

Research Articles: Cellular/Molecular

P2Y₆ receptor-dependent microglial phagocytosis of synapses during development regulates synapse density and memory

<https://doi.org/10.1523/JNEUROSCI.1089-23.2023>

Cite as: J. Neurosci 2023; 10.1523/JNEUROSCI.1089-23.2023

Received: 15 May 2023

Revised: 27 July 2023

Accepted: 31 August 2023

This Early Release article has been peer-reviewed and accepted, but has not been through the composition and copyediting processes. The final version may differ slightly in style or formatting and will contain links to any extended data.

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1 **P2Y₆ receptor-dependent microglial phagocytosis of synapses**
2 **during development regulates synapse density and memory**

3
4 Running title: P2Y₆ receptor and brain development

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14
15 Number of pages: 28

16 Number of figures: 12 (9 in main, 3 extended)

17
18 Abstract: 181 words

19 Introduction: 547 words

20 Discussion: 1,370 words

21
22 **Conflict of interest statement**

23 The authors declare no competing interests.

24
25 **Acknowledgments**

26 We thank Bernard Robaye for the P2Y₆R knockout mice, and Stefan Milde for
27 carrying out the behavioral tests on 9-12-month-old mice. The authors gratefully
28 acknowledge the Cambridge Advanced Imaging Centre for their support &
29 assistance in this work. This work was supported by: the Medical Research Council
30 UK [MR/L010593]; the Biotechnology and Biological Sciences Research Council
31 [BB/T508160/1]; and Eli Lilly [ARUK-DC2017-4 and RG99704].

32 **Abstract**

33 During brain development, excess synapses are pruned (i.e. removed), in part by
34 microglial phagocytosis, and dysregulated synaptic pruning can lead to behavioral
35 deficits. The P2Y₆ receptor (P2Y₆R) is known to regulate microglial phagocytosis of
36 neurons, and to regulate microglial phagocytosis of synapses in cell culture and *in*
37 *vivo* during aging. However, currently it is unknown whether P2Y₆R regulates
38 synaptic pruning during development. Here, we show that P2Y₆R knockout mice of
39 both sexes had strongly reduced microglial internalization of synaptic material,
40 measured as Vglut1 within CD68-staining lysosomes of microglia at postnatal day 30
41 (P30), suggesting reduced microglial phagocytosis of synapses. Consistent with this,
42 we found an increased density of synapses in the somatosensory cortex and the
43 CA3 region and dentate gyrus of the hippocampus at P30. We also show that adult
44 P2Y₆R knockout mice have impaired short- and long-term spatial memory and
45 impaired short- and long-term recognition memory compared to wild-type mice, as
46 measured by novel location recognition, novel object recognition, and Y-maze
47 memory tests. Overall, this indicates P2Y₆R regulates microglial phagocytosis of
48 synapses during development, and this contributes to memory capacity.

49

50 **Significance statement**

51 The P2Y₆ receptor (P2Y₆R) is activated by UDP released by neurons, inducing
52 microglial phagocytosis of such neurons or synapses. We tested whether P2Y₆R
53 regulates developmental synaptic pruning in mice and found that P2Y₆R knockout
54 mice have reduced synaptic material within microglial lysosomes, and increased
55 synaptic density in the brains of postnatal day 30 mice, consistent with reduced
56 synaptic pruning during development. We also found that adult P2Y₆R knockout
57 mice had reduced memory, consistent with persistent deficits in brain function,
58 resulting from impaired synaptic pruning. Overall, the results suggest that P2Y₆R
59 mediates microglial phagocytosis of synapses during development, and the absence
60 of this results in memory deficits in the adult.

61 **Introduction**

62 During development, synapse formation in the human brain is thought to peak at 1
63 million synapses per second (Tang et al., 2001). Excess synapses are generated
64 compared to what remains in adulthood, and some of these synapses are then
65 pruned to shape neuronal networks in an activity-dependent manner (Hong, Dissing-
66 Olesen and Stevens, 2016; Faust, Gunner and Schafer, 2021). This synaptic pruning
67 is by multiple mechanisms but is partly mediated by microglial phagocytosis of the
68 synapses (Stevens et al 2007; Tremblay, Lowery, and Majewska, 2010; Paolicelli et
69 al., 2011; Schafer et al., 2012; Mordelt and de Witte, 2023).

70

71 Microglia are macrophages, resident central nervous system, that play a variety of
72 roles in maintaining a healthy brain. One such role involves the phagocytosis (i.e.,
73 engulfment and degradation) of synapses, neurons, debris, bacteria, and aggregated
74 proteins (Sierra et al., 2013; Wolf, Boddeke and Kettenmann, 2017; Tay et al., 2018;
75 Gabandé-Rodríguez, Keane and Capasso, 2020). Microglial phagocytosis of
76 synapses is involved in learning and memory in adults (Miyanishi et al., 2021), but
77 excessive microglial phagocytosis of the synapses may contribute to memory loss
78 with aging and neurodegeneration (Hong et al., 2016; Dundee et al., 2023). Signals
79 regulating microglial phagocytosis of synapses include fractalkine and the fractalkine
80 receptor, complement components C1q, C3 and C4 and complement receptor 3,
81 adenosine 2A, TREM2, GPR56, phosphatidylserine, CD47, TGF β , and the P2Y₆
82 receptor (P2Y₆R) (Faust, Gunner and Schafer, 2021; Dundee et al., 2023).

83

84 P2Y₆R is a G-protein-coupled receptor for extracellular uridine diphosphate (UDP),
85 expressed from the *P2ry6* gene by myeloid and other cells in the body, but in the
86 brain, is almost exclusively expressed by microglia (Koizumi et al., 2007). Koizumi et
87 al. (2007) found that kainite-stressed neurons released uridine triphosphate, which
88 was hydrolyzed into UDP and induced microglia to phagocytose neurons via
89 activating P2Y₆R on microglia. We subsequently showed that extracellular UDP
90 could induce microglia to phagocytose live neurons, and that inhibition of P2Y₆R
91 prevented neuronal loss induced by LPS in glial-neuronal cultures and in mouse
92 brains *in vivo* (Neher et al., 2014; Milde et al., 2021). P2Y₆R knockout also prevented
93 microglial phagocytosis of neurons and loss of neurons and memory induced beta
94 amyloid and TAU *in vivo* (Puigdellívol et al., 2021).

95

96 Recently, we reported that P2Y₆R regulated microglial phagocytosis of synapses
97 (Dundee et al., 2023). In culture, we showed that P2Y₆R inhibition or knockout
98 strongly reduced microglial phagocytosis of isolated synapses (synaptosomes), and
99 that glial-neuronal cultures from P2Y₆R knockout mice had reduced inflammatory
100 loss of synapses (Dundee et al., 2023). *In vivo*, we found that P2Y₆R knockout
101 prevented an aging-associated increase in microglial phagocytosis of synapses, and
102 reduced loss of synapses and memory with age (Dundee et al., 2023).

103

104 Here, we investigated whether P2Y₆R regulated microglial phagocytosis of synapses
105 during development, and whether this affected synaptic density and memory. We
106 found that microglia from P2Y₆R knockout mice had reduced internalization of
107 synaptic material at postnatal day 30 compared to wild-type mice, coupled with an
108 increase in synaptic density in three different regions of P2Y₆R knockout mouse
109 brains. P2Y₆R knockout mice also performed worse than wild-type mice in both long-
110 and short-term memory tests. These findings are significant as they support the
111 hypothesis that microglial phagocytosis of synapses during development contributes
112 to a healthy brain and this is, in part, through activation of the P2Y₆R.

113 **Materials and Methods**

114

115 **Animals**

116

117 All animal work was carried out in accordance with the Animals (Scientific
118 Procedures) Act 1986 Amendment Regulations 2012 following ethical review by the
119 University of Cambridge Animal Welfare and Ethical Review Body (AWERB). P2Y₆R
120 knockout mice were kindly provided by Bernard Robaye (ULB Brussels) and
121 maintained on a C57BL/6 background (Charles River Laboratories). P2Y₆R knockout
122 mice and wild-type littermates were used to establish homozygous P2Y₆R wild-type
123 and knockout sub-lines. Postnatal day 30 (P30) mice included 4 male and 4 female
124 wild-type mice and 4 male and 4 female P2Y₆R knockout mice. 4-month-old mice
125 included 6 male and 5 female wild-type mice and 8 male and 3 female P2Y₆R
126 knockout mice. 9-12-month-old mice included 9 female wild-type mice and 7 female
127 P2Y₆R knockout mice. Mice utilized in short-term memory tests received
128 intracerebroventricular injections of 4 μ L PBS 3 or 14 days prior to testing.

129

130 **Fixation and tissue sectioning**

131

132 P30 mice were sacrificed following cervical dislocation and decapitation. Brains were
133 removed and fixed for 48 hours in 4% paraformaldehyde and cryoprotected by
134 immersion in an increased 10%–30% sucrose solution until sectioning. Serial coronal
135 sections (25 μ m) through the whole brain were collected using a sliding microtome
136 and placed in PBS with 0.025% sodium azide as free-floating sections.

137

138 **Immunohistochemistry of free-floating brain slices**

139

140 All steps were carried out at room temperature, with shaking, and rinsing thrice with
141 PBS after each incubation unless stated otherwise. Five to six free-floating 25 μ m
142 sections were taken every 12th brain section of 8 P2Y₆R wild-type and knockout
143 mice at P30 for immunohistochemistry. Sections were rinsed three times in PBS and
144 incubated with 50 mM ammonium chloride in PBS for 30 min to quench free
145 aldehyde groups from fixation. Sections were then incubated in 0.1% Sudan Black B

146 in 70% ethanol for 20 min to reduce autofluorescence, permeabilized using 1%
147 Triton X-100 in PBS for 30 min to facilitate antibody penetration, and blocked for 1 h
148 with blocking solution (2% bovine serum albumin, 3% goat serum, and 0.03% Triton
149 X-100 in PBS). Subsequently, sections were incubated with mouse anti-Vglut1
150 (1:200, Thermo Fisher, MA5-31373), rabbit anti-Homer1 (1:500, Synaptic Systems,
151 160003), rabbit anti-Iba1 (1:200, Wako, 019-19741), and rat CD68 (1:200, Thermo
152 Fisher, 14-0681-82) antibodies in blocking solution for 2 h at 37°C (Xiao et al., 2017).
153 Sections were then rinsed three times with PBS and then incubated with Alexa-Fluor
154 568 goat anti-mouse (1:200, Thermo Fisher, A5054), Alexa-Fluor goat 488 anti-
155 rabbit (1:200, Thermo Fisher, A11008), and Alexa-Fluor 647 goat anti-rat (1:200,
156 Thermo Fisher, A21247) antibodies for 2 h at 37°C. Sections were then rinsed three
157 times with PBS and mounted on poly-l-lysine-treated glass slides and dried at 37°C.
158 Sections were then mounted using Vectashield mounting medium with DAPI (Vector
159 Laboratories, H1500) and imaged using confocal microscopy.

160

161 **Synaptic internalization analysis of free-floating brain sections**

162

163 Imaging was carried out on a Nikon C2si confocal microscope with a 63×, 1.35 NA
164 oil immersion objective using 488, 561, and 640 nm laser lines. Microglia were
165 imaged and analyzed following Schafer et al. (2014). Briefly, Z-stacks (0.5 μm step
166 intervals) were collected being 2 μm from the surface of the section at each region of
167 interest. Fourteen to fifteen microglia were analyzed across three sections 300 μm
168 apart per mouse. Background subtraction (six pixels/0.2 μm rolling ball) and intensity
169 normalization (2%) across the sections was carried out using Fiji (Schindelin et al.,
170 2012). Microglial structures were analyzed using 3D Morph analysis (York et al.,
171 2018). Microglial (Iba1), lysosomal (CD68), and synaptic (Vglut1) surface rendering
172 was carried out using Imaris 9.1.2. The results for the surface-rendered objects were
173 represented as volume (μm³).

174

175 **Microglial density of free-floating brain sections**

176

177 Imaging was carried out on an EVOS M5000 fluorescent microscope with a 20×
178 objective. Nine images in the somatosensory cortex were taken across three

179 sections 300 μm apart per mouse. Total microglia (Iba1- and DAPI-positive) per field
180 were counted using Fiji (Schindelin et al., 2012).

181

182 **Synaptic density analysis of free-floating brain sections**

183

184 Imaging was carried out on a Nikon C2si confocal microscope with a 63 \times , 1.35 NA
185 oil immersion objective using 405, 488, and 561 nm laser lines. A 2- μm Z-stack
186 (0.125 μm step intervals) was collected being 2–5 μm from the surface of the section
187 at each region of interest. Nine images were taken across three sections 300 μm
188 apart per mouse. Background subtraction (six pixels/0.2 μm rolling ball) and intensity
189 normalization (2%) across the sections were carried out using Fiji (Schindelin et al.,
190 2012). We developed a custom script for Fiji to map Vglut1 and Homer1 puncta
191 positions in 3D and analyze their distributions. In order to detect puncta 0.2 μm in
192 diameter (Moreno Manrique et al., 2021), the script applies a Laplacian of Gaussian
193 filter with the standard deviation set from this estimated diameter and detects local
194 maxima as puncta candidates. Candidate points are then clustered into final puncta
195 by merging points within one punctum width of each other, excluding points on
196 image edges and below the global Otsu intensity threshold (Otsu et al., 1979). The
197 number of puncta “colocalized” between the two channels is counted as the number
198 of C2 (Homer1) coordinates having at least one C3 (Vglut1) coordinate within one
199 punctum width. Local density is calculated for each punctum using a Gaussian
200 kernel density estimate (Davis, Lii and Politis, 1956), and overall density in each
201 channel is calculated as puncta per μm^3 . The puncta densities were normalized to
202 the mean of wild-type mice.

203

204 **Long-term novel object location test and novel object recognition test**

205

206 To test long-term location and recognition memory, object location recognition
207 testing (OLRT) and novel object recognition testing (NORT) was performed in a
208 30 \times 44 cm arena with opaque sides, with a 24 h retention time (Murai et al., 2007;
209 Puigdemívol et al., 2021). Briefly, 4-month-old mice were first habituated to the arena
210 in the absence of objects on two consecutive days (15 min/day), when spontaneous
211 locomotor activity (total distance traveled) and anxiety/motivation (distance traveled
212 in periphery versus center of the open field) were measured. On the third day, visual

213 clues were included on the walls of the open field and two similar objects were
214 presented for 10 min (A and A' objects). Twenty-four hours later, the same animals
215 were retested for 5 min retention test in the arena with a familiar (A) and a new (B)
216 object location. The object location preference was measured as the time exploring
217 each object \times 100/time exploring both objects. On the fifth day, visual clues were
218 removed and two new identical objects were presented to the animals for 10 min (C
219 and C'). Twenty-four hours later, the same animals were retested for 5 min retention
220 test in the arena with a familiar (C) and a new (D) object. The object preference was
221 measured as the time exploring each object \times 100/time exploring both objects.
222 Animals were tracked and recorded with SMART Junior software (Panlab). Objects
223 and arena were cleaned thoroughly with 70% ethanol and dried after each trial to
224 eliminate odor cues. Experimenter was blinded to the genotype of the individual
225 animals. NORT data was reanalyzed from Dundee et al. (2023).

226

227 **Short-term novel object recognition memory test**

228

229 To test short-term recognition memory, novel object recognition testing (NORT) was
230 performed in a 30 \times 44 cm arena with opaque sides, with a 2 min retention time
231 (Puigdemívol et al., 2021). Briefly, 9-12-month-old mice were first habituated to the
232 arena in the absence of objects for two 10 min sessions 2 h apart. The next day, two
233 identical objects were presented for 10 min (A and A' objects). 2 minutes later, the
234 same animals were retested for 5 min in the arena with a familiar (A) and a novel (B)
235 object. The order of testing of mice from different experimental groups was
236 randomized on day 1 and maintained in the same order on day 2. Object interaction
237 times and ratios were extracted from digital recordings of the trials using modified
238 "Autotyping" software. Experimenter was blinded to the genotype of the individual
239 animals. This data was reanalyzed from Puigdemívol et al. (2021).

240

241 **Y-maze memory test**

242

243 To test short-term spatial memory, spontaneous alternation performance in a Y-
244 maze was tested as described previously (Prieur and Jadavji, 2019). Briefly, 9-12-
245 month-old mice were placed in the center of the symmetrical Y-maze and were
246 allowed to explore freely through the maze during an 8 min session. The sequence,

247 total number of arms entered, and total distance travelled was recorded.
248 Spontaneous alternations (%) are as follows: number of arm entries that were not
249 identical to the previous arm entry/total number of arm entries \times 100. Arm entries
250 and distance travelled were extracted from digital recordings of the trials using
251 modified “Autotyping” software. Experimenter was blinded to the genotype of the
252 individual animals.

253

254 **Statistical analysis**

255

256 Bars represent mean \pm SEM, and each data point represents one animal (Fig. 1d-f,
257 2b,d,e-g, 3b, 4d, 5d, 6d, 7d, 8b,c,d,e,f, 9b-d,f,g, and Extended Fig. 4-1a-l, 4-2).
258 Statistical differences were calculated using unpaired *t* tests and one-sample *t* tests.
259 Statistical correlation was carried out using paired Pearson correlation coefficients.
260 All experiments were analyzed using GraphPad Prism 9 (GraphPad software).
261 Graphical data were shown as individual data points, including mean values with
262 error bars indicating SEM. *p*-values of $*p < 0.05$, $**p < 0.01$ indicated significant
263 differences between groups. For each experiment and graph, statistical details
264 including the statistical test used, the exact value of *n*, what *n* represents (number of
265 animals per genotype) as well as dispersion and precision measures (mean, SEM,
266 etc.) can be found in each figure legend.

267 **Results**

268

269 **P2Y₆R knockout results in reduced synaptic internalization during**
270 **development**

271

272 During development, synaptic levels are regulated in part by the phagocytosis of
273 synapses by microglia (Schafer et al., 2012). We have previously shown that P2Y₆R
274 knockout microglia have reduced synaptic internalization during aging (Dundee et
275 al., 2023). To test whether P2Y₆R is involved in the microglial phagocytosis of
276 synapses during development, we analyzed the internalization of synaptic material
277 within microglial lysosomes in the somatosensory cortex of P30 wild-type and
278 knockout mice by confocal microscopy (Fig. 1a). Coronal brain slices were
279 immunostained using antibodies to Iba1 (microglial marker), CD68 (lysosomal
280 marker), and Vglut1 (pre-synaptic marker), and internalized Vglut1 volume within
281 lysosomal microglia was analyzed by generating Z-projection surface renderings
282 using Imaris software (Fig. 1b,c). We chose a pre-synaptic marker (Vglut1), rather
283 than a post-synaptic marker, as there is clearer evidence for microglial phagocytosis
284 of pre-synaptic markers during development (Weinhard et al., 2018; Mordelt and de
285 Witte, 2023). We chose to examine synapse phagocytosis at P30 because synaptic
286 pruning has been reported to still be actively occurring in somatosensory cortex of
287 mice at this timepoint (Cong et al., 2020). We chose the somatosensory cortex
288 because we have previously found P2Y₆R affects synaptic density in this area during
289 aging (Dundee et al., 2023) and the somatosensory cortex affects working memory
290 (Long and Zhang, 2021).

291

292 There was a small but significant increase in microglial Iba1 volume in the P2Y₆R
293 knockout mice compared to the wild-type mice (Fig. 1d, n=7-8, unpaired t-test, WT:
294 282, KO: 332, p=0.024, $\eta^2=0.269$), however there were no other observable effects
295 on morphology (York et al., 2018), such as the number of branches (n=8, unpaired t-
296 test, WT: 63, KO: 60, p=0.376, $\eta^2=0.007$), the number of branch points (n=8,
297 unpaired t-test, WT: 52, KO: 50, p=0.391, $\eta^2=0.006$), the average branch length
298 (n=8, unpaired t-test, WT: 32, KO: 33 p=0.327, $\eta^2=0.015$), or the maximum (n=8,
299 unpaired t-test, WT: 70, KO: 71, p=0.393, $\eta^2=0.005$) and minimum (n=8, unpaired t-

300 test, WT: 4.1, KO: 4.4, $p=0.178$, $\eta^2=0.061$) branch length (Fig. 2). Microglial density
301 was subtly but significantly reduced in P2Y₆R knockout mice suggesting a small
302 effect on microglial proliferation at a young age (Fig. 3, $n=8$, unpaired t-test, WT: 54,
303 KO: 49, $p=0.007$, $\eta^2=0.357$). There were no significant differences in microglial CD68
304 volume between wild-type and knockout mice (Fig. 1e, $n=7-8$, unpaired t-test, WT:
305 4.2, KO: 3.4, $p=0.210$, $\eta^2=0.051$), indicating no difference in microglial lysosomal
306 volume. Note that virtually all of the CD68 staining was within Iba1-stained microglia
307 (Extended Fig. 1-1), indicating that this marker of phagocytic lysosomes was
308 relatively specific to microglia. Importantly, the volume of Vglut1 internalized within
309 microglial CD68 was strongly and significantly lower in the knockout mice, indicating
310 that the microglial phagocytosis of synapses is strongly reduced in knockout mice
311 (Fig. 1f, $n=7-8$, unpaired t-test, WT: 0.028, KO: 0.007, $p=0.001$, $\eta^2=0.506$). This
312 suggests that P2Y₆R mediates the phagocytosis of synapses during development.

313

314 **P2Y₆R knockout results in increased synaptic density during development**

315

316 In order to investigate whether this decrease in microglial phagocytosis of synapses
317 had an effect on synapse density during development, we examined the synaptic
318 density of wild-type and P2Y₆R knockout mice at P30. Synaptic density was
319 analyzed in the somatosensory cortex, the hippocampal CA1 and CA3 stratum
320 radiatum, and the dentate gyrus molecular layer of P30 P2Y₆R wild-type and
321 knockout mice using confocal microscopy (Fig. 4a, 5a, 6a, and 7a). Coronal brain
322 slices were immunostained using antibodies to Vglut1 (pre-synaptic marker) and
323 Homer1 (post-synaptic marker), and synaptic density was measured as colocalized
324 (<200 nm) puncta of both synaptic markers in an entire Z-stack.

325

326 In the somatosensory cortex, the density of Vglut1 puncta ($n=8$, unpaired t-test, WT:
327 100, KO: 105, $p=0.030$, $\eta^2=0.229$) and the colocalization of Vglut1 and Homer1
328 puncta ($n=8$, unpaired t-test, WT: 100, KO: 109, $p=0.039$, $\eta^2=0.206$) was significantly
329 increased in knockout mice compared to wild-type mice, with no difference in
330 Homer1 puncta density (Fig. 4b-d, $n=8$, unpaired t-test, WT: 100, KO: 101, $p=0.418$,
331 $\eta^2=0.003$). No difference in density was observed in the hippocampal CA1 stratum
332 radiatum for Vglut1 puncta ($n=8$, unpaired t-test, WT: 100, KO: 98, $p=0.335$,

333 $\eta^2=0.013$), Homer1 puncta (n=8, unpaired t-test, WT: 100, KO: 102, p=0.215,
334 $\eta^2=0.045$), or the colocalization of Vglut1 and Homer1 puncta (Fig. 5b-d, n=8,
335 unpaired t-test, WT: 100, KO: 101, p=0.394, $\eta^2=0.005$). In the hippocampal CA3
336 stratum radiatum, the density of Vglut1 puncta (n=8, unpaired t-test, WT: 100, KO:
337 108, p=0.038, $\eta^2=0.208$) and the colocalization of Vglut1 and Homer1 puncta (n=8,
338 unpaired t-test, WT: 100, KO: 115, p=0.048, $\eta^2=0.184$) was significantly increased in
339 knockout mice compared to wild-type mice, with no difference in Homer1 puncta
340 (Fig. 6b-d, n=8, unpaired t-test, WT: 100, KO: 108, p=0.092, $\eta^2=0.123$). In the
341 molecular layer of the dentate gyrus, the density of Homer1 puncta (n=8, unpaired t-
342 test, WT: 100, KO: 114, p=0.005, $\eta^2=0.388$) and the colocalization of Vglut1 and
343 Homer1 puncta (n=8, unpaired t-test, WT: 100, KO: 113, p=0.027, $\eta^2=0.241$) was
344 significantly increased in knockout mice compared to wild-type mice, with no
345 difference in Vglut1 puncta (Fig. 7b-d, n=8, unpaired t-test, WT: 100, KO: 100,
346 p=0.485, $\eta^2=0.0001$). We did not examine synapse density in any other areas of the
347 brain. Raw values for the Vglut1 puncta density, Homer1 puncta density, and the
348 colocalization density are visible in Extended Figure 4-1. Furthermore, the Pearson
349 correlation coefficient was calculated comparing the internalization of synaptic
350 material per mouse in the somatosensory cortex to their corresponding synaptic
351 density. We found a significant, negative correlation between the volume of
352 internalized synaptic material by microglia and the synaptic density in the
353 somatosensory cortex (Extended Fig. 4-2, n=15, Pearson correlation coefficient, r=-
354 0.532, p=0.021). Altogether, these data indicate that P2Y₆R knockout increases
355 synapse density and decreases microglial phagocytosis of synapses during
356 development, and the data are consistent with P2Y₆R-mediated microglial
357 phagocytosis of synapses causing synaptic pruning during development.

358

359 **P2Y₆R knockout results in reduced memory**

360

361 In the previous section, we found that P2Y₆R knockout increased synapse density in
362 the somatosensory cortex and hippocampus, regions that are associated with
363 various forms memory (Rebola, Carta and Mulle, 2017; Hainmueller and Bartos,
364 2020; Long and Zhang, 2021). Dysregulated synaptic pruning during development
365 has been associated with memory deficits (Wang et al., 2022). And we have

366 previously observed a role for P2Y₆R-dependent memory loss in aging, and both an
367 acute amyloid model and chronic tau model of neurodegeneration (Puigdemívol et al.,
368 2021; Dundee et al., 2023).

369

370 To investigate whether the reduced synaptic pruning in P2Y₆R knockout mice had a
371 long-term effect on memory, we tested the memory of adult wild-type and P2Y₆R
372 knockout mice using the object location recognition test (OLRT), the novel object
373 recognition test (NORT), and the Y-maze. The NORT tests object recognition
374 memory (Ennaceur and Delacour, 1988) whilst the OLRT and Y-maze test spatial
375 memory (Vogel-Ciernia and Wood, 2014; Kraeuter, Guest and Sarnyai, 2019). We
376 first tested long-term spatial memory using the OLRT. 4-month-old mice were
377 habituated in an open field for two consecutive days, and on the third day, mice were
378 allowed to explore two identical objects for 10 min in the presence of visual clues on
379 the walls of the open field. 24 h after this training, one of the objects remained in the
380 same location (familiar) whilst the other was moved (novel), and the relative time
381 spent with novel and familiar objects was quantified (Fig. 8a). During the training
382 session, wild-type mice had no overall preference for the objects at either location
383 (n=11, one-sample t-test to 50%, WT: 52, p=0.356, $\eta^2=0.086$) whilst P2Y₆R knockout
384 mice had a slight object preference (Fig. 8b, n=11, one-sample t-test to 50%, KO: 54,
385 p=0.015, $\eta^2=0.459$). 24 h after training, wild-type mice spent more time with the
386 object at the novel location, meaning they had long-term memory of the object
387 locations (Fig. 8c, n=11, one-sample t-test to 50%, WT: 72, p=0.005, $\eta^2=0.556$).
388 However, P2Y₆R knockout mice did not show a preference for either object location
389 (n=11, one-sample t-test to 50%, KO: 55, p=0.305, $\eta^2=0.105$) and showed reduced
390 preference compared to wild-type mice (n=11, unpaired t-test, WT: 72, KO: 55,
391 p=0.023, $\eta^2=0.186$), indicating that knockout mice have impaired long-term spatial
392 memory.

393

394 Using the same animals, we then tested long-term recognition memory using the
395 Novel Object Recognition Test (NORT). The animals had previously been
396 habituated in the open field (day 1 and 2) followed by the OLT (day 3 and 4). Visual
397 clues were removed from the walls of the open field and on the fifth day mice were
398 allowed to explore two new identical objects for 10 min. 24 h after training, one of the

399 objects (familiar) was replaced by a new object (novel), and the relative time spent
400 with novel and familiar objects was quantified (Fig. 8d). During the training session,
401 both wild-type (n=11, one-sample t-test to 50%, WT: 56, $p=0.033$, $\eta^2=0.379$) and
402 P2Y₆R knockout (n=11, one-sample t-test to 50%, KO: 55, $p=0.049$, $\eta^2=0.335$) mice
403 had a slight preference for one object (Fig. 8e). During the test, both wild-type (n=11,
404 one-sample t-test to 50%, WT: 74, $p=0.0002$, $\eta^2=0.770$) and P2Y₆R knockout (n=11,
405 one-sample t-test to 50%, KO: 64, $p=0.005$, $\eta^2=0.564$) mice spent noticeably more
406 time with the novel object, meaning they had long-term memory of the familiar
407 object. However, P2Y₆R knockout mice showed a significantly reduced preference
408 for the novel object compared to wild-type mice (Fig. 8f, n=11, unpaired t-test, WT:
409 74, KO: 64, $p=0.042$, $\eta^2=0.142$), indicating that knockout mice have impaired long-
410 term recognition memory. These data suggest that long-term memory is impaired in
411 P2Y₆R knockout mice.

412

413 We tested short-term spatial memory of 9-12-month old mice using the Y-maze,
414 where mice were placed in the center of the symmetrical Y-maze and were allowed
415 to explore freely for 8 min (Fig. 9a). The sequence, total number of arms entered,
416 and distance traveled was recorded to determine the spontaneous alterations, which
417 is a measure of short-term spatial memory. There was a non-significant increase in
418 the number of total arm entries (Fig. 9b, n=7-9, unpaired t-test, WT: 34, KO: 46,
419 $p=0.064$, $\eta^2=0.157$), and a significant increase in total distance travelled for P2Y₆R
420 knockout mice compared to wild-type mice (Fig. 9c, n=7-9, unpaired t-test, WT: 675,
421 KO: 901, $p=0.027$, $\eta^2=0.239$), suggesting mildly increased motility of P2Y₆R
422 knockout mice. Both wild-type (n=9, one-sample t-test to 50%, WT: 72, $p=0.0003$,
423 $\eta^2=0.819$) and P2Y₆R knockout (n=7, one-sample t-test to 50%, KO: 60, $p=0.010$,
424 $\eta^2=0.691$) mice carried out spontaneous alterations in the Y-maze, meaning both
425 groups of mice had short-term spatial memory. However, there was a significant
426 decrease in spontaneous alterations for P2Y₆R knockout mice compared to wild-type
427 mice, reduced from 72% in wild-type mice to 60% in P2Y₆R knockout mice, where
428 50% indicates random arm entry and zero memory (Fig. 9d, n=7-9, unpaired t-test,
429 WT: 72, KO: 60, $p=0.013$, $\eta^2=0.304$). This data indicates that P2Y₆R knockout mice
430 have impaired short-term spatial memory.

431

432 Finally, we tested short-term recognition memory of 9-12-month-old mice using the
433 NORT. Mice were habituated in an open field for two consecutive days, and on the
434 third day, mice were allowed to explore two identical objects for 10 min. 5 min after
435 training, one of the objects (familiar) was replaced by a new object (novel), and the
436 relative time spent with the novel and familiar objects was quantified (Fig. 9e). During
437 the training session, both wild-type (n=7, one-sample t-test to 50%, WT: 50, p=0.919,
438 $\eta^2=0.002$) and P2Y₆R knockout (n=7, one-sample t-test to 50%, KO: 52, p=0.638,
439 $\eta^2=0.039$) mice had no overall preference for either object (Fig. 9f). By contrast,
440 during the test, both wild-type (n=7, one-sample t-test to 50%, WT: 72, p=0.001,
441 $\eta^2=0.860$) and P2Y₆R knockout (n=7, one-sample t-test to 50%, KO: 62, p=0.112,
442 $\eta^2=0.685$) mice spent more time with the novel object, meaning they had short-term
443 memory of the familiar object. However, P2Y₆R knockout mice showed a significantly
444 reduced preference for the novel object compared to wild-type mice (Fig. 9g, n=7,
445 unpaired t-test, WT: 72, KO: 63, p=0.044, $\eta^2=0.222$), indicating that knockout mice
446 have impaired short-term recognition memory. Overall, our data indicates that adult
447 P2Y₆R knockout mice have reduced memory, consistent with the reduced synaptic
448 pruning during development resulting in long-term deficits in brain function.

449 **Discussion**

450

451 We had previously found that the P2Y₆ receptor (P2Y₆R) regulates microglial
452 phagocytosis of synapses during aging in mice, by measuring microglial
453 phagocytosis of isolated synapses, synaptic loss in culture, and synaptic
454 internalization and loss *in vivo* (Dundee et al., 2023). We were therefore interested
455 here in whether P2Y₆R regulates microglial phagocytosis of synapses during
456 development in mice, and if so, whether this affects adult brain function. We found
457 here that P2Y₆R knockout mice had greatly reduced synaptic material within
458 microglial lysosomes, and increased synaptic density at postnatal day 30 (P30),
459 consistent with P2Y₆R regulating synaptic pruning. We found that adult P2Y₆R
460 knockout mice had impaired long- and short-term memory, consistent with reduced
461 synaptic pruning causing memory dysfunction. Thus, it appears that P2Y₆R mediates
462 synaptic pruning via regulating microglial phagocytosis of synapses during
463 development, and if this does not occur, adult brain dysfunction results. However,
464 there are a number of uncertainties and limitations of our study that are outlined
465 below.

466

467 We measured microglial phagocytosis of synapses as the internalization of the pre-
468 synaptic marker Vglut1 into the CD68-stained lysosomes of Iba1-stained microglia in
469 fixed sections of the somatosensory cortex of wild-type and P2Y₆R knockout mice at
470 P30. We found a large reduction in Vglut1 within microglial lysosomes in the
471 knockout mice, consistent with reduced phagocytosis. This is consistent with
472 previous research identifying a role for microglial phagocytosis in developmental
473 synaptic pruning (Tremblay, Lowery, and Majewska, 2010; Paolicelli et al., 2011).
474 However, we did not visualize/image phagocytosis over time, and it is conceivable
475 that the reduced Vglut1 within microglial lysosomes in the knockout mice was due to
476 increased digestion of the synaptic material within the knockout lysosomes. To
477 directly test this possibility, in our previous publication, we fed isolated synapses
478 (synaptosomes) to immortalized microglia and measured the rate of Vglut1
479 degradation with and without the P2Y₆R inhibitor MRS2578 and found no difference
480 in Vglut1 degradation rate (Dundee et al., 2023). Moreover, we showed that
481 microglia isolated from P2Y₆R knockout mice had a reduced rate of microglial

482 phagocytosis of the isolated synapses (Dundee et al., 2023). Thus, the reduced
483 Vglut1 within microglial lysosomes in the knockout mice found here is most likely due
484 to reduced microglial phagocytosis of synapses. However, further experiments
485 should be carried out on P2Y₆R knockout microglial lysosomes to test for possible
486 functional deficits, such as lysosomal acidity and enzyme composition.

487

488 We chose a pre-synaptic marker (Vglut1), rather than a post-synaptic marker, as
489 there is clearer evidence for microglial phagocytosis of pre-synaptic markers during
490 development (Weinhard et al., 2018; Mordelt and de Witte, 2023). However, it would
491 be interesting to know whether P2Y₆R also affects microglial phagocytosis of post-
492 synaptic elements, such as dendritic spines, or the microglial phagocytosis of
493 inhibitory synapses. We chose to examine synapse phagocytosis and density at P30
494 because synaptic pruning has been reported to still be actively occurring in
495 somatosensory cortex of mice at this timepoint (Cong et al., 2020). However, it would
496 be useful to know whether P2Y₆R affects microglial phagocytosis at other
497 developmental timepoints, including adults. We chose somatosensory cortex
498 because there is evidence for a role in working memory and we have previously
499 identified P2Y₆R-dependent phagocytosis occurring in this region (Long and Zhang,
500 2021; Dundee et al., 2023), but it would be informative to look at other areas of the
501 brain.

502

503 We measured synapse density as the density of Vglut1-staining puncta, Homer1-
504 staining puncta, and their co-localization (<200 nm) within the Z-stack of 25 μm thick
505 brain sections using confocal microscopy. Thick sections have limitations with
506 confocal microscopy, especially in relation to the z-plane (Avila and Henstridge,
507 2022), so ideally our data should be verified using array tomography of ultrathin
508 sections. Vglut1 and Homer1 puncta density is high in the hippocampal stratum
509 radiatum and molecular layer, as well as the somatosensory cortex, due to the high
510 density of glutamatergic synapses. However, in the hippocampal stratum lacunosum-
511 moleculare, Vglut1 puncta density is substantially lower compared to Vglut2 puncta
512 density as Vglut2 is found primarily on output neurons in the hippocampus (Wozny et
513 al., 2018). It may be beneficial to utilize excitatory synaptic markers that are less
514 region specific for future studies, such as synaptophysin and PSD-95, as well as
515 investigate inhibitory synapses. Note that we analyzed synapses at a single time

516 point (P30) in mouse development, and it would be useful to analyze other time
517 points in order to understand how P2Y₆R affects changes in synapses during
518 development.

519

520 We found a subtle increase in synaptic density in P2Y₆R knockout mice, consistent
521 with reduced microglial phagocytosis of synapses. However, this finding is also
522 consistent with increased synaptogenesis or reduced retraction of synapses in the
523 knockout mice. On the other hand, as there is clear evidence from us and others that
524 P2Y₆R mediates microglial phagocytosis (Koizumi et al., 2007; Puigdellívol et al.,
525 2021; Dundee et al., 2023), but there is no evidence that P2Y₆R affects
526 synaptogenesis or synapse retraction. The most parsimonious explanation of our
527 finding of increased synaptic density in the P2Y₆R knockouts is reduced microglial
528 phagocytosis of synapses, for which we have independent evidence. However, it
529 would be useful to test the effect of P2Y₆R knockout on other potential regulators of
530 synaptic density. It would also be interesting to look at the effect of P2Y₆R knockout
531 on the density of particular synaptic types, such as inhibitory synapses.

532

533 We found that adult P2Y₆R knockout mice had reduced short- and long-term
534 memory, consistent with reduced synaptic pruning during development causing
535 memory dysfunction. However, the memory defects of adult mice could result from
536 reduced microglial phagocytosis of synapses in the adult mice, because such
537 phagocytosis has been found to contribute to learning and memory in the adult brain
538 (Miyanishi et al., 2021). This might be clarified by testing memory at earlier ages, or
539 using conditional knockout of P2Y₆R. However, if learning and memory is regulated
540 by P2Y₆R-dependent microglial phagocytosis of synapses, it may be difficult to
541 disentangle a developmental effect from an adult effect. We have attributed the
542 reduced memory of P2Y₆R knockout mice to reduced synaptic pruning during
543 development, however, it cannot be ruled out that the memory deficits result from
544 other effects of P2Y₆R yet to be discovered. We have previously reported that P2Y₆R
545 knockout mice have improved memory when 17 months old relative to wild-type
546 mice, which have strongly degraded memory at this age (Dundee et al., 2023). We
547 attributed this protection to P2Y₆R mediating the excessive microglial phagocytosis
548 of synapses occurring at this age (Dundee et al., 2023). Thus, P2Y₆R regulated

549 phagocytosis of synapses may potentially protect memory in the young and degrade
550 memory in the old.

551

552 Disruption of synaptic pruning can lead to behavioral deficits in mice, reminiscent of
553 neurodevelopmental disorders, such as autism and schizophrenia, suggesting that
554 these disorders result from disrupted microglial phagocytosis of synapses (Faust,
555 Gunner and Schafer, 2021; Mordelt and de Witte, 2023). It would therefore be
556 interesting to test young P2Y₆R knockout mice for behaviors, such as socialization,
557 related to neurodevelopmental disorders. However, socialization deficits have been
558 linked to specific neuronal-microglial signaling, such as via fractalkine (Corona et al.,
559 2010), rather than microglia per se (Mordelt and de Witte, 2023). It is unclear what
560 behavioral effects may result from loss of UDP-P2Y₆R signaling but it would be
561 interesting to investigate further.

562

563 P2Y₆R is a receptor on microglia for extracellular UDP, and extracellular UDP
564 acutely induces microglial phagocytosis (Koizumi et al., 2007). Kainate-stressed
565 neurons have been shown to release UTP that is converted to UDP in the mouse
566 brain (Koizumi et al., 2007), and amyloid beta-stressed neurons have also been
567 shown to release UDP that activates P2Y₆R (Puigdellivol et al., 2021). UTP is known
568 to be released from stressed or apoptotic cells via pannexin or connexin channels
569 (Elliott et al., 2009; Lazarowski, 2012). UDP activation of P2Y₆R induces formation of
570 the phagocytic cup in microglia, which is a late stage in the engulfment process,
571 normally preceded by recognition of other phagocytic signals on the target. Thus,
572 UDP-P2Y₆R signaling may potentially combine with other signals to recognize
573 synapses that should be engulfed (Cockram et al., 2021). It would be useful to test
574 whether and in what conditions synapses release UDP (although this is currently
575 difficult to do), and how UDP might combine with other signals on the synapse to
576 regulate microglial phagocytosis of synapses.

577

578 **Author contributions**

579

580 JMD and MP managed the animal colonies. JMD analyzed the brain sections. RB
581 wrote the puncta distribution analysis script. MP performed the 4-month-old

582 behavioral studies. GCB conceived and managed the research. JMD and GCB wrote
583 the manuscript. All authors reviewed and approved the manuscript.

584

585 **Data availability statement**

586

587 All code used to analyze synaptic puncta can be found at [https://github.com/gurdon-](https://github.com/gurdon-institute/Synaptic-Density-Analysis)
588 [institute/Synaptic-Density-Analysis](https://github.com/gurdon-institute/Synaptic-Density-Analysis). Experimental data reported in this paper will be
589 shared by the corresponding authors upon request.

590

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731

732 **Figure Legends**

733 **Figure 1: P2Y₆R deficiency reduces synaptic pruning during development. (a)**
 734 Representative confocal microscopy image of mice stained for Iba1 (green,
 735 microglial marker), CB68 (blue, lysosomal marker), and Vglut1 (red, synaptic marker)
 736 in the somatosensory cortex. Scale bar = 2 μm. Virtually all of the CD68 staining was
 737 within Iba1-stained microglia, as shown in Extended Figure 1-1. **(b)** Representative
 738 surface-rendered microglia (from a). Scale bar = 3 μm. **(c)** Enlarged inset of Vglut1
 739 colocalization within CD68, denoted by the white dotted line (from b). Scale bar = 0.5
 740 μm. Microglial volume **(d)**, CD68 volume within microglia **(e)**, and Vglut1
 741 colocalization within CD68 **(f)** across P30 wild-type and P2Y₆R knockout mice (n=7-
 742 8). Each point represents one animal comprised of 14–15 microglia analyzed across
 743 three equidistant sections. Statistical comparisons were made via unpaired t-tests.
 744 Error bars represent ±SEM, ns=p≥0.05, *p<0.05, **p<0.01.

745 **Figure 2: P2Y₆R deficiency does not affect microglial branching in the**
 746 **somatosensory cortex at P30. (a)** Microglial branch end points identified by
 747 3DMorph analysis (York et al., 2018). **(b)** Mean number of branch end points per
 748 microglia (n = 8). **(c)** Microglial branching points identified by 3DMorph analysis

749 (York et al., 2018). **(d)** Mean number of branching points per microglia ($n = 8$). **(e)**
750 Mean branch length (μm) per microglia ($n = 8$). **(f)** Mean of the maximum branch
751 length (μm) per microglia ($n = 8$). **(g)** Mean of the minimum branch length (μm) per
752 microglia ($n = 8$). Each point represents one animal. Statistical comparisons were
753 made via unpaired t-tests. Error bars represent $\pm\text{SEM}$.

754 **Figure 3: P2Y₆R deficiency reduces microglial density in the somatosensory**
755 **cortex at P30. (a)** Representative fluorescent microscopy images of mice stained for
756 Iba1 (green, microglial marker) and DAPI (blue, nuclei marker) in the somatosensory
757 cortex. Scale bar = 50 μm . **(b)** Mean number of microglia per 20x field ($n = 8$). Each
758 point represents one animal. Statistical comparison was made via an unpaired t-test.
759 Error bars represent $\pm\text{SEM}$, $**p < 0.01$.

760 **Figure 4: P2Y₆R deficiency results in increased synaptic density in the**
761 **somatosensory cortex. (a)** Nissl stain from the Allen Mouse Brain Atlas and Allen
762 Reference Atlas—Mouse Brain of a coronal section of the mouse brain, with the
763 somatosensory cortex labeled. Available from: mouse.brain-map.org/static/atlas. **(b)**
764 Colocalization of Vglut1 and Homer1 puncta. Scale bar = 0.5 μm . **(c)** Representative
765 confocal microscopy images of P30 wild-type and P2Y₆R knockout mice stained for
766 Vglut1 (red, pre-synaptic marker), Homer1 (green, post-synaptic marker), and with
767 DAPI (blue, nuclear stain) in the somatosensory cortex. Scale bar = 5 μm . **(d)** Vglut1
768 puncta density, Homer1 puncta density, and synaptic density of the somatosensory
769 cortex ($n = 8$, 3 equidistant planes 300 μm apart per mouse). Synaptic density
770 determined as colocalized Vglut1 and Homer1 puncta ($< 200 \text{ nm}$). Raw values for
771 synaptic puncta density across brain regions are available in Extended Figure 4-1.
772 Synaptic internalization by microglia negatively correlates with synaptic density in the
773 somatosensory cortex, as shown in Extended Figure 4-2. Each point represents one
774 animal. Statistical comparisons were made via unpaired t-tests. Error bars represent
775 $\pm\text{SEM}$, $*p < 0.05$.

776 **Figure 5: P2Y₆R deficiency does not affect synaptic density in the CA1**
777 **hippocampus. (a)** Nissl stain from the Allen Mouse Brain Atlas and Allen Reference
778 Atlas—Mouse Brain of a coronal section of the mouse brain, with the hippocampal
779 CA1 stratum radiatum labeled. Available from: mouse.brain-map.org/static/atlas. **(b)**
780 Colocalization of Vglut1 and Homer1 puncta. Scale bar = 0.5 μm . **(c)** Representative
781 confocal microscopy images of P30 wild-type and P2Y₆R knockout mice stained for
782 Vglut1 (red, pre-synaptic marker), Homer1 (green, post-synaptic marker), and with

783 DAPI (blue, nuclear stain) in the hippocampal CA1 stratum radiatum. Scale bar = 5
784 μm . **(d)** Vglut1 puncta density, Homer1 puncta density, and synaptic density of the
785 hippocampal CA1 stratum radiatum ($n = 8$, 3 equidistant planes $300\ \mu\text{m}$ apart per
786 mouse). Synaptic density determined as colocalized Vglut1 and Homer1 puncta
787 ($<200\ \text{nm}$). Each point represents one animal. Statistical comparisons were made via
788 unpaired t-tests. Error bars represent $\pm\text{SEM}$.

789 **Figure 6: P2Y₆R deficiency results in increased synaptic density in the CA3**
790 **hippocampus.** **(a)** Nissl stain from the Allen Mouse Brain Atlas and Allen Reference
791 Atlas—Mouse Brain of a coronal section of the mouse brain, with the hippocampal
792 CA3 stratum radiatum labeled. Available from: mouse.brain-map.org/static/atlas. **(b)**
793 Colocalization of Vglut1 and Homer1 puncta. Scale bar = $0.5\ \mu\text{m}$. **(c)** Representative
794 confocal microscopy images of P30 wild-type and P2Y₆R knockout mice stained for
795 Vglut1 (red, pre-synaptic marker), Homer1 (green, post-synaptic marker), and with
796 DAPI (blue, nuclear stain) in the hippocampal CA3 stratum radiatum. Scale bar = 5
797 μm . **(d)** Vglut1 puncta density, Homer1 puncta density, and synaptic density of the
798 hippocampal CA3 stratum radiatum ($n = 8$, 3 equidistant planes $300\ \mu\text{m}$ apart per
799 mouse). Synaptic density determined as colocalized Vglut1 and Homer1 puncta
800 ($<200\ \text{nm}$). Each point represents one animal. Statistical comparisons were made via
801 unpaired t-tests. Error bars represent $\pm\text{SEM}$, * $p < 0.05$.

802 **Figure 7: P2Y₆R deficiency results in increased synaptic density in the dentate**
803 **gyrus.** **(a)** Nissl stain from the Allen Mouse Brain Atlas and Allen Reference Atlas—
804 Mouse Brain of a coronal section of the mouse brain, with the hippocampal dentate
805 gyrus molecular layer labeled. Available from: mouse.brain-map.org/static/atlas. **(b)**
806 Colocalization of Vglut1 and Homer1 puncta. Scale bar = $0.5\ \mu\text{m}$. **(c)** Representative
807 confocal microscopy images of P30 wild-type and P2Y₆R knockout mice stained for
808 Vglut1 (red, pre-synaptic marker), Homer1 (green, post-synaptic marker), and with
809 DAPI (blue, nuclear stain) in the hippocampal dentate gyrus molecular layer. Scale
810 bar = $5\ \mu\text{m}$. **(d)** Vglut1 puncta density, Homer1 puncta density, and synaptic density
811 of the hippocampal dentate gyrus molecular layer ($n = 8$, 3 equidistant planes
812 $300\ \mu\text{m}$ apart per mouse). Synaptic density determined as colocalized Vglut1 and
813 Homer1 puncta ($<200\ \text{nm}$). Each point represents one animal. Statistical
814 comparisons were made via unpaired t-tests. Error bars represent $\pm\text{SEM}$, * $p < 0.05$.

815 **Figure 8: P2Y₆R deficiency results in long-term memory loss.** **(a)** Schematic
816 representation of the training and testing sessions in the NORT. **(b)** Object

817 preference of each animal as percentage of time spent exploring two identical
818 objects (n = 11). **(c)** Object preference of each animal as percentage of time spent
819 exploring the novel object 24 h after training (n = 11). **(d)** Schematic representation
820 of the training and testing sessions in the OLRT. **(e)** Object preference of each
821 animal as percentage of time spent exploring two identical objects (n = 11). **(f)** Object
822 preference of each animal as percentage of time spent exploring the novel object
823 24 h after training (n = 11). Data from (b) and (c) was reanalyzed from Dundee et al.
824 (2023). Each point represents one animal. Dashed lines indicate a 50% chance
825 level. Statistical comparisons were made via unpaired t-tests. Error bars represent
826 \pm SEM, * $p < 0.05$.

827 **Figure 9: P2Y₆R deficiency results in short-term memory loss.** **(a)** Schematic
828 representation of spontaneous alterations in the Y-maze. **(b)** Total number of arm
829 entries by each animal in the Y-maze (n = 7-9). **(c)** Distance travelled (inches) by
830 each animal in the Y-maze (n = 7-9). **(d)** Spontaneous alterations by each animal in
831 the Y-maze (n = 7-9). **(e)** Schematic representation of the training and testing
832 sessions in the NORT. **(f)** Object preference of each animal as percentage of time
833 spent exploring two identical objects (n = 7). **(g)** Object preference of each animal as
834 percentage of time spent exploring the novel object 24 h after training (n = 7). Data
835 from (f) and (g) was reanalyzed from Puigdellívol et al. (2021). Each point represents
836 one animal. Dashed lines indicate a 50% chance level. Statistical comparisons were
837 made via unpaired t-tests. Error bars represent \pm SEM, * $p < 0.05$.

838 **Extended Figure 1-1: CD68 staining is predominantly found within Iba1-**
839 **positive microglia in the somatosensory cortex at P30.** **(a)** Representative
840 confocal microscopy images of mice stained for Iba1 (green, microglial marker) and
841 CD68 (blue, lysosomal marker) in the somatosensory cortex. Scale bar = 10 μ m. **(b)**
842 Enlarged inset of CD68 colocalization within Iba1, denoted by the white dotted line
843 (from a). White arrows indicate CD68 colocalization. Scale bar = 5 μ m.

844 **Extended Figure 4-1: Raw values for synaptic densities of the somatosensory**
845 **cortex, CA1 hippocampal, CA3 hippocampal, and dentate gyrus regions of P30**
846 **mice.** Vglut1 **(a)**, Homer1 **(b)**, and colocalization of Vglut1 and Homer1 **(c)** puncta
847 densities in the somatosensory cortex (From Figure 4, n = 8, 3 equidistant planes
848 300 μ m apart per mouse). Vglut1 **(d)**, Homer1 **(e)**, and colocalization of Vglut1 and
849 Homer1 **(f)** puncta densities in the hippocampal CA1 stratum radiatum (From Figure
850 5, n = 8, 3 equidistant planes 300 μ m apart per mouse). Vglut1 **(g)**, Homer1 **(h)**, and

851 colocalization of Vglut1 and Homer1 **(i)** puncta densities in the hippocampal CA3
852 stratum radiatum (From Figure 6, n = 8, 3 equidistant planes 300 μ m apart per
853 mouse). Vglut1 **(j)**, Homer1 **(k)**, and colocalization of Vglut1 and Homer1 **(l)** puncta
854 densities in the hippocampal dentate gyrus molecular layer (From Figure 7, n = 8, 3
855 equidistant planes 300 μ m apart per mouse). Synaptic density determined as
856 colocalized Vglut1 and Homer1 puncta (<200 nm). Each point represents one
857 animal. Statistical comparisons were made via unpaired t-tests. Error bars represent
858 \pm SEM, *p < 0.05, **p < 0.01.

859 **Extended Figure 4-2: Synaptic internalization by microglia negatively**
860 **correlates with synaptic density in the somatosensory cortex.** Synaptic density
861 of the somatosensory cortex (normalized to *P2ry6*^{+/+} mice) plotted against volume of
862 internalized synaptic material within microglia in the somatosensory cortex (n=15,
863 WT=7, KO=8). Each point represents one animal. Statistical correlation was made by
864 calculating the Pearson correlation coefficient.

Figure 1

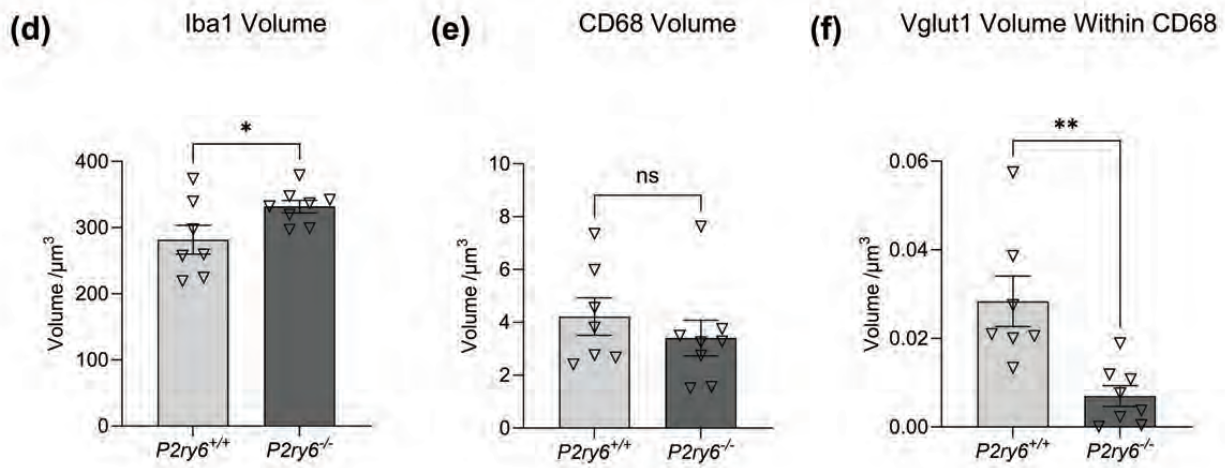
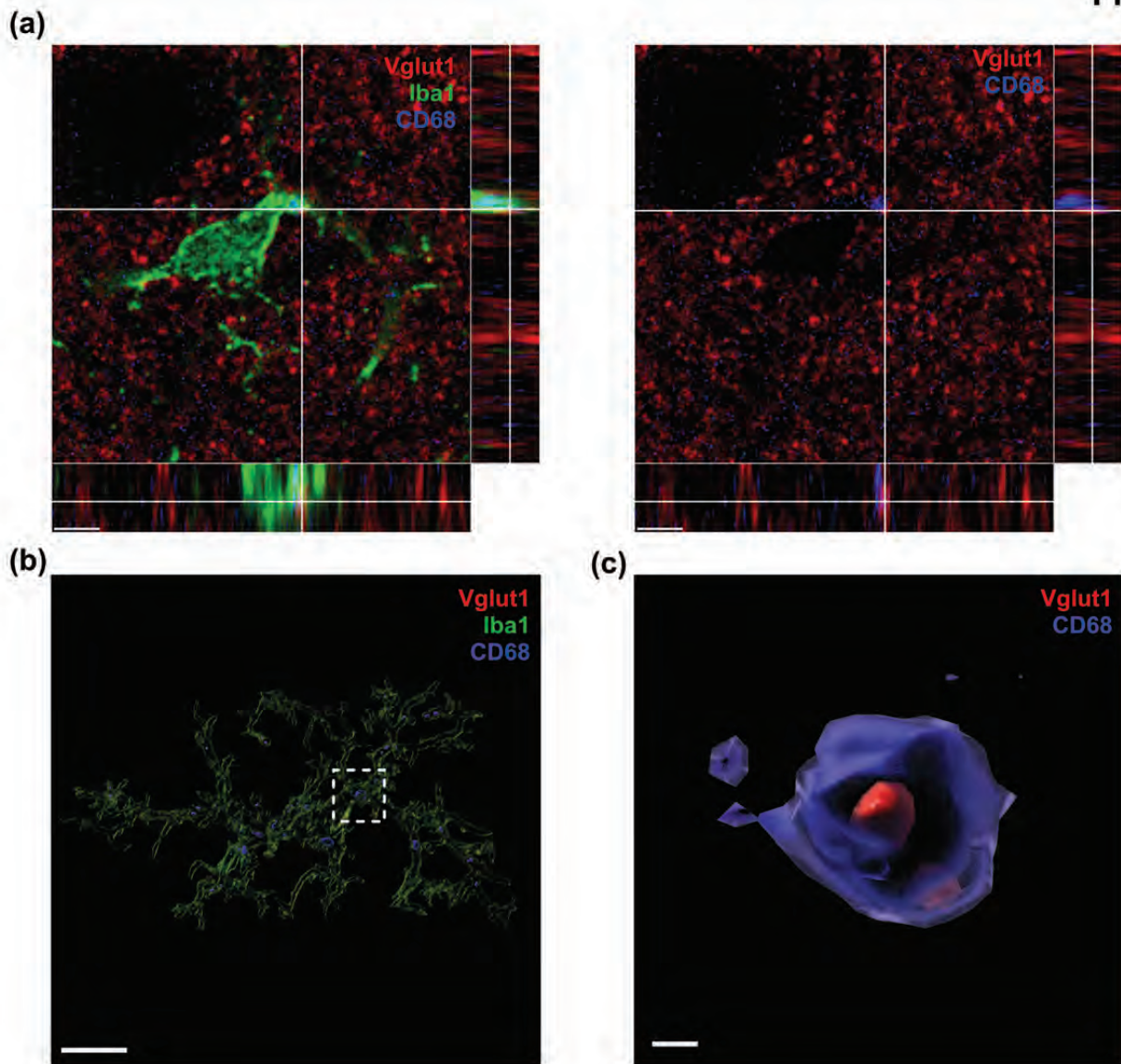
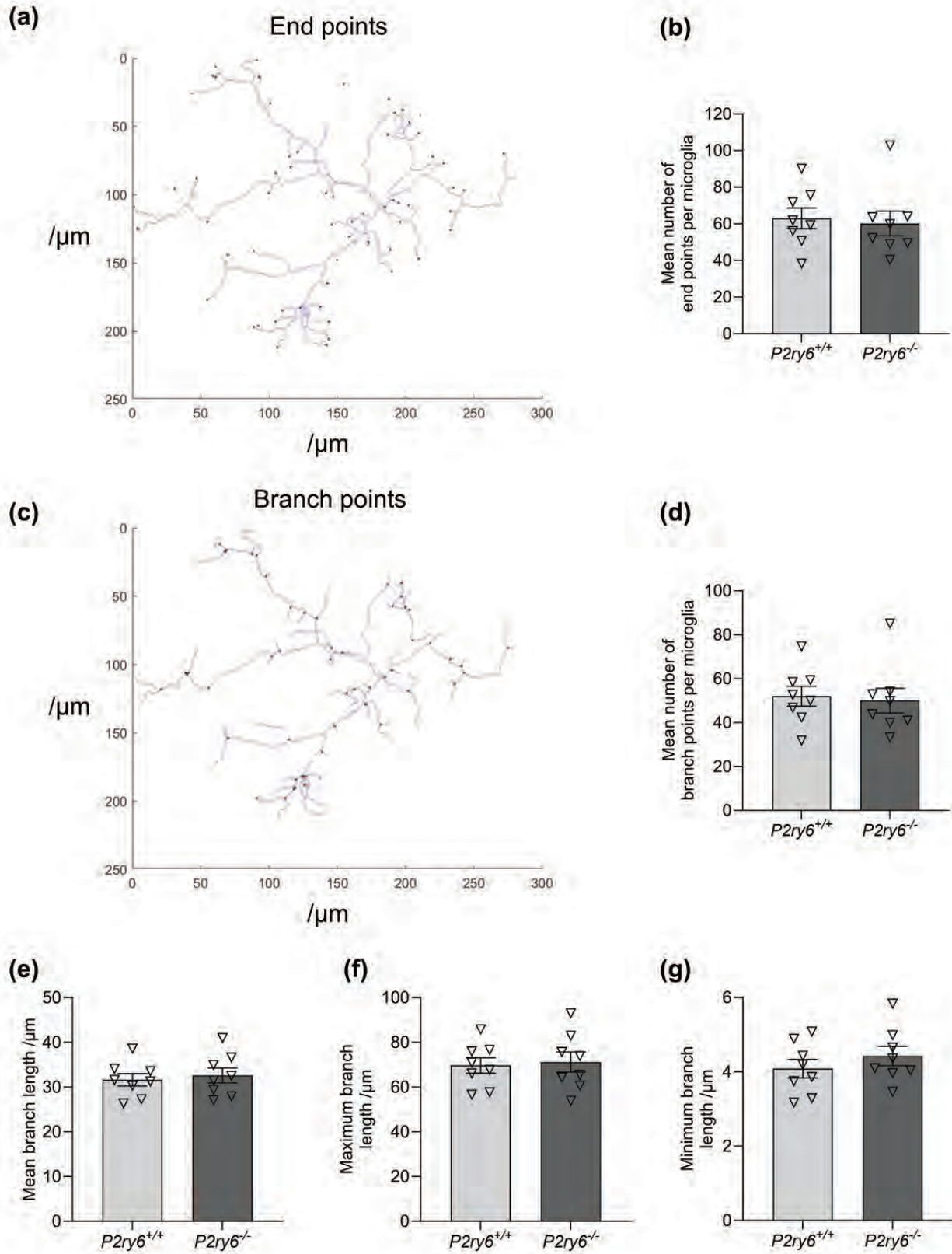


Figure 2



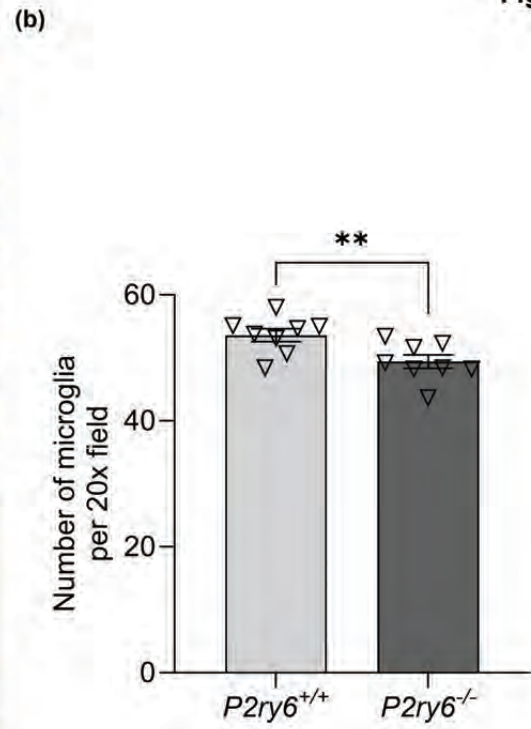
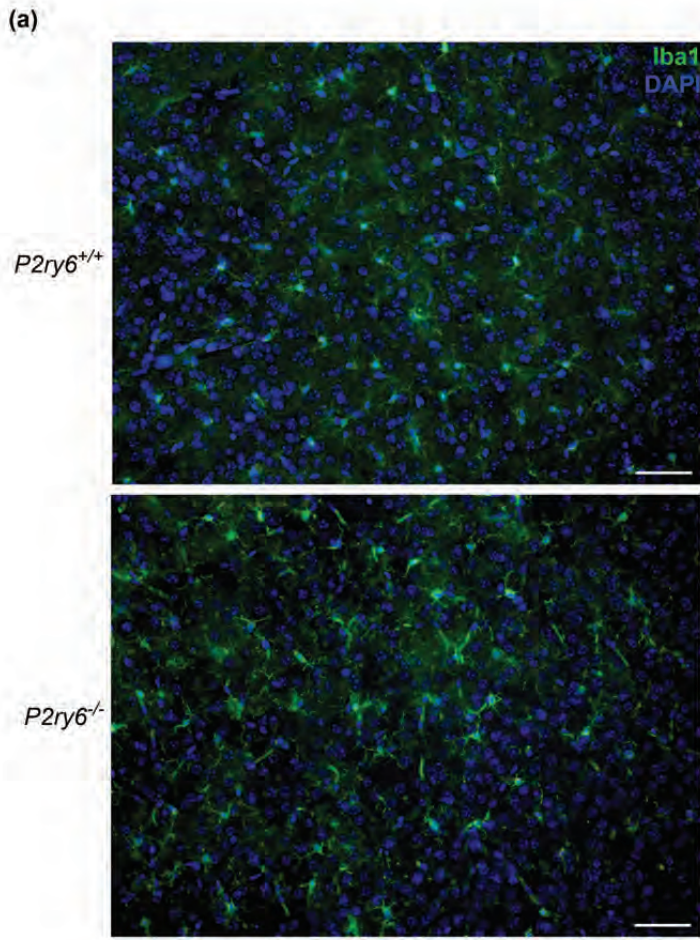


Figure 3

Figure 4

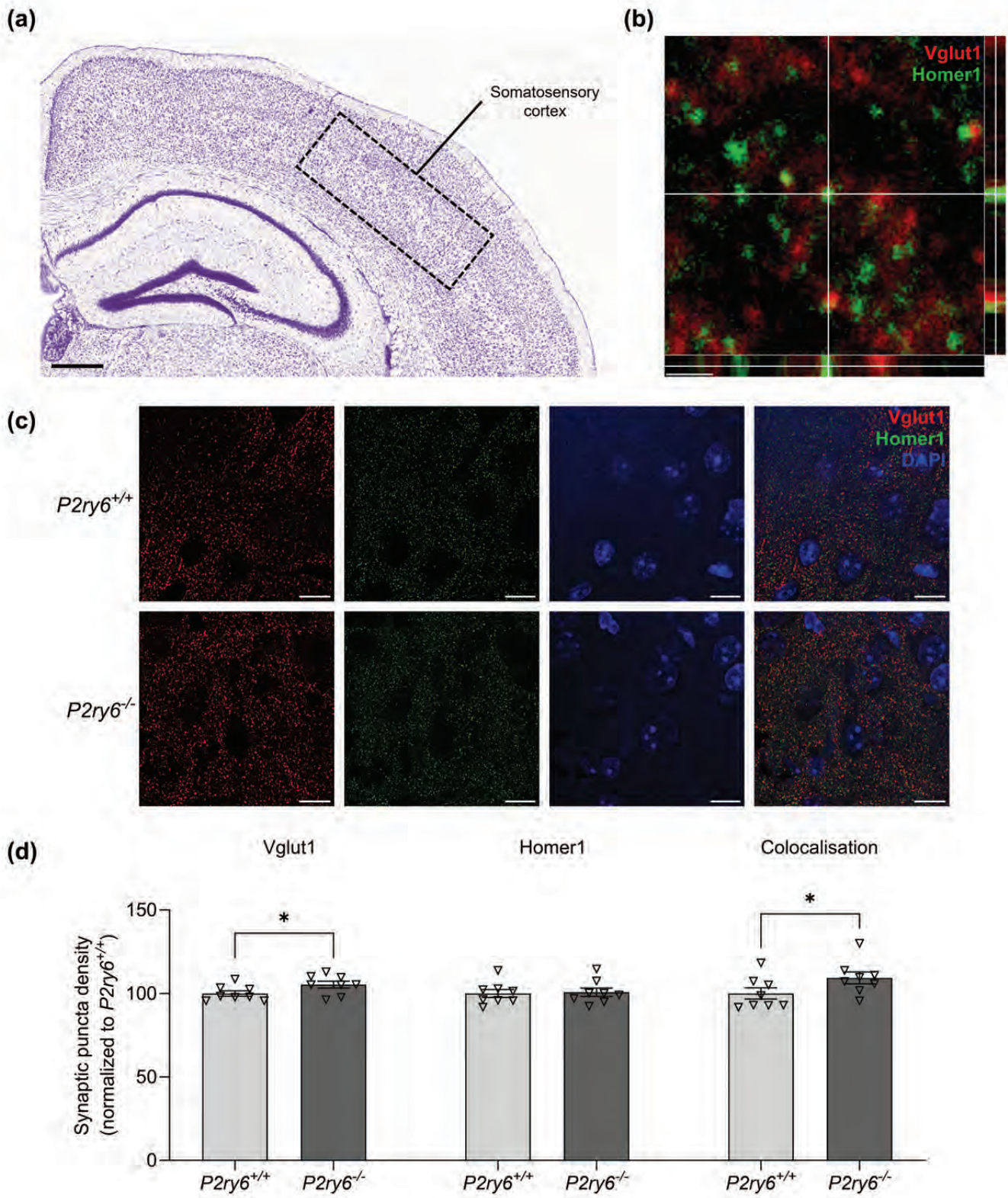


Figure 5

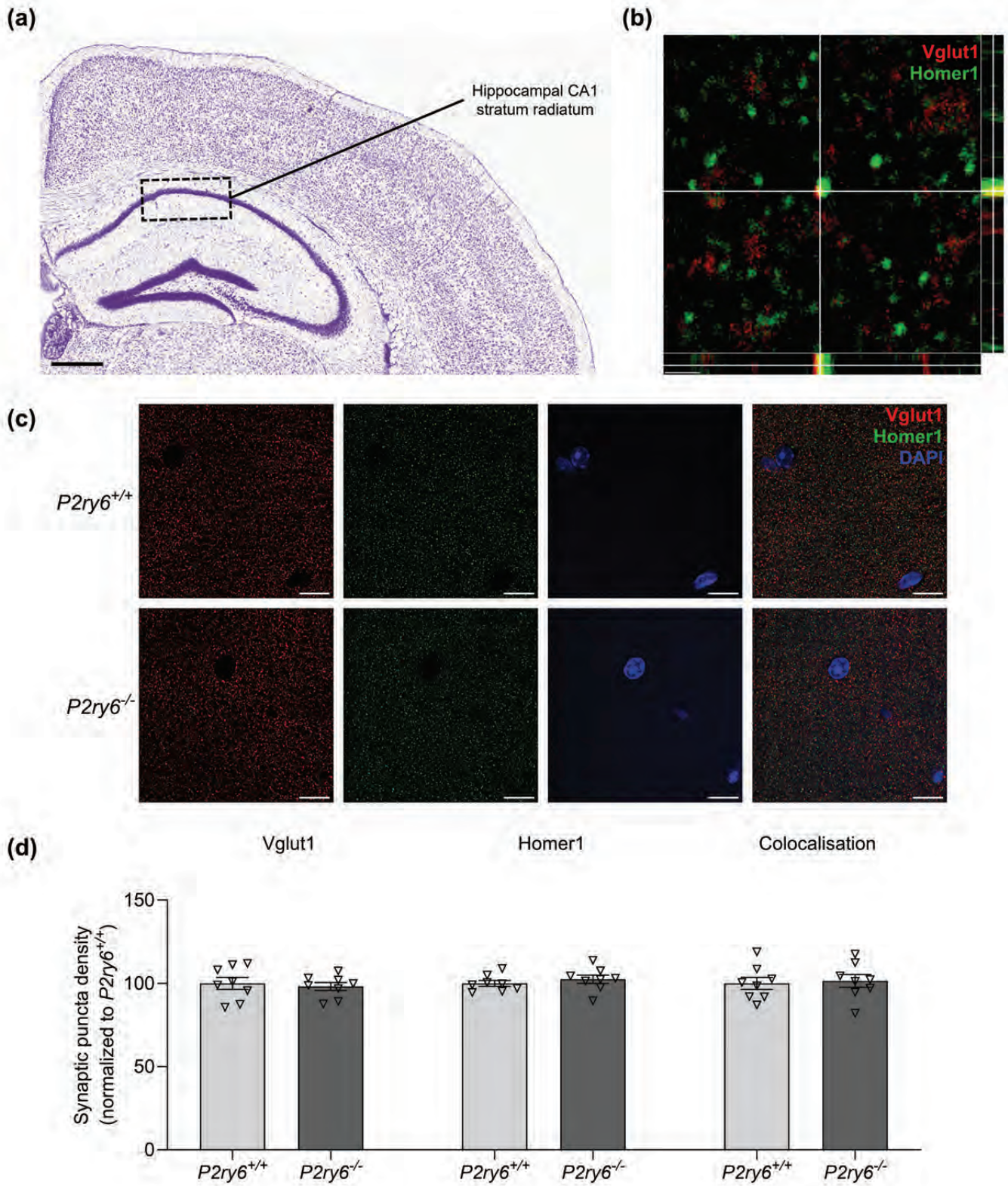


Figure 6

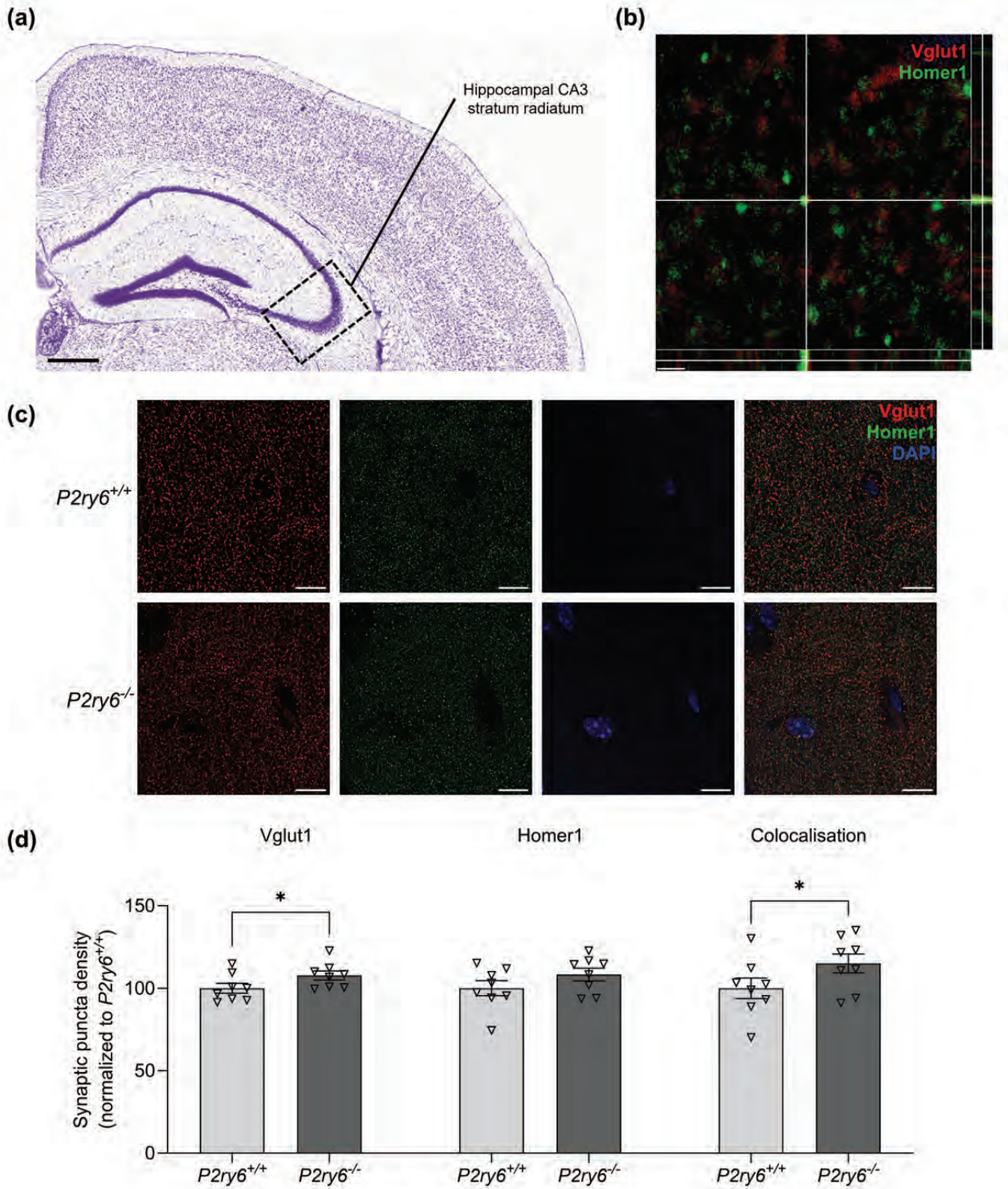


Figure 7

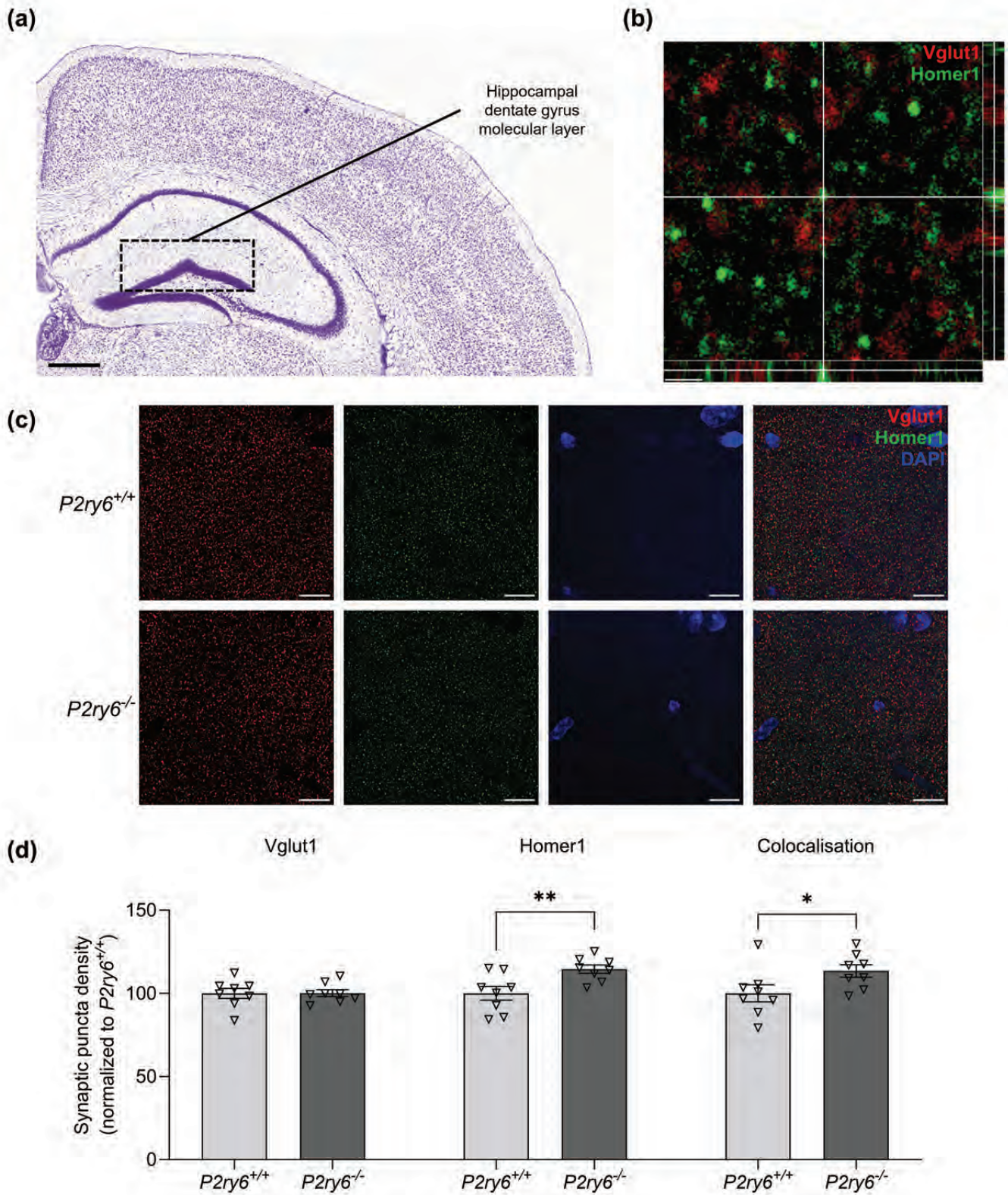
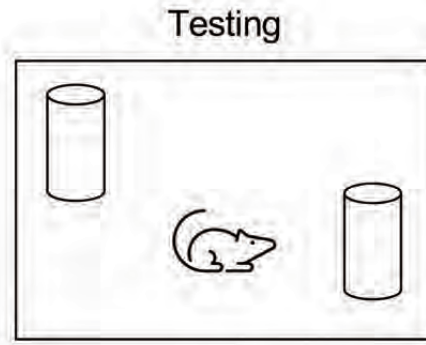
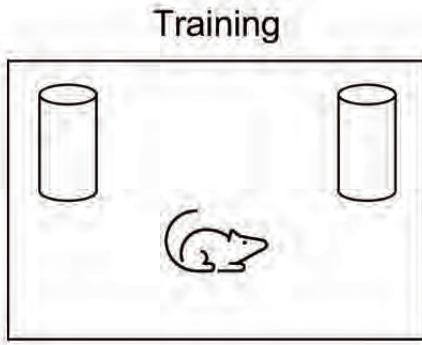
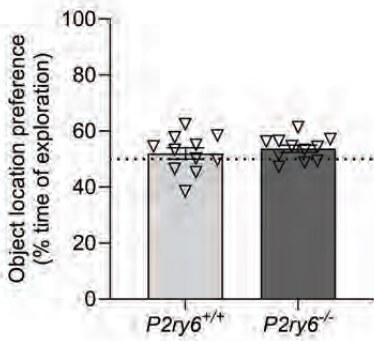


Figure 8

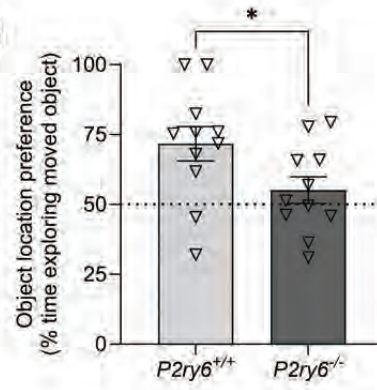
(a)



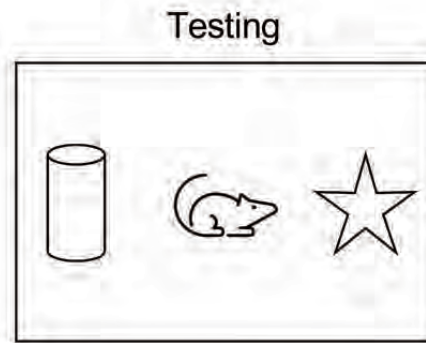
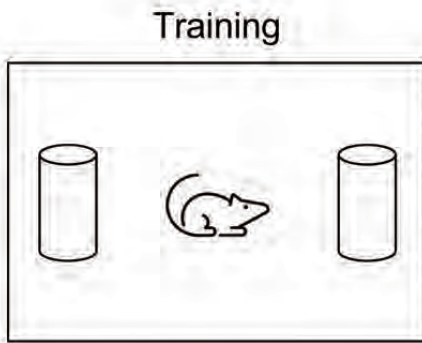
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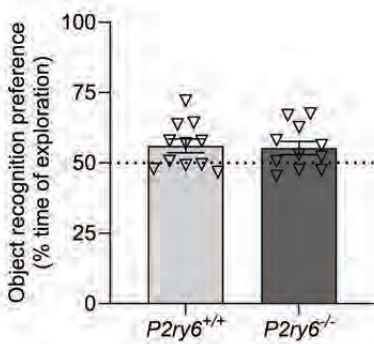
(c)



(d)



(e)



(f)

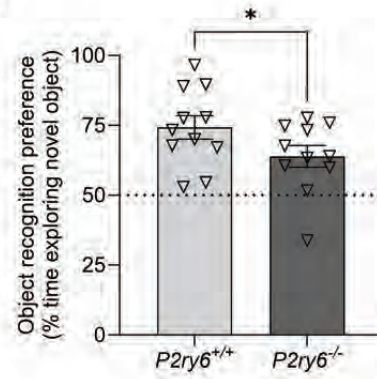


Figure 9

