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Short-term Dendritic Dynamics of Neonatal Cortical Neurons Revealed by *in vivo* Imaging with Improved Spatiotemporal Resolution

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1 **Short-term Dendritic Dynamics of**
2 **Neonatal Cortical Neurons Revealed by *in vivo* Imaging**
3 **with Improved Spatiotemporal Resolution**

4
5 Abbreviated title:

6 Short-term Dendritic Dynamics of Neonatal Neurons

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44 **Abstract:**

45 Individual neurons in sensory cortices exhibit specific receptive fields based on their dendritic
46 patterns. These dendritic morphologies are established and refined during the neonatal period
47 through activity-dependent plasticity. This process can be visualized using two-photon *in vivo*
48 time-lapse imaging, but sufficient spatiotemporal resolution is essential. We previously examined
49 dendritic patterning from spiny stellate (SS) neurons, the major type of layer 4 (L4) neurons, in
50 the mouse primary somatosensory cortex (barrel cortex), where mature dendrites display a strong
51 orientation bias toward the barrel center. Longitudinal imaging at 8-h intervals revealed the long-
52 term dynamics by which SS neurons acquire this unique dendritic pattern. However, the
53 spatiotemporal resolution was insufficient to detect the more rapid changes in SS neuron dendrite
54 morphology during the critical neonatal period. In the current study, we imaged neonatal L4
55 neurons hourly for 8 h and improved the spatial resolution by uniform cell-surface labeling. The
56 improved spatiotemporal resolution allowed detection of precise changes in dendrite morphology
57 and revealed aspects of short-term dendritic dynamics unique to the neonatal period. Basal
58 dendrites of barrel cortex L4 neurons were highly dynamic. In particular, both barrel-inner and
59 barrel-outer dendrites (trees and branches) emerged/elongated and disappeared/retracted at
60 similarly high frequencies, suggesting that SS neurons acquire biased dendrite patterns through
61 rapid trial-and-error emergence, elongation, elimination, and retraction of dendritic trees and
62 branches. We also found correlations between morphology and behavior (elongation/retraction)
63 of dendritic tips. Thus, the current study revealed short-term dynamics and related features of
64 cortical neuron dendrites during refinement.

65

66

67 **Significance Statement:**

68 The formation of proper dendritic patterns during early postnatal development is essential for
69 normal neuronal circuit function in adulthood. To elucidate the mechanisms responsible for this

70 refinement, *in vivo* imaging with high spatiotemporal resolution is useful. Our previous long-term
71 *in vivo* imaging studies have clarified aspects of dendritic refinement mechanism; however, due
72 to the long intervals (8-h) between image acquisitions, rapid changes in dendritic morphology
73 were missed. Here hourly *in vivo* time-lapse imaging of neonatal mouse barrel cortex over 8 h
74 revealed the rapid changes in the dendrite morphology of layer 4 neurons, thereby providing a
75 more comprehensive record of dendritic refinement during postnatal development for mechanistic
76 analysis.

77

78

79 **Introduction:**

80 Complex yet precise connectivity of neurons underlies higher brain functions in mammals. The
81 specificity of neuronal connections depends largely on dendritic patterns as they determine the
82 inputs received by individual neurons. Mature spiny stellate (SS) neurons in the mouse primary
83 somatosensory cortex (barrel cortex) layer 4 (L4) show highly biased basal dendritic patterns
84 oriented toward the barrel center, where thalamocortical axons (TCAs) transmit single-whisker
85 inputs. This unique asymmetric basal dendritic pattern, which underlies the precise one-to-one
86 functional relationship between whiskers and barrels, is formed during neonatal stages in a TCA
87 input-dependent manner (Datwani et al., 2002; Espinosa et al., 2009; Mizuno et al., 2014). Thus,
88 barrel cortex SS neuron basal dendrites are an ideal model for studying the mechanisms of
89 dendritic refinement.

90 To reveal the mechanisms of dendritic refinement, it is critical to elucidate the dynamic
91 features of dendrites during early postnatal development. Two-photon microscopy has been used
92 extensively for imaging of spine dynamics *in vivo* during late postnatal development and
93 adulthood (Holtmaat & Svoboda, 2009; Lendvai et al., 2000; Yang et al., 2009). Some studies
94 have also addressed axon dynamics *in vivo* (Carrillo et al., 2013; Portera-Cailliau et al., 2005;
95 Ruthazer et al., 2003). On the other hand, understanding dendrite dynamics is largely delayed

96 (Mizuno et al., 2014; Nakazawa et al., 2018), which is partly because visualization of dendrite
97 morphology *in vivo* is difficult in the neonatal brain (Ako et al., 2011; Espinosa et al., 2009; Luo
98 et al., 2016; Young et al., 2008). We previously achieved that by developing the Supernova
99 system, a versatile sparse cell labeling method (Luo et al., 2016; Mizuno et al., 2014), and
100 successfully performed two-photon *in vivo* imaging of individual L4 neurons in the neonatal
101 barrel cortex at 4 time points within 18 h between postnatal day 5 (P5) and P6 (Mizuno et al.,
102 2014). This study analyzed morphological changes in barrel cortex L4 neuron basal dendrites in
103 the neonatal cortex and detected dendritic motility *in vivo* for the first time. Subsequently, we
104 achieved longitudinal *in vivo* imaging of morphological changes in basal dendrites from P3 to P6
105 (Nakazawa et al., 2018).

106 Although the aforementioned study revealed the long-term dynamics that generate
107 orientation bias, the spatiotemporal resolution was not sufficient to capture short-term dendritic
108 dynamics during neonatal dendritic refinement. In that study, to cover the whole refinement
109 processes between P3 and P6, we set acquisition intervals of 8 or 24 hours. However, the changes
110 in dendrite morphology over 8 h are much larger than expected (Nakazawa et al., 2018), which
111 hindered the detection of rapid changes over shorter periods. For example, if a dendritic branch is
112 eliminated and a new branch emerges at a similar position within 8 h, it could be mistakenly
113 recognized as the same. Thus, greater temporal resolution is required to reduce such potential
114 errors.

115 To characterize the short-term dynamics of SS neuron dendrites more precisely, we here
116 conducted hourly imaging for 8 h. We also improved the spatial resolution by labeling cell
117 surfaces with a membrane-binding fluorescent protein, which was helpful to visualize more
118 precise dendrite morphology. In the current study, we describe the short-term dynamics of
119 cortical neuron dendrites during the critical period of dendritic refinement. Also, we report that
120 the elongation and retraction of dendrites were positively correlated with dendrite tip thickness.

121

122

123 **Materials and Methods:**

124 **Animals**

125 All experiments were performed according to guidelines for animal experimentation of National
126 Institute of Genetics (NIG) and were approved by the animal experimentation committee of NIG.
127 Thalamocortical axon (TCA)-green fluorescent protein (GFP) transgenic (Tg) mice (Mizuno et al.,
128 2014) were backcrossed from C57BL/6J to ICR more than four times and were intercrossed to
129 obtain TCA-GFP Tg homozygous male mice. Homozygosity of the TCA-GFP Tg mice was
130 determined by quantitative polymerase chain reaction genotyping (qPCR) using the primers
131 [KS89] TTCTCGTTGGGGTCTTTGCT and [KS90] ACTTCTCAAGTCCGCCATG. To obtain
132 pups, ICR female mice were mated with the TCA-GFP Tg homozygous male mice. The day at
133 which the vaginal plug was detected (between 10 AM to 11 AM) was designated as embryonic
134 day (E) 0 and E19 was defined as postnatal day (P) 0.

135 **Plasmids**

136 Flpe/FRT-based Supernova vectors: pK036 (TRE-Flpe-WPRE) (Luo et al., 2016); pK037 [CAG-
137 FRT-STOP-FRT-turbo red fluorescent protein (turboRFP)-ires-tTA-WPRE] (Luo et al., 2016),
138 and pK300 (CAG-FRT-STOP-FRT-GAP43tagRFP-ires-tTA-WPRE) were used for sparse
139 labeling. pK302 [CAG-blue fluorescent protein (BFP)-WPRE] (Nakazawa et al., 2020) was used
140 for dense labeling. The plasmid DNA was purified by a Midi-prep kit (Macherey-Nagel). For the
141 generation of pK300, the GAP43tagRFP sequence was obtained from pK190 (CAG-loxP-STOP-
142 loxP-GAP43tagRFP-ires-tTA-WPRE) vector by Sall/EcoRV restriction digestion and inserted
143 into Sall/EcoRV sites of pK068 (CAG-FRT-STOP-FRT-EGFP-ires-tTA-WPRE vector) (Luo et
144 al., 2016). For the generation of pK190, GAP43 sequence (Chiaramello et al., 1996) and tagRFP
145 sequence from pTagRFP-N vector (Haas et al., 1996; Kozak, 1987) were fused and PCR

146 amplified using the primers [HM67]
147 CTAGTGTGCGACATGCTGTGCTGTATGAGAAGAACCAAACAGGTTGAAAAGAATGAT
148 GAGGACCAAAAGATCGAG and [HM75]
149 CCGATATCTTCAATTAAGTTTGTGCCCCAGTTTGCTAGG, and then inserted into the
150 SalI/EcoRV sites of pK038 (CAG-loxP-STOP-loxP-EGFP-ires-tTA-WPRE) vector (Luo et al.,
151 2016).

152 ***In utero* electroporation**

153 *In utero* electroporation (IUE) was performed on TCA-GFP Tg heterozygous mice at E14
154 between 10 AM and noon to label L4 neurons in the barrel cortex. The pregnant mothers were
155 anesthetized via intraperitoneal injection of triple anesthesia (11 $\mu\text{g/g}$ body weight). The triple
156 anesthesia contains medetomidine hydrochloride (0.75 $\mu\text{g/g}$), midazolam (4 $\mu\text{g/g}$), and
157 butorphanol tartrate (5 $\mu\text{g/g}$). A total of 0.5 μl of DNA solution (mixed with 5% methylene blue)
158 was injected into the right lateral ventricle of embryos via a pulled glass capillary (Drummond),
159 and square electric pulses (40 V; 50 ms) were delivered 5 times at 1 Hz by a CUY21EDIT
160 electroporator (NepaGene). The electric pulses were given 3 times/embryo, and the current was
161 70–100 mA/electric pulse. After the IUE, pregnant mothers were injected with an antagonist (11
162 $\mu\text{g/g}$ body weight), which contained atipamezole hydrochloride (0.75 $\mu\text{g/g}$), to wake them up
163 from anesthesia and kept on a 37°C heater until they recovered from anesthesia.

164 For Supernova-cytosolic RFP (cRFP) labeling, a DNA solution containing pK036 (20
165 ng/ μl) and pK037 (1 $\mu\text{g}/\mu\text{l}$) was used. For Supernova-membrane-bound RFP (mRFP) labeling, a
166 DNA solution containing pK036 (20 or 40 ng/ μl), pK300 (1 $\mu\text{g}/\mu\text{l}$), and pK302 (200 ng/ μl) was
167 used.

168 ***In vivo* imaging with two-photon microscopy**

169 For *in vivo* imaging of L4 neurons in the barrel cortex, TCA-GFP Tg heterozygous pups at P4
170 were used, in which L4 neurons were labeled by IUE-based BFP and Supernova-mRFP. Mice in
171 which L4 neurons in the barrel cortex were labeled appropriately were identified by BFP
172 fluorescence over the skull after removing the skin from the corresponding areas. Then, the
173 craniotomy was performed on the right half of the head while the pups were under isoflurane
174 anesthesia (3%) as described previously (Mizuno et al., 2014; Nakazawa et al., 2018). Briefly, the
175 skull above the IUE-labeled area in the barrel cortex was carefully removed by a razor blade with
176 the dura remaining intact, and a small round cover glass (2.5 mm in diameter, Matsunami) was
177 placed to cover the open skull area. Gelfoam (Pfizer) was used to stop the occasional minor
178 bleeding and cortex buffer (Holtmaat et al., 2009) (125 mM NaCl, 5 mM KCl, 10 mM glucose,
179 10mM Hepes, 2 mM CaCl₂, and 2 mM MgSO₄; pH 7.4) was applied to the skull-removed area to
180 keep the brain moist during surgery. A customized titanium bar (~30 mg) (Nakazawa et al., 2018)
181 was attached to the adjacent area to the cranial window. No apparent surgery-induced discomfort
182 was observed. After 1 h of recovery on a 37°C heater with littermates, pups with cranial windows
183 were used for *in vivo* imaging. Pups were anesthetized with isoflurane (0.9%–1.4%) and head
184 fixed to the microscope stage using the titanium bar during *in vivo* imaging. A heating pad was
185 used to keep the pups warm. Images were acquired using a two-photon microscope (LSM 7MP,
186 Zeiss) with a W Plan-Apochromat 20x/1.0 DIC objective lens (Zeiss) and an LSM BiG detector
187 (Zeiss). A HighQ-2 laser (Spectra-Physics) at 1,045 nm was used in all experiments. GFP and
188 RFP were simultaneously excited, and the emitted fluorescence was filtered (500–550 nm for
189 GFP and 575–620 nm for RFP). *In vivo* imaging was performed as follows: h0, h1, h2, h3, h4, h5,
190 h6, h7, and h8 sessions started around 13:00, 14:00, 15:00, 16:00, 17:00, 18:00, 19:00, 20:00, and
191 21:00 at P4, respectively. Each imaging session took about 20 min. Between two imaging
192 sessions, pups were returned to their littermates and kept on a 37°C heater. The pups usually start
193 to respond to touch from littermates and move a few minutes after the end of the imaging session.

194 **Histology and confocal microscopy**

195 After *in vivo* imaging, brain samples were obtained and fixed with ice-cold 4% paraformaldehyde
196 in 0.1 M phosphate buffer (PB). The brain samples were kept at 4°C overnight and protected from
197 light. For tangential sectioning, the right hemispheres were flattened and transferred to 30%
198 sucrose in 0.1 M PB and kept overnight at 4°C. Tangential sections (100 μm thick) of the
199 flattened cortices were made using a freezing microtome (ROM-380; Yamato).

200 Three-dimensional (3D) fluorescence images were acquired using a confocal microscope
201 (TCS SP5; Leica). Images of the barrel map and labeled neurons were taken with a 10x/0.4 CS2
202 objective lens with 2x digital zoom at a step size around 2.5 μm and the total thickness was \leq 100
203 μm . The 20x images were taken with 1024 \times 1024 pixels. Images of fine structures of dendrites
204 were taken with a 63x/1.4 oil immersion lens with 2x digital zoom at a step size around 0.1 μm
205 and the total thickness was \leq 100 μm . The 126x images were taken with 2048 \times 2048 pixels. Step
206 sizes were determined according to the optimized setting for each objective lens.

207 **Image analysis and quantification**

208 The Image J/FIJI (NIH) (version 1.53t) software was used for histological analysis. The TCA
209 signal intensities were used to distinguish barrels and septa areas in *post hoc* analysis with
210 confocal images (taken by a 10x lens with 2 times zoom). TCA cluster boundaries were defined
211 as barrel edges. *In vivo* imaged L4 neurons in barrels were identified in confocal images
212 according to their relative positions with each other and their dendritic patterns. The distance of
213 neurons to barrel edges was defined as the distance from the center of the cell body to the nearest
214 barrel edge, and within 20 μm were classified as edge-located neurons. Neurons located barrel
215 inside and distances to the barrel edge longer than 20 μm were classified as barrel-center neurons
216 and not included in the analysis of the current study.

217 Acquired 3D neuron images from two-photon microscopy were analyzed using the
218 IMARIS Filament Tracer (Bitplane, version 9.5.1) software to reconstruct dendritic patterns and
219 quantify basal dendritic morphologies. To keep the accuracy and efficiency of data analysis, we
220 selected *in vivo* imaged neurons for analysis with the following three criteria: I, basal dendritic
221 morphologies were clearly visible both in *in vivo* imaging (all 9 imaging sessions) and *post hoc*
222 confocal imaging; II, neurons were located on edges of main barrels (rows A~E, arcs 1~5); III,
223 neurons were surrounded by two (or more) main barrels. In total, 19 neurons from 3 pups
224 satisfied the above criteria. The same “dendritic trees” and “dendritic segments” were identified
225 from different imaging sessions. A “dendritic tree” is an extension originating from the cell body.
226 A “dendritic segment” was defined as a segment between two branching points, or between a
227 branching point and the origin of the dendrite (or the distal end). A primitive dendritic tree with
228 no branches is also a dendritic segment. Newly formed and eliminated dendrite segments were
229 detected by comparing the dendrite morphologies of the same neuron between two consecutive
230 sessions. A 180°-barrel boundary was used to separate barrel inner and outer sides as in our
231 previous study (Nakazawa et al., 2018). When more than half the length of a dendritic segment of
232 a basal dendrite was located in the barrel inner side, it was defined as an “inner segment”.
233 “Orientation bias index” (OBI) = total inner segment length/total basal dendritic length. The
234 average OBI of 9 imaging sessions from the same neuron was used as the OBI of the neuron.

235 For the quantification of basal dendritic tree and tip dynamics, the number of emerged,
236 eliminated, and transient basal dendritic trees/tips from each neuron in 8 h of imaging was
237 calculated. These dynamic events of dendritic tips were normalized with their corresponding
238 neuron’s total basal dendritic length or total inner or outer segment length (#/millimeter, mm).
239 We classified length changes (μm) of “tip-segments” into 3 subgroups: retraction (R, length
240 change $< -3 \mu\text{m}$), elongation (E, length change $> 3 \mu\text{m}$), and stable (length change within $3 \mu\text{m}$),
241 as previously reported (Mizuno et al., 2014). A “tip-segment” is defined as a dendritic segment

242 whose one end is a dendritic tip. A primitive dendritic tree with no branches is also a tip-segment.
243 For the analysis of dynamics in two consecutive hours, tip-segments that were continuously
244 existing (no formation or elimination) were selected.

245 To analyze the correlation of dendrite tip morphologies with its dynamics, we first
246 investigated all basal dendritic tips of 19 neurons (h0 to h7 of imaging session) and identified 64
247 tips that were apparently thick. Most of these thick parts of the dendritic tips were between 5–10
248 μm in length. Therefore, we used the most distal 5 μm part of the dendritic tip as the tip part (T),
249 skipped the adjacent 5 μm , and then used the subsequent 5 μm as the shaft part (S). By excluding
250 5 μm of the intermediate area from the analyses, we avoided overlaps between the T and S.
251 Regions of interest (ROIs) of T and S were then determined based on the mRFP-signals at the
252 resolution of the single pixel level. Initially, we picked out 126 tips that showed tip morphologies
253 distinguishable from their surrounding environment with a segment length longer than 15 μm and
254 had no overlapping with other dendrites or imaging noise. 79 out of 126 tips were continuously
255 presented, without branching or branch elimination, in two consecutive sessions [between 1 h
256 before (Time -1) and 1 h after (Time +1) of the TTI-checking session (Time 0)]. The mRFP
257 signal intensity (gray value) of ROIs was analyzed with the Image J/FIJI software. The dendritic
258 tip's "tip thickness index" (TTI) = (gray value of T)/(gray value of S). We compared TTI and the
259 length change (μm) between Time -1 and Time +1.

260 **Statistical analysis**

261 The quantification of L4 neurons' basal dendritic dynamics was done with Microsoft Excel
262 (Microsoft 365). The statistical analyses of dynamics comparisons were performed with
263 GraphPad Prism (version 9.3.1). The significance of the comparisons was assessed by the Mann-
264 Whitney test, Wilcoxon matched-pairs signed rank test, or paired t-test. Values are given as mean
265 \pm standard error of the median (SEM). The asterisks in the figures indicate the following: * $p <$
266 0.05, ** $p <$ 0.01, *** $p <$ 0.001, and **** $p <$ 0.0001. When $p >$ 0.05, it indicates ns.

267 **Reanalysis of the data of previous longitudinal *in vivo* imaging**

268 We reanalyzed the P4 *in vivo* imaging data of OBIs from our previous study (Nakazawa et al.,
269 2018). We compared the OBI difference of L4 neurons at P4_M (noon) and P4_L (8 PM) since that
270 time window fits with the time at which we performed *in vivo* imaging in the current study (1 PM
271 to 9 PM at P4). We also analyzed the OBI differences between SS and SP neurons at P4 (the
272 average OBI of P4_M and P4_L). See Table 2.

273

274

275 **Results:**

276 ***In vivo* imaging of developing dendrites with improved spatiotemporal resolution**

277 To visualize the dendritic morphology of neonatal cortical neurons *in vivo*, we used the *in utero*
278 electroporation (IUE)-based Supernova system (Luo et al., 2016; Mizuno et al., 2014), which
279 labels cortical neurons sparsely and brightly for improved signal to background (Figures 1A, 1B).
280 As a fluorescent label, in our previous studies we employed regular RFP (cytosolic RFP, cRFP)
281 (Mizuno et al., 2014; Nakazawa et al., 2018), but the much stronger fluorescence in the cell body
282 than dendrites (Figure 1C) necessitated multiple tracings of the same dendrite at different image
283 contrast settings for complete and accurate reconstruction. To solve this problem and facilitate
284 faster dendrite reconstruction, in the current study we used membrane-bound RFP (mRFP) (Liu et
285 al., 1994; Moriyoshi et al., 1996), which reduced somal relative to dendrite emission and enabled
286 clearer visualization of dendritic morphologies *in vivo* (Figure 1D). This enhanced accuracy and
287 efficiency even enabled detailed detection of dendritic tips, including growth cone-like and
288 filopodium-like structures (Figures 1D¹–1D³), which were subsequently confirmed by *post hoc*
289 high-magnification confocal imaging in brain slices (Figures 1E–1E³).

290 To visualize the barrel map, we used TCA-GFP Tg mice (Mizuno et al., 2014). We
291 labeled L4 neurons in the barrel cortex of these mice with the IUE-based Supernova-mRFP. We
292 then performed craniotomy at P4 (morning) and conducted hourly time-lapse imaging for a total
293 of 8 h [or 9 imaging sessions, hour 0 (h0) to h8, Figures 2A, 2B]. Brain samples were then
294 collected for *post hoc* histological analysis (Figure 2C). For quantitative analysis of dendritic
295 dynamics, we considered only L4 neurons located on barrel edges with clear basal dendritic
296 patterns for all 9 imaging sessions (see Materials and Methods). Tangential views of an example
297 neuron from h0 to h8 sessions and *post hoc* histological imaging are shown in Figures 2D and 2E,
298 respectively. In total, we obtained dendritic images from 19 neurons (Table 1) in three mice
299 satisfying these criteria.

300 This improved spatiotemporal resolution revealed rapid changes in dendrite morphology,
301 including the emergence (*e.g.*, Figures 3A, 3C, 3D) and elimination (*e.g.*, Figures 3B, 3C, 3D) of
302 dendritic trees and branches between two consecutive imaging sessions. These rapid dendrite
303 dynamics detected with hourly imaging were difficult to detect using our previous 8-h interval
304 imaging strategy. For example, in the case of Figure 3C, a dendritic branch that emerged between
305 the first and second (h0 and h1) sessions disappeared between the h2 and h3 sessions. This kind
306 of transient branches were likely missed by our previous 8-h-interval imaging. In the case of
307 Figure 3D, after a branch was eliminated between the h1 and h2 sessions, another branch
308 emerged at a similar position between h2 and h3 sessions. It is highly likely that these two
309 branches at similar locations were recognized as the same branch by lower time-resolution
310 imaging (Figure 3D'). Thus, the high-spatiotemporal resolution achieved using mRFP and shorter
311 acquisition interval in the current study allowed us to conduct more comprehensive mechanistic
312 analyses.

313

314 **Three-dimension (3D) reconstruction of L4 neuron dendritic patterns.**

315 To reveal the detailed dendritic dynamics of barrel cortex L4 neurons at P4, we reconstructed the
316 3D morphology of all dendrites (basal and apical dendrites) from 19 neurons (the example neuron
317 in Figure 4 is the same neuron as shown in Figure 2D). After tracing and reconstructing dendritic
318 patterns of the same neuron from the 9 imaging sessions, we simplified the temporal patterns by
319 drawing schematics for more efficient evaluation of morphological changes (Figure 4).

320 We identified the same dendritic trees, which are extensions originating from the cell
321 body, across imaging sessions by comparing morphologies in the 3D view (Figure 4A). Then,
322 tree IDs (tr X) were assigned to the individual basal dendritic trees of each neuron (Figure 4B).
323 The same trees from each imaging session were arranged in the same rows in the schematics (left
324 to right, h0 to h8 imaging sessions). The same “dendritic segments” (see Materials and Methods)
325 were also identified from different imaging sessions and placed at the same angles. We then
326 classified individual dendritic segments into barrel-inner and -outer according to location as
327 previously described (Nakazawa et al., 2018). Inner and outer dendritic segments accounted for
328 all or the majority of segments belonging to the barrel-inner and outer half, respectively. If a
329 dendritic tree extends along the barrel border, it is possible that it includes both inner and outer
330 dendritic segments (*e.g.*, tr 3 in Figure 4B).

331 By comparing the same dendritic trees and dendritic segments in schematics, we were
332 able to easily identify newly emerged (arrowheads in schematics) and eliminated dendritic trees
333 and branches. These schematics were useful for characterizing detailed changes in dendritic
334 patterns during the 8 h of *in vivo* imaging.

335

336 **Basal dendritic dynamics of L4 neurons in neonatal barrel cortex.**

337 We first analyzed the general properties of dendritic tree dynamics *in vivo*. The total number of
338 basal dendritic trees per neuron was rather stable during the 8 h of imaging (Figure 5A). To

339 investigate dynamics of L4 neuron dendritic trees systematically, we quantified emerged,
340 eliminated and transiently appearing (emerged and then eliminated within 8 h) trees across the 9
341 imaging sessions. In the temporal window, 3.32 ± 0.45 trees emerged and 3.32 ± 0.46 were
342 eliminated. Notably, a substantial number (1.68 ± 0.28) were transient (Figure 5B). Further,
343 newly emerged trees were often short lived, and about 20% (6 out of 33) disappeared by the next
344 imaging session (Figure 5C), underscoring the importance of the improved temporal resolution.

345 Next, we analyzed dendritic tip dynamics of L4 neurons. The total number of basal
346 dendritic tips from each neuron did not change substantially over 8 h (Figure 5D). We then
347 normalized the numbers of basal dendritic tips that emerged, were eliminated, or appeared
348 transiently in 8 h by the total basal dendritic length (see Materials and Methods) and again found
349 similar densities of emergence and elimination (34.62 ± 5.01 vs. 32.44 ± 3.73 per mm) over 8 h
350 (Figure 5E). There were 21.50 ± 3.00 (per mm) transient tips and about half (109 out of 214) of
351 the newly emerged tips disappeared by the next imaging session (Figure 5F).

352 We then grouped basal dendritic tips into three categories, retraction (R: shortened more
353 than $3 \mu\text{m}$), elongation (E: extended longer than $3 \mu\text{m}$), and stable (S: length changes less than 3
354 μm) (see Materials and Methods). Roughly 28% of tips retracted during the time window, 28%
355 elongated, and 43% were stable (Figure 5G). An important parameter of dendritic dynamics is
356 how often dendritic tips switch between E and R behavior. We quantified these events for two
357 consecutive imaging sessions as follows. When a tip continued to elongate across two
358 consecutive sessions, it was categorized as “Elongation-Elongation (EE)”, and when a tip became
359 shorter across two consecutive hours it was categorized as “Retraction-Retraction (RR)”. Further,
360 when a tip was elongated at one measurement time and shorter at the next, it was categorized as
361 “Elongation-Retraction (ER)” and vice versa. The overall frequency of switching events (ER &
362 RE) was higher than the frequency of stability (EE & RR) over two consecutive sessions (Figure
363 5H). This result further highlights the highly dynamic nature of L4 neuron basal dendrites at P4.

364

365 **Comparison of short-term dynamics between the inner and outer basal dendrites.**

366 Although L4 neurons in barrel cortex have both inner and outer basal dendrites, these neurons
367 receive inputs primarily onto inner basal dendrites because TCAs densely innervate the barrel
368 center but not the septa (Figure 2C). Dendritic refinement largely relies on TCA inputs (Harris &
369 Woolsey, 1981; Li et al., 2013; Mizuno et al., 2018; Nakazawa et al., 2018; Nakazawa et al.,
370 2020; Narboux-Nême et al., 2012), so the spatial bias of TCA inputs may drive dendritic
371 dynamics. The numbers of inner dendritic trees and tips did not change markedly during 8 h of
372 imaging (Figures 6A, 6B). Similarly, the number of outer dendritic trees and tips did not change
373 substantially (Figures 6A, 6B). Moreover, the numbers of newly emerged, eliminated, and
374 transient trees were similar between the inner and outer of barrels over 8 h (Figure 6C). The
375 numbers of dendritic tips (per total inner or outer dendritic length) that emerged, were eliminated,
376 and transiently appeared within 8 h did not differ significantly between inner and outer dendritic
377 fields (Figure 6D). Furthermore, the change in tip length across two consecutive imaging sessions
378 was also similar between inner and outer fields (Figure 6E). Thus, basal dendritic trees and tips
379 demonstrated comparable short-term dynamics between inner and outer dendrites of barrel cortex
380 L4 neurons at P4.

381 In the mouse barrel cortex L4, excitatory neurons are classified as SS or star pyramid
382 (SP) neurons. Our IUE-based Supernova labels both types of neurons, although the majority of
383 labeled neurons are of the SS type (Nakazawa et al., 2018). The barrel center-oriented
384 asymmetric basal dendritic pattern is a characteristic of SS neurons but not SP neurons,
385 suggesting potential differences in short-term dynamics between inner and outer basal dendrites,
386 especially those of SS neurons. In the adult brain, SP neurons are distinguished by the presence of
387 apical dendrites, which are lacking in SS neurons (Scala et al., 2019; Staiger et al., 2004). On the
388 other hand, the majority of SS neurons at P4 still have the apical dendrite (Callaway & Borrell,

389 2011; Nakazawa et al., 2018) and thus are difficult to distinguish from SP neurons. However, our
390 previous longitudinal *in vivo* imaging study conducted between P3 and P6 revealed that SS
391 neurons exhibit a higher orientation bias index (OBI, see Materials and Methods) than SP neurons
392 even before the initiation of apical dendrite retraction (Nakazawa et al., 2018). Therefore, we here
393 re-evaluated all imaged neurons (14 SS and 7 SP neurons) in our previous longitudinal imaging
394 study and found that OBIs were significantly higher in SS than SP neurons at P4 [0.417–0.858
395 (median = 0.730) vs 0.416–0.580 (median = 0.473)] ($p = 0.006$, Mann-Whitney test) (Table 2).
396 Based on this finding, we selected only neurons with High-OBI (> 0.600 : 9 neurons) as
397 prospective SS neurons from 19 neurons that were imaged in the current study (Table 1). Notably,
398 4 out of 9 High-OBI neurons had no apical dendrite or an extremely short ($\sim 30 \mu\text{m}$) apical
399 dendrite, which is a characteristic of mature SS neurons. We then compared the dynamics
400 between inner and outer basal dendrites of these High-OBI neurons and found that the numbers of
401 trees and tips of basal dendrites that emerged, disappeared, or were transient were still similar
402 between inner and outer dendrites. Thus, even within the High-OBI neuron group (*i.e.*,
403 prospective SS neurons), there were no differences in the dynamics between inner and outer
404 dendrites (Figures 6F, 6G). Furthermore, the length change in 1 h was also similar between inner
405 and outer tips of High-OBI neurons (Figure 6H).

406 We also compared basal dendritic dynamics between High- and Low-OBI neurons
407 (defined as lower than 0.500, 8 out of 19 neurons, Table 1), while neurons with intermediate
408 OBIs (0.500–0.600, 2 neurons) were excluded. All Low-OBI neurons had substantially longer
409 apical dendrites, suggesting that these were SP or less mature SS neurons. High-OBI neurons
410 showed fewer newly emerged and transient basal dendritic trees, compared with Low-OBI
411 neurons (Figure 7A). A similar tendency was observed in the number of eliminated dendritic trees,
412 although the difference was not significant ($p = 0.106$). These results indicate that High-OBI
413 neurons produce fewer trees than Low-OBI neurons in a short time window. In other words,

414 High-OBI neurons were more stable than Low-OBI neurons. We next compared the dynamics of
415 basal dendritic tips between High- and Low-OBI neurons. There were fewer emergent and
416 transient dendritic tips (per mm) among High-OBI neurons compared to Low-OBI neurons
417 (Figure 7B). The number of eliminated dendritic tips also showed a similar tendency for lower
418 dynamics, although the difference was not significant ($p = 0.059$). It is likely that both of
419 prospective SP and less mature SS neurons are categorized as Low-OBI neurons at P4. In fact, in
420 the re-evaluation of our previous longitudinal imaging data, 1 out of 14 SS neurons and 5 out of 7
421 SP neurons showed low OBI (< 0.500) at P4. Thus, these results from the current imaging may
422 suggest that mature SS neurons have less dynamic basal dendrites than SP and less mature SS
423 neurons.

424 In addition to these quantitative analyses, we also carefully examined the behavior of
425 individual basal dendritic trees located around barrel edges in High-OBI neurons, including
426 whether the growth direction of dendritic tips is re-oriented from outward to inward at the barrel
427 edge and if basal dendrites form more branches inward than outward. Examination of all basal
428 dendritic trees (43 trees from 9 High-OBI neurons) expanded around the barrel edges revealed
429 neither reorientation at the barrel edge nor more frequent inward orientation (see example in
430 Figure 7C). There were no detectable differences in dynamics or turning direction between inner
431 and outer tips. Both inner and outer tips demonstrated similar behaviors such as retraction,
432 elongation, elimination, and emergence. These results suggest that basal dendritic asymmetry is
433 generated not through directed outgrowth or reorientation of dendrites but through trial-and-error
434 emergence, elongation, elimination, and retraction of dendritic branches and trees.

435

436 **The tip features of basal dendrites are correlated with behavior.**

437 Taking advantage of the high spatiotemporal resolution of our *in vivo* imaging system, we
438 examined if basal dendrite tip thickness represents its behavior. Indeed, elongating dendritic tips
439 were often thicker (red arrow in Figure 8A), while retracting tips were often thinner (red arrow in
440 Figure 8B). We analyzed these associations quantitatively by calculating correlations. For this
441 analysis, we considered only dendritic tips with clear morphology and measured the mRFP signal
442 intensity in the tip (T) and shaft (S) parts to calculate a “tip thickness index” (TTI) by dividing T
443 by S (examples as in Figure 8C, also see Materials and Methods). We found that TTI was
444 positively correlated with tip-segment length changes in 2 h (between 1 h before and 1 h after the
445 tip thickness quantification) (Figure 8D). In other words, dendritic tips with smaller TTIs, which
446 showed thinner tips in morphology, tended to be retracting while thicker tips with larger TTIs
447 tended to be elongating. We also found that the continuously elongating (EE) tips in two
448 consecutive hours had significantly larger TTIs than those of continuously retracting (RR) tips at
449 the middle time point of extension or retraction (Figure 8E).

450 In *Drosophila* larvae, dendritic pruning of C4da sensory neurons is mediated by severing
451 and degeneration, in which the proximal part of a dendrite is severed and subsequently the parts
452 of dendrites that are distal from the severed point are degenerated and disappeared (Jan & Jan,
453 2010; Kanamori et al., 2013; Kuo et al., 2005). To examine whether a similar
454 severing/degeneration mechanism is involved in basal dendrite retraction of mouse neonatal
455 cortical neurons, we focused on dendritic tips that were retracted between h7 and h8 sessions. We
456 carefully observed both *in vivo* images of neurons at the h8 session and high magnification
457 confocal images of the same neurons in the *post hoc* fixed brain slices and examined whether
458 there are debris-like structures at sites where the dendritic tip disappeared between the h7 and h8
459 sessions. No such debris-like structures were found in 41 instances of late tip disappearance
460 (including 10 cases from the outer basal dendrites of High-OBI neurons) (examples as in Figure

461 9). Thus, we found no evidence for the possibility that dendritic retraction in barrel cortex L4
462 neurons involves dendrite severing and degradation.

463

464

465 **Discussion:**

466 **Short-term dynamics of dendrites in the neocortex during developmental refinement.**

467 In the current study, we analyzed the short-term dendritic dynamics of L4 neurons in the mouse
468 barrel cortex during neonatal development. *In vivo* imaging at 1-h interval allowed us to precisely
469 identify the same dendritic trees and branches across imaging sessions (Figures 2–4), which were
470 often difficult in previous 8-h interval imaging (Nakazawa et al., 2018). In addition to greater
471 temporal resolution, this study used a membrane-bound RFP, which enabled the precise
472 visualization of dendrite morphology *in vivo* without laborious contrast adjustments (Figure 1).
473 These improvements in spatiotemporal resolution allowed us to detect detailed morphological
474 changes at the level of “individual” dendritic trees and branches in the neonatal cortex (Figures 3
475 and 4). We found that many dendritic trees and branches (tip-segments) emerged and were
476 eliminated within 8 h (Figures 5A–5F). In addition, dendritic tips often changed direction from
477 elongation to retraction or from retraction to elongation between two consecutive imaging
478 sessions (Figures 5G, 5H). Thus, basal dendrites of barrel cortex L4 neurons are highly dynamic
479 during the refinement period in neonatal development.

480 Traditional studies of dendrite dynamics in the developing mammalian brain have used *in*
481 *vitro* systems such as pyramidal neurons in acute slices of the mouse neocortex or cultured
482 cerebellar Purkinje cells of the mouse (Fujishima et al., 2012; Jontes & Smith, 2000; Portera-
483 Cailliau et al., 2003). *In vivo* studies mostly used optic tectal neurons of *Xenopus* tadpole and
484 zebrafish larvae, whose brains are transparent, and focused primarily on how dendrites arborize

485 (Nagel et al., 2015; Niell et al., 2004; Rajan et al., 1999; Sanchez et al., 2006; Sin et al., 2002; Wu
486 & Cline, 1998; Wu et al., 1999). The optic tectal neuron has a single dendrite. In contrast, SS
487 neurons, the major type of L4 glutamatergic neurons (65%–80% of the total) (Fox, 2008), in the
488 mouse barrel cortex have multiple basal dendrites that exhibit an asymmetric pattern strongly
489 oriented toward the barrel center in adulthood, which is the basis of the precise one-to-one
490 relationship between individual whisker stimulation and barrel activation (Lübke et al., 2000;
491 Staiger et al., 2004). Therefore, it is particularly important to determine how the asymmetric
492 dendritic patterns of SS neurons are generated.

493 It is generally assumed that SS neuron dendrites show symmetric patterns in early
494 neonatal stages and are refined to acquire asymmetric patterns by simply eliminating outer
495 dendrites and adding new inner dendrites and/or elaborating existing inner dendrites during later
496 neonatal stages (Emoto, 2011; Espinosa et al., 2009; Greenough & Chang, 1988; Iwasato, 2020).
497 We have challenged this conventional view by developing *in vivo* imaging approaches for the
498 neonatal mouse cortex to analyze dynamic processes of SS neuron dendritic refinement (Iwasato,
499 2020). As a first step, we previously conducted *in vivo* imaging of L4 neuron dendrites starting at
500 P5 (9-h interval for 18 h) (Mizuno et al., 2014), which provides the first observation of dendritic
501 motility in the mammalian brain *in vivo*. This study revealed that branches of inner dendrites are
502 not only elongated but also often retracted. Similarly, branches of outer dendrites show both
503 elongation and retraction. Our recent longitudinal *in vivo* imaging of L4 neurons in the mouse
504 barrel cortex revealed that SS neurons establish basal dendritic orientation bias through two
505 phases (Nakazawa et al., 2018). During Phase I (by P3), SS neurons produce a larger number of
506 inner basal dendritic trees than outer trees, although both inner and outer trees are similarly
507 primitive in morphology at P3. During Phase II (between P3 and P6), the ratio of inner to outer
508 trees does not change. But individual dendritic trees show extensive turnover. Both inner and
509 outer dendritic trees often disappear quickly. Meanwhile, only a few trees highly elaborate. These

510 “winner” trees emerged specifically from inner trees, which generates a strong orientation bias of
511 SS neuron basal dendrites. Although that study revealed the long-term dynamics that generate
512 orientation bias, the spatiotemporal resolution was not sufficient to capture short-term dynamics.
513 In that study, to cover the whole refinement processes between P3 and P6, we set acquisition
514 intervals of 8 or 24 hours. However, the changes in dendrite morphology over 8 h are larger than
515 we initially expected, which hindered the detection of rapid changes over shorter periods. For
516 example, temporal resolutions of these studies were not high enough to quantitatively analyze the
517 frequency of emergence and elimination of dendritic branches.

518 In the current study, we focused on P4, which is the peak time of Phase II (Nakazawa et
519 al., 2018), and asked whether there were differences in short-term dynamics between inner and
520 outer basal dendrites. On the other hand, at this age, it is often difficult to distinguish SS and SP
521 neurons because many SS neurons still have apical dendrites (Nakazawa et al., 2018). Therefore,
522 we also compared dynamics between inner and outer basal dendrites of High-OBI neurons by
523 assuming that the majority of these neurons is prospective SS neurons (see Results for the detail).
524 In either of all-L4-excitatory-neuron population or High-OBI neuron population, both inner and
525 outer trees emerged and were eliminated with no marked difference in frequency (Figure 6).
526 Similarly, inner and outer branches showed comparable frequencies of emergence and
527 elimination and similar behaviors of elongation and retraction (Figure 6). Thus, although basal
528 dendritic trees and branches of SS neurons were highly dynamic during Phase II, rapid changes in
529 basal dendritic structure did not directly contribute to the formation of asymmetric basal dendritic
530 patterns of SS neurons. Taken all the data from our current and previous *in vivo* imaging studies
531 together, we propose that SS neurons establish highly asymmetric dendritic patterns through
532 extensive trial-and-error emergence, elongation, elimination, and retraction of dendritic trees and
533 branches rather than simple emergence/elongation of inner dendrites and elimination/retraction of
534 outer dendrites.

535 In the current study, we also found that the basal dendritic trees and tips of High-OBI (>
536 0.600) neurons were less dynamic (*i.e.*, exhibited fewer emergent and transient events over the 8-
537 h imaging period) compared to Low-OBI (< 0.500) neurons (Figures 7A, 7B). It is likely that the
538 majority of High-OBI neurons are of the SS type while Low-OBI neurons may include SP
539 neurons and less mature SS neurons (See Results for the detail). Our previous study shows that
540 dendritic dynamics at P5 are increased in the absence of *N*-Methyl-D-aspartic acid (NMDA)
541 receptor activity (Mizuno et al., 2014). Because NMDA receptor is a key player in dendritic
542 refinement (Datwani et al., 2002; Espinosa et al., 2009; Mizuno et al., 2014), it is possible that
543 dendritic dynamics are reduced with the progress of dendritic refinement.

544 A caveat of our short-interval imaging is possible effects of anesthesia on activity-
545 dependent dendritic refinement. During early postnatal stages, sensory cortices show spontaneous
546 activities characterized by correlated firing that is arranged spatially according to the modality,
547 and this correlated activity is important for neuronal circuit refinement (Nakazawa & Iwasato,
548 2021). In the neonatal barrel cortex, spontaneous activity shows a barrel-corresponding
549 patchwork-typed spatial pattern, and this activity is blocked by general anesthesia with isoflurane
550 (Mizuno et al., 2018). Patchwork spontaneous activity is detected again in 5~7 minutes after
551 stopping the isoflurane supply and fully recovers within 10 minutes (See Fig. S1D of (Mizuno et
552 al., 2018)). Similarly, spontaneous activity in the neonatal mouse visual cortex is recovered
553 within 10 minutes after stopping isoflurane inhalation (Ackman et al., 2012). In the experiment
554 setting of the current study, we started and stopped isoflurane supply 2 minutes before the start of
555 each imaging session and 2 minutes before the end of the imaging session, respectively. After
556 finishing each imaging session (20 minutes), we confirmed that pups woke up and started to
557 interact with littermates within 5 minutes. Therefore, pups were anesthetized for nearly 30
558 minutes in one hour between two consecutive imaging sessions. Thus, it is likely that short-term
559 dendritic dynamics of mouse cortical neurons is even faster than what we observed here.

560

561 **Specific dendritic features during development revealed by improved spatiotemporal**
562 **resolution of *in vivo* imaging.**

563 The high spatiotemporal-resolution imaging achieved in the current study also enabled us to
564 examine correlations between morphological features and dendritic tip dynamics (Figure 8). We
565 found a positive correlation between dendritic tip thickness and its behavior (Figure 8D). Thick
566 tips, which may have dendritic growth cones (Figure 9), were more likely to be extending (Figure
567 8A) and thin tips were more likely to be retracting (Figure 8B). Conversely, extending and
568 retracting tips are more likely to be thick and thin, respectively (Figure 8E). Similar correlations
569 between dendritic tip morphology and dendrite behavior are observed in optic tectal neurons in
570 *Xenopus* tadpoles (Hossain et al., 2012; Wu & Cline, 1998) and cortical pyramidal neurons in
571 mouse acute slices (Portera-Cailliau et al., 2003). Thus, it is possible to assume dendritic tip
572 behavior based on dendritic tip morphology revealed by two-photon *in vivo* imaging or confocal
573 imaging of fixed brain slices.

574 As mentioned above, before *in vivo* time-lapse imaging was achieved, it was generally
575 believed that SS neurons initially have radially oriented basal dendrites and during critical period
576 of dendritic refinement these neurons specifically prune the outer dendrites to acquire highly
577 asymmetric dendritic patterns (Emoto, 2011; Mizuno et al., 2014). If so, it could be similar to
578 class IV dendritic arborizing (C4da) sensory neurons in *Drosophila*, which prune their dendritic
579 arbors during the early metamorphosis. In this pruning process, dendrites are severed and
580 fragmented, and fragments are finally cleared by engulfment (Kanamori et al., 2013; Kuo et al.,
581 2005; Williams et al., 2006; Williams & Truman, 2005). It was wondered whether similar
582 mechanisms may be involved in SS neuron dendritic refinement. By taking advantage of high
583 spatiotemporal resolution of the current imaging, we here carefully observed loci where dendritic
584 segments disappeared within the last 1 h (h7 to h8) combined with *post hoc* confocal imaging of

585 the same loci with high magnification. However, we found no evidence of dendrite severing and
586 fragmentation. This result suggests that the outer basal dendritic trees are unlikely to be pruned
587 from SS neurons in a process akin to that observed for *Drosophila* class IV C4da neurons. In
588 accord with this finding, our previous longitudinal imaging study reveals that SS neurons
589 establish a highly oriented basal dendritic pattern between P3 and P6 without selective
590 elimination of outer dendritic trees. In fact, the numbers of both inner and outer basal dendrites do
591 not differ substantially between P3 and P6 (Nakazawa et al., 2018). Rather, SS neurons establish
592 unique basal dendritic patterns highly oriented toward the barrel center through selection and
593 elaboration of a few inner dendritic trees as “winners”.

594 In the current study, we focused our analyses on the short-term dynamics of barrel cortex
595 L4 neurons at P4, the time of peak refinement. Taking our current results and previous
596 longitudinal imaging together, we have revealed highly dynamic features of basal dendrites in the
597 early postnatal period. Rapid dendritic dynamics do not directly contribute to the highly
598 asymmetric basal dendritic pattern of L4 neurons. Instead, dendritic refinement progresses
599 through massive trial-and-error emergence, elongation, elimination, and retraction of dendritic
600 tips and trees.

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770 **Figure legends**

771 **Figure 1: Improvement in the spatial resolution of cortical neuron *in vivo* imaging.**

772 **(A)** The Supernova membrane-bound RFP (mRFP) vector set (Vector 1 and Vector 2) used for *in*
773 *utero* electroporation (IUE)-based sparse labeling of barrel cortex layer 4 (L4) neurons. Vector 3
774 was used to identify mice in which neurons in the barrel field were appropriately labeled before
775 the cranial window surgery because dense BFP labeling by Vector 3, but not Supernova-mRFP,
776 was detectable over the skull.

777 **(B)** Tangential stacks of Supernova-mRFP-labeled L4 neurons in the thalamocortical axon
778 (TCA)-GFP transgenic (Tg) mouse at postnatal day (P) 4. Left to right, the barrel map visualized
779 with TCA-GFP Tg labeling, Supernova-mRFP-labeled L4 neurons, BFP-labeled L4 neurons, and
780 the merged image. A: anterior; L: lateral. Scale bar: 100 μm .

781 **(C)** *In vivo* images of an L4 neuron labeled with Supernova-cytosolic RFP (cRFP) at P4 (z-
782 stacked). Scale bar: 50 μm .

783 **(D)** Tangential stacks of *in vivo* images of a Supernova-mRFP-labeled L4 neuron. Scale bar: 50
784 μm .

785 **(D'-D''')** Higher magnification images of the red box regions in (D) showed growth cone-like
786 structures (arrows) and filopodium-like protrusions (arrowheads). Scale bar: 10 μm .

787 **(E)** Tangential stacks of a high magnification confocal image of the neuron shown in (D). Images
788 were taken by a 63x lens with a 2x zoom. Scale bar: 50 μm .

789 **(E'-E''')** Higher magnification images of corresponding areas shown in (D'-D'''). The growth
790 cone-like and filopodium-like morphologies were confirmed (arrows and arrowheads,
791 respectively). Scale bar: 10 μm .

792

793 **Figure 2: One-hour interval *in vivo* time-lapse imaging of L4 neurons in the neonatal barrel**
794 **cortex.**

795 **(A)** Experimental design of time-lapse *in vivo* imaging. TCA-GFP Tg mice were used for
796 visualizing the barrel map. L4 neurons in the TCA-GFP mouse barrel cortex were sparsely
797 labeled with Supernova-mRFP via IUE at embryonic day (E) 14. The cranial window was made
798 at postnatal day (P) 4 morning, and the 1-h-interval *in vivo* imaging started 1 h after the surgery.
799 In every imaging session, the pup for *in vivo* imaging was head-fixed to the imaging stage under a
800 two-photon microscope with light anesthesia. The brain of the *in vivo* imaged mouse was
801 collected after 9 (h0 to h8) imaging sessions, and then fixed. Tangential sections were obtained
802 for *post-hoc* confocal analyses.

803 **(B)** An example of 1-h-interval *in vivo* imaging of Supernova-mRFP-labeled L4 neurons at P4. Z-
804 stacked images from the top view. Scale bar: 100 μm .

805 **(C)** The *post-hoc* confocal image (z-stacked) of the tangential section of the same area in (B). The
806 neuron pointed with a yellow arrow in (C) is the same neuron pointed in h8 session of (B). The
807 same neurons were identified by their relative positions with each other and dendritic patterns.
808 The *in vivo* imaged neurons were located on C1–C4 barrels in this example mouse. The sample
809 orientations in (B) and (C) were similar. A, anterior; L, lateral. Scale bar: 100 μm .

810 **(D)** An example of time-lapse images of dendritic morphologies of an L4 neuron in 8 h of
811 imaging sessions. Top views of an L4 neuron (z-stacked), which was located at the edge of the
812 C1 barrel. Snapshots were taken from the 3D reconstruction software Imaris. The neuron was
813 rotated to the same angle and at the same magnification from 9 imaging sessions. The green line
814 in the h8 imaging session showed the C1 barrel edge. Scale bar: 20 μm .

815 **(E)** The corresponding confocal image (z-stacked) of the neuron (yellow arrow) shown in (D) at a
816 similar angle and magnification. An example neuron is located at the C1 barrel edge. Scale bar:
817 20 μm .

818

819 **Figure 3: Detection of rapid dendritic dynamics with improved spatiotemporal resolution *in***
820 ***vivo* imaging.**

821 **(A)** A basal dendritic tree (arrowhead) that newly emerged between h7 (left) and h8 (right)
822 sessions. Snapshot images taken in the Imaris software. Scale bar: 5 μm .

823 **(B)** A basal dendritic tree that disappeared between h0 (left) and h1 (right) sessions. Snapshot
824 images taken in the Imaris software. Scale bar: 5 μm .

825 **(C)** A transient basal dendritic branch (arrowheads) emerged between h0 and h1 sessions and
826 disappeared between h2 and h3 sessions. Scale bar: 10 μm .

827 **(D, D')** After a branch (red arrowheads) was eliminated between h1 and h2 sessions, another
828 branch (yellow arrowheads) emerged at a similar position between h2 and h3 sessions **(D)**. It is
829 highly likely that these two branches (red and yellow arrowheads) are recognized as an identical
830 dendrite if only h0 and h5 images are available **(D')**. Snapshot images taken in the Imaris
831 software. Scale bar: 5 μm .

832 Examples in (A-C) and (D, D') are from Neuron #1 and Neuron #3 (Table 1), respectively.
833

834 **Figure 4: Reconstruction of dendritic patterns imaged *in vivo*.**

835 **(A)** A 2-dimensional view of 3-dimensionally (3D) reconstructed dendritic morphology from the
836 neuron shown in Figure 2D. Representative basal dendritic trees (tr 2 and tr 7) that originated
837 from the barrel-side half (IN) of the cell body and the other half (OUT), respectively, are labeled
838 in red. Scale bars: 20 μm .

839 **(B)** A schematic that enables simple tracking of dendritic pattern changes during imaging
840 sessions. The representative schematic for the example neuron in (A) is shown. Imaging session
841 numbers (h0 to h8) are shown on the top. The same basal dendritic trees (tr 1–8) were arranged in
842 the same rows in schematics. The same dendritic segments from different imaging sessions were
843 placed at the same angles. Individual dendritic segments were classified into barrel-inner (IN,
844 cyan) and -outer (OUT, magenta) according to their location. The newly emerged inner and outer

845 branches (and trees) were marked with cyan and magenta arrowheads, respectively.
846 The lengths of individual dendritic trees and segments are roughly proportional. Gray lines, apical
847 dendrite. Scale bar: 20 μm .

848

849 **Figure 5: General properties of the dendritic dynamics of L4 neurons at P4.**

850 **(A)** Changes in basal dendritic tree numbers from each neuron during 8 h of imaging. The gray
851 and black lines represent data of individual neurons and the average, respectively. $n = 19$ neurons,
852 3 mice.

853 **(B)** Numbers of emerged, eliminated, and transient trees from each neuron in 8 h of imaging. The
854 “+” represents the mean value and the horizontal line in the boxplot represents the median. Each
855 dot represents one neuron. $n = 19$ neurons, 3 mice.

856 **(C)** Survival ratio of newly emerged trees. In this analysis, only dendritic trees that were first
857 detected at h1, h2, h3, h4, or h5 imaging sessions were used. The imaging session at which the
858 tree was first detected was defined as the time point 0 ($n = 33$ trees). 27 (81.8%), 19 (57.6%), and
859 16 (48.5%) of the 33 trees were still present at time points 1, 2, and 3, respectively. $n = 16$
860 neurons, 2 mice. 3 out of 19 *in vivo* imaged neurons have no trees that emerged between h1 and
861 h5 sessions.

862 **(D)** Changes in dendritic tip numbers from each neuron during 8 h of imaging. The gray lines and
863 black lines represent data of individual neurons and the average, respectively. $n = 19$ neurons, 3
864 mice.

865 **(E)** Numbers of emerged, eliminated, and transient tips (per mm) from each neuron in 8 h of
866 imaging. The “+” represents the mean value and the line in the boxplot represents the median.
867 Each dot represents one neuron. $n = 19$ neurons, 3 mice.

868 **(F)** Survival ratio of the newly emerged dendritic tips. Only dendritic tips that emerged between
869 h1 to h5 of the *in vivo* imaging sessions were used. At the time point 0, $n = 214$ tips; 105 (49.1%),
870 74 (34.6%), and 61 (28.5%) out of 214 dendritic tips were still present at time points 1, 2, and 3,

871 respectively. n = 19 neurons, 3 mice.

872 **(G)** Histogram of dendritic tip length changes in 1 h. x-axis shows the length change (μm), and y-
873 axis shows the event number of corresponding length changes. Length changes larger than 3 μm ,
874 smaller than -3 μm , and between -3 μm and +3 μm were classified as elongation (E, 28.1%),
875 retraction (R, 28.5%), and stable (S, 43.4%), respectively. n = 2089 dendritic tips (from 19
876 neurons, 3 mice).

877 **(H)** Frequencies of tip behavior in which individual tips continued to elongate or retract in
878 consecutive 2 imaging sessions (EE&RR) and those of tip behavior in which dendritic tips
879 changed the motility direction from E to R or R to E (ER&RE) were compared. Other behaviors
880 (ES, RS, SE, SR & SS) were excluded from comparison. The “+” represents the mean value and
881 the horizontal line in the boxplot represents the median. Each dot represents one neuron. p =
882 0.022, Mann-Whitney test. n = 19 neurons, 3 mice.

883

884 **Figure 6: Comparison of dynamic properties between the inner and outer dendrites.**

885 **(A, B)** Changes in the numbers of inner (IN) and outer (OUT) trees (A) and tips (B) from each
886 neuron across 9 imaging sessions. Gray lines represent data for individual neurons. Black lines
887 show the average of all neurons. n = 19 neurons, 3 mice.

888 **(C)** There were no significant differences between IN and OUT trees in the numbers of emerged,
889 eliminated, and transient trees. p = 0.796, 0.556, 0.234, respectively; Wilcoxon matched-pairs
890 signed rank tests; n = 19 neurons, 3 mice. Red lines represent the average.

891 **(D)** There were no significant differences between IN and OUT tips in the numbers of emerged,
892 eliminated, and transient tips (per mm basal dendritic length) (p = 0.738, 0.568, and 0.441,
893 respectively; Wilcoxon matched-pairs signed rank tests; n = 19 neurons, 3 mice). The dendritic
894 tip numbers were normalized with the IN or OUT basal dendritic length of the neuron (average
895 length from 9 imaging sessions). Red lines represent the average.

896 **(E)** There were no significant differences between IN and OUT dendritic tips in 1 h length
897 changes (μm). $p = 0.304$; Mann-Whitney test; $n = 945$ IN and 852 OUT tips (from 19 neurons, 3
898 mice).

899 **(F)** There were no significant differences between IN and OUT basal dendritic trees of "High-
900 orientation bias index (OBI)" neurons in the numbers of emerged, eliminated, and transient trees.
901 $p = 0.648, 0.750, \text{ and } > 0.999$, respectively; Wilcoxon matched-pairs signed rank tests. $n = 9$
902 High-OBI neurons, 3 mice. Red lines represent the average.

903 **(G)** There were no significant differences between IN and OUT dendritic tips of High-OBI
904 neurons in the numbers of emerged, eliminated, and transient tips (per mm IN or OUT basal
905 dendritic length) ($p = 0.820, 0.570, \text{ and } 0.250$, respectively; Wilcoxon matched-pairs signed rank
906 tests. $n = 9$ High-OBI neurons, 3 mice. Red lines represent the average.

907 **(H)** There were no significant differences between IN and OUT tips of High-OBI neurons in 1 h
908 length changes (μm). $p = 0.260$; Mann-Whitney test; $n = 568$ High-IN and 340 High-OUT tips
909 (from 9 High-OBI neurons, 3 mice).

910

911 **Figure 7: Comparison of dynamic properties of dendrites between High- and Low-OBI**
912 **neurons.**

913 **(A)** Numbers of basal dendritic trees that emerged ($p = 0.024$), eliminated ($p = 0.106$), and were
914 transient ($p = 0.030$) were compared between High- and Low-OBI neurons. The "+" represents
915 the mean value and the horizontal line in the boxplot represents the median. Each dot represents
916 one neuron. Mann-Whitney tests, $n = 9$ High- and 8 Low-OBI neurons.

917 **(B)** Numbers of basal dendritic tips that emerged ($p = 0.047$), eliminated ($p = 0.059$), and
918 transient ($p = 0.036$) were compared between High- and Low-OBI neurons. The dendritic tip
919 number was normalized with the total basal dendritic length of the neuron (average length from 9
920 imaging sessions). The "+" represents the mean value and the horizontal line in the boxplot

921 represents the median. Each dot represents one neuron. Mann-Whitney tests. $n = 9$ High- and 8
922 Low-OBI neurons.

923 **(C)** Reconstructed dendritic pattern for an example dendritic tree (yellow) from a High-OBI
924 neuron, which is located near the barrel edge (green). An outward basal dendritic tip did not
925 change its direction or start to retract at the barrel edge and elongated outward further (arrow at
926 h8). Two inward tips were generated near the barrel-edge, but both were eliminated immediately
927 (arrowheads at h1 and h6 sessions). Scale bar: 20 μm .

928

929 **Figure 8: Dendrite tip features represent their behavior.**

930 **(A)** An example image of a dendritic branch (tip-segment) whose tip appeared thick at Time 0
931 (red arrow) was becoming longer and longer between the Time -1 and Time +1 sessions. Red and
932 yellow arrows indicate the distal end of the branch. Snapshot images taken in the Imaris software.
933 Scale bar: 10 μm .

934 **(B)** An example image of a dendritic branch whose tip appeared thin at Time 0 (red arrow) was
935 becoming shorter and shorter between the Time -1 and Time +1 sessions. Red and yellow arrows
936 indicate the distal end of the branch. Snapshot images taken in the Imaris software. Scale bar: 10
937 μm .

938 **(C)** Example diagrams of quantitative analysis for “tip thickness index” (TTI). Left panel: an
939 example for a thin tip, $\text{TTI} = 0.26$. Right panel: an example for a thick tip, $\text{TTI} = 1.68$. The distal
940 part of the tip-segment was divided into 3 units: tip (T), middle, and shaft (S). The length of each
941 unit is 5 μm . Regions of interest (ROIs, the magenta areas) of T and S units were determined
942 according to their morphologies. Snapshot images taken in the Imaris software. Scale bar: 10 μm .

943 **(D)** Quantification of correlations of TTIs and length changes of tip-segments in 2 h (between
944 Time -1 and Time +1). The x-axis is in \log_2 units. $y = 3.689 * x - 4.826$. The black curve represents
945 the fitted curve, and the dashed line represents the 95% confidence interval. Each inverted
946 triangle represents individual dendritic tips. $n = 79$ tips, from 14 neurons, 3 mice.

947 **(E)** TTIs of tip-segments that continued to extend (EE) were compared with TTIs of tip-segments
948 that continued to retract (RR) in 2 consecutive hours. In this analysis, TTIs were calculated from
949 the middle time points of continuous elongation or retraction. The “+” represents the mean value
950 and the horizontal line in the boxplot represents the median. Each dot represents one neuron. $p =$
951 0.003, Mann-Whitney test. $n = 5$ EE and 7 RR tip-segments, from 5 neurons, 2 mice.

952

953 **Figure 9: Observation of loci immediately after dendrite retraction.**

954 **(A)** Three representative basal dendritic tips retracted between h7 and h8 sessions. The red and
955 yellow arrowheads show positions of basal dendritic tips in h7 and h8 sessions, respectively.

956 Snapshot images taken in the Imaris software. Scale bars: 10 μm .

957 **(B)** *Post-hoc* (126x) confocal z-stack images corresponding to images in (A). The red and yellow
958 arrowheads show positions of dendritic tips in h7 and h8 sessions, respectively. Brain slices: 100

959 μm -thick. Scale bars: 10 μm .

960 **(B')** The higher magnification images of red box areas in (B). Scale bars: 5 μm .

961 **Table 1. Summary of 19 neurons that were used for statistical analyses.**

Neuron ID	Barrel ^{a)}	Distance ^{b)}	OBI ^{c)}	Type ^{d)}	AD ^{e)}	AD length (μm)
#1	C1	12.8	0.689	High	–*	29.7
#2	B2	10.6	0.415	Low	+	157.6
#3	C3	2.7	0.464	Low	+	190.1
#4	C2	9.9	0.475	Low	+	178.0
#5	C2	0	0.386	Low	+	486.8
#6	C4	5.8	0.747	High	+	234.6
#7	C3	2.4	0.714	High	–	0
#8	B1	11.8	0.666	High	+	187.4
#9	B2	11.7	0.616	High	+	189.6
#10	B4	3.3	0.545	Medium	+	267.1
#11	C5	4.2	0.562	Medium	+	124.7
#12	B3	9.9	0.465	Low	+	435.8
#13	B3	19.7	0.757	High	–	0
#14	B3	2.7	0.454	Low	+	679.8
#15	B2	16.1	0.693	High	–	0
#16	C2	1.7	0.644	High	+	225.7
#17	B2	7.4	0.381	Low	+	229.8
#18	C2	19.3	0.681	High	+	447.5
#19	C2	14.0	0.367	Low	+	207.8

962 a) The barrel column which the neuron belongs to.

963 b) The distance (μm) from the center of cell body to the barrel edge.

964 c) Mean OBI (orientation bias index: see Materials and Methods) of 9 imaging sessions of the

965 neuron.

966 d) High: High-orientation bias index (OBI) neuron, whose OBI was >0.6 , Low: Low-OBI neuron,
 967 whose OBI was <0.5 . Medium: Neurons whose OBI were >0.5 and <0.6 were not included in
 968 either High- or Low-OBI groups.

969 e) AD: apical dendrite. “+” and “-” mean presence and absence of the apical dendrite,
 970 respectively. *Note that “Neuron #1” has only an extremely short ($\sim 30 \mu\text{m}$) apical dendrite.

971

972 **Table 2. Reanalysis of neurons that were analyzed in previous longitudinal *in vivo* imaging.**

Neuron ID	OBI	Neuron type
#a	0.712	SS
#b	0.869	SS
#c	0.857	SS
#d	0.808	SS
#e	0.795	SS
#f	0.752	SS
#g	0.747	SS
#h	0.753	SS
#i	0.629	SS
#j	0.575	SS
#k	0.560	SS
#l	0.516	SS
#m	0.503	SS
#n	0.417	SS
#o	0.580	SP
#p	0.563	SP
#q	0.497	SP
#r	0.473	SP

#s	0.439	SP
#t	0.430	SP
#u	0.416	SP

973

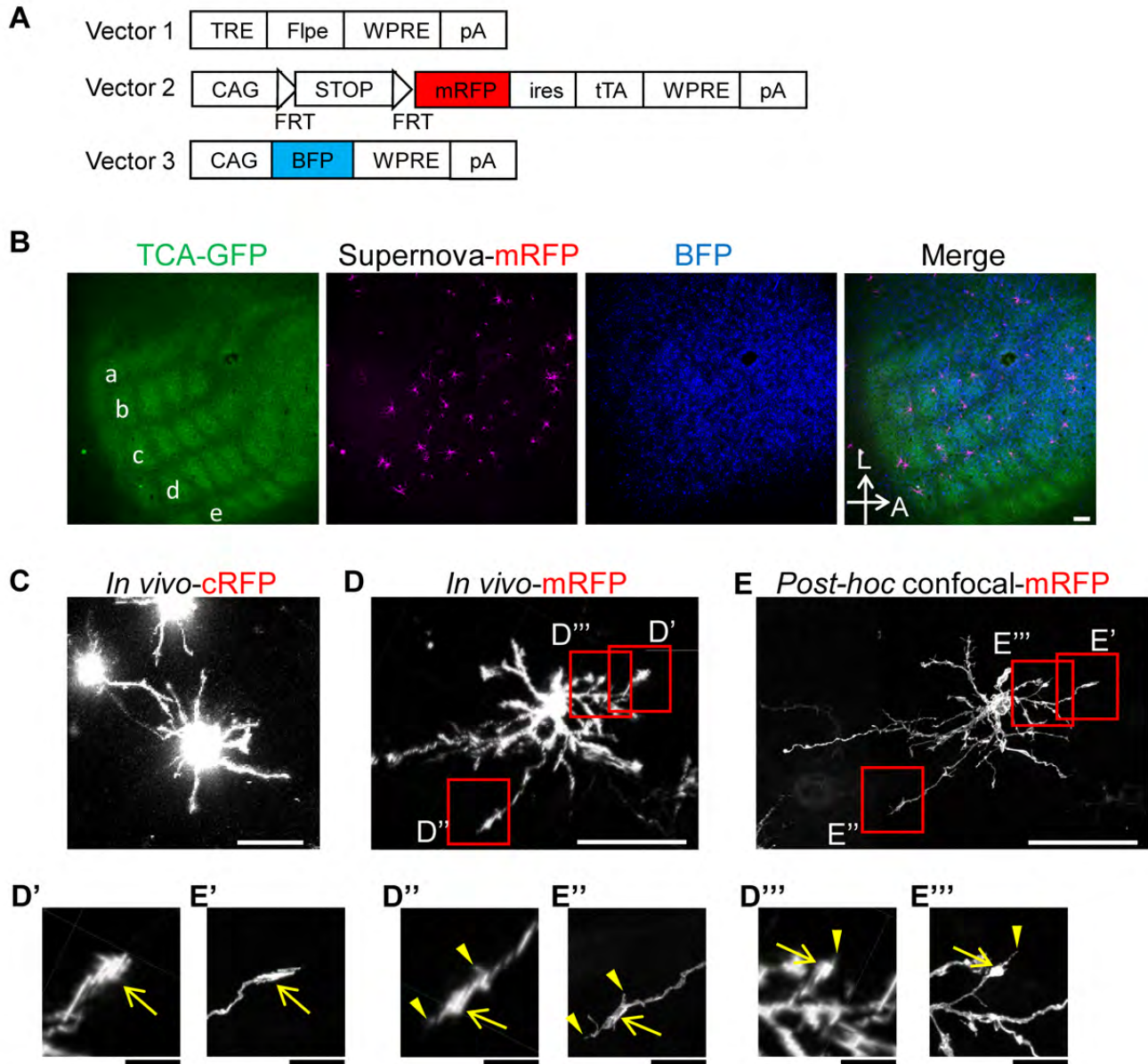


Figure 1

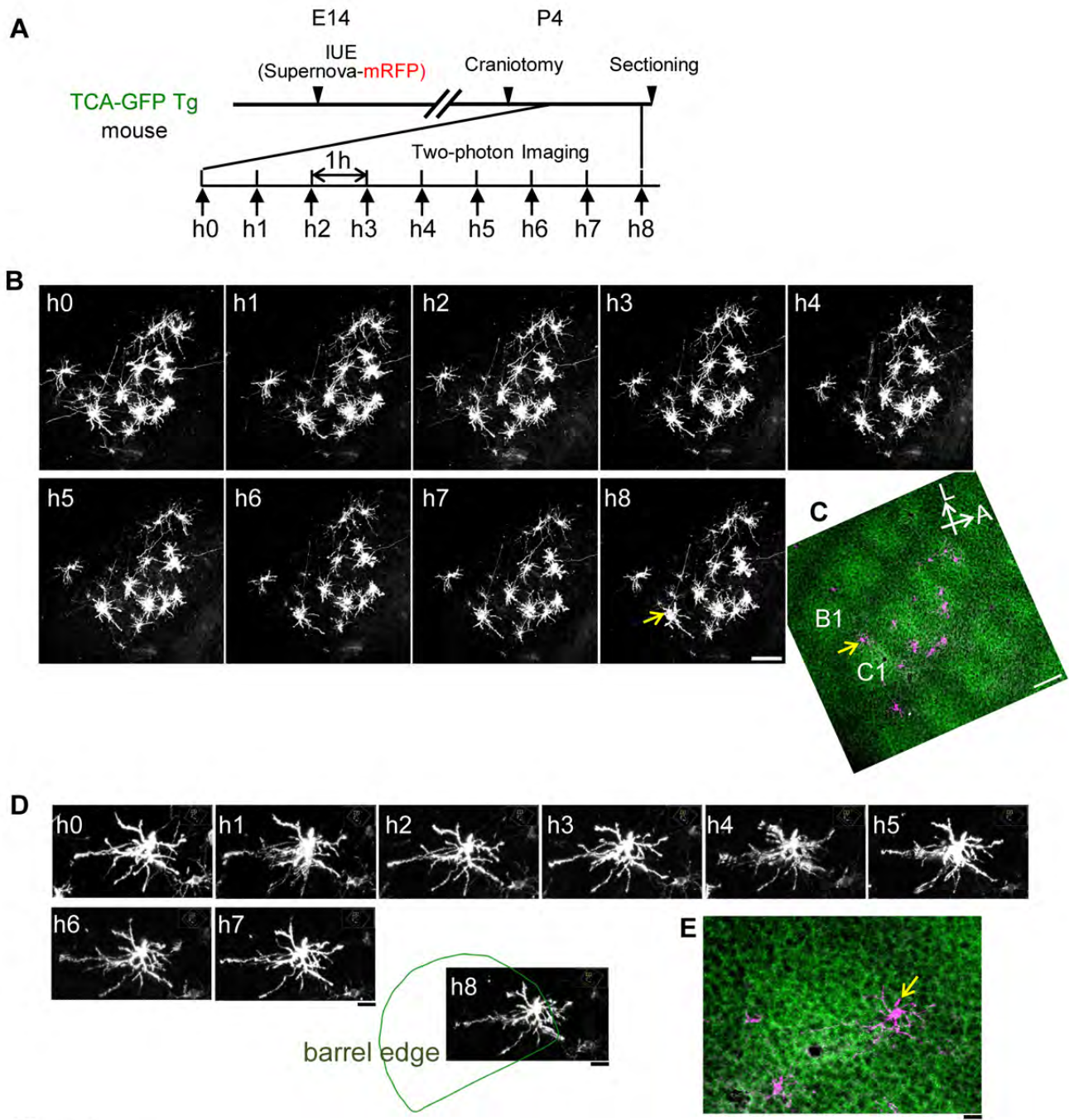


Figure 2

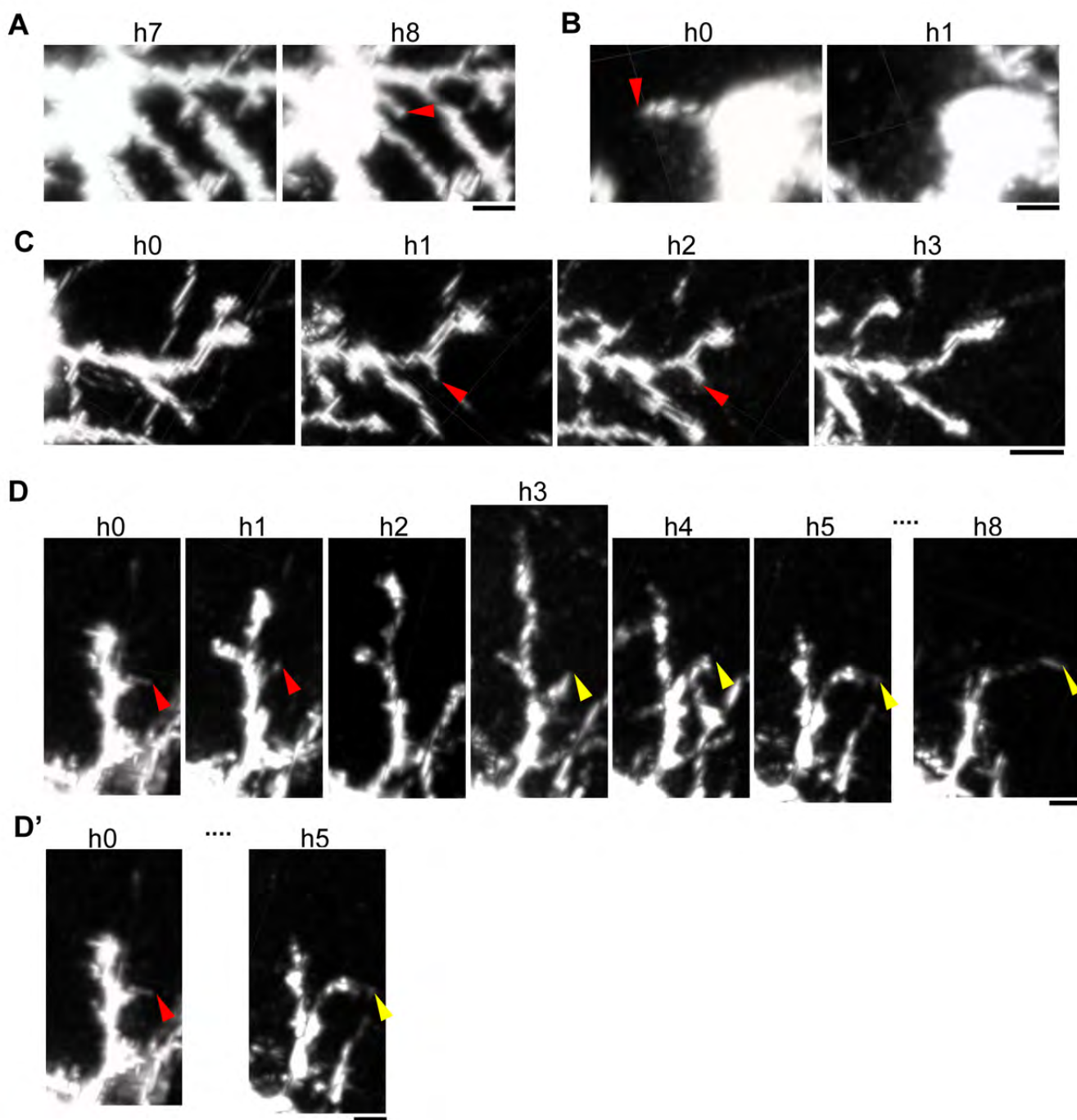


Figure 3

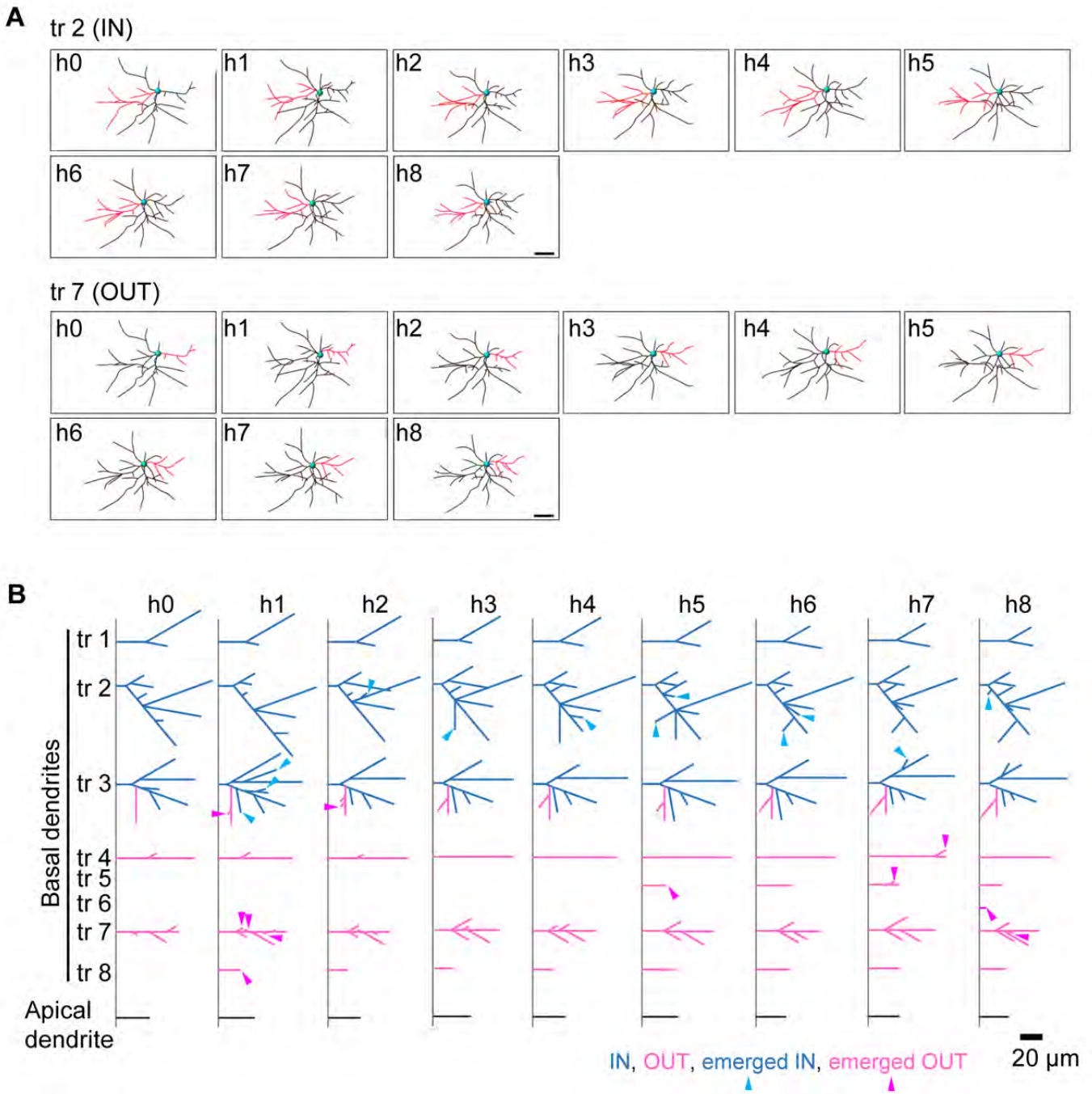


Figure 4

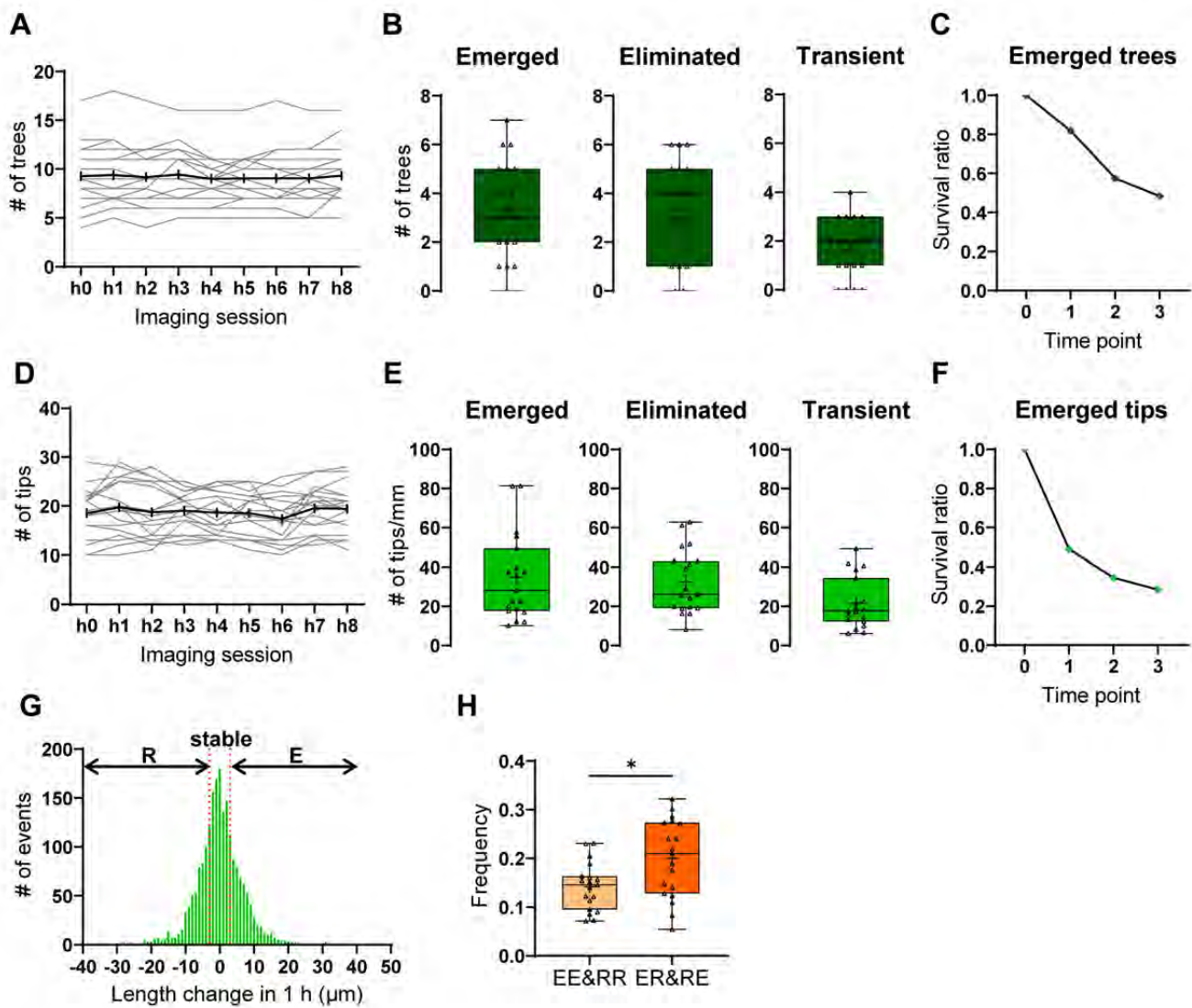


Figure 5

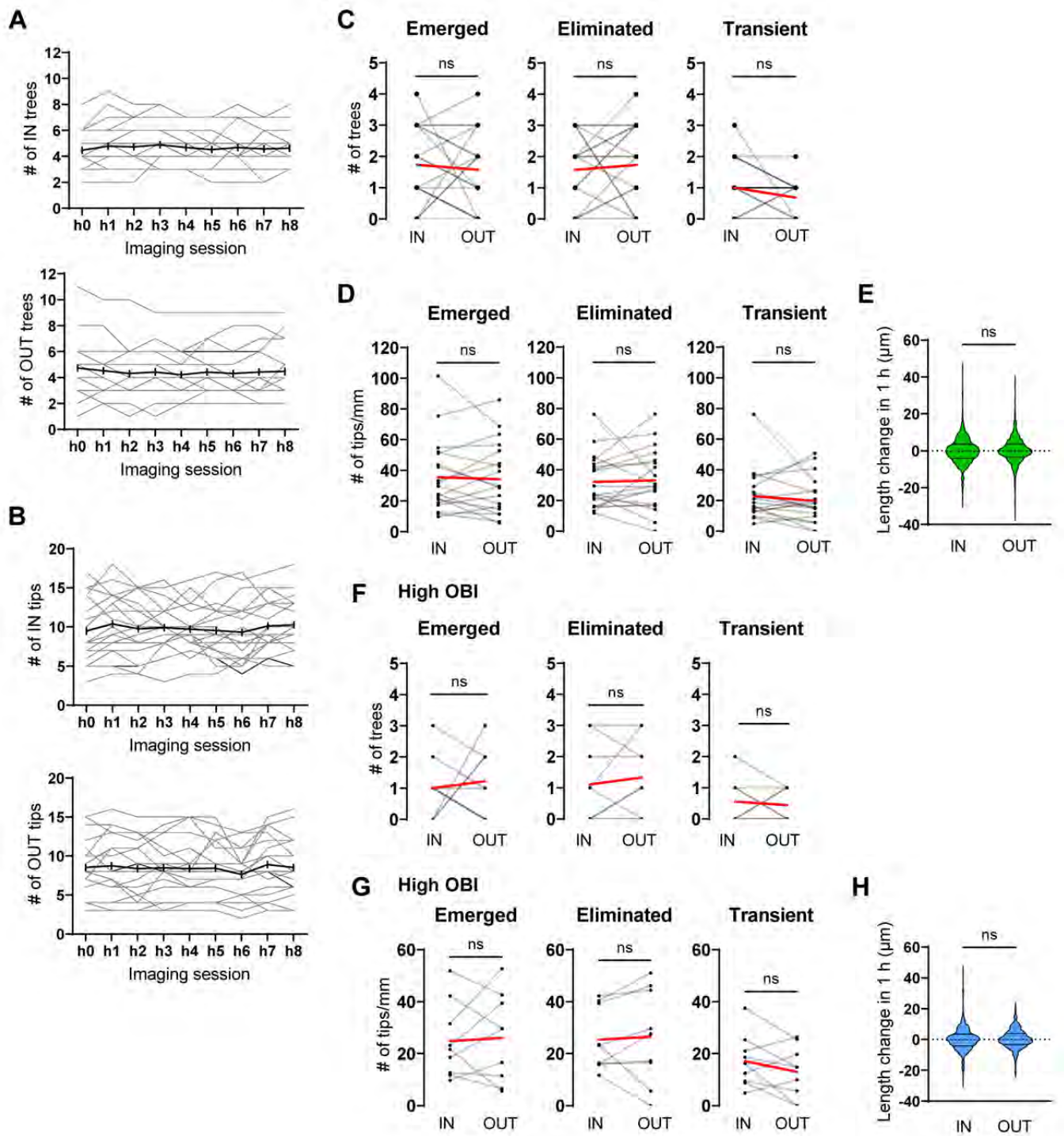


Figure 6

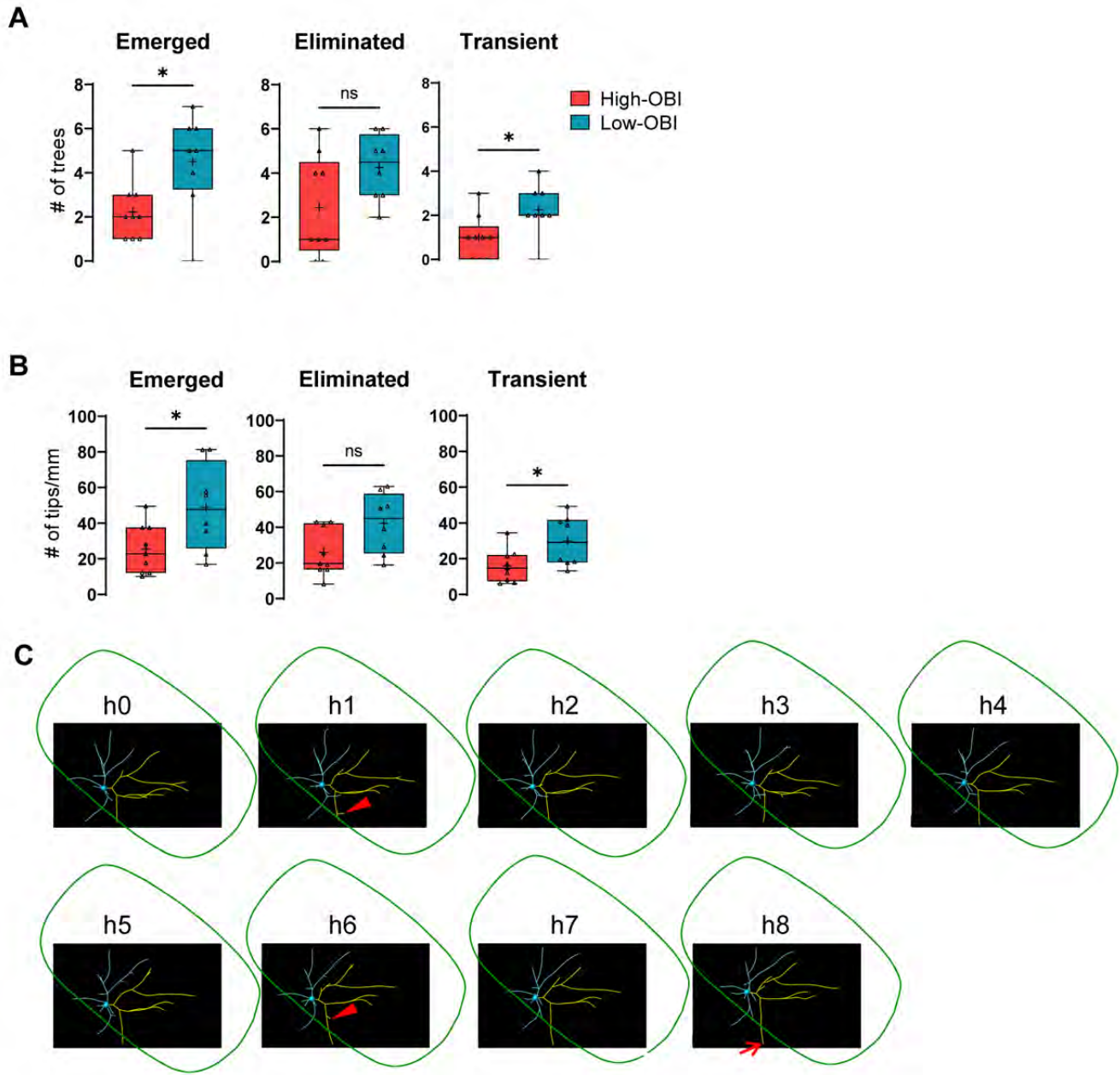


Figure 7

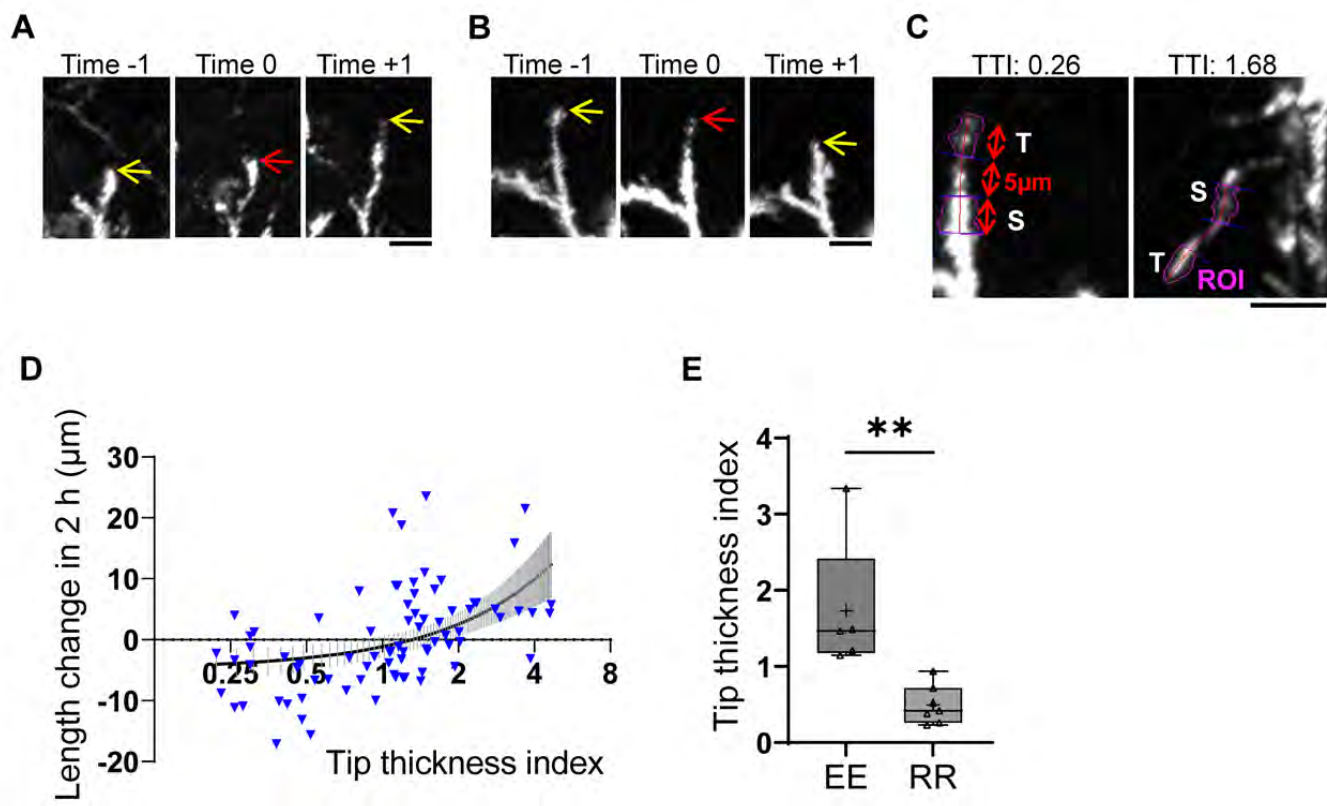


Figure 8

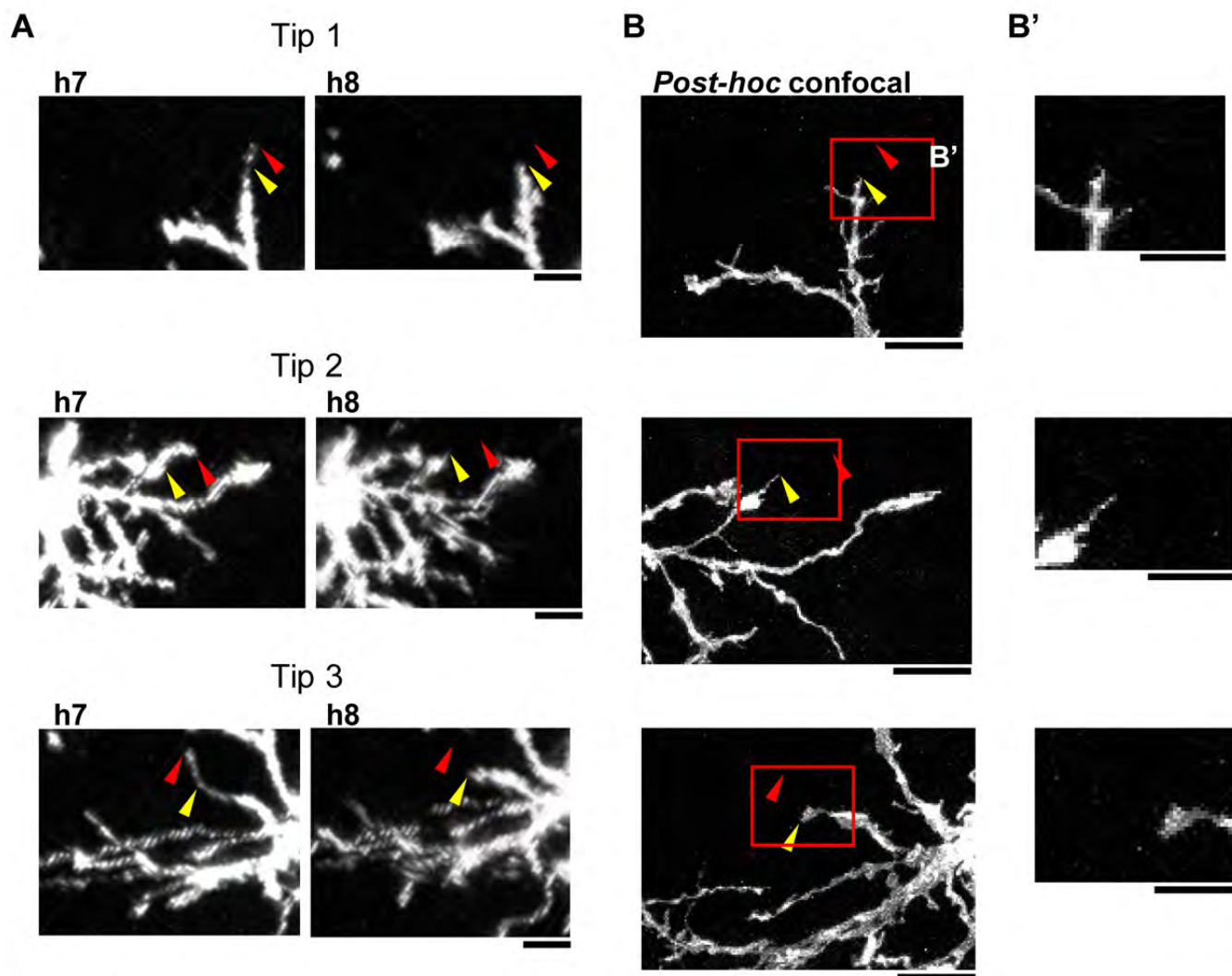


Figure 9