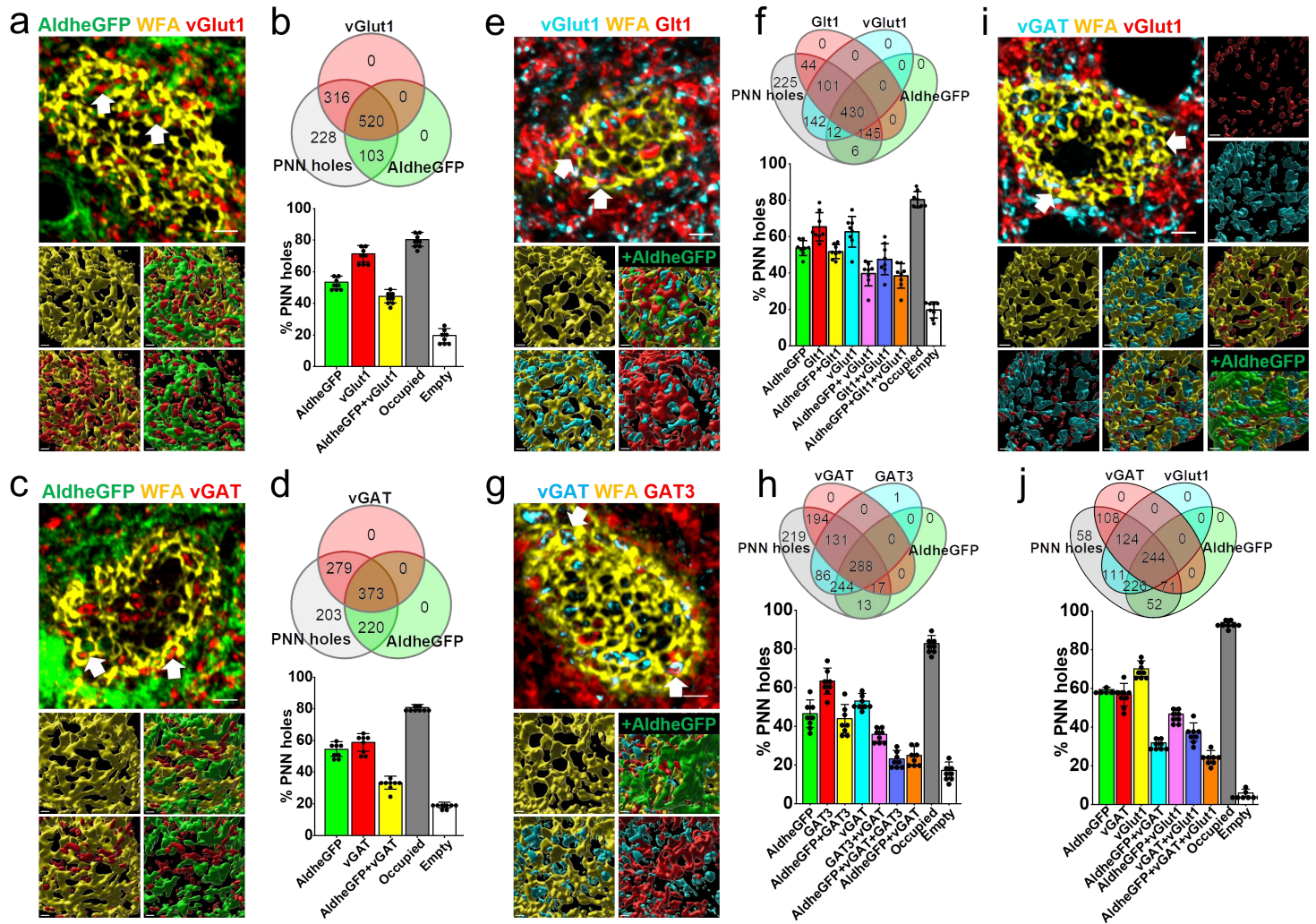


Figure 8

Figure 2

## Supplementary Files

This is a list of supplementary files associated with this preprint. Click to download.

- [Table1AntibodiesAstroPNN.pdf](#)
- [FigS6AstroPNN.tif](#)
- [FigS5AstroPNN.tif](#)
- [Table2Summaryofstatistics.xlsx](#)
- [FigS2AstroPNN.tif](#)
- [FigS1AstroPNN.tif](#)
- [FigS4AstroPNN.tif](#)
- [FigS3AstroPNN.tif](#)

- [FigS1AstroPNN.tif](#)
- [FigS2AstroPNN.tif](#)
- [FigS3AstroPNN.tif](#)
- [FigS4AstroPNN.tif](#)
- [FigS5AstroPNN.tif](#)
- [FigS6AstroPNN.tif](#)
- [Table1AntibodiesAstroPNN.pdf](#)
- [Table2Summaryofstatistics.xlsx](#)