TITLE

Afadin-deficient mouse retinas exhibit severe neuronal lamination defects but preserve visual functions

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ABSTRACT

Neural lamination is a common feature of the central nervous system (CNS), with several subcellular structures, such as adherens junctions (AJs), playing a role in this process. The retina is also heavily laminated, but it remains unclear how laminar formation impacts retinal cell morphology, synapse integrity, and overall retinal function. In this study, we demonstrate that the loss of afadin, a key component of AJs, in mice leads to significant pathological changes. These include the disruption of outer retinal lamination and a notable decrease as well as mislocalization of photoreceptors, their outer segments, and photoreceptor synapses. Interestingly, despite these severe impairments, we recorded small local field potentials, including the a- and b-waves. We also classified retinal ganglion cells (RGCs) into ON, ON-OFF, and OFF types based on their firing patterns in response to light stimuli. Additionally, we successfully characterized the receptive fields of certain RGCs. Overall, these findings provide evidence that retinal circuit function can be partially preserved even when there are significant disruptions in both retinal lamination and photoreceptor synapses. Our results indicate that retinas with severely altered morphology still retain some capacity to process light stimuli.

INTRODUCTION

A common feature of the central nervous system (CNS) in vertebrates is highly laminated structures, which are established by well-ordered neuronal migration (Ayala et al., 2007; Liu, 2011; Veeraval et al., 2020). Lamination defects are observed in some patients with psychiatric disorders, such as autism spectrum disorder, and these rodent models suggest that lamination may be associated with neural circuit formation and function (Miao et al., 1994; Pan et al., 2019; Romero et al., 2018; Stouffer et al., 2016). However, several studies report that orderly lamination is dispensable for the assembly of direction-selective tectal circuits as well as for several cortical circuit functions and synaptic connections (Dräger, 1981; Guy et al., 2015; Guy and Staiger, 2017; Nikolaou and Meyer, 2015). Therefore, the full extent of the association between lamination and the integrity of neural circuit function remains elusive.

Adherens junctions (AJs), known to be correlated with CNS lamination, are adhesive intercellular junctions composed of protein complexes that mediate strong cell-cell adhesion (Harris and Tepass, 2010; Masai et al., 2003; Meng and Takeichi, 2009; Park et al., 2002; Takai et al., 2008b). Cadherin-catenin complex, nectin-afadin complex, and actin cytoskeleton are the major components of AJs (Gil-Sanz et al., 2014; Harris and Tepass, 2010; Ikeda et al., 1999; Meng and Takeichi, 2009; Tachibana et al., 2000; Takai et al., 2008b, 2008a). Cell-cell adhesion of AJs is mainly mediated by the cadherin-catenin complex, whereas nectin and afadin are required for the initial establishment of AJs (Takai and Nakanishi, 2003). A recent report shows that afadin in intestinal

cells is essential for epithelial cell-cell adhesion which was previously assumed to be primarily regulated by cadherin and catenin (Mangeol et al., 2024). Following this result, the functional importance of afadin and nectin in AJs is being reconsidered (Sebbagh and Schwartz, 2024). Animals with mutations in AJ molecules exhibit cell migration defects in the developing stage, resulting in disruption of lamination, which might be associated with brain dysfunctions, such as intellectual disability (Baum and Georgiou, 2011; de Ligt et al., 2012). However, how the defects in AJs affect the CNS function also remains unclear.

The retina, one of the best-understood mammalian neural circuits in the CNS, is highly laminated and organized into three nuclear layers and two synaptic layers (Amini et al., 2017). In the retina, incoming light is converted into neural signals in photoreceptors, processed in parallel through a network of interneurons, and finally sent to the brain via the optic nerve, the bundle of retinal ganglion cell (RGC) axons (Masland, 2012). Photoreceptors, bipolar cells (BCs), and RGCs are connected sequentially, and horizontal cells and amacrine cells (ACs) modulate the information flow in the circuit (Wässle, 2004). The mechanisms underlying retinal circuit formation and function have been actively studied, but the effects of retinal lamination disruption on retinal neural circuits remain unclear (Duan et al., 2018, 2014; Gollisch and Meister, 2010; Zapp et al., 2022). In mammals, most of the major AJ molecules belong to protein families, possibly resulting in functional redundancy with no apparent lamination phenotype in a single knockout (KO) of a member of this complex. Indeed, the loss of central AJ molecules, nectin-1 and nectin-3, does not cause severe

106 defects in retinal lamination (Inagaki et al., 2005). To clarify the effects of AJ
107 molecule depletion on retinal neural circuits and cell morphology while avoiding
108 the issue of functional redundancy, we focused on afadin, which is a scaffolding
109 protein for nectin and has no paralog in mice. Afadin regulates the localization
110 of nectin, which initiates cell–cell adhesion and promotes AJ formation by
111 recruiting the cadherin–catenin complex. (Ohama et al., 2018; Takai and
112 Nakanishi, 2003). In addition, afadin interacts with various cell adhesion and
113 signaling molecules, as well as the actin cytoskeleton, and contributes to the
114 accumulation of β-catenin, αE-catenin, and E-cadherin at AJs (Sakakibara et
115 al., 2018; Sato et al., 2006). *Afadin* KO mice exhibit severe disruption of AJs in
116 the ectoderm, along with other developmental defects, leading to embryonic
117 lethality (Ikeda et al., 1999; Zhadanov et al., 1999). Conditional deletion of
118 afadin in RGCs leads to disruption of dendrites in ON-OFF direction-selective
119 RGCs (Duan et al., 2018). However, the effect of afadin loss on retinal
120 lamination, circuit formation, and function is poorly understood.
121
122 In this study, using the *Afadin* conditional KO (cKO) mouse, we revealed that
123 the *Afadin*-deficient retinas exhibit severe pathological defects, such as outer
124 retinal lamination disruption, as well as decrease and mislocalization of
125 photoreceptors and their synapses. In contrast to severe disruption of the outer
126 retina, the inner plexiform layer (IPL) structure in the inner retina was relatively
127 intact. Using the retinas isolated from the *Afadin* cKO mouse, we could record
128 local field potentials (micro electoretinograms: mERGs) and RGC firings. Based
129 on the light-evoked firing pattern, RGCs could be classified into ON, ON-OFF,

OFF, and other types as previously established. Furthermore, the receptive field (RF) could be mapped in some RGCs. Consistent with this, the cKO mice responded to moving stimuli in visual behavioral tests. Our results suggest that neural circuits in the retina with severe defects of the outer retinal lamination and photoreceptor synapses can mediate some visual information processing and partially preserve visual functions.

RESULTS

Afadin is localized to adherens junctions, synaptic regions, and the surface of bipolar cells in the retina

To elucidate afadin localization in mature retina (1-month-old: 1M), we immunostained retinal sections with anti-afadin antibody. Afadin signals were observed in the outer segment (OS), outer limiting membrane (OLM), outer plexiform layer (OPL), inner nuclear layer (INL), and inner plexiform layer (IPL) (Figure 1A *Top*). These signals were not observed in retinal sections immunostained using anti-afadin antibody pre-absorbed with the antigen peptide (Figure 1A *Bottom*). Thus, we confirmed that the signals are specific to afadin. To explore whether afadin is co-localized with nectin in the retina, we analyzed afadin and nectin localization in the retina by immunostaining using antibodies against afadin and nectins1-3. Afadin signals were aligned and overlapped with all three nectin signals in the OLM (Figure 1B, C, Figure 1-figure supplement 1A), consistent with the previous finding that AJs are formed in the OLM in the retina (Willbold and Layer, 1998). Nectin-1, unlike nectin-2 and nectin-3, was partially co-localized with afadin at the OPL and IPL, in addition to the OLM (Figure 1B, Figure 1-figure supplement 1B). Nectin-1 was co-localized with afadin in cone synapses immunostained with Arrestin3 (Arr3, cone photoreceptor marker), suggesting that puncta adherent junctions may be formed between cone pedicles and BC dendritic tips. In the retina at postnatal day 0 (P0), afadin and nectin-1 signals were observed in the outer retinal surface where AJs are formed (Figure 1-figure supplement 1C) (Koike et al., 2005). Immunostaining with anti-afadin, anti-SCGN (Type 2-6 cone BC marker)

161 (Puthussery et al., 2010), and anti-PKCα (rod BC marker) revealed that afadin
162 localizes to the dendritic tips, cell bodies, and axons of ON- BCs and at least
163 cell bodies of OFF-BCs (Figure 1D).
164
165 **Afadin is required for proper retinal lamination**
166 To investigate the role of afadin in the retina, we generated retina-specific
167 *Afadin* cKO mice by crossing *Afdn*^{flox} mice with *Dkk3-Cre* mice (Majima et al.,
168 2009; Sato et al., 2007; Yamamoto et al., 2020). Cre activity in *Dkk3-Cre* mice
169 was confirmed by crossing with R26R-H2B-EGFP reporter mice (Abe et al.,
170 2011), in which Cre-dependent excision of a *loxP*-flanked STOP cassette drives
171 nuclear EGFP expression from the *Rosa26* locus (Figure 2-figure supplement
172 1A, B). Afadin was not detected at the protein level by Western blotting in the
173 *Afadin* cKO retina (Figure 2-figure supplement 1C). The *Afadin* cKO mouse was
174 viable and showed no gross morphological abnormalities. To examine retinal
175 lamination and morphology, using the cKO retina at 1M, we performed toluidine
176 blue staining. We found that there were no clear outer and inner segments of
177 photoreceptors in the outer retina, no distinct OPL, and no clear
178 compartmentalization of the ONL and INL (Figure 2A). Rod photoreceptors,
179 identified by inverted nuclear architecture (Solovei et al., 2009) (Figure 2A
180 insets), were widely scattered in the cKO retina, whereas the IPL and GCL
181 appeared intact. Furthermore, rosette-like structures, which lacked cell bodies
182 centrally, were sometimes observed in the cKO outer retina (Figure 2A, B, C:
183 arrowhead).
184

185 To investigate the defects in the OS of the cKO retina in more detail, we stained
186 retinal sections with rhodamine-labeled peanut agglutinin (PNA, a marker of
187 cone photoreceptor outer segment and axon terminals) or antibodies against
188 Rom1 (rod OS marker), Rhodopsin (rod OS marker), S-opsin (S-cone OS
189 marker), and M-opsin (M-cone OS marker). Rhodopsin, Rom1, S-opsin, and M-
190 opsin play central roles in phototransduction in the OS (Clarke et al., 2000;
191 Greenwald et al., 2014; Tanimoto et al., 2015; Xu et al., 2020). Distribution of
192 Rom1, Rhodopsin, S-opsin, M-opsin, and PNA signals in the cKO retina differed
193 significantly from those of the *Afadin* conditional heterozygous (cHet) retina. In
194 the cKO retina, Rom1, Rhodopsin, S-opsin, M-opsin, and PNA signals were
195 remarkably decreased in number and mislocalized in the retina (Figure 2B, C).
196 These markers were not observed in the rosette-like structure of the cKO retina,
197 suggesting that the rosette-like structure is different from the photoreceptor
198 rosette as reported (Stuck et al., 2012). Electron microscopic analysis also
199 demonstrated that only a few OS structures were observed between retinal cells
200 in the disrupted ONL and INL regions of the cKO retina (Figure 2-figure
201 supplement 1D). These findings indicate that rod and cone phototransduction
202 may be severely impaired in the cKO retina. To investigate the distribution of
203 cone and rod photoreceptors in the cKO retina, we performed wholemount
204 immunostaining with antibodies against Arr3, Rhodopsin, and Rom1. Cell
205 bodies of cone photoreceptors were arranged regularly in the cHet retina,
206 whereas cone photoreceptor arrangements and morphology were significantly
207 disordered in the cKO retina (Figure 2D). Rom1, Rhodopsin, and Arr3 signals
208 were decreased but relatively dense in some areas (Figure 2D). These results

209 suggest that photoreceptor distribution is disrupted, but photoreceptors remain
210 in some areas heterogeneously.
211
212 We further investigated AJs in the cKO retina by immunostaining with OLM
213 adherens junction markers β-catenin, N-cadherin, and nectin-1. We found that
214 these signals were scattered in the cKO retina (Figure 2-figure supplement 1E).
215 This data suggests that AJs in the OLM are disrupted in mature cKO retina.
216
217 As toluidine blue staining of the cKO retina suggested mislocalization of each
218 cell type, we investigated the position of each retinal cell body in the cKO retina
219 by immunostaining with retinal cell type-specific markers; Otx2 (a photoreceptor
220 and BC marker), Chx10 (a BC marker), Arr3, Calbindin (a horizontal cell and AC
221 marker), AP2α (an AC maker), RBPMS (a RGC marker), and Lhx2 (a Müller
222 glial cell marker) (Figure 2E). All retinal cell types were aligned in restricted and
223 orderly regions in the cHet retina. In the cKO retina, the spatial organization of
224 rod photoreceptors, BCs, cone photoreceptors, horizontal cells, and Müller glial
225 cells appeared particularly disrupted, whereas positions of ACs and RGCs
226 remained relatively intact, except for the immediately surrounding region of
227 rosette-like structures. To quantitatively assess the retinal cell positioning in the
228 cKO retina, the distance from the soma of each cell type to the outer or inner
229 retinal surface was measured (Figure 2F-L). The distance for each cell type in
230 the cKO retina was significantly different from that in the cHet retina. In the cKO
231 retina, BCs, horizontal cells, Müller glial cells were positioned close to the outer
232 retinal surface and ACs were close to the inner retinal surface, whereas

233 photoreceptors and RGCs were positioned away from the outer and inner
234 retinal surfaces, respectively.
235
236 Notably, BCs in the cKO retina were broadly distributed throughout the retina
237 (Figure 2H). Rod photoreceptors also tended to be dispersed throughout the
238 cKO retina. However, approximately 75% of them in the cKO retina were
239 located within 60 μm of the outer retinal surface, which corresponds to the
240 region where all rods are positioned in the cHet retina (Figure 2G). Cone
241 photoreceptors, horizontal cells, and Müller glial cells were similarly spread, but
242 their distribution was largely confined to the area distal to the IPL (Figure 2F, I,
243 J). Although ACs and RGCs were mainly located in the inner retina, a sparse
244 distribution was also observed throughout the retina, which may be attributed to
245 the cells surrounding the rosette-like structure (Figure 2K, L). The reduction in
246 the distance from ACs to the inner retinal surface in the cKO retina is
247 conceivably due to a shortening of the IPL (Figure 2K), though there was no
248 significant difference in retinal thickness between cHet and cKO retinas (Figure
249 2M). The cKO mice appeared to show greater inter-individual variability in
250 retinal layer structure and cell positioning than the cHet mice.
251
252 Immunostaining with an anti-GS antibody (a Müller glial cell marker) showed
253 that Müller glial cell morphology was affected, especially at the distal part.
254 However, a distinct inner limiting membrane was retained in the cKO retina
255 (Figure 2-figure supplement 1F). It has been reported that defects in the distal
256 processes of Müller glia are associated with abnormal retinal vasculature (Shen

257 et al., 2012). Thus, we immunostained the cKO retina with anti-CD31, a blood
258 vessel marker, but no apparent vascular abnormalities were detected (Figure 2-
259 figure supplement 1G). These results indicate that afadin is essential for normal
260 retinal lamination.
261
262 **Ablation of afadin results in the remarkable decrease and mislocalization**
263 **of photoreceptor-BC synapses**
264 No distinct OPL structure in the cKO retina suggests that synapses between
265 photoreceptor and BC may be disrupted. To assess their integrity, we
266 immunostained retinal sections of the cKO mouse (1 M) with antibodies against
267 Bassoon (a photoreceptor synaptic ribbon marker), PSD95 (a photoreceptor
268 synaptic terminal marker), mGluR6 (an ON-BC dendritic tip marker), GluR5 (an
269 OFF-BC dendritic tip marker), PKCα, and PKARIIβ (a type 3b OFF cone BC
270 marker). In the cHet retina, photoreceptor synapses containing synaptic ribbons
271 formed a row in the OPL (Figure 3A, B, Figure 3-figure supplement 1A). On the
272 other hand, in the cKO retina, the row of photoreceptor synapses was not
273 observed, and photoreceptor-BC synapses were remarkably reduced in
274 number. Bassoon and mGluR6 were aberrantly distributed, and the position of
275 the soma and dendritic tips of rod BCs was noticeably disturbed (Figure 3A).
276 Only a few ribbon synapses remained (Figure 3A arrowheads). The 3D
277 reconstructions of the confocal image stacks showed more clearly that
278 photoreceptor synapses decreased in number and mislocated in the cKO retina
279 (Figure 3C, D). The contact between the axon terminal of photoreceptors and
280 the dendrite of ON- and OFF-BCs was also significantly reduced in number,

281 suggesting that photoreceptor-BC synapses were considerably lost in the cKO
282 retina (Figure 3B, D, Figure 3-figure supplement 1A, B). Along with these
283 defects, GluR5 signals with a branch-like pattern were detected in the IPL of the
284 cKO retina, though no GluR5 signals were found in the IPL of the cHet retina
285 (Figure 3B, D). Photoreceptor-ON BC ribbon synapses and photoreceptor-OFF
286 BC synapses in the outer retinal region beneath 1 mm^2 of the surface area of
287 the cKO retina were significantly reduced to about 5% and 15% of those in the
288 cHet retina, respectively (Figure 3E, F, Figure 3-figure supplement 1C). These
289 results indicate that afadin may regulate the formation and localization of
290 photoreceptor synapses.
291
292 **Afadin is required for proper localization of GluR5 to OFF BC dendrites**
293 GluR5, a kainate receptor subunit that forms functional homomeric and
294 heteromeric channels, is specifically expressed in OFF BCs and localizes to
295 their dendrites (Haverkamp et al., 2003). To investigate GluR5 localization in
296 more detail, we immunostained the cKO retina with anti-GluR5, anti-SCGN, and
297 anti-PKCα antibodies. We found that ectopic GluR5 signals in the IPL
298 overlapped not only with SCGN but also PKCα in the cKO retina (Figure 4A).
299 These data indicate that afadin may be required for specific localization of
300 GluR5 in OFF BC dendrites. It seems likely that ectopically localized GluR5 may
301 receive glutamate released from surrounding cells in the IPL, resulting in
302 depolarization of ON- and OFF-BCs. It should be noted that there were no
303 remarkable differences between cKO and cHet retinas in the localization of
304 glutamate transporter EAAT5 and vesicular glutamate transporter vGlut1 to the

305 axon terminal of photoreceptors and BCs (Figure 4B, Figure 3-figure
306 supplement 1D, E). It has been reported that mGluR6 localization to ON BC
307 dendritic tips requires trans-synaptic interaction with ELFN1 in the synaptic cleft
308 and that ELFN1 ablation results in a remarkable reduction of mGluR6 in ON BC
309 dendritic tips (Cao et al., 2015). As GluR5 localization may also be regulated by
310 trans-synaptic interaction with photoreceptor synaptic terminal proteins, it
311 seems possible that the defect of photoreceptor-OFF BC contacts prevents this
312 trans-synaptic interaction, resulting in GluR5 mislocalization in the cKO retina.
313
314 Then, we investigated the IPL structure in the cKO retina by immunostaining it
315 with anti-ChAT, Calbindin, and Calretinin antibodies. Distinct ON- and OFF
316 ChAT bands, Calbindin bands, and Calretinin bands were observed in the cKO
317 retina (Figure 3-figure supplement 1F). However, type 2-6 BCs immunostained
318 with SCGN were slightly disturbed in the axon terminal position (Figure 4A, B,
319 Figure 3-figure supplement 1D). These results suggest that
320 stratification patterns in the IPL may be slightly impaired but almost intact in the
321 cKO retina. Immunostaining analysis with anti-HPC-1 (also known as syntaxin,
322 an AC marker), anti-Calbindin, anti-RBPMS, and anti-Tuj1 (an RGC marker)
323 antibodies revealed that the rosette-like structure contains AC and RGC
324 processes inside (Figure 4C). These data indicate that the rosette-like structure
325 in the cKO may be an ectopic IPL, termed “acellular patches” (Nahar and Cho,
326 2022).
327
328 **Retinal cell fate is altered in the cKO retina**

329 To examine whether afadin loss affects retinal cell fate determination, we
330 counted the number of retinal cells immunostained with each retinal cell marker.
331 In the cKO retina at 1M, BCs significantly increased to about 150% of the cHet
332 retina, but rods, cones, and Müller glial cells were significantly decreased to
333 about 30%, 40%, and 80% of the cHet retina, respectively (Figure 4D). PKCα-
334 positive cells and SCGN-positive cells were significantly increased (Figure 4-
335 figure supplement 1A), indicating that both rod- and OFF-BCs were increased in
336 the cKO retina. In contrast, the number of rod BC processes in the IPL was
337 significantly decreased in the cKO retina (Figure 4-figure supplement 1B).
338 These data suggest that ectopically located BCs may increase in the cKO retina
339 at 1M. A significant increase in BCs and a significant decrease in
340 photoreceptors were also observed in the cKO retina at P14, a stage at which
341 retinal differentiation is complete (Figure 4-figure supplement 1C). Consistently,
342 the expression of marker genes for photoreceptors and BCs was significantly
343 decreased and increased, respectively, in the cKO retina (Figure 4-figure
344 supplement 1D). In contrast, the number of Müller glial cells was unaltered in
345 the cKO retina at P14 (Figure 4-figure supplement 1C), suggesting that the
346 reduction in Müller cells observed at 1M may result from cell death after P14
347 rather than a differentiation defect. To assess whether retinal cell fate changes
348 occur during differentiation, we examined the number of BCs and the
349 expression of rod marker genes in the cKO and cHet retinas at P1, P3, and P5.
350 No significant differences were observed between genotypes at P1 and P3,
351 whereas the number of BCs was significantly increased, and the expression of
352 rod marker genes was significantly decreased in the cKO retina at P5 (Figure 4-

353 figure supplement 1E, F). These results suggest that defects in cell fate
354 determination of BCs and rod photoreceptors begin to emerge between P3 and
355 P5. Based on these data and ectopic localization of GluR5 to ON-BCs in the
356 cKO retina (Figure 4A), we propose that afadin may be required for normal
357 retinal cell fate determination and gene expression.
358

359 **Retinal structure is severely disrupted during development in *Afadin* cKO**
360 **mice**

361 To investigate whether the aberrant lamination in the cKO retina can be
362 attributed to a defect in development or maintenance, we immunostained the
363 developing cHet and cKO retinas using markers for individual retinal cell types,
364 outer segments, and photoreceptor synapses (Figure 4E, Figure 4-figure
365 supplement 2A-D). In the cKO retina at P1, retinal cells appeared to be
366 organized into distinct multicellular compartments with apparent boundaries
367 (Figure 4E, Figure 4-figure supplement 2A, B). RGCs, ACs, and their dendritic
368 processes, which are components of the IPL, were localized along the
369 boundaries of these compartments (Figure 4-figure supplement 2A, B).
370 Furthermore, RGCs and ACs were observed in the outer part of the retina. At
371 P3 and P5, the interspaces between compartments appeared to be reduced,
372 coinciding with an increase in RGCs and ACs positioned on the inner side of the
373 retina (Figure 4-figure supplement 2A, B). By comparison, the positioning of
374 BCs and rod photoreceptors appears to remain disorganized throughout
375 postnatal development (Figure 4E). Contrary to our expectation that retinal
376 structure would become increasingly disorganized as development progresses,

377 laminar disruption in the cKO retina appeared more severe at earlier postnatal
378 stages than at later ones, particularly in the IPL, ACs, and RGCs (Figure 4-
379 figure supplement 2A, B). Aberrant acellular regions were observed in the cKO
380 retina from P1 through P14 (Figure 4E, Figure 4-figure supplement 2A-C).
381 Some of these regions contained AC processes and were surrounded by AC
382 and RGC somata from P3 onward (Figure 4-figure supplement 2A, B).
383 Correspondingly, clear signals of rod outer segment markers were observed in
384 the interior of some acellular regions at P8, P11, and P14 (Figure 4-figure
385 supplement 2C). At 1M, these structures were not observed in the cKO retina
386 (Figure 2B). These data suggest that both acellular patch and photoreceptor
387 rosette may be formed in the developing retina, but photoreceptor rosette may
388 degenerate after P14. The outer segment marker signals were disorganized in
389 the cKO retina at P8, P11, and P14 (Figure 4-figure supplement 2C). Moreover,
390 the row of rod and cone photoreceptor synapses was not observed in the cKO
391 retina at P11 (Figure 4-figure supplement 2D). These data indicate that proper
392 development of retinal lamination and synapse formation is impaired in the cKO
393 retina.
394
395 **Small local field potentials are recorded from the cKO retina**
396 To examine whether the cKO mouse retina can respond to light stimulation, we
397 first recorded electoretinogram (ERG) from anaesthetized animals. Under the
398 scotopic condition, a flashlight was applied to eyes in the dark (Figure 5A *left*).
399 Scotopic ERG from the cHet mouse showed a negative a-wave and a following
400 positive b-wave, whereas both waves from the cKO mouse were almost flat.

401 Under the photopic condition, the mouse was adapted to the background light
402 for 10 min, then a flashlight was superimposed on the adapting background
403 (Figure 5A *right*). Photopic ERGs from the cHet mouse showed a small a-wave
404 and a prominent b-wave, whereas those from the cKO mouse were almost flat.
405 These results may be attributed to the scattered distribution of rod and cone
406 photoreceptors, a lack of a uniform photoreceptor layer, and the decrease of
407 outer segments and photoreceptor synapses in the cKO retina (see Figures 2
408 and 3).
409
410 Then, to examine whether local field potentials can be detected from the cKO
411 retina, we performed multielectrode array (MEA) recordings from the retinas
412 isolated from the cKO and cHet mice. Surprisingly, a flashlight could
413 occasionally evoke micro-ERGs (mERGs) from the cKO retina (SD of the
414 amplitude during 300 ms after the flashlight onset; >50 μV, 13/590 electrodes, n
415 = 10 retinas from 6 mice) (Figure 5B *right*). However, the amplitude of mERGs
416 from the cKO retina was smaller than that from the cHet retina (SD of the
417 amplitude during 200 ms after the flashlight onset; >1,000 μV, 292/413
418 electrodes, n = 7 retinas from 4 mice) (Figure 5B *left*). As the a-wave represents
419 hyperpolarization of the photoreceptors and the b-wave arises from the inner
420 retina, predominantly from ON BCs with a contribution from Müller glial cells
421 (Perlman, 2020), the first negative and following positive waves of mERGs may
422 correspond to the a- and b-waves, respectively. To confirm this, L-AP4, an
423 agonist of mGluR6, was bath-applied to suppress the response of ON BCs. In
424 the presence of L-AP4, the second positive wave disappeared, whereas the first

425 negative wave was enhanced in the cHet retina and remained in the cKO retina
426 (Figure 5B), indicating that the first negative and following positive waves are
427 the a- and b-waves, respectively. We compared the amplitude and implicit time
428 (time-to-peak latency) of the a- and b-waves and found that both a- and b-
429 waves were significantly smaller in amplitude (Figure 5C) and longer in implicit
430 time in the cKO retina than those in the cHet retina (Figure 5E). The ratio of b-
431 wave amplitude to a-wave amplitude (b/a) was significantly smaller in the cKO
432 retina than in the cHet retina (Figure 5D), suggesting that the efficiency of
433 synaptic transmission from photoreceptors to ON BCs may be decreased in the
434 cKO retina. However, the difference of the implicit time between a- and b-waves
435 was similar in the cKO and cHet retinas (Figure 5F), suggesting that signals
436 from photoreceptors to ON BCs may be transmitted properly. These results
437 indicate that despite severe disorganization of the cKO outer retina,
438 photoreceptors can still respond to light stimulation and that the synaptic
439 transmission from photoreceptors to ON-BCs remains functional.
440

441 **RGCs in the cKO retina evoke firings to light stimulation**

442 Using the MEA system, we recorded spike discharges from RGCs in retinas
443 isolated from the cKO and cHet mice. The frequency of spontaneous firings in
444 the dark was not different significantly between cKO and cHet retinas (cHet;
445 12.5 ± 19.2 Hz, n = 434, cKO; 10.8 ± 14.6 Hz, n = 388, p = 0.79, Mann-Whitney
446 U test). We examined whether RGCs in the cKO retina respond to light
447 stimulation. The isolated retina was stimulated by a flash seven times every 8 s,
448 and the peristimulus time histogram (PSTH) was calculated for each RGC after

449 spike sorting. RGCs could be classified into “ON”, “ON-OFF”, “ON-OFF
450 inhibition”, “OFF”, and “None” types (Figure 6A, B; see **MATERIALS and**
451 **METHODS**). All types were found in both cKO and cHet retinas, and the
452 composition of RGC types was similar in both retinas (Figure 6C). We also
453 investigated whether RGCs received both rod and cone inputs and found that >
454 60% of RGCs among successfully spike sorted RGCs in both the cKO and cHet
455 retinas received both inputs (Figure 6D).
456
457 **Receptive fields are characterized in some RGCs**
458 Using pseudorandom checkerboard pattern stimulation, we examined the
459 properties of spatiotemporal RF of RGCs by calculating the spike-triggered
460 average. We could identify RFs in the cKO retina. To check the morphology of
461 the retina where RFs were identified, the retina was removed from the MEA
462 after recording, and sliced retinal sections were immunostained with anti-
463 PKARIIβ, anti-PSD95, and anti-PKCα antibodies (Figure 7A). The sectioned
464 region clearly showed impairment of the outer retina, where it was difficult to
465 identify or locate photoreceptors. On the other hand, the inner retina was
466 apparently normal, and both the IPL and the proximal part of the INL could be
467 recognized. Even though the outer retina was severely impaired, we could
468 identify RF of RGCs (Figure 7B). However, the cells with clear RF were fewer in
469 the cKO retina (69/388 cells, n = 4 retinas) than those in the cHet retina
470 (195/434 cells, n = 4 retinas). By fitting ellipse to each RF, we measured the
471 short and long axes and calculated the RF area (Figure 7C, D, E). The RF area
472 in the cKO retina was significantly smaller than that in the cHet retina. We tried

473 to find an antagonistic surround of RFs, but it was difficult to detect a clear
474 surround in both cKO and cHet retinas.
475
476 The temporal profile of RFs was calculated, and the principal component
477 analysis was performed. Then, using Uniform Manifold Approximation and
478 Projection (UMAP) for clustering, we could recognize 5 clusters (Figure 7F,
479 cKO; n = 4 retinas, cHet; n = 4 retinas). The temporal profile belonged to each
480 cluster was superimposed separately (Figure 7G). Among 5 clusters, clusters 1
481 and 2 did not include the temporal profiles calculated from RGCs in the cKO
482 retina. The temporal profiles of cluster 1 and cluster 2 were ON and OFF types,
483 respectively, each of which had a shorter time-to-peak latency than the other
484 temporal profiles obtained from RGCs in both cKO and cHet retinas (Figure
485 7H). These results indicate that the light-evoked responses of RGCs in the cKO
486 retina are not as refined as cHet. Actually, the rising time of transient responses
487 (determined by an upward deflection soon after light stimulation obtained from
488 the cumulative curve of each PSTH) was significantly longer in the cKO retina
489 (58.5 ± 19.8 ms, n = 17) than in the cHet retina (43.0 ± 17.8 ms, n = 23) (***p <
490 0.001, Student's test).
491
492 It has been reported that degenerated retinas, such as the *rd1* retinas (Poria
493 and Dhingra, 2015), show oscillatory spontaneous firings in RGCs. Thus, we
494 checked whether RGCs in the cKO retina show RGC oscillations. RGCs in the
495 cKO retina showed oscillations (P30; 1.55% of 382 RGCs, 14.0 ± 2.4 Hz, n = 4
496 retinas, P40; 15.0% of 333 RGCs, 9.2 ± 3.8 Hz, n = 4 retinas). On the other

497 hand, RCGs in the cHet retina did not show obvious RGC oscillations (0% of
498 432 RGCs, n = 4 retinas).
499
500 ***Afadin* cKO mice retain partial visual function**
501 As the MEA analyses suggested partial preservation of retinal functions in the
502 cKO retina, we further examined whether cKO mice retained visual function,
503 using visual behavior tests. We first conducted the optomotor response (OMR)
504 test, a robust method for assessing contrast-based visual function and
505 estimating visual acuity in mice (Thaung et al., 2002; Ye et al., 2021). The OMR
506 test measures head-tracking movements in response to a vertical square wave
507 grating stimulus (Figure 8A). Mice capable of detecting contrasting motion
508 passing through their visual field exhibit head-tracking movements in response
509 to stimuli (Kretschmer et al., 2015; Mitchiner et al., 1976; Schmucker et al.,
510 2005). Apparent head-tracking movements were observed in the cHet and cKO
511 mice during presentation of drifting grating stimuli at spatial frequencies of 0.056
512 and 0.1 cycles/degree, but not in those with optic nerve crush (ONC), in which
513 the optic nerve had been mechanically crushed, resulting in vision loss (Figure
514 8B, C). The number of head-tracking movements in response to these stimuli
515 was significantly higher in the cHet and cKO mice than in the ONC mice. At 0.27
516 cycles/degree, the cKO mice hardly exhibited head-tracking movements, in
517 contrast to the cHet mice, and no significant difference was observed between
518 the cKO and ONC mice under this condition. The number of movements was
519 significantly lower in the cKO mice than in the cHet mice at all three spatial
520 frequencies (Figure 8C). There was no significant difference in the number of

521 head-tracking movements across the three spatial frequencies in the cHet mice,
522 whereas in the cKO mice, the number tended to decrease with increasing
523 spatial frequency (Figure 8D). These data suggest that the cKO mice can
524 recognize moving visual stimuli, but their ability is inferior to that of the cHet
525 mice.
526
527 We further conducted the visual cliff test to evaluate visual performance in a
528 perceptual task. This test exploits rodents' innate tendency to avoid the deep
529 side of the cliff—a plexiglass surface suspended 80 cm above a checkerboard
530 on the floor—and prefer the shallow side, where the plexiglass covered directly
531 atop the checkerboard (Figure 8E) (Fox, 1965). The cHet mice spent
532 approximately 80% of their time on the shallow side, significantly more time
533 than the cKO and ONC mice, while the cKO mice did not differ significantly from
534 the ONC mice in time spent on the shallow side (Figure 8F). Nonetheless, some
535 cKO mice exhibited a behavior in which they looked down toward the bottom of
536 the deep side and quickly backed away (Video 1). This behavior, observed in
537 the cHet mice but not in the ONC mice, may imply that they can visually
538 recognize the deep side (Video 2). Furthermore, the cKO mice, but not the ONC
539 mice, showed a significant preference for the shallow side over the deep side
540 (57.35%, p = 0.048, one-sample t-test, figure not shown). These results indicate
541 that visual information from the retina is transmitted to the brain and visual
542 function is partially maintained in *Afadin* cKO mice.
543
544 **Ectopic AJs and aberrant localization of progenitor cells are observed in**

545 **the developing *Afadin* cKO retina**

546 As RGC oscillations were observed in the cKO retina, like the *rd1* retinas with
547 photoreceptor degeneration (Choi et al., 2014), we examined whether RGC
548 oscillations were related to cell death in the cKO retina. Using anti-active
549 caspase3 (an apoptotic cell marker), we counted the number of dead cells
550 immunohistochemically. We found that dead cells were significantly increased in
551 the cKO retina at around P30 (Figure 9-figure supplement 1A, B). These
552 findings suggest that the progressive cell death observed in the cKO retina may
553 lead to a further decrease of photoreceptors, resulting in an increase of
554 oscillatory RGCs at P40. Apoptotic cells were significantly increased in the cKO
555 retina in the developing stage at P1, P3, P5, and P14 (Figure 9-figure
556 supplement 1B). In the cKO retina at 1 M, the number of rod photoreceptors
557 and Müller glial cells was significantly reduced to approximately 60% and 70%
558 of that at P14, respectively (P14; rod 98.3 ± 3.8, Müller glial cell 17.2 ± 3.3, n =
559 3, 1M; rod 57.3 ± 17.3, Müller glial cell 12.4 ± 1.5, n = 4. rod p = 0.011, Müller
560 glial cell p = 0.046 by Student's t-test), in contrast to the cHet retina, in which
561 the number of rod photoreceptors remained unchanged (P14; rod 176.5± 14.3,
562 Müller glial cell 17.3 ± 1.4, n = 4, 1M; rod 185.9 ± 17.5, Müller glial cell 15.7 ±
563 2.3, n = 4, rod p = 0.43, Müller glial cell p = 0.27 by Student's t-test). This
564 reduction may result from ongoing rod cell death even after the completion of
565 differentiation.

566

567 Then we investigated the mechanism underlying lamination defects in the cKO
568 retina. We previously demonstrated that the photoreceptor-cell specific *aPKCλ*

569 KO retina shows disruption of AJs and mislocalization of mitotic progenitors in
570 developing stage, followed by severe lamination defects in adults (Koike et al.,
571 2005). In the developing retina, AJs are observed in the outer retinal surface,
572 where progenitor cells divide and migrate toward the inner retina. To investigate
573 the possibility that disruption of AJs may lead to abnormal cell migration and
574 lamination, we immunostained the cKO retina at P0 with anti-β-catenin, N-
575 cadherin, nectin-1, and phospho-histone H3S10 (pHH3, a marker of the mitotic
576 cell) (Figure 9A). N-cadherin, nectin-1, and β-catenin signals were obviously
577 detected, but these signals were localized in the inner retina with aberrant
578 patterns. These signals were slightly observed in the outer retina. These data
579 indicate that ectopic AJs are present in the cKO retina. Mitotic cells
580 immunostained by pHH3 were localized near the ectopic AJs. The number of
581 pHH3-positive cells was significantly increased to approximately 150% of that in
582 the cHet (Figure 9B). At P1, P3, and P5, mitotic cells were also mislocalized to
583 the middle retinal region in the cKO mice, and their distance from the outer
584 retinal surface was significantly larger —approximately 45 μm, 70μm, and
585 75 μm, respectively— than that in the cHet (Figure 9C, D). Mitotic cells showed
586 a broader distribution in the cKO retina than in the cHet. We further examined
587 the distribution of retinal progenitor cells during development. In the cHet retina,
588 progenitor cells labeled with BrdU were scattered but relatively enriched in the
589 middle region at P1 (Figure 9E, F). As the development progressed, the number
590 of progenitor cells decreased, and their distribution was progressively confined
591 to the middle region of the retina. In the cKO retina, progenitor cells were more
592 broadly distributed (Figure 9E, F). At P5, no significant difference in the distance

593 was observed between the cHet and cKO retinas However, progenitor cells in
594 the cKO retina showed a trend toward a broader distribution compared to the
595 cHet retina. Retinal thickness and the number of progenitor cells did not differ
596 significantly between the cHet and cKO retinas at P1, P3, and P5 (Figure 9-
597 figure supplement 1C, D). These data suggest that progenitor cells may divide
598 at ectopic regions and migrate aberrantly into the cKO retina, resulting in
599 lamination defects.
600
601

602 **DISCUSSION**

603 In the current study, we showed that the *Afadin* cKO retina exhibits various
604 severe pathological defects, including photoreceptor morphological defects,
605 mislocalization, reduced number of photoreceptors and their synapses,
606 disruption of the outer retinal lamination, ectopic localization of GluR5, aberrant
607 rosette-like structure in the outer retina (Figures 2-4). These results suggest that
608 afadin is required for normal retinal lamination, photoreceptor cell morphology,
609 photoreceptor synapse formation, and proper GluR5 localization to OFF bipolar
610 cell dendrites. Conditional loss of AJ molecules in the retina has been reported
611 to cause partial or complete disruption of retinal lamination (Erdmann et al.,
612 2003; Fu et al., 2006; Masai et al., 2003). However, the effect of AJ molecule
613 loss on the retinal cell morphology and synapses was unclear. We revealed that
614 loss of afadin significantly but not completely reduces photoreceptor synapses.
615

616 In the *Afadin* cKO mouse retina with severe outer retinal disorganization, we
617 detected the a- and b-waves of mERG and the RF of RGCs (Figures 2, 5, 7).
618 These results suggest that some visual information processing may be
619 performed by newly formed and preserved circuits even when the retinal
620 structure is severely impaired and the number of photoreceptors is remarkably
621 decreased. To our knowledge, this is the first report that the RF of RGCs could
622 be formed if several functional photoreceptors connect to BCs, even when the
623 outer retinal lamination is severely disrupted (Figure 7). The number and size of
624 RFs observed in the cKO retina were smaller than those in the cHet retina,
625 suggesting that the precise arrangement of retinal cells and lamination is

626 necessary for normal visual function. It is important to note that light stimulation
627 evoked firing responses in RGCs around the rosette-like structure, and that
628 such RGC firings could not be detected by a large distance between RGC soma
629 and MEA electrode.
630
631 We would like to speculate why the RF of RGCs was observed in the *Afadin*
632 cKO retina. First, visual processing may be partially preserved without any
633 remodeling if only a small number of functional photoreceptor-BC-RGC
634 pathways remain intact. It has been shown that the RF of RGCs is detected
635 even when approximately half of cone photoreceptors are stimulated (Care et
636 al., 2019; Lee et al., 2022). Thus, in the *Afadin* cKO retina, where cone
637 photoreceptors are reduced to about 40% of the cHet retina, such reduction *per*
638 *se* may not critically affect the RF organization. Furthermore, RFs of RGCs are
639 also detected in several mouse models of retinitis pigmentosa, in which rod
640 photoreceptors are degenerated and surviving cone photoreceptors lose their
641 OS discs and pedicles, instead forming abnormal processes resembling
642 synaptic dendrites (Barhoum et al., 2008; Ellis et al., 2023; Scalabrino et al.,
643 2022). A few photoreceptor-BC-RGC pathways (vertical pathways of the retina)
644 are inferred to be maintained in the cKO retina. In some regions, the density of
645 photoreceptors was high enough to make functional synapses between
646 photoreceptor and BC (Figure 2D), possibly resulting in the RF of RGCs.
647 Second, synaptic rewiring of interneurons such as BCs may recover retinal
648 function in the *Afadin* cKO retina. Various studies show partial loss of
649 photoreceptors or BCs, or partial silencing of BCs causes rewiring in the

650 developing and mature retinas (Johnson and Kerschensteiner, 2014; Jones et
651 al., 1995; Okawa et al., 2014; Shen et al., 2020; Strettoi et al., 2022). Also, in
652 the *rd1* retina, neural connections between inner retinal neurons via gap
653 junctions may be enhanced, potentially resulting in the widespread distribution
654 of electrically evoked RGC responses (Ahn et al., 2022). These reports suggest
655 that reduced input from photoreceptors and BCs may lead to synaptic
656 reorganization among surviving cells. It is possible that synaptic reorganization
657 triggered by reduced photoreceptor and BC inputs may occur in the *Afadin* cKO
658 retina with severe defects of the photoreceptor synapses and outer segments in
659 the developing stages, and that this might contribute to RF formation. Synaptic
660 rewiring may occur between BC dendrite and their closely located
661 photoreceptor terminal in the cKO retina. It may be necessary to count the
662 number of BC dendritic connections to each cone pedicle to verify the synaptic
663 rewiring between photoreceptor and BC, but this was quite difficult because of
664 the remarkably aberrant cone morphology in the cKO retina. As the IPL was
665 relatively intact and BC axon terminals were surrounded by many RGC
666 dendrites, synaptic reconnections between BC axon terminal and RGC dendrite
667 are presumed to be more easily established than those between photoreceptor
668 terminal and BC dendrite in the cKO retina. Thus, several BCs connected with
669 multiple functional photoreceptors may rewire to RGC dendrites and transmit
670 signals efficiently. Our findings imply that a very few BCs receiving input from
671 functional photoreceptors responded to light stimulation, and that the output
672 signal can be amplified downstream. The above two possibilities could both
673 contribute to the formation of RGC RFs. Furthermore, we cannot rule out the

674 possibility that gene expression may be changed in the *Afadin* cKO retina,
675 resulting in facilitation of RF formation.
676
677 The ratio of responding to non-responding RGCs to light stimulation in the
678 *Afadin* cKO retina was comparable to that in the cHet retina despite notable
679 reduction of photoreceptors (Figures 4D, 5, 6). It seems likely that the number
680 of photoreceptors connected to spreading dendrite tips of one BC may be
681 increased, and that glutamate spillover from BC terminals may contribute to
682 increasing the responsiveness of RGCs in the cKO retina because glutamate
683 transporter EAAT5 and vesicular glutamate transporter vGlut1 were similar in
684 both cKO and cHet retinas (Figure 4B, Figure 3-figure supplement 1D, E).
685 Glutamate released from BC terminals may spread and bind GluR5, which is
686 ectopically localized in BC processes in the IPL, enhancing the response and
687 expanding the responding area (Figure 4A, B). These factors may contribute to
688 the emergence of RF, but the RF size in the cKO retina was still smaller than
689 that in the cHet retina (Figure 7). Photoreceptors in the cKO retina were
690 reduced to 30 to 40% of the cHet retina and scattered at 1M (Figures 2, 4D).
691 The random checkerboard pattern used for STA was projected onto the
692 assumed photoreceptor layer of the normal retina. However, as the morphology
693 and position of photoreceptors were disturbed in the cKO retina (Figure 2,
694 Figure 2-figure supplement 1D), the remaining photoreceptors would be
695 stimulated by an unfocused image with decreased intensity. Accordingly,
696 photoreceptors possibly responded weakly to unfocused images, and
697 thresholding of noise (increment >+4 SDs, decrement <-2 SDs, determined by

698 the spontaneous firing rate, see **MATERIALS and METHODS**) may result in
699 reduction of the RF size. Rod-driven responses were observed in RGCs in the
700 cKO retina (Figures 5, 6). However, it is unlikely that the rod photoreceptors
701 transmit signals to cone photoreceptors through gap junctions in the disrupted
702 cKO outer retina, and thus, rod signals may be transmitted to RGCs via the rod
703 BC-AII amacrine-ON cone BC pathway.
704
705 In the cKO retina, the temporal profile of RFs with short time-to-peak latency
706 was not included (Clusters 1 and 2; Figure 7), indicating that RGCs in the cKO
707 retina cannot respond to rapid changes in luminance. There are several ways to
708 explain this result. For example, disruption of the outer retinal lamination may
709 make it difficult to synchronize photoreceptor responses without gap junctions,
710 resulting in dull responses. BCs with fast response kinetics may not be present
711 or not functional in the cKO retina (Kuo et al., 2024). Furthermore, glutamate
712 spillover from BC terminals may amplify the signals, but at the same time, the
713 time-to-peak of RGC responses may be delayed. These possibilities should be
714 investigated in the future.
715
716 Ectopic continuous AJs formed in the developing *Afadin* cKO retina (Figure 9A)
717 are possibly associated with partial disruption of retinal lamination. The *β-*
718 *catenin* cKO retina also shows ectopic continuous AJs in developing stage and
719 the disruption of the outer retinal lamination with relatively intact IPL and GCL
720 structure in adult (Fu et al., 2006), suggesting that afadin and β-catenin are
721 required for normal continuous AJ formation at the outer retinal surface but

722 dispensable for maintaining AJs. The photoreceptor-specific *aPKCλ* cKO retina
723 in which AJs are dispersed in the developing stage exhibits disruption of whole
724 retinal lamination despite aPKCλ remaining in cells excluding photoreceptors.
725 Individual cells contain AJs, but the AJs are not continuous in the *aPKCλ* cKO
726 retina (Koike et al., 2005). These findings, based on the studies using each
727 mutant mouse, imply that continuous AJs are important for the formation of the
728 outer retinal structure. The ONL was severely disrupted in the cKO retina,
729 despite the fact that afadin is not localized in the layer. The disruption is
730 presumed to be caused by defects in AJs in the outer retina during
731 development. In the developing *Afadin* cKO retina, ectopic AJs and mislocalized
732 progenitor cells were observed (Figure 9). These defects may lead to abnormal
733 cell migration and positioning, ultimately resulting in ONL disorganization. In
734 contrast, the IPL structure appeared relatively intact in the cKO retina, possibly
735 due to functional redundancy between afadin and other cell adhesion
736 molecules.
737
738 In the *Afadin* cKO retina, significant increase in BCs and significant decrease in
739 photoreceptors and Müller glial cells were observed at 1M (Figure 4D). The
740 changes in the number and/or marker gene expression of photoreceptors and
741 BCs were also observed in the developing cKO retina (Figure 4-figure
742 supplement 1C-F). These results suggest that retinal cell fate was affected in
743 the *Afadin* cKO mice. As retinal cell fate is not altered in the *β-catenin* cKO
744 retina (Fu et al., 2006), the phenotypic difference between *β-catenin* cKO and
745 *Afadin* cKO retinas may imply distinct functional properties of the in AJ

746 molecules.

747

748 One of the major concerns of retinal regeneration therapy is that while
749 transplanted photoreceptors can form synapses with donor BCs, the
750 transplanted tissue often shows aberrant structures, such as rosettes
751 (Yamasaki et al., 2022). Our study demonstrated that RGCs show RFs and
752 visual behaviors are partially preserved even when the outer retinal lamination
753 is severely disrupted, provided that synaptic connectivity is maintained to a
754 certain extent. Therefore, mice with photoreceptor degeneration that have
755 undergone retinal transplantation might recognize stimulus patterns to some
756 extent, even if aberrant structures are formed after retinal transplantation.
757 Moreover, retinal regeneration therapy could be more effective in restoring
758 vision than previously anticipated. In recent studies, distinct RFs of RGCs were
759 detected in some mouse lines with progressive photoreceptor degeneration or
760 substantial loss of cone photoreceptors or BCs, indicating the robustness of
761 retinal circuitry (Care et al., 2019; Ellis et al., 2023; Johnson and
762 Kerschensteiner, 2014; Lee et al., 2022; Scalabrino et al., 2022). However,
763 whether visual function is retained in these models remains unclear. Behavioral
764 visual responses are partially preserved in the *Afadin* cKO mice, in which RFs
765 of RGCs are reduced in number and size but clearly detectable. Given that
766 RGCs are intact in the mouse lines used in the aforementioned studies, it is
767 possible that visual functions are preserved to a substantial extent in those
768 models. Although the *Afadin* cKO mice responded to certain visual stimuli, the
769 extent of what visual patterns or stimuli they can perceive remains unclear.

770 Further detailed analyses of visual functions are required in the future.

771

772 **MATERIALS and METHODS**

773

Key resources table				
Reagent type (species) or resource	**Designation**	**Source or reference**	**Identifiers**	**Additional information**
strain, strain background (Mus musculus)	*Afdn^{flox}*	Majima et al., 2009		
strain, strain background (Mus musculus)	*Dkk3-Cre*	Sato et al., 2007, Yamamoto et al., 2020	RBRC05427	
strain, strain background (Mus musculus)	R26R-H2B-EGFP	Abe et al., 2011	CDB0203K	
antibody	anti-α-afadin (mouse monoclonal)	Abcam	Cat# ab90809	IF (1:100), WB(1:1000)
antibody	anti-nectin1 (Rat monoclonal)	MBL	Cat# D146-3, RRID:AB_590847	IF (1:100)
antibody	anti-nectin2 (Rat monoclonal)	MBL	Cat# D083-3, RRID:AB_590848	IF (1:100)
antibody	anti-nectin3 (Rat monoclonal)	MBL	Cat# D084-3	IF (1:100)
antibody	anti-PKCα (Mouse monoclonal,Rabbit polyclonal)	Sigma-Aldrich	Cat# P5704, RRID:AB_477375 Cat# P4334, RRID:AB_477345	IF (1:1000, 1:10000)
antibody	anti-SCGN (Goat polyclonal)	R&D Systems	Cat# AF4878, RRID:AB_2269934	IF (1:2000)

antibody	anti-Arr3 (Rabbit polyclonal)	Millipore	Cat# AB15282, RRID:AB_1163387	IF (1:1000)
antibody	anti-Calbindin (Rabbit polyclonal)	Millipore	Cat# PC253L-100, RRID:AB_213554	IF (1:200)
antibody	anti-ChAT (Goat polyclonal)	Millipore	Cat# AB144P, RRID:AB_2079751	IF (1:50)
antibody	anti-Bassoon (Mouse monoclonal)	Enzo	RRID:AB_1860018	IF (1:1000)
antibody	anti-mGluR6 (Guinea pig polyclonal)	Current study		IF (1:3000)
antibody	anti-PSD95 (Mouse monoclonal)	Synaptic Systems	Cat# 124 014, RRID:AB_2619800	IF (1:3000)
antibody	anti-GluR5 (Goat polyclonal)	Steve H. DeVries		IF (1:2000)
antibody	anti-PKARIIβ (Mouse monoclonal)	BD Biosciences	Cat# 610625, RRID:AB_397957	IF (1:500)
antibody	anti-Calretinin (Mouse monoclonal)	Millipore	Cat# PC235L-100UCN	IF (1:5000)
antibody	anti-EAAT5 (Rabbit polyclonal)	Sigma-Aldrich	Cat# HPA049124	IF (1:100)
antibody	anti-vGlut1 (Guinea pig polyclonal)	Millipore	Cat# AB5905, RRID:AB_2301751	IF (1:6000)
antibody	anti-HPC-1 (Mouse monoclonal)	Sigma-Aldrich	Cat# S0664, RRID:AB_477483	IF (1:10000)
antibody	anti-active caspase-3	R&D Systems	Cat# AF835, RRID:AB_2243952	IF (1:300)

	(Rabbit polyclonal)			
antibody	anti-Chx10 (Rabbit polyclonal)	Hori et al., 2019		IF (1:100)
antibody	anti-Otx2 (Goat polyclonal)	R&D Systems	Cat# AF1979, RRID:AB_2157172	IF (1:200)
antibody	anti-Glutamine synthetase (Mouse monoclonal)	Millipore	Cat# MAB302, RRID:AB_2110656	IF (1:500)
antibody	anti-CD31 (Rat monoclonal)	BD Biosciences	Cat# 557355, RRID:AB_396660	IF (1:100)
antibody	anti-Lhx2 (Goat polyclonal)	Santa Cruz Biotechnology	Cat# sc-19344, RRID:AB_2135660	IF (1:100)
antibody	anti-RBPMS (Rabbit polyclonal)	GeneTex	Cat# GTX118619, RRID:AB_10720427	IF (1:500)
antibody	anti-Ki67 (Mouse monoclonal)	BD Biosciences	Cat# 556003, RRID:AB_396287	IF (1:100)
antibody	anti-Tuj1 (Mouse monoclonal)	BioLegend	Cat# 801201, RRID:AB_2313773	IF (1:500)
antibody	anti-AP2α (Mouse monoclonal)	DSHB	Clone 3B5-c	IF (1:1000)
antibody	anti-Rom1 (Rabbit polyclonal)	Robert Molday		IF (1:10)
antibody	anti-Rhodopsin (Rabbit polyclonal)	St John's Laboratory	Cat# STJ95452	IF (1:100)
antibody	anti-S-opsin (Goat polyclonal)	Santa Cruz Biotechnology	Cat# sc-14363, RRID:AB_2158332	IF (1:500)

antibody	anti-M-opsin (Rabbit polyclonal)	Millipore	Cat# AB5405, RRID:AB_177456	IF (1:500)
antibody	anti-GFP (Chicken polyclonal)	Aves Labs	Cat# GFP-1010, RRID:AB_2307313	IF (1:1000)
antibody	anti-β-catenin (Mouse monoclonal)	BD Biosciences	Cat# 610153, RRID:AB_397554	IF (1:1000)
antibody	anti-N-cadherin (Mouse monoclonal)	BD Biosciences	Cat# 610920, RRID:AB_2077527	IF (1:500)
antibody	anti-BrdU (Mouse monoclonal)	BD Biosciences	Cat# 347580, RRID:AB_400326	IF (1:100)
antibody	anti-phospho-histone H3 (Rabbit polyclonal)	Millipore	Cat# 06-570, RRID:AB_310177	IF (1:2000)
software, algorithm	MATLAB (MathWorks)	MathWorks	RRID:SCR_001622	
software, algorithm	Psychtoolbox-3	http://psychtoolbox.org/	RRID:SCR_002881	
software, algorithm	Python3	https://www.python.org/	RRID:SCR_008394	

774
775 *Animals*

776 All animal experimental protocols were conducted in accordance with local
777 guidelines and the ARVO statement on the use of animals in ophthalmic and
778 vision research. These procedures were approved by the Institutional Safety
779 Committee on Recombinant DNA Experiments (approval ID R4016) and Animal
780 Experimental Committees (approval ID BKC2022-017) of Ritsumeikan
781 University. Mice were kept in the temperature-controlled room at 25°C with a 12
782 h/12 h day/night cycle. Fresh water and rodent diets were always provided.

783

784 WT, *Afadin* cHet, and cKO mice on 129S6/SvEvTac background, and R26R-
785 H2B-EGFP mice on C57BL/6J background were used in this study. The *Afdn*^{flox}
786 mice (Majima et al., 2009), *Dkk3-Cre* mice (Sato et al., 2007; Yamamoto et al.,
787 2020), and R26R-H2B-EGFP mice (Abe et al., 2011,CDB Accession number.
788 CDB0203K), have been described previously. Exon 2 of the *Afadin* gene is
789 flanked by loxP sites in the *Afdn*^{flox} mice, and Cre activity is detected in the
790 retina of the *Dkk3-Cre* mice from embryonic day 10.5 (E10.5). In R26R-H2B-
791 EGFP mice, Cre-dependent, nuclear localized green fluorescent protein
792 targeted to the Rosa26 locus. Mice of either sex were used for all animal
793 experiments. The stages used for individual experiments are described in figure
794 legends.

795

796 *Immunohistochemistry*

797 Immunohistochemical analysis was performed as described previously (Hori et
798 al., 2019; Kubo et al., 2021). In brief, the retina used for MEA recordings and
799 isolated mice eyes were fixed with 4% PFA in PBS for 30 min at room
800 temperature. After three-time washes, retinas were cryoprotected by 30%
801 sucrose in PBS overnight, embedded in an OCT compound (Sakura, Japan),
802 frozen on dry ice, and sectioned at 20 μm of thickness. Whole-mount
803 immunostaining was performed as previously described with some
804 modifications (Ueno et al., 2018). The retinas were gently peeled off from the
805 sclera, fixed with 4%PFA in PBS for 30 min at room temperature, and washed
806 three times. Retinal sections and whole retinas were soaked in blocking buffer

807 (5% NDS, 0.1% Triton X-100, in 1x PBS) for 1-2 h at room temperature, and
808 incubated with primary antibodies in blocking buffer 1 or 2 overnight at 4°C. The
809 sections were washed with PBS three times and incubated with fluorescent dye-
810 conjugated secondary antibodies and DAPI (1:1000) for more than 2 h at room
811 temperature or overnight at 4°C under the light-shielded condition. The
812 specimens were observed using a laser confocal microscope (LSM900; Carl
813 Zeiss, Germany). For BrdU staining, mice were given an intraperitoneal
814 injection of 20 mg/kg BrdU and sacrificed 2 h later. Retinal sections were then
815 pretreated with 2 N HCl at 37 °C for 30 min before blocking. The antibodies and
816 dilution ratios were as follows; anti-l-afadin (ab90809, Abcam, UK, 1:100), anti-
817 nectin1 (D146-3, MBL, Japan, 1:100), anti-nectin2 (D083-3, MBL, Japan,
818 1:100), anti-nectin3 (D084-3, MBL, Japan, 1:100), anti-PKCα (P5704, Sigma,
819 USA, 1:1000, P4334, Sigma, 1:10000), anti-SCGN (AF4878, R&D systems,
820 USA, 1:2000), anti-Arr3 (AB15282, Millipore, USA, 1:1000), anti-Calbindin
821 (PC253L-100, Millipore, USA,1:200), anti-ChAT (AB144P, Millipore, USA, 1:50),
822 anti-Bassoon (SAP7F407, Enzo, USA, 1:1000), anti-mGluR6 (current study,
823 1:3000), anti-PSD95 (#124 014, Synaptic Systems, Germany, 1:3000), anti-
824 GluR5 (Grik1, gift from Steve H DeVries, 1:2000), anti-PKARIIβ (#610625, BD
825 biosciences, USA, 1:500), anti-Calretinin (PC235L-100UCN, Millipore, USA,
826 1:5000), anti-EAAT5 (HPA049124, Sigma, USA, 1:100), anti-vGlut1 (AB5905,
827 Millipore, USA, 1:6000), anti-HPC-1 (S0664, Sigma, USA, 1:10000), anti-active
828 caspase3 (AF835, R&D Systems, USA, 1:300), anti-Chx10 (Hori et al., 2019,
829 1:100), anti-Otx2 (AF1979, R&D systems, USA, 1:200), anti-Glutamine
830 synthetase (GS, MAB302, Millipore, USA, 1:500), anti-CD31 (#557355, BD

831 Biosciences, USA, 1:100), anti-Lhx2 (sc-19344, Santa Cruz Biotechnology,
832 USA, 1:100), anti-RBPMS (GTX118619, GeneTex, USA, 1:500), anti-Ki67
833 (#556003, BD Biosciences, USA, 1:100), anti-Tuj1 (#801201, BioLgend, USA,
834 1:500), anti-AP2α (3B5-c, DSHB, USA, 1:1000), anti-Rom1 (gift from Robert
835 Moldey, 1:10), anti-Rhodopsin (STJ95452, ST John's Laboratory, UK, 1:100),
836 anti-S-opsin (sc-14363, Santa Cruz Biotechnology, USA, 1:500), anti-M-opsin
837 (AB5405, Sigma, USA, 1:500), anti-GFP (GFP-1010, Aves labs, USA, 1:1000),
838 anti-β-catenin (#610153, BD Biosciences, USA, 1:1000), anti-N-cadherin
839 (#610920, BD Transduction Laboratories, USA, 1:500), anti-BrdU (#347580, BD
840 Biosciences, USA, 1:100), and anti-phospho-histone H3S10 (#06-570, Millipore,
841 USA, 1:2000) antibodies.
842
843 *Counting of retinal cells and synapses*
844 The cell number of each retinal cell type was counted around the area 500 μm
845 away from the optic nerve using immunostained retinal sections, and we
846 calculated the cell number per 100 μm width of retinal section. Each retinal cell
847 type was identified as follows: rod (Otx2 signal at the nuclear periphery), cone
848 (Arr3+, Otx2 signal in the soma), BC (Chx10+), horizontal cell (Calbindin+ and
849 AP2α-), AC (AP2α+), RGC (RBPMS+), and Müller glial cell (Lhx2+).
850
851 It was difficult to count the number of synapses between photoreceptor and BC
852 in 2D images of the cKO retina because the direction of the photoreceptor
853 synapse was aberrant. Thus, the number of synapses was counted using 3D
854 images reconstituted from immunostained vertical sections. The image of

855 vertical sections was acquired with a width (x-axis) of 126.8 μm, a length (y-
856 axis) of 126.8 μm, and a thickness (z-axis) of 12 μm. Then, the number of
857 synapses per 1 mm^2 of retinal surface was calculated. The regions distal to the
858 IPL were analyzed because photoreceptor synapses were not observed below
859 the IPL. Since the synapses between the rod photoreceptor and BC in the cHet
860 retina were highly dense, we counted them every 1.8 μm using 2D images and
861 integrated them. We confirmed that the number of synapses was not different
862 between 2D and 3D images.
863
864 *Processing of tissues for electron microscopy*
865 The eyeball was removed, cut in the cornea, and fixed with 2.5% glutaraldehyde
866 in PBS for 2.5 h on ice, and with 2% osmium tetroxide for 2 h. Retinas were
867 washed with 1x PBS, and dehydrated with a graded series of ethanol, followed
868 by propylene oxide, and embedded in Epon 812. Seventy-nm ultrathin sections
869 were cut by an ultramicrotome (Ultracut E; Reichert-Jung, Germany), mounted
870 on nickel grids, and stained with 2% uranyl acetate for 4 h and with nitrate for 5
871 min. Retinal sections were observed by transmission electron microscope (H-
872 7500; HITACHI Co, Japan).
873
874 *Reverse transcription real-time quantitative PCR (RT-qPCR)*
875 Total retinal RNA was isolated using Isogen II reagent (Nippon Gene, Japan)
876 following the manufacturer's instructions, and reverse transcribed into cDNA
877 using SuperScript II Reverse Transcriptase (Thermo Fisher Scientific, USA)
878 with random hexamers and oligo (dT) 12-18 primers. Diluted cDNA was used as

879 a template for qPCR using a TB Green® Premix Ex Taq™ II (Tli RNaseH Plus)
880 (Takara Bio, Japan) according to the manufacturer's instructions, on Thermal
881 Cycler Dice® Real Time System II (Takara Bio, Japan). The primer sequences
882 are listed in **Supplementary File 1**.
883
884 *Western blot analysis*
885 Retinal tissues were lysed in RIPA buffer (50 mM Tris-HCl pH7.6, 150 mM NaCl,
886 1mM EDTA, pH8.0, 1% Nonided-P40, 1% Sodium Deoxycholate, 0.1% SDS),
887 cooled for 10 min on ice, and centrifuged at 14,000 rpm for 10 min at 4ºC. Then,
888 a 3x sample buffer was added to the supernatant, and it was boiled for 10 min
889 at 95ºC. Samples were separated by SDS-PAGE and transferred to PVDF
890 membranes (Millipore, USA) using a wet transfer cell (Bio-Rad, USA). The
891 membranes were soaked in blocking buffer (5% skim milk (W/V) and 0.1%
892 Tween 20 in TBS) and incubated with anti-l-afadin antibody (1:1000) overnight
893 at 4°C or horseradish peroxidase-conjugated anti-GAPDH antibody (1:5000) for
894 2 h at room temperature. After washing with TBS/0.1% Tween-20 four times for
895 10 min each, the membranes were incubated with horseradish peroxidase-
896 conjugated anti-mouse IgG antibody for 2 h at room temperature. Signals were
897 detected using ImmunoStar LD (Fujifilm Wako, Japan).
898
899 *Antibody production*
900 cDNA fragments encoding a C-terminal portion of mouse mGluR6 (853-871
901 residues) were subcloned into *pGEX4T-1* plasmid (Amersham, UK). The GST-
902 mGluR6 fusion protein was expressed in *Escherichia coli* strain BL21 and

903 purified with glutathione Sepharose 4B (GE Healthcare, USA) according to the
904 manufacturer's instructions. The anti-mGluR6 antibody was obtained by
905 immunizing guinea pigs with the purified GST-mGluR6.
906
907 *ERG recordings*
908 Mice were dark-adapted for more than 1 h and then anesthetized deeply by
909 intraperitoneally injecting mixed anesthesia consisting of (in mg/kg) 0.3
910 medetomidine, 4 midazolam, and 5 butorphanol tartrate. We also administered
911 0.1% tropicamide and 0.1% phenylephrine to the eyes to dilate the pupil. ERGs
912 were recorded via contact lens electrodes (Mayo Corporation, Japan), and data
913 were sampled at 1,250 Hz by using the PuREC system (Mayo Corporation,
914 Japan). Reference and ground electrodes were placed in the mouth and on the
915 tail, respectively. Full-field stimulation was presented via an optical stimulator
916 equipped with white LED (RMG; Mayo Corporation, Japan), and its light
917 intensity and duration were controlled by an LED visual stimulator (LS-100;
918 Mayo Corporation, Japan). Scotopic ERGs were recorded by applying 5-ms
919 flash (1.0×10^4 cd/m^2) three times in the dark. After 10 min light adaptation
920 (31.6 cd/m^2), photopic ERGs were recorded by superimposing 5-ms flash ($1.0 \times$
921 10^4 cd/m^2) sixteen times on the adapting background. After noise reduction by a
922 program installed in the PuREC system, ERGs were lowpass-filtered (<50 Hz)
923 on Python3 and then analyzed.
924
925 *MEA recordings*
926 The retinal preparation was made as described previously (Takeuchi et al.,

927 2018). In brief, a dark-adapted (>1 h) mouse was sacrificed by cervical
928 dislocation under a dim red light, and its eyes were enucleated. Under a
929 stereomicroscope equipped with an infrared (IR) image converter (V6833P,
930 Hamamatsu Photonics, Japan) and IR illumination (HVL-IRM, Sony, Japan), the
931 retina was isolated from the eye, and the dorsal retina was placed ganglion cell
932 layer side down onto the MEA (Multichannel Systems, Germany,
933 60pMEA200/30iR-Ti: 60 electrodes, electrode size 30x30 μm, inter-electrode
934 distance 200 μm) and attached to the MEA by suction using a vacuum pump
935 (Constant Vacuum Pump, Multichannel Systems, Germany). When another type
936 of MEA (Multi Channel Systems, Germany, 60MEA200/30iR-Ti-gr: 60
937 electrodes, electrode size 30×30 μm, inter-electrode distance 200 μm) was
938 used, a piece of anodisc (13 mm, 0.2 μm, 6809-7023, Whatman, UK) and a
939 weight with nylon fibers were placed on the retina to improve the contact with
940 MEA. Then, the MEA chamber was constantly superfused with Ames' medium
941 (A1372-25, United States Biological, USA) bubbled with 95% O_2/5% CO_2 at the
942 rate of 6.0 mL/min at 32°C. After >30 min superfusion, we started recordings.
943 Signals were amplified, sampled at 20 kHz, and stored using MEA2100-Lite-
944 System (Multichannel Systems, Germany). In pharmacological experiment, L-
945 (+)-2-amino-4-phosphonobutyric acid (10 μM; L-AP4, 23052-81-5, Tocris, UK)
946 dissolved in Ames' medium was bath-applied to the retina.
947
948 Light stimulus generated by Psychtoolbox-3 on MATLAB was presented on a
949 monitor display (P2314H, Dell, Japan), and projected through optics on the
950 photoreceptor layer. Flashes (24.2 or 27.3 cd/m^2, 2 s in duration, 1,600 μm in

diameter) were presented seven times every 8 s. In some experiments, the retina was stimulated by 5-ms diffuse green LED light (2.5 x 10^4 photons/s/μm^2 for mERG recordings, 4.8 x 10^3 photons/s/μm^2 for rod stimulation: LED; λ_{max} = 518 nm, IF filter; 510 nm, FWHM 10 nm, #65-697, ND filter; OD 1.0 #47-207, Edmund Optics, USA) or 5-ms diffuse UV LED light (3.3 x 10^4 photons/s/μm^2 for cone stimulation; LED; λ_{max} = 370 nm, KED365UH, IF filter; 360 nm, FWHM 10 nm #67-827, Edmund Optics, USA) controlled by a function generator (WF1973, NF Corporation, Japan) three times every >10 s.

Spike sorting

For spike sorting, the algorithm described on Python3 was used. The data were high pass-filtered (>300 Hz), and minimum values were detected by using scipy.signal.argrelmin function. To decide whether each detected minimum value reflects the peak of spike event, we used a threshold value defined by the following formula (Quiroga et al., 2004).

$$threshold = -4 \times median\ (|x|\ /0.6745),$$

where x represents each data point*, as* our data were recorded extracellularly, the threshold polarity was reversed. The detected values (> threshold) were judged as spike events, and the waveform of spikes was obtained from the data points between -1 and +2 ms from each peak. The 1st and 2nd differences (numpy.diff) of each waveform were joined, and then principal component analysis (PCA) was carried out for feature compression to two dimensions. Then, clustering was performed using template matching method (Zhang et al., 2004) and UMAP. To check the accuracy of spike sorting, we performed

975 autocorrelation analysis to confirm the presence of a refractory period (the
976 number of spike events at ± 1 ms bin was less than 1% of total spike events).
977
978 *Classification of RGCs based on PSTH*
979 Using the MEA system, we recorded firing responses of RGCs to light
980 stimulation (0.24 or 0.27 cd/m^2 in intensity, 2 s in duration) presented 7 times
981 from the monitor and calculated the PSTHs with 20-ms bin. The PSTH was
982 smoothed by a Gaussian filter (kernel size: 3 bins, $\sigma = 1$). The light-evoked
983 response was defined when the firing rate of the PSTH exceeded the threshold
984 (increment >+4 SDs, decrement <-2 SDs) determined by the spontaneous firing
985 rate for 2 s before light stimulation. RGCs were classified into “ON” (increment
986 within 2 s after light onset), “ON-OFF” (increment within 2 s after light onset and
987 offset), “ON-OFF inhibition” (decrement within 2 s after light onset and offset),
988 “OFF” (decrement within 2 s after light onset and/or increment within 2 s after
989 light offset), and “None” (no clear change in firing rate) types.
990
991 To determine the type of photoreceptor inputs, a 2-s flash of green LED (λ_{max} =
992 510 nm, 4.8x10^3 photons/s/μm^2) and 2-s flash of UV LED (λ_{max} = 360 nm,
993 3.3x10^4 photons/s/μm^2) were applied seven times at 8-s interval serially. We
994 classified RGCs into “Rod (+)” (responded to green LED), “Cone (+)-Rod (+)”
995 (responded to both green and UV LEDs), “Cone (+)” (responded to UV LED),
996 and “None” responsive types.
997
998 *Receptive Field*

The reverse correlation method was used to detect the receptive field (RF). Pseudorandom checkerboard patterns (32×32 pixels; black/white = 1; pixel size 50×50 μm) were applied to the retina at 60 Hz. The checkerboard images that preceded each spike event (20 frames) were averaged (spike-triggered average: STA). The intensity of each pixel was +1 or 0, and thus the area with pixels that exceeded +5 SDs of the noise intensity (~0.5: the SD calculated from the frame at t = 0) were judged to be an RF, and the pixel with maximal intensity was defined as the RF center. An ellipse was fitted to the area, and its long and short axes, as well as the area, were calculated. To estimate the temporal profile of RF, the mean intensity of 5×5 pixels in the RF central region was obtained from a series of averaged frames. The temporal profile of RF was normalized, assuming that the mean of the frame at t = 0 was 0. The normalized temporal profiles were compressed into two dimensions by using UMAP, and then k-means clustering was performed.

Visual behavioral tests

We examined the visual behavior of the *Afadin* cHet, *Afadin* cKO, and ONC (optic nerve crush) mice. The optic nerve crush was performed as previously described (Cameron et al., 2020), and successful nerve damage was confirmed by absence of a pupillary light reflex.

The optomotor response (OMR) test was performed as described previously with slight modifications (Kretschmer et al., 2015, 2013; Warwick et al., 2024). Mice were placed on a central platform surrounded on all four sides by

monitors, with a mirror positioned on the floor beneath the platform. Stimuli were comprised of vertical square wave gratings (white; 116.2 cd/m^2, black; 1.66 cd/m^2, Michelson contrast; 97.2) with spatial frequency (0.056, 0.10, or 0.27 cycles/degree) manipulated to induce the illusion of a virtual cylinder with identical spatial frequency across the entire display at a constant speed of 12 degree/second. Each trial consisted of 2 min of drifting gratings, followed by either 30 s or 2 min of a uniform white screen (88.6 cd/m^2), and then another 2 min of drifting gratings in the opposite direction. This trial was repeated three or four times. Head movements were tracked using a camera (29.97 fps, AN-S093, KEIYO, Japan) and the number of head movements consistent with the stimulus direction and speed was counted in a blind manner.

For the visual cliff test, mice were placed along the centerline of a transparent plane (60x60 cm, width x length), positioned 80 cm high above the floor. The half of the arena was immediately beneath the platform covered by a black-and-white checkered sheet (pixel size; 2x2 cm) (shallow side) and the other half was a transparent plane, below which the identical checkerboard pattern was located on the floor (deep side). Mouse behavior was recorded using the camera, and the percentage of time spent on each side during the first 5 minutes was calculated by an observer in a blind manner.

Statistical Analysis

In Figures 2H-L, 3E and 3F, a generalized linear mixed model (GLMM) was

used. In Figures 2F, 2G, 9D, and 9F, a log1p-transformation followed by a linear mixed model (LMM) was applied. In Figures 2M, 9B, Figure 4-figure supplement 1A-C and E, Student's t-test was used. In Figure 8C and 8D, Dunn's test was used. In Figure 8F, the one-way ANOVA followed by Tukey's multiple comparisons test was applied. In Figure 3-figure supplement 1C, LMM was used. In Figures 5C-F and 7D, the Mann-Whitney U test was used. In Figure 6A, the Kolmogorov-Smirnov test was used. In Figure 4-figure supplement 1D, F, and Figure 9-figure supplement 1B–D, in which multiple stages and genes were analyzed, and data distribution and variance differed across conditions. Statistical tests (Student's t-test, Welch's t-test, or Mann–Whitney U test) were chosen accordingly. The specific statistical test used for each stage or gene is described in the corresponding Figure legends. Error bars denote standard error. The following asterisks in Figures indicate *p* values: * <0.05, ** <0.01, *** <0.001.

Data availability

Source data for Figures 2F–M; 3E, F; 4D; 5A-F; 6C, D; 7B–H; 8C, D, F; 9B, D, F; 3-figure supplement 1C; 4-figure supplement 1A-F; and 9-figure supplement 1B–D, together with full-size blot images of Figure 2-figure supplement 1C, are available at https://zenodo.org/records/17037918.

Acknowledgments

We thank Dr. Takahisa Furukawa for providing *Dkk3*-Cre mice, Dr. Toshihiko

Fujimori for providing R26R-H2B-EGFP mice, Takefumi Yamamoto for technical assistance in electron microscopy, Drs. Akishi Onishi, Katsunori Kitano, and Kiyo Sakagami for helpful comments, and Y. Shibata for technical assistance.

Funding

This work was supported by Grant-in-Aid for Scientific Research (B) (24390019), Fund for the Promotion of Joint International Research (22KK0137), Early-Career Scientists (23K15920), and Research Activity Start-up (22K20698) from the Japan Society for the Promotion of Science (JSPS), Takeda Science Foundation, the Kobayashi Foundation, JST PRESTO (08062795), and R-GIRO.

1084 **FIGURE LEGENDS**

1085 **Figure 1 Afadin is localized to the OS, OLM, OPL, INL, and IPL.**

1086 **A**. Immunostaining of the wild-type (WT) mouse retina (1M) with anti-afadin
1087 antibody. The afadin signals disappeared in the pre-absorbed (peptide+)
1088 sample. The signal observed in the peptide+ image represents the background
1089 and non-specific staining. Nuclei were stained with DAPI (blue).

1090 **B**. Immunostaining of the WT retinal section (1M) using anti-afadin (green)
1091 combined with anti-Nectin-1 (red) antibody. Necin-1 was partially co-localized
1092 with afadin in the OLM, OPL, and IPL.

1093 **C**. Afadin was colocalized with nectin-1, nectin-2, and nectin-3 at the OLM. The
1094 1M WT retinal sections were immunostained with anti-afadin (green), anti-
1095 nectin-1 (red, upper panels), anti-nectin-2 (red, middle panels), and anti-nectin-
1096 3 (red, lower panels) antibodies.

1097 **D**. Immunostaining of 1M WT retinal sections using anti-afadin (green), anti-
1098 SCGN (red), and anti-PKCα(white) antibodies. Afadin signals overlapped with
1099 SCGN and PKCα in the OPL and INL, and PKCα in the IPL.

1100

1101 OS, the outer segment; IS, the inner segment; OLM, the outer limiting
1102 membrane; ONL, the outer nuclear layer; OPL, the outer plexiform layer; INL,
1103 the inner nuclear layer; IPL, the inner plexiform layer; GCL, the ganglion cell
1104 layer.

1105

1106

1107 **Figure 1-figure supplement 1 Afadin is localized to AJs in developing and**

1108 **mature retinas.**

1109 **A**. Immunostaining of the WT retinal section (1 M) with anti-afadin (green)
1110 antibody combined with nectin-2 (red, upper and middle panels) and nectin-3
1111 (red, lower panels) antibodies. Nectin-2 and nectin-3 were localized in the OLM.
1112 The nectin-2 signal in the IPL was insufficient for reliable assessment of its
1113 localization and colocalization.

1114 **B**. Immunostaining with anti-afadin (green), anti-nectin-1 (red), and anti-Arr3 (a
1115 cone marker, white) antibodies. Afadin was co-localized with nectin-1 in cone
1116 synapses.

1117 **C**. Immunostaining of the WT retinal section (P0) with anti-afadin (green) and
1118 anti-nectin-1 (red) antibodies.

1119

1120

1121 **Figure 2 Outer retinal lamination is severely disrupted in the *Afadin* cKO**
1122 **retina.**

1123 **A**. Toluidine blue staining of the cHet and cKO mouse retinal sections at 1M.
1124 The retinal layer structure was disrupted in the cKO retina. Rod photoreceptor
1125 nuclei (insets) were deeply stained. The red asterisk indicates a rod
1126 photoreceptor. The arrowhead indicates a rosette-like structure. Scale bar in the
1127 inset, 2.5 μm.

1128 **B**. Representative images of the cHet and cKO retinas (1M) stained with anti-
1129 Rom1 (green) and anti-Rhodopsin (Rho, red) antibodies and PNA-rhodamine
1130 (white). Rom1, Rhodopsin, and PNA signals were remarkably decreased and
1131 scattered in the cKO mice. The arrowhead indicates the rosette-like structure.

1132 **C**. Representative images of the cHet and cKO retinas (1M) stained with anti-S-
1133 opsin (green) and anti-M-opsin (white) antibodies and PNA-rhodamine (red).
1134 Cone OSs were aberrantly located in the cKO retina. The arrowhead indicates
1135 the rosette-like structure.
1136 **D**. Retinal flat-mount immunostaining of the cHet and cKO retinas
1137 immunostained with anti-Arr3 (green, left panels), anti-Rom1 (green, right
1138 panels), and anti-Rhodopsin (red, right panels) antibodies. The inset, a
1139 magnified view of the area enclosed by the dashed box, shows an area with a
1140 relatively high density of photoreceptors in the cKO retina.
1141 **E**. Immunofluorescent analysis of the cHet and cKO retinas (1M) using anti-Arr3
1142 (cone marker, green in the left panels), anti-Otx2 (photoreceptor and BC
1143 marker, red in the left panels), anti-RNA binding protein with multiple splicing
1144 (RBPMS, RGC marker, green in second left panels), anti-Chx10 (BC marker,
1145 red in second left panels), anti-Lhx2 (Müller glia marker, green in third left and
1146 third right panels), anti-AP2α (AC marker, red in third left and second right
1147 panels), and anti-Calbindin (horizontal cell and AC marker, white in third left and
1148 right panels) antibodies. Arrowheads indicate RGCs and ACs near the rosette-
1149 like structure.
1150 **F-L**. Distribution of each retinal cell type based on their distance from the retinal
1151 surface in the cHet and cKO mice. The shortest distance between each cell
1152 body and either the outer or inner retinal surface was measured using ImageJ.
1153 For cones (**F**), rods (**G**), bipolar cells (**H)**, horizontal cells (**I**), and Müller glia (**J**),
1154 the distance was measured from the outer retinal surface. For amacrine cells
1155 (**K**) and retinal ganglion cells (**L**), the distance was measured from the inner

retinal surface. *Note: In **K** and **L**, the X-axis is reversed such that short distance appears on the right.* The outer retinal surface was defined as the region immediately above the somata of the outer retinal cells. Bin width was 5 μm (0–5 up to 175–180 μm). Inset box plots show the distribution of raw values, with whiskers extending to 1.5 times the interquartile range and dots indicating outliers (cHet; Cone 3.45 ± 2.72 μm, n = 141 cells from 4 mice, Rod 23.1 ± 13.8 μm, n = 3055 cells from 4 mice, Bipolar cell 59.4 ± 6.74 μm, n = 432 cells from 3 mice, Horizontal cell 57.5 ± 6.45 μm, n = 20 cells from 3 mice, Müller glial cell 69.9 ± 7.61 μm, n = 191 cells from 3 mice, Amacrine cell 51.8 ± 18.6 μm, n = 362 cells from 3 mice, Ganglion cell 8.71 ± 3.71 μm, n = 93 cells from 3 mice, cKO; Cone 14.9 ± 14.2 μm, n = 42 cells from 4 mice, Rod 35.5 ± 30.5 μm, n = 716 cells from 4 mice, Bipolar cell 56.6 ± 32.7 μm, n = 1236 cells from 5 mice, Horizontal cell 37.2 ± 31.2 μm, n = 14 cells from 4 mice, Müller glial cell 45.9 ± 22.9 μm, n = 95 cells from 4 mice, Amacrine cell 63.8 ± 34.6 μm, n = 265 cells from 4 mice, Ganglion cell 38.9 ± 33.6 μm, n = 116 cells from 5 mice). ***$p < 0.001$, **$p < 0.01$ by a generalized linear mixed model (GLMM) or log1p-transformation followed by a linear mixed model (LMM). Statistical tests: Cone, Rod, log1p-transformation followed by a LMM; Bipolar cell, Horizontal cell, Müller glial cell, Amacrine cell, Ganglion cell, GLMM.

M. The thickness of the cHet and cKO retinas at 1M. (cHet; 137.9 ± 11.1 μm, n = 4, cKO; 135.5 ± 9.1 μm, n = 5). Error bars, mean ± SD. No significant difference was observed between the two groups ($p = 0.735$ by Student's t-test).

1180 **Figure 2-figure supplement 1 AJs in the OLM are disrupted in the *Afadin***
1181 **cKO retina.**
1182 **A, B.** Immunostaining of *Dkk3*-Cre; R26R-H2B-EGFP and R26R-H2B-EGFP
1183 retinas at P0 (**A**) and 1M (**B**) using an anti-GFP antibody (green). The Cre
1184 recombinase activity was detected in the *Dkk3*-Cre; R26R-H2B-EGFP retina at
1185 both stages.
1186 **C**. Western blot analysis of the cHet and cKO retinas using anti-afadin (upper
1187 panel) and anti-GAPDH (lower panel) antibodies. No significant afadin band
1188 was detected in the cKO retina.
1189 **D.** Electron microscopic analysis of the cHet and cKO retinas (1M). A few
1190 ectopic disc structures were observed in the cKO retina. The inset, a magnified
1191 view of the area enclosed by the dashed box, shows the ectopic disc structure
1192 at high magnification. The arrowhead indicates the outer segment disc
1193 structure. RPE, the retinal pigment epithelium.
1194 **E.** Immunostaining of the cHet and cKO retinal sections at 1M using anti-β-
1195 catenin (green, left panels), anti-nectin-1 (red, left panels), and anti-N-cadherin
1196 (green, right panels). Signals of these markers were not observed at the outer
1197 retinal surface in the cKO mice.
1198 **F.** Immunostaining of the cHet and cKO retinal sections at 1M using anti-
1199 glutamine synthetase (GS, green) antibody. Obvious GS signals were observed
1200 in the inner side of the retina but not at the outer retinal surface in the cKO
1201 mice.
1202 **G.** Retinal sections from the cHet and cKO mice at 1M were stained with an
1203 anti-CD31 antibody (red). No notable vascular differences were observed in the

1204 cKO retina.

1205

1206

1207 **Figure 3 Photoreceptor-BC synapses were severely impaired and**

1208 **mislocalized in the *Afadin* cKO retina.**

1209 **A**. Immunostaining of the cHet and cKO retinal sections at 1M with anti-

1210 Bassoon (green), anti-mGluR6 (red), and anti-PKCα (white) antibodies. The

1211 arrowhead indicates the synapse between rod and rod-BC. Small panels are

1212 magnified views of the area enclosed by the dashed box.

1213 **B**. Immunostaining of the cHet and cKO retinas at 1M with anti-GluR5 (green),

1214 anti-PSD95 (red), and anti-PKARIIβ (white) antibodies. The arrowhead indicates

1215 the synapse between the cone and OFF-BC. Small panels are magnified views

1216 of the area enclosed by the dashed box.

1217 **C**. 3D projection of confocal image immunostaining with anti-Bassoon (green),

1218 anti-mGluR6 (red), and anti-PKCα (white) using the cHet and cKO retina (1M).

1219 The inset, a magnified view of the area enclosed by the dashed box, shows the

1220 synapse between the rod and BC at high magnification. The arrowhead

1221 indicates the ribbon synapse between rod and rod-BC.

1222 **D**. 3D projection of confocal image immunostaining with anti-GluR5 (green),

1223 anti-PSD95 (red), and anti-PKARIIβ (white) antibodies using the cHet and cKO

1224 retina (1M).

1225 **E**. Quantification of the number of synapses between rod photoreceptor and

1226 rod-BC under 1mm^2 of retinal surface in the cHet and cKO mice immunostained

1227 with anti-Bassoon, anti-mGluR6, and anti-PKCα antibodies (cHet; 599.6 ± 49.7,

1228 n = 3, cKO; 29.1 ± 12.75, n = 4, 10 images from each mouse. Error bars, mean
1229 ± SD. ***$p < 0.001$ by a GLMM). The number of synapses decreased to about
1230 5 % of the cHet in the cKO retina.
1231 **F.** Quantification of the number of synapses between photoreceptor cells and
1232 type 3 OFF-BC under 1mm^2 of the retinal surface in the cHet and cKO mice
1233 immunostained with anti-GluR5, anti-PSD95, and anti-PKARIIβ antibodies
1234 (cHet; 24.0 ± 2.8, n = 3, cKO; 4.1 ± 1.3, n = 3, 10 images from each mouse.
1235 Error bars, mean ± SD. ***$p < 0.001$ by a GLMM). The number of synapses
1236 between the photoreceptor and Type 3 OFF-BC in the cKO retina decreased to
1237 about 15% of the cHet retina.
1238
1239
1240 **Figure 3-figure supplement 1 The OS and photoreceptor-BC synapses are**
1241 **affected in the developing *Afadin* cKO retina.**
1242 **A**. Immunostaining of the cHet and cKO retinas at 1M with anti-PKCα (green),
1243 anti-PSD95 (red) antibodies. Small panels are magnified views of the area
1244 enclosed by the dashed box.
1245 **B**. 3D projection of confocal image immunostaining with anti-PKCα (green),
1246 anti-PSD95 (red) antibodies.
1247 **C**. Quantification of the number of contacts between photoreceptor and rod ON-
1248 BC under 1mm^2 of retinal surface in the cHet and cKO mice immunostained
1249 with anti-GluR5, anti-PSD95, and anti-PKARIIβ antibodies (cHet; 550.5 ± 29.1,
1250 n = 3, cKO; 65.2 ± 7.9, n = 3, 10 images from each mouse. Error bars, mean ±
1251 SD ***$p < 0.001$ by a LMM). The number of synapses between the rod and ON

1252 BC decreased to about 10% of the cHet in the cKO retina.

1253 **D, E.** The cHet and cKO retinas (1M) stained with anti-EAAT5 (green), anti-
1254 SCGN (red in **D**), and anti-PKCα (white in **D**), anti-PSD95 (red in **E**) antibodies
1255 and PNA-rhodamine (white in **E**). Glutamate transporter EAAT5 localization to
1256 ON and OFF BC axon terminals and photoreceptor axon terminals were
1257 unaffected in the cKO retina. Small panels are magnified views of the area
1258 enclosed by the dashed box.

1259 **F.** Immunostaining of the cHet and cKO retinal sections at 1M with anti-ChAT
1260 (green), anti-Calretinin (red), and anti-Calbindin (white) antibodies. Obvious
1261 ChAT bands, Calretinin bands, and Calbindin bands were observed in the IPL of
1262 the cKO mice.

1263

1264

1265 **Figure 4 GluR5 is ectopically localized to ON and OFF BC processes.**

1266 **A.** Immunostaining of the cHet and cKO retinal sections at 1M with anti-GluR5
1267 (green), anti-SCGN (red), and anti-PKCα antibodies. GluR5 signal was
1268 obviously observed in the IPL of the cKO retina. Yellow and white arrowheads
1269 indicate overlap of GluR5 and PKCα, and GluR5 and SCGN, respectively. Small
1270 panels are magnified views of the area enclosed by the dashed box.

1271 **B.** Representative images of the immunostained cHet and cKO retinas (1M)
1272 with anti-GluR5 (green), anti-vGlut1 (red), and anti-SCGN (white) antibodies.

1273 **C.** Immunofluorescent analysis of the cHet and cKO retinas (1M) with anti-HPC-
1274 1 (AC marker, green in the left panels), anti-Calbindin (red in the left panels),
1275 anti-Tuj1 (RGC marker, green in the right panels), and anti-RBPMS (red in the

1276 right panels) antibodies. Arrowhead indicates acellular patches.

1277 **D.** The number of each retinal cell type per 100 μm width of retinal section at

1278 1M (cHet; Rod 185.9 ± 17.5, Cone 8.4 ± 1.4, Bipolar cell 45.5 ± 5.2, Ganglion

1279 cell 7.4 ± 1.5, Horizontal cell 2.1 ± 0.4, Amacrine cell 30.1 ± 3.2, Müller glial cell

1280 15.7 ± 2.3, n = 4, cKO; Rod 57.3 ± 17.3, Cone 3.5 ± 1.3, Bipolar cell 64.6 ± 7.8,

1281 Horizontal cell 1.9 ± 0.7, Amacrine cell 27.6 ± 6.3, Ganglion cell 5.7 ± 0.9,

1282 Müller glial cell 12.4 ± 1.5, n = 4. Error bars, mean ± SD. Horizontal cell p =

1283 0.58, AC p = 0.52, RGC p =0.10. ***p < 0.001, **p < 0.01, *p < 0.05 by Student's

1284 t test. In the cKO retina, rod and cone photoreceptors and Müller glial cells

1285 significantly decreased to about 30, 40, and 80% of the cHet retina,

1286 respectively, and BCs significantly increased to about 150% of the cHet retina.

1287 **E.** Immunofluorescent analysis of the cHet and cKO retinas at P1, P3, P5, P8,

1288 P11, and P14 using anti-Chx10 (green), anti-Otx2 (red), and anti-Ki67

1289 (progenitor cell marker, white) antibodies. NBL, the neuroblastic layer.

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1291

1292 **Figure 4-figure supplement 1 Retinal cell fate determination is altered**

1293 **during development in the *Afadin* cKO mice.**

1294 **A.** The number of PKCα+ and SCGN+ cells per 100 μm width of retinal sections

1295 at 1M (cHet; PKCα+ 15.8 ± 2.6, SCGN+ 24.7 ± 4.8, n = 4, cKO; PKCα+ 23.6 ±

1296 5.2, SCGN+ 37.4 ± 7.7, n = 4. Error bars, mean ± SD. *p < 0.05 by Student's t-

1297 test). PKCα+ cells and SCGN+ cells significantly increased to 150% of the cHet.

1298 **B.** The number of PKCα+ cell processes in the IPL per 100 μm width of retinal

1299 sections at 1M (cHet; 18.6 ± 0.5, n = 4, cKO; 14.7 ± 1.2, n = 4. Error bars, mean

± SD. ***$p < 0.01$ by Student's t-test). The number of PKCα+ cell processes significantly decreased to 80% of the cHet.

C. The number of each retinal cell types per 100 μm width of retinal sections at P14 (cHet; Rod 176.5± 14.3, Cone 10.6 ± 1.3, Bipolar cell 49.2 ± 1.9, Ganglion cell 8.1 ± 0.8, Horizontal cell 2.5 ± 0.7, Amacrine cell 30.3 ± 2.1, Müller glial cell 17.3 ± 1.4, n = 4, cKO; Rod 98.3 ± 3.8, Cone 4.7 ± 0.9, Bipolar cell 76.8 ± 10.3, Horizontal cell 1.9 ± 0.8, Amacrine cell 32.2 ± 8.9, Ganglion cell 8.5 ± 1.0, Müller glial cell 17.2 ± 3.3, n = 3. Error bars, mean ± SD. Horizontal cell p = 0.36, Amacrine cell p = 0.68, Ganglion cell p =0.57, Müller glial cell p = 0.94. ***$p < 0.001$ by Student's t-test). Rod and cone photoreceptors were significantly reduced to approximately 55% and 45%, respectively, of those in the cHet retina, while BCs significantly increased to approximately 150% in the cKO retina.

D. Expression of photoreceptor and BC marker genes in the cHet and cKO at P14 (Opn1sw and Opn1mw; cone maker, Nrl and Rhodopsin; rod marker, Chx10; BC marker, Trpm1 and mGluR6; ON BC marker). Each gene expression level was normalized to GAPDH, a housekeeping gene. The expression of Nrl, Rhodopsin, Opn1sw, and Opn1mw significantly decreased, whereas the expression of mGluR6, Trpm1, and Chx10 significantly increased in the cKO retina. Error bars, mean ± SD. ***$p < 0.001$, **$p < 0.01$. Statistical tests: Opn1sw Welch's t-test; Opn1mw Student's t-test; Nrl Welch's t-test; Rho Student's t-test; Chx10 Welch's t-test; Trpm1 Student's t-test; mGluR6 Student's t-test.

E. The number of BCs per 100 μm width of retinal sections at P1, P3, and P5 (cHet; P1 4.42 ± 1.09, n = 5, P3 17.2 ± 2.47, n = 3, P5 59.9 ± 7.42, n = 4, cKO;

1324 P1 4.71 ± 1.75, n = 4, P3 18.7 ± 1.74, n = 4, P5 96.4 ± 6.91, n = 3). BCs
1325 (Chx10-positive and Ki67-negative) were significantly increased in the cKO
1326 retina at P5, but not at P1 or P3. Error bars, mean ± SD. P1 p = 0.77, P3 p =
1327 0.56, **p < 0.01 by Student's t-test.
1328 **F.** Expression of rod photoreceptor cell marker genes (Rho and Nrl) in the cHet
1329 and cKO at P1, P3, and P5. Each gene expression was normalized to GAPDH,
1330 a housekeeping gene. The expression of Nrl and Rhodopsin significantly
1331 decreased in the cKO retina at P5. No significant difference was observed at P1
1332 and P3. Error bars, mean ± SD. P1: Nrl p = 0.50, Rho p = 0.81, P3: Nrl p = 0.97,
1333 Rho p = 0.81, *p < 0.05. Statistical tests: P1 Rho, Mann–Whitney U test; P1 Nrl,
1334 P3 Rho, P3 Nrl, P5 Nrl, and P5 Rho, Student's t-test.
1335
1336
1337 **Figure 4-figure supplement 2 Retinal lamination is disputed in the *Afadin***
1338 **cKO retina during development.**
1339 **A**. Immunostaining of the cHet and cKO retinal sections at P1, P3, P5, P8, P11,
1340 and P14 with anti-HPC-1 (green), anti-RBPMS (red) antibodies. Yellow and
1341 white arrowheads indicate acellular patches and aberrant acellular regions not
1342 containing IPL components, respectively.
1343 **B**. Immunostaining of the cHet and cKO retinal sections at P1, P3, P5, P8, P11,
1344 and P14 with anti-AP2α (green), anti-Calbindin (red) antibodies. Yellow and
1345 white arrowheads indicate acellular patches and aberrant acellular regions not
1346 containing IPL components, respectively.
1347 **C**. Immunostaining of the cHet and cKO retinal sections at P8, P11, and P14

with anti-Rom1 (green), anti-Rho (red) antibodies. Yellow and white arrowheads indicate the rosette-like structure and aberrant acellular regions not containing Rom1 and Rho signals, respectively.

D. Representative images of the cHet and cKO retinas (P11) stained with antibodies against PKARIIβ (green, left panels), Arr3 (red, left panels), Bassoon (green, right panels), mGluR6 (red, right panels), PKCα (white, right panels). The inset, a magnified view of the area enclosed by the dashed box, shows the synapse between Cone and Type 3 OFF-BC (left panels), and ribbon synapses (right panels) at high magnification.

Figure 5 Reduction of the a- and b-waves in the cKO retina.

A. *Left*, Scotopic ERGs from the anesthetized cHet (n = 3; black) and cKO (n = 3; magenta) mice. A flashlight (duration; 5 ms, white LED, intensity; 1.0×10^4 cd/m^2) was applied to eyes in the dark three times. Traces: averaged (dark) and SD (pale). *Right*, Photopic ERGs. The anesthetized cHet (black) and cKO (magenta) mice were light (31.6 cd/m^2) adapted for 10 min, and then, a flashlight (duration; 5 ms, intensity; 1.0×10^4 cd/m^2) was superimposed on the adapted light to eyes 16 times.

B. mERGs recorded by MEA from the cHet (black) and cKO (magenta) retinas. A flashlight (duration; 5 ms, green LED; λ_{max} = 510 nm, intensity; 2.5×10^4 photon/s/μm^2) was applied to the isolated whole retina 3 times every 10 s. Control (solid line), L-AP4 (10 μM) (pale), and after washout (dotted line). *Left*, cHet; SD of the amplitude for 200 ms after the flash onset; >1,000 μV, 292/413

electrodes, n = 7 retinas from 4 mice. *Right*, cKO; SD of the amplitude for 300 ms after the flash onset; >50 μV, 13/590 electrodes, n = 10 retinas from 6 mice.

C. *Left*, Amplitude of the a-wave (cHet; 155 ± 58.7 μV, n = 292 electrodes, n = 7 retinas from 4 mice, cKO; 44.1 ± 21.7 μV, n = 13 electrodes, n = 10 retinas from 6 mice, Error bars, mean ± SD. ***$p < 0.001$, Mann-Whitney U test). *Right*, Amplitude of the b-wave (cHet; 163 ± 76.5 μV, n = 292 electrodes, cKO; 28.4 ± 19.6 μV, n = 13 electrodes, $p = 7.08 \times 10^{-9}$, Mann-Whitney U test).

D. Ratio of b-wave amplitude / a-wave amplitude (b / a) (cHet; 1.19 ± 0.67, cKO; 0.59 ± 0.23, Error bars, mean ± SD. ***$p < 0.001$, Mann-Whitney U test).

E. *Left*, Implicit time of the a-wave (cHet; 62.6 ± 10.8 ms, n = 292 electrodes, cKO; 149 ± 16.0 ms, n = 13 electrodes, $p = 5.34 \times 10^{-10}$, Mann-Whitney U test). *Right*, Implicit time of the b-wave (cHet; 127 ± 25.3 ms, n = 292 electrodes, cKO; 229 ± 60.7 ms, n = 13 electrodes, Error bars, mean ± SD. ***$p < 0.001$, Mann-Whitney U test).

F. Difference between implicit time of the b-wave and that of the a-wave (cHet; 64.4 ± 22.0 ms, n = 292 electrodes, cKO; 80.4 ± 50.9 ms, n = 13 electrodes, Error bars, mean ± SD. $p = 0.40$, Mann-Whitney U test).

Figure 6 RGC classification based on the light-evoked responses.

A, B. Representative light-evoked responses. **A**, The cHet retina. **B**, The cKO retina. A flash (duration; 2 s, intensity; 24.2 or 27.3 cd/m^2) was applied seven times every 8 s. After spike sorting, the raster plots (*top*) and the PSTHs (*bottom*) were created (20 ms/bin). Based on the PSTHs, RGC responses were

classified into “ON”, “ON-OFF”, “ON-OFF inhibition”, “OFF”, and “None” types.

C. Ratio of responded to non-responded types (cHet; n = 434, cKO; n = 388).

D. Rod and cone inputs to RGCs (cHet; n = 434, cKO; n = 388). 2-s green LED flash (λ_{max} = 510 nm, intensity; 4.8 x 10^3 photons/s/μm^2) and UV LED flash (λ_{max} = 360 nm, intensity; 3.3 x 10^4 photons/s/μm^2) were applied 7 times every 8 s to stimulate mainly rods and S cones, respectively.

Figure 7 Spatiotemporal properties of the RF of RGCs.

A. *Left*, An isolated retina on the MEA. Electrodes (black square), mapped RFs (colored oval), and the region where immunohistochemical examination was performed after recording (magenta square). Scale bar 200 μm. *Right*, Immunohistochemical staining with anti-PKARIIβ (green), anti-PSD95 (red), and anti-PKCα (white) antibodies. IPL: the inner plexiform layer.

B. Number of cells whose RF was clearly observed (cHet; 195/434 cells, n = 4 retinas, cKO; 69/388 cells, n = 4 retinas).

C. The spatial profile of RFs in the cHet (*Top*) and cKO (*Bottom*) retinas. Color scale illustrates high (red) to low (blue) relative to the average (green). Scale bar 100 μm.

D. RF size of the cKO retina was significantly smaller than that of the cHet retina (cHet; 0.0307 ± 0.0210 mm^2 n = 195, n = 4 retinas, cKO; 0.0161 ± 0.00839 mm^2 n = 69, n = 4 retinas, Error bars, mean ± SD. ***$p < 0.001$, Mann-Whitney U test).

E. Short and long axes of the oval fitted to each RF.

1420 **F**. Clustering of RGCs based on the temporal profile of RFs. Five clusters were
1421 visualized (cHet; circle, n = 4 retinas, cKO; x, n = 4 retinas).
1422 **G**. Temporal profiles from each cluster were superimposed separately (cHet;
1423 black, cKO; magenta, mean; solid line, SD: pale line).
1424 **H**. Time-to-peak latency of the temporal profile (cHet; filled bar, cKO; open bar).
1425 Mean ± SD. (cHet; cluster 1; -113 ± 15 msec, n = 31, cluster 2; -111 ± 13 msec,
1426 n = 36, cluster 3; -200 ± 31 msec, n = 51, cluster 4; -171 ± 19 msec, n = 37,
1427 cluster 5; -178 ± 28 msec, n = 40, cKO; cluster 3; -163 ± 21 msec, n = 10,
1428 cluster 4; -238 ± 34 msec, n = 36, cluster 5; -217 ± 41 msec, n = 23).
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1430
1431 **Figure 8 Visual function is partially retained in the *Afadin* cKO mice.**
1432 **A.** Schematic diagram of the experimental setup for the OMR test. Drifting
1433 square-wave gratings were displayed on monitors surrounding the mouse, and
1434 head-tracking movements were recorded under photopic conditions.
1435 **B.** Representative traces of head-tracking movements in the cHet, cKO, and
1436 ONC mice. Blue lines indicate head movement trajectories, and red segments
1437 denote the periods when the head movement aligned with the direction and
1438 timing of the visual stimulus.
1439 **C.** Summary of head-tracking movements in response to drifting gratings with
1440 spatial frequencies of 0.056, 0.10, and 0.27 cycles/degree, as well as under no-
1441 stimulus conditions, in the cHet, cKO, and ONC mice (cHet; 0.056
1442 cycles/degree 7.30 ± 1.94, 0.10 cycles/degree 7.89 ± 2.00, 0.27 cycles/degree
1443 7.16 ± 1.39, n = 8 mice, cKO; 0.056 cycles/degree 2.36 ± 1.59, 0.10

cycles/degree 1.64 ± 1.29, 0.27 cycles/degree 0.0268 ± 0.103, n = 14 mice, ONC; 0.056 cycles/degree 0.0833 ± 0.121, 0.10 cycles/degree 0.0714 ± 0.161, 0.27 cycles/degree 0.0476 ± 0.128, n = 7 mice). Error bars, mean ± SD. No stimulus; cHet vs cKO $p > 0.9999$, cHet vs ONC $p > 0.9999$, cKO vs ONC $p > 0.9999$, 0.27 cycles/degree; cKO vs ONC $p > 0.9999$. ***$p < 0.001$, Dunn's test.

D. Comparison of head-tracking responses across spatial frequencies (0.056, 0.10, and 0.27 cycles/degree). Statistical comparisons were conducted within each mouse group. No significant differences were detected across spatial frequencies in cHet and ONC mice. In contrast, the cKO mice exhibited a trend toward decreased head-tracking responses with increasing spatial frequency. Error bars, mean ± SD. cHet; 0.056 cycles/degree vs 0.1 cycles/degree $p = 0.3566$, 0.1 cycles/degree vs 0.27 cycles/degree $p = 0.2606$, 0.056 cycles/degree vs 0.27 cycles/degree $p > 0.9999$, cKO; 0.056 cycles/degree vs 0.1 cycles/degree $p = 0.2577$. ***$p < 0.001$, Dunn's test.

E. Schematic diagram of the visual cliff test. The apparatus consists of a single transparent platform spanning two arenas with identical checkerboard patterns: one immediately beneath the platform (the shallow side) and the other on the floor (the deep side), creating a visual cliff. Mice are placed at the boundary between the two sides, and their location preference is evaluated.

F. Time spent on the shallow side during the visual cliff test. The percentage of time spent on the shallow side was significantly higher in the cHet mice than in the cKO and ONC mice. No significant difference was observed between the cKO and ONC mice (cHet; 81.1 ± 14.6, n = 19 mice, cKO; 57.4 ± 17.6, n = 26 mice, ONC; 51.1 ± 6.38, n = 6 mice). Error bars, mean ± SD. ***$p < 0.001$, one-

1468 way ANOVA followed by Tukey's multiple comparisons test.

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1471 **Figure 9 Ectopic AJs are observed in the developing *Afadin* cKO retina.**

1472 **A.** Immunostaining of the cHet and cKO retinas at postnatal day 0 (P0) with
1473 anti-phospho-histone H3S10 (pHH3, mitotic cell marker, green), anti-N-
1474 Cadherin (red in left, second left, and second right panels), anti-Nectin-1 (white
1475 in left, second left, and third right panels), anti-β-catenin (red in the right panels)
1476 antibodies. Ectopic N-cadherin, nectin-1, and β-catenin signals were observed
1477 in the cKO retinas, and pHH3-positive cells were localized near these ectopic
1478 signals.

1479 **B.** Quantification of pHH3-positive cells per 100 μm width of retinal sections in
1480 the cHet and cKO retinas at P0 (cHet; 4.40 ± 0.937, n = 3, cKO; 6.69 ± 0.507, n
1481 = 4). pHH3 positive cells significantly increased in the cKO retina. Error bars,
1482 mean ± SD. $^{**}p < 0.01$ by Student's t-test).

1483 **C.** Retinal sections from the cHet and cKO mice immunostained with anti-pHH3
1484 antibody (red) at P1 (upper panels), P3 (middle panels), and P5 (lower panels).

1485 **D.** Distribution of mitotic cells based on their distance from the outer retinal
1486 surface in the cHet and cKO mice at P1 (top), P3 (middle), and P5 (bottom).
1487 The shortest distance between mitotic cell body and the outer retinal surface
1488 was measured using ImageJ. Bin width was 5 μm (0–5 up to 245–250 μm).
1489 Inset box plots show the distribution of raw values, with whiskers extending to
1490 1.5 times the interquartile range and dots indicating outliers (cHet; P1 3.76 ±
1491 17.5 μm, n = 300 cells from 5 mice, P3 6.73 ± 28.8 μm, n = 469 cells from 3

1492 mice, P5 16.7 ± 44.7 μm, n = 180 cells from 4 mice, cKO; P1 48.4 ± 24.1 μm, n
1493 = 326 cells from 4 mice, P3 76.0 ± 41.9 μm, n = 902 cells from 4 mice, P5 94.0
1494 ± 44.8 μm, n = 163 cells from 4 mice). ***$p < 0.001$ by log1p-transformation
1495 followed by a LMM.
1496 **E.** Retinal sections from the cHet and cKO mice immunostained with anti-BrdU
1497 antibody (green) at P1 (upper panels), P3 (middle panels), and P5 (lower
1498 panels).
1499 **F.** Distribution of progenitor cells based on their distance from the outer retinal
1500 surface in the cHet and cKO mice at P1 (top), P3 (middle), and P5 (bottom). Bin
1501 width was 5 μm (0–5 up to 245–250 μm). Inset box plots show the distribution of
1502 raw values, with whiskers extending to 1.5 times the interquartile range and
1503 dots indicating outliers (cHet; P1 58.0 ± 22.1 μm, n = 2527 cells from 4 mice, P3
1504 83.5 ± 31.7 μm, n = 1904 cells from 4 mice, P5 95.3 ± 32.4 μm, n = 620 cells
1505 from 5 mice, cKO; P1 62.7 ± 35.2 μm, n = 1713 cells from 4 mice, P3 96.5 ±
1506 48.5 μm, n = 2025 cells from 3 mice, P5 90.7 ± 40.5 μm, n = 630 cells from 3
1507 mice). P5 $p = 0.49$, ***$p < 0.001$, **$p < 0.01$ by log1p-transformation followed by
1508 a LMM.
1509
1510
1511 **Figure 9-figure supplement 1 The number of apoptotic cells decreased in**
1512 **the developing cKO retina.**
1513 **A.** Immunostaining of the cHet and cKO retinas (1M) with anti-active caspase 3
1514 (AC3, green) antibody. Arrowhead indicates AC3 positive cell.
1515 **B.** Quantification of apoptotic cells per 1mm^2 in the cHet and cKO retinas at P1,

P3, P5, P14, and 1M (cHet; P1 176.9 ± 31.0, n = 4, P3 164.7 ± 33.4, n = 3, P5 112.1 ± 25.8, n = 4, P14 13.7 ± 4.5, n = 5, 1M 0.577 ± 0.699, n = 4, cKO; P1 322.1 ± 65.5, n = 4, P3 cKO; 267.4 ± 29.1, n = 3, P5 177.1 ± 36.0, n = 4, P14 86.4 ± 17.3, n = 3, 1M 66.7 ± 8.2, n = 5). Apoptotic cells significantly increased in the cKO retina. Error bars, mean ± SD. *$p < 0.05$, **$p < 0.01$. Statistical tests: P1, P3, P5 Student's t-test; P14 Mann–Whitney U test; 1M Welch's t-test.

C. Quantification of progenitor cells labeled by BrdU per 100 μm width of retinal section in the cHet and cKO retinas at P1, P3, and P5 (cHet; P1 70.8 ± 7.64, n = 4, P3 66.6 ± 17.0, n = 4, P5 17.6 ± 8.87, n = 3. cKO; P1 63.2 ± 7.31, n = 4, P3 65.1 ± 2.15, n = 3, P5 31.8 ± 5.03, n = 3). There was no significant difference between the cHet and cKO retinas. Error bars, mean ± SD. P1 $p = 0.20$, P3 $p = 0.87$, P5 $p = 0.074$. Statistical tests: P1, P5 Student's t-test; P3 Welch's t-test.

D. Quantification of the retinal thickness in the cHet and cKO mice at P1, P3, and P5 (cHet; P1 145.1 ± 16.6, n = 5, P3 190.2 ± 9.81, n = 4, P5 203.9 ± 26.6, n = 5. cKO; P1 139.2 ± 26.1, n = 4, P3 203.7 ± 21.6, n = 5, P5 200.0 ± 26.7, n = 4). There was no significant difference between the cHet and cKO retinas. Error bars, mean ± SD. P1 $p = 0.54$, P3 $p = 0.29$, P5 $p = 1.0$. Statistical tests: P1, P3 Student's t-test; P5 Mann–Whitney U test.

Video 1: Behavior of the *Afadin* cKO mouse in the visual cliff test.

Excerpt of the cKO mouse behavior during the visual cliff test.

Video 2: Behavior of the *Afadin* cHet mouse in the visual cliff test.

Excerpt of the cHet mouse behavior during the visual cliff test.

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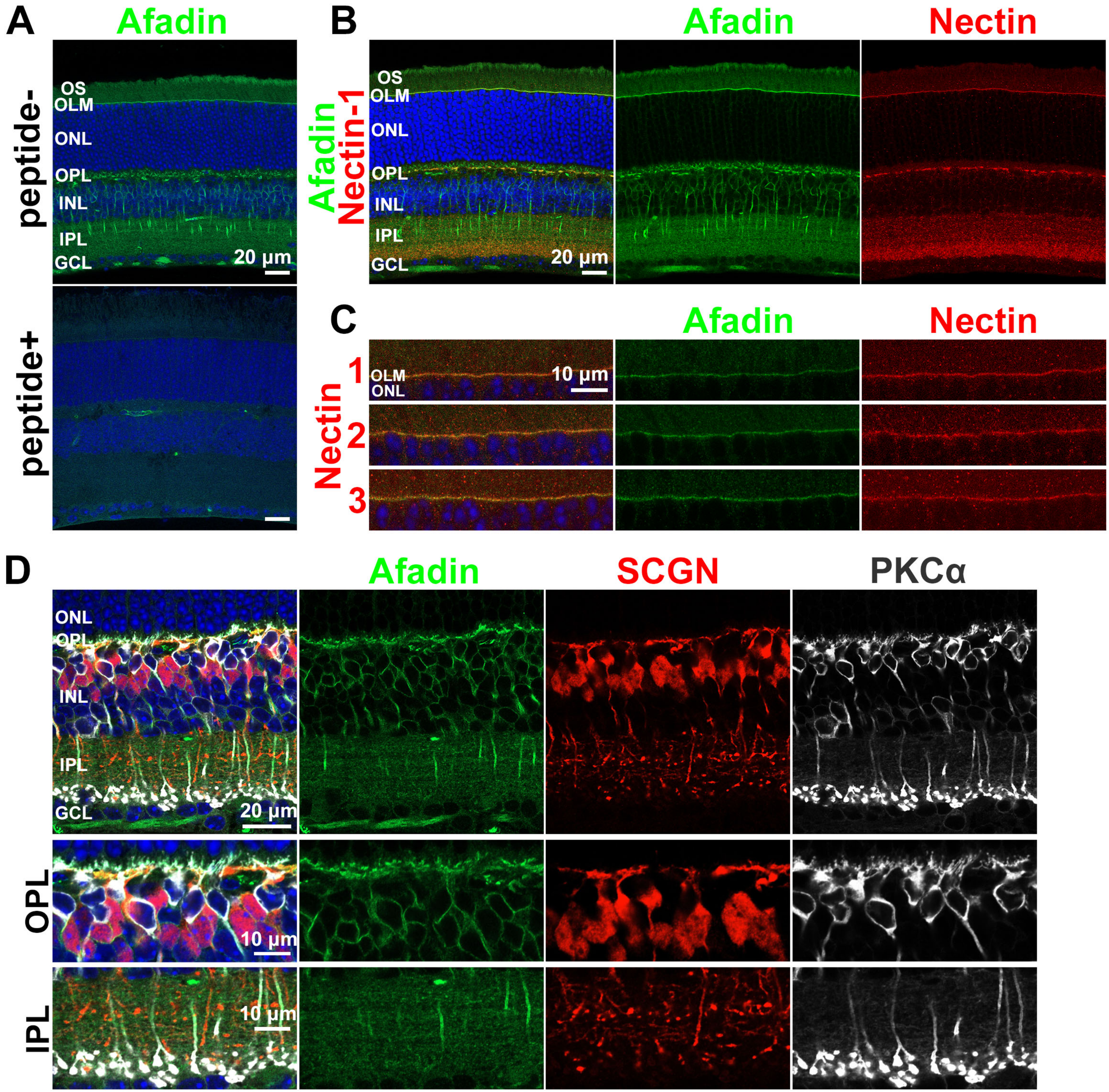
A
Afadin
peptide-
peptide+
OS
OLM
ONL
OPL
INL
IPL
GCL
20 µm
B
Afadin
Nectin
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OS
OLM
ONL
OPL
INL
IPL
GCL
20 µm
C
Afadin
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Nectin
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OLM
ONL
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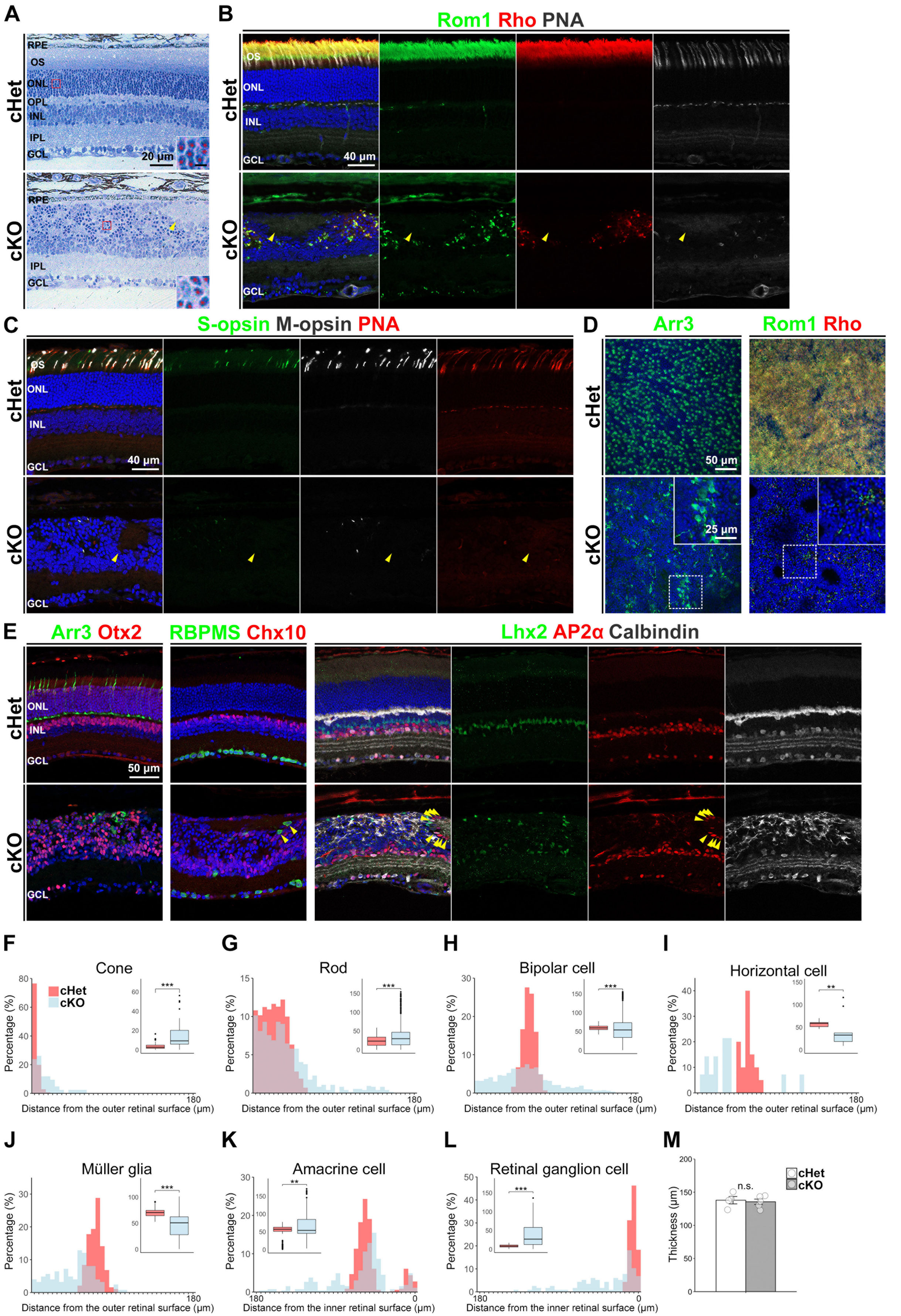
A
cHet
cKO
RPE
OS
ONL
OPL
INL
IPL
GCL
20 μm
B
Rom1 Rho PNA
40 μm
C
S-opsin M-opsin PNA
40 μm
D
Arr3
Rom1 Rho
50 μm
25 μm
E
Arr3 Otx2
RBPMS Chx10
Lhx2 AP2α Calbindin
50 μm
F
Cone
G
Rod
H
Bipolar cell
I
Horizontal cell
J
Müller glia
K
Amacrine cell
L
Retinal ganglion cell
M
cHet
cKO
Percentage (%)
Distance from the outer retinal surface (μm)
Distance from the inner retinal surface (μm)
Thickness (μm)
n.s.
***
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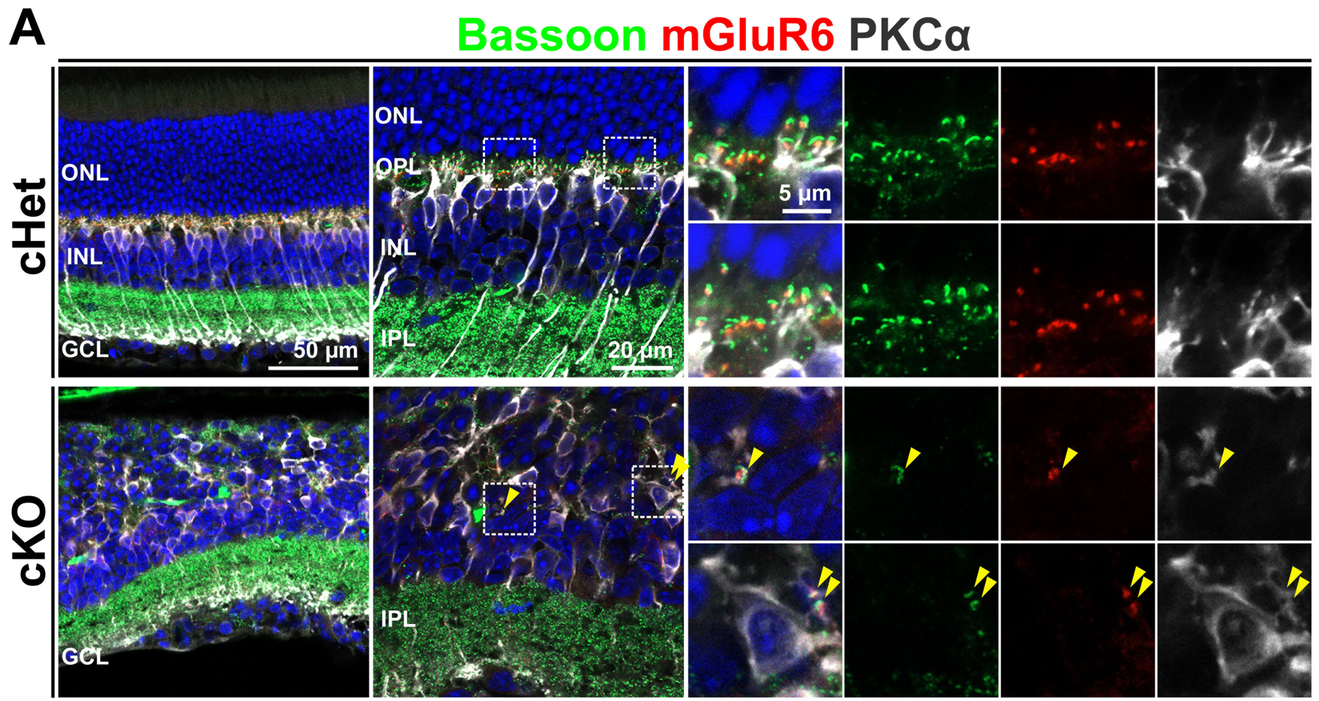
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Bassoon mGluR6 PKCα
cHet
cKO
ONL
INL
GCL
50 µm
OPL
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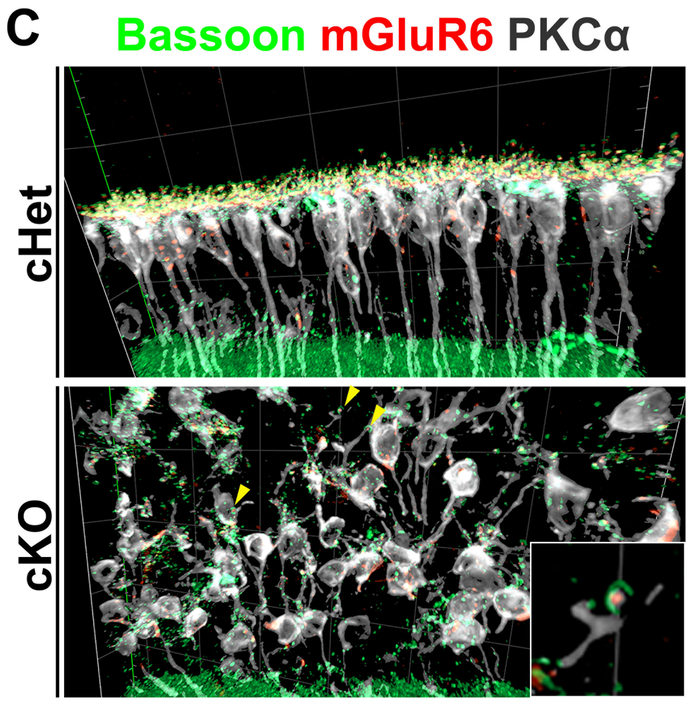
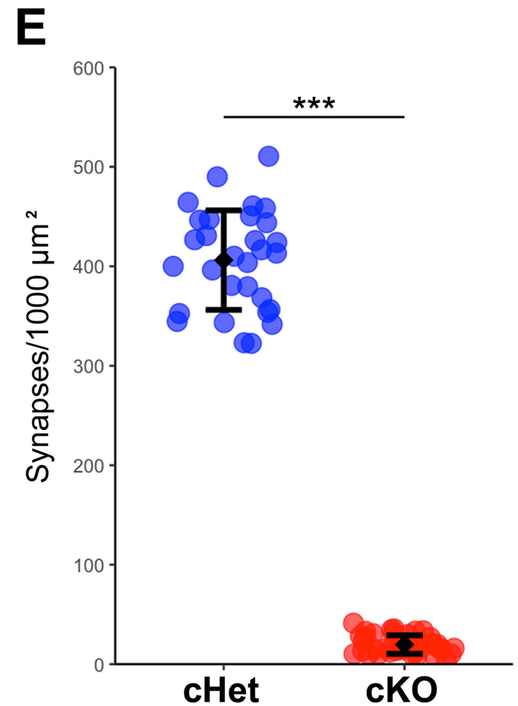
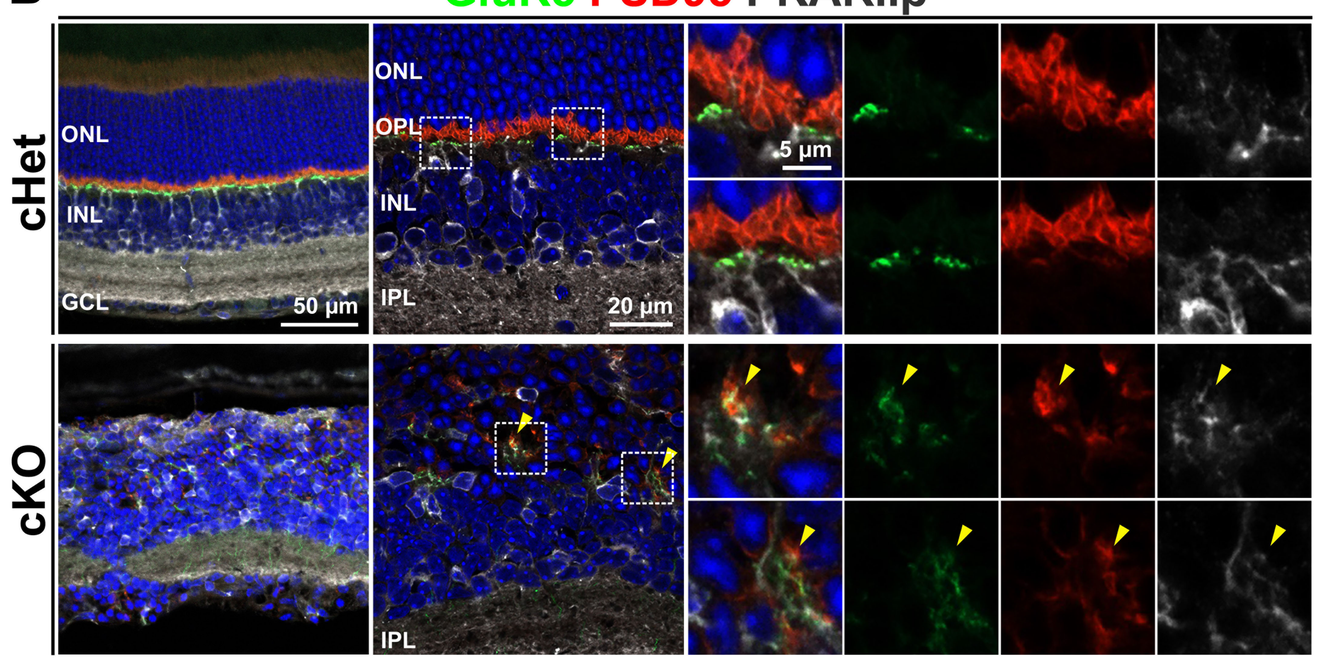
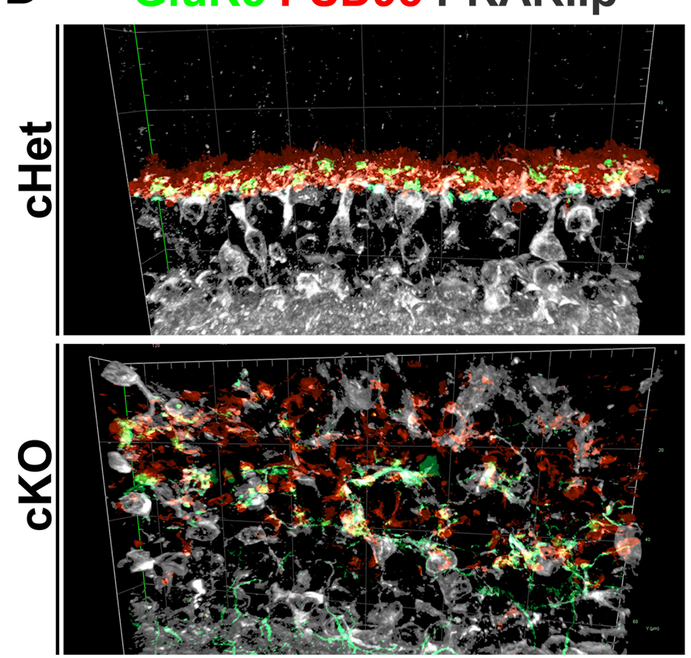
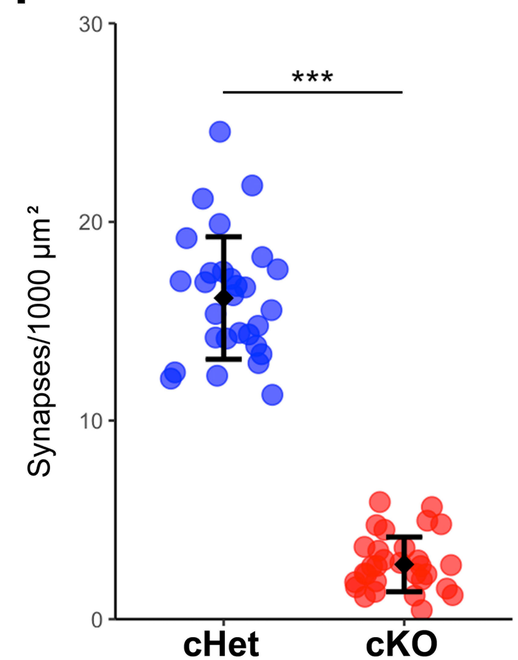
20 µm
5 µm

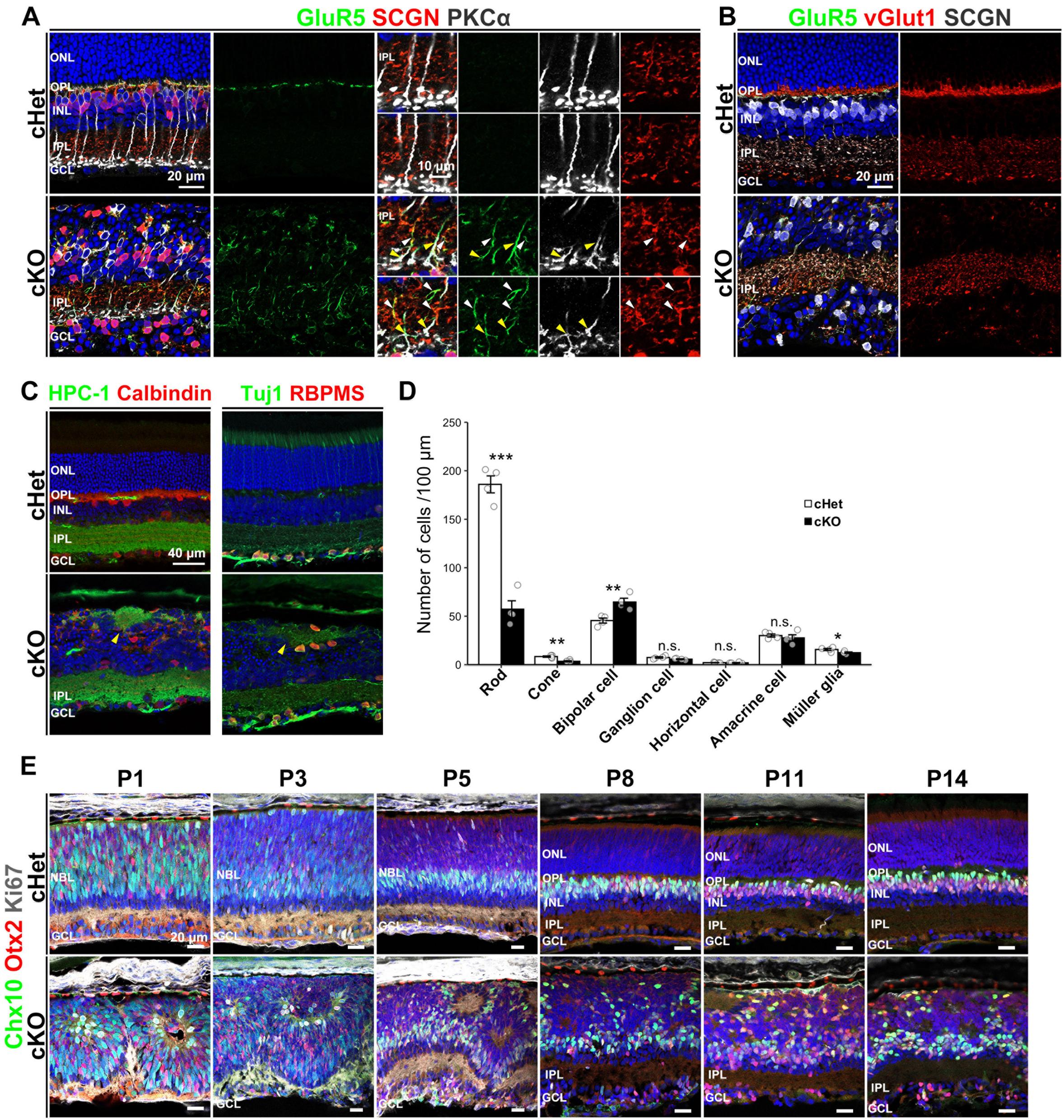

C
Bassoon mGluR6 PKCα
cHet
cKO


E
Synapses/1000 µm²
600
500
400
300
200
100
0
***
cHet
cKO


B
GluR5 PSD95 PKARIIβ
cHet
cKO
ONL
INL
GCL
50 µm
OPL
IPL
20 µm
5 µm


D
GluR5 PSD95 PKARIIβ
cHet
cKO


F
Synapses/1000 µm²
30
20
10
0
***
cHet
cKO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GluR5 SCGN PKCα
cHet
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ONL
OPL
INL
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GCL
20 μm
10 μm
B
GluR5 vGlut1 SCGN
C
HPC-1 Calbindin
Tuj1 RBPMS
40 μm
D
Number of cells /100 μm
250
200
150
100
50
0
***
**
n.s.
*
cHet
cKO
Rod
Cone
Bipolar cell
Ganglion cell
Horizontal cell
Amacrine cell
Müller glia
E
P1
P3
P5
P8
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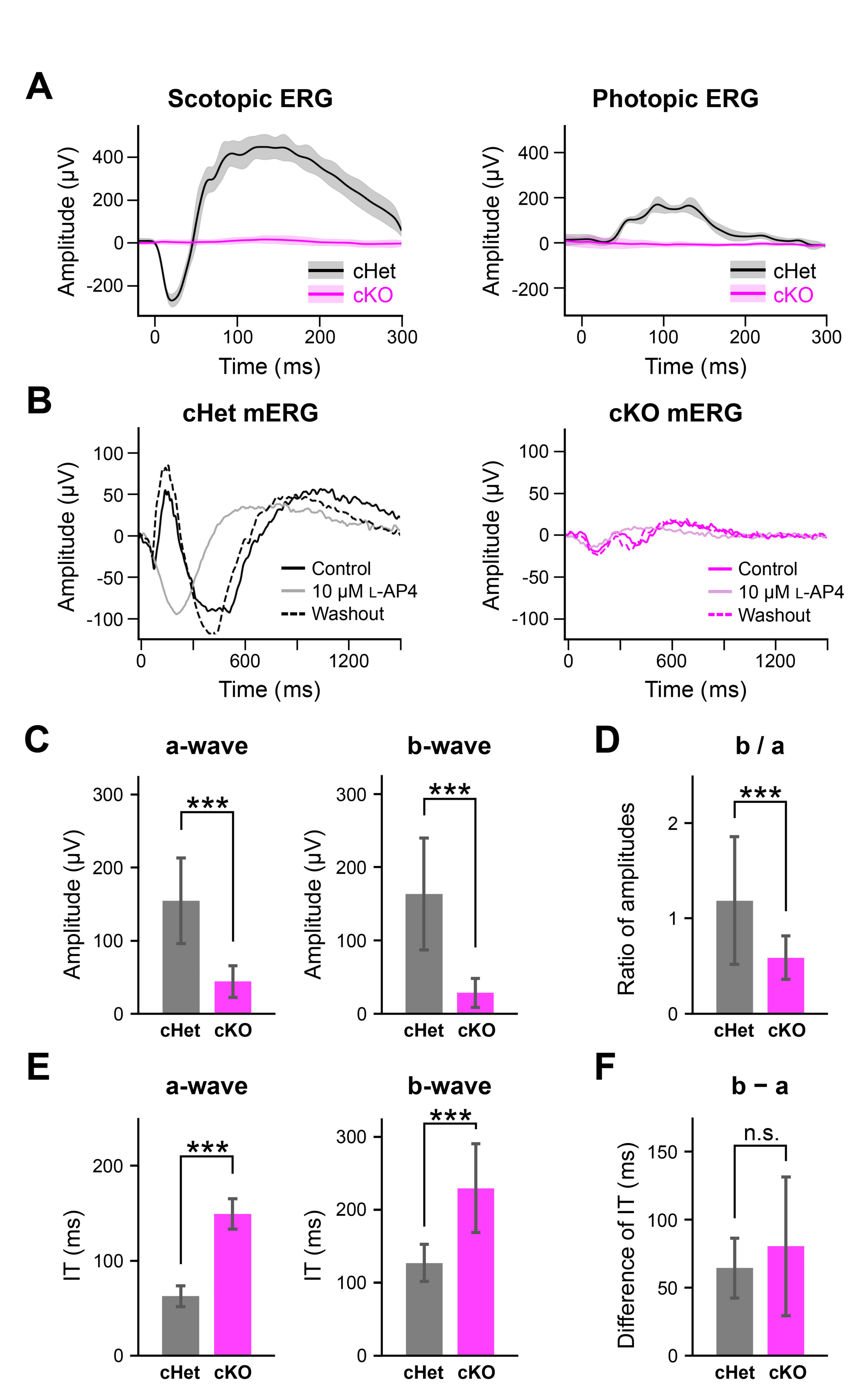
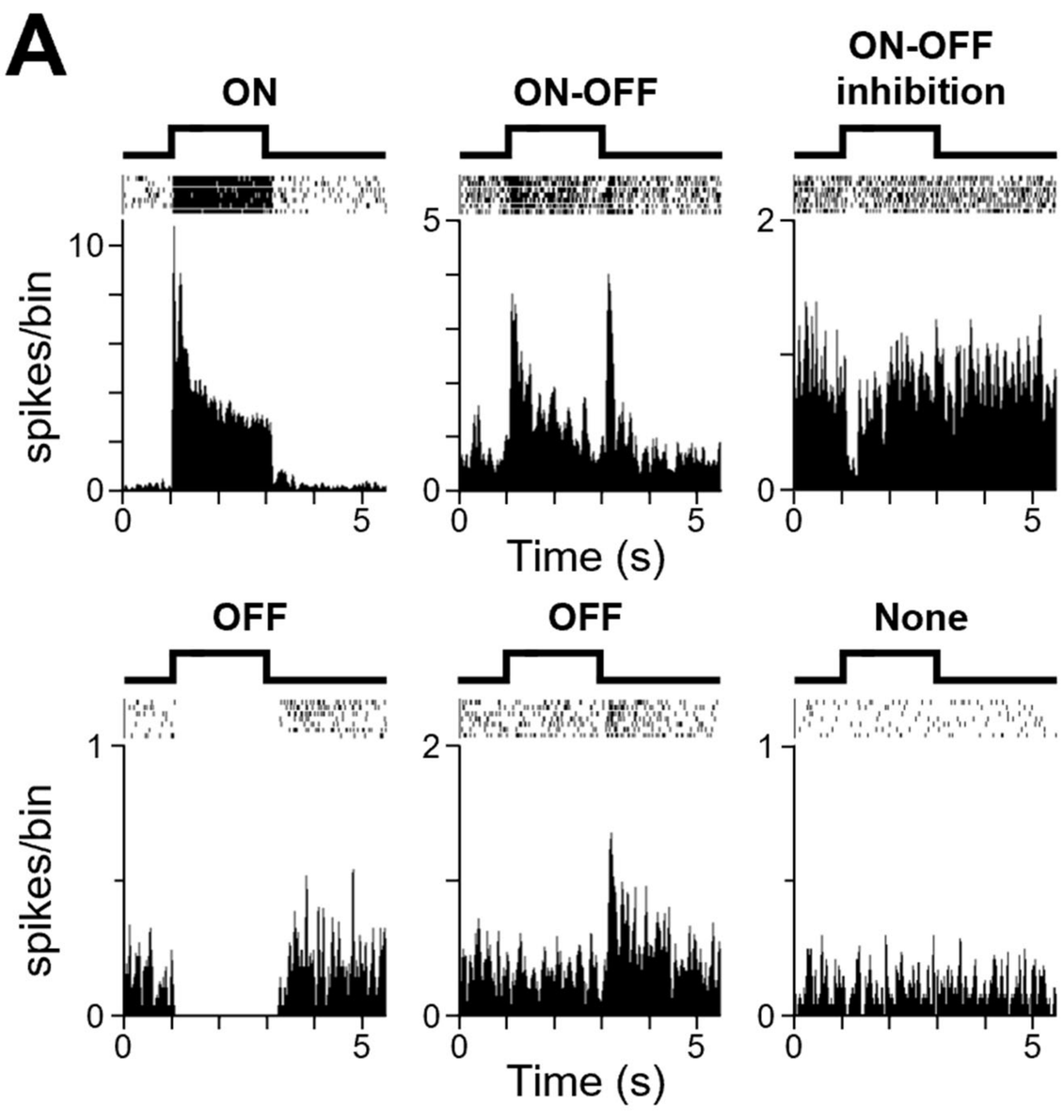
Chx10 Otx2 Ki67
NBL


A
Scotopic ERG
Photopic ERG
Amplitude (μV)
400
200
0
-200
0
100
200
300
Time (ms)
cHet
cKO
B
cHet mERG
cKO mERG
100
50
0
-50
-100
600
1200
Control
10 μM L-AP4
Washout
C
a-wave
b-wave
300
200
100
0
***
cHet
cKO
D
b / a
Ratio of amplitudes
2
1
E
IT (ms)
F
b − a
Difference of IT (ms)
150
50
n.s.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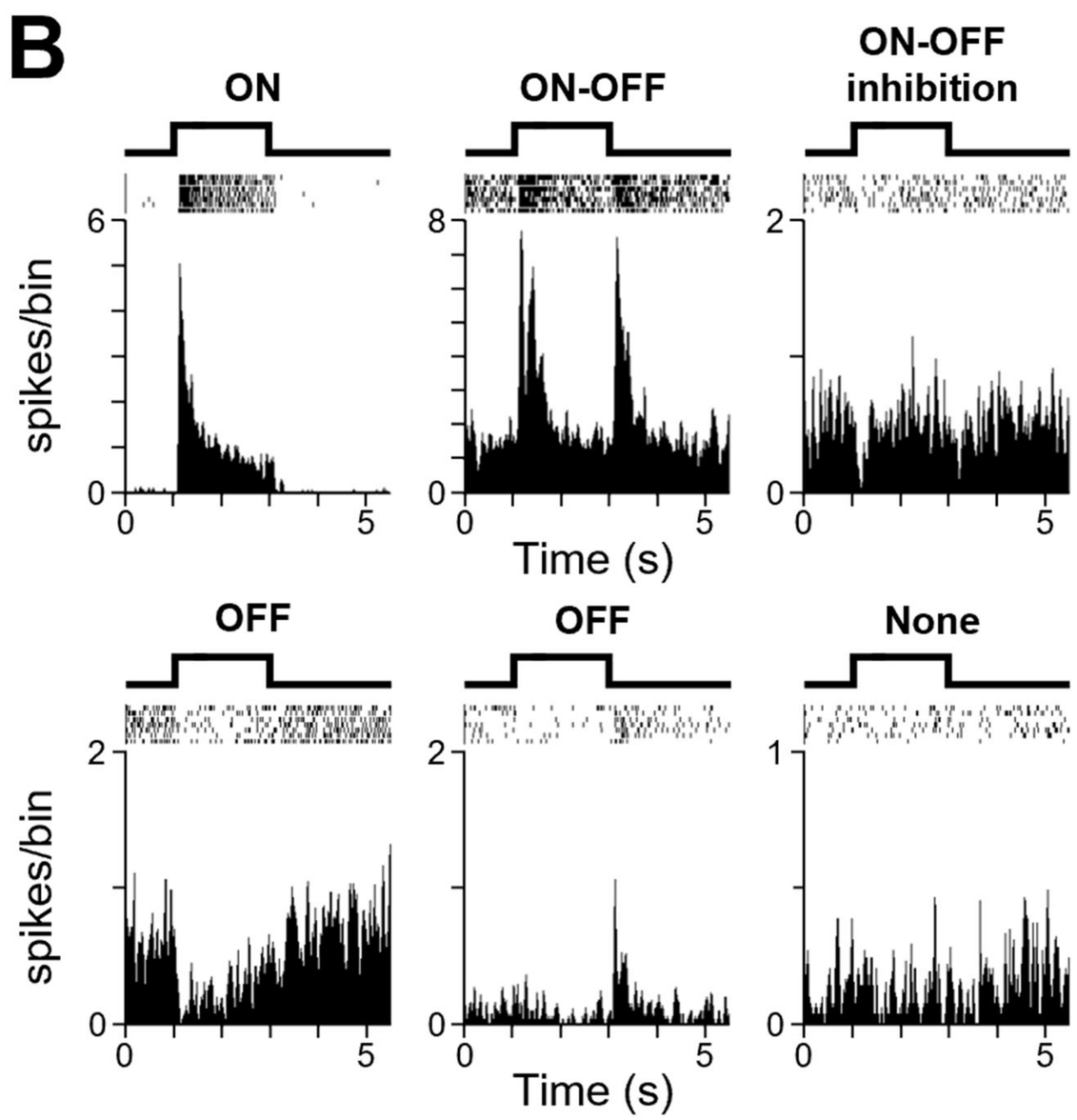
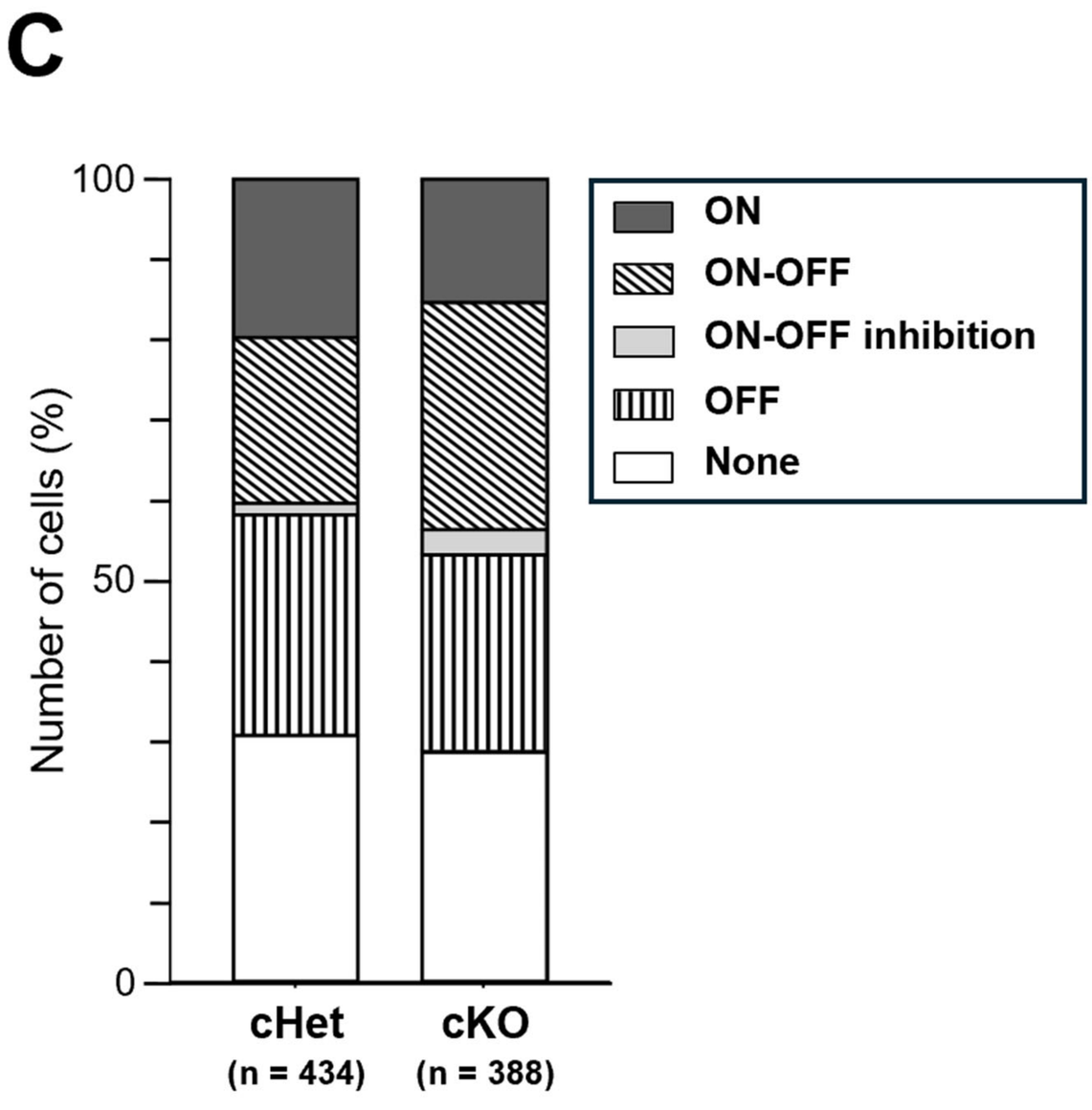
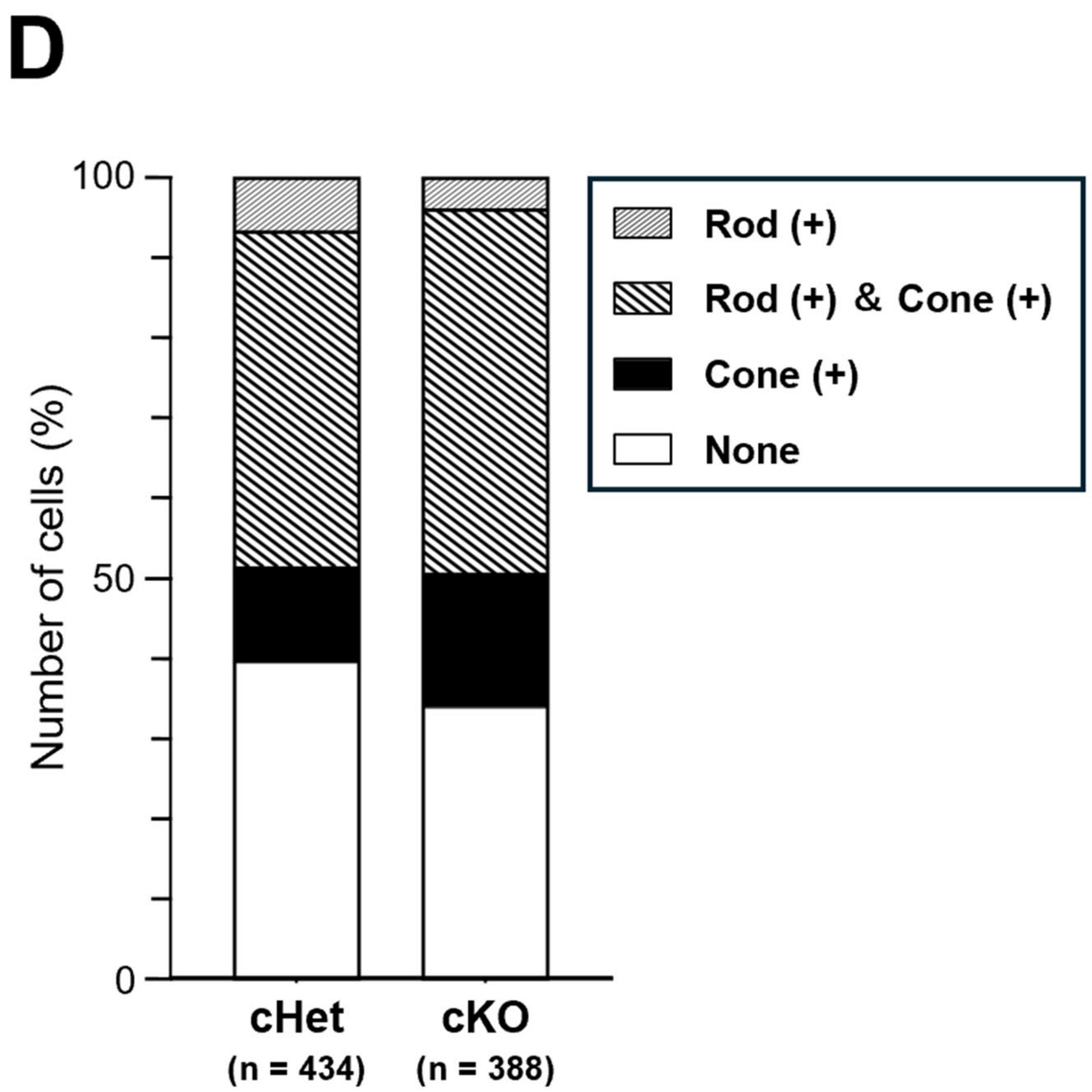
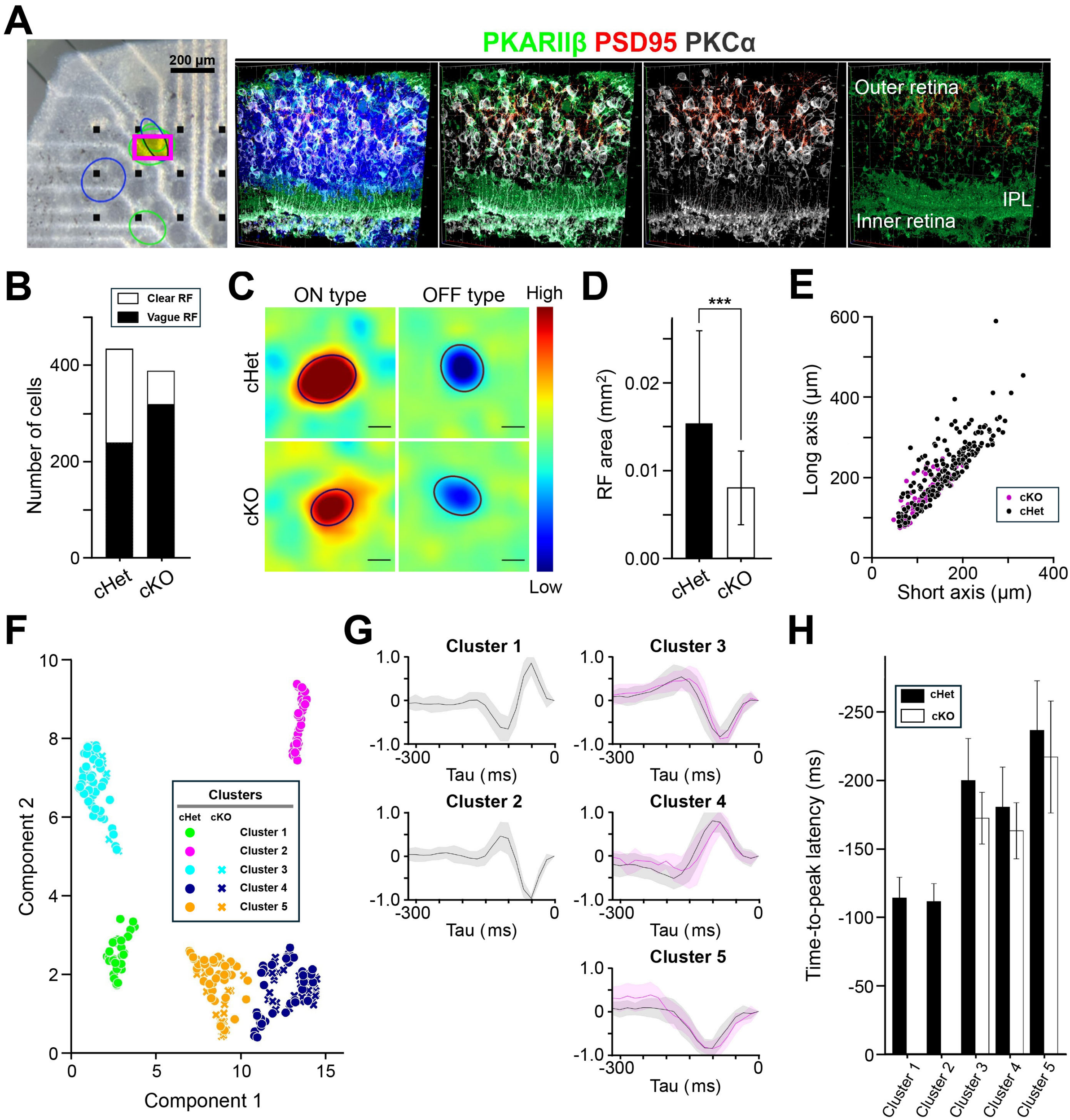
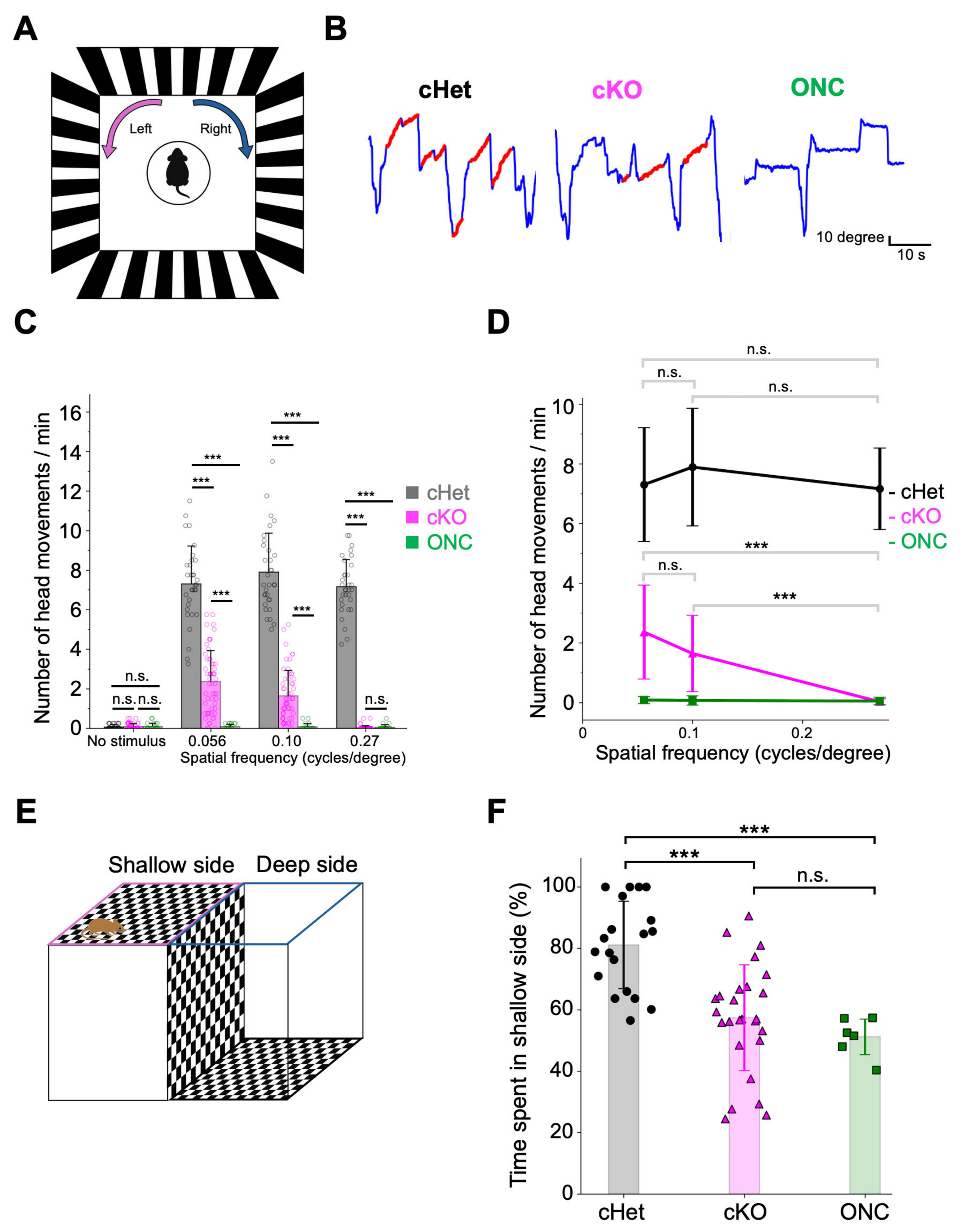
ON
ON-OFF
ON-OFF inhibition
OFF
OFF
None
spikes/bin
Time (s)

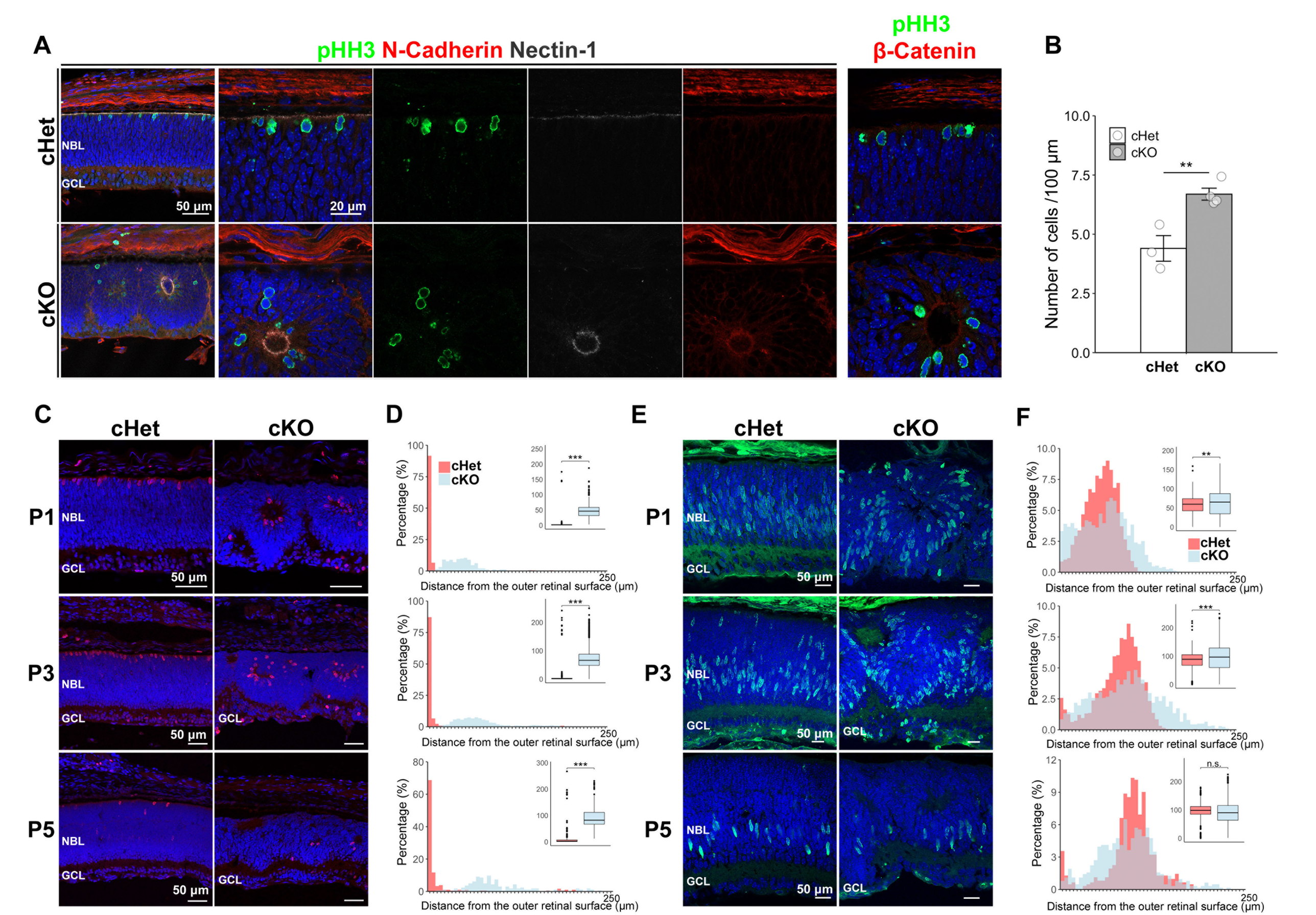

B
ON
ON-OFF
ON-OFF inhibition
OFF
OFF
None
spikes/bin
Time (s)


C
Number of cells (%)
ON
ON-OFF
ON-OFF inhibition
OFF
None
cHet
(n = 434)
cKO
(n = 388)

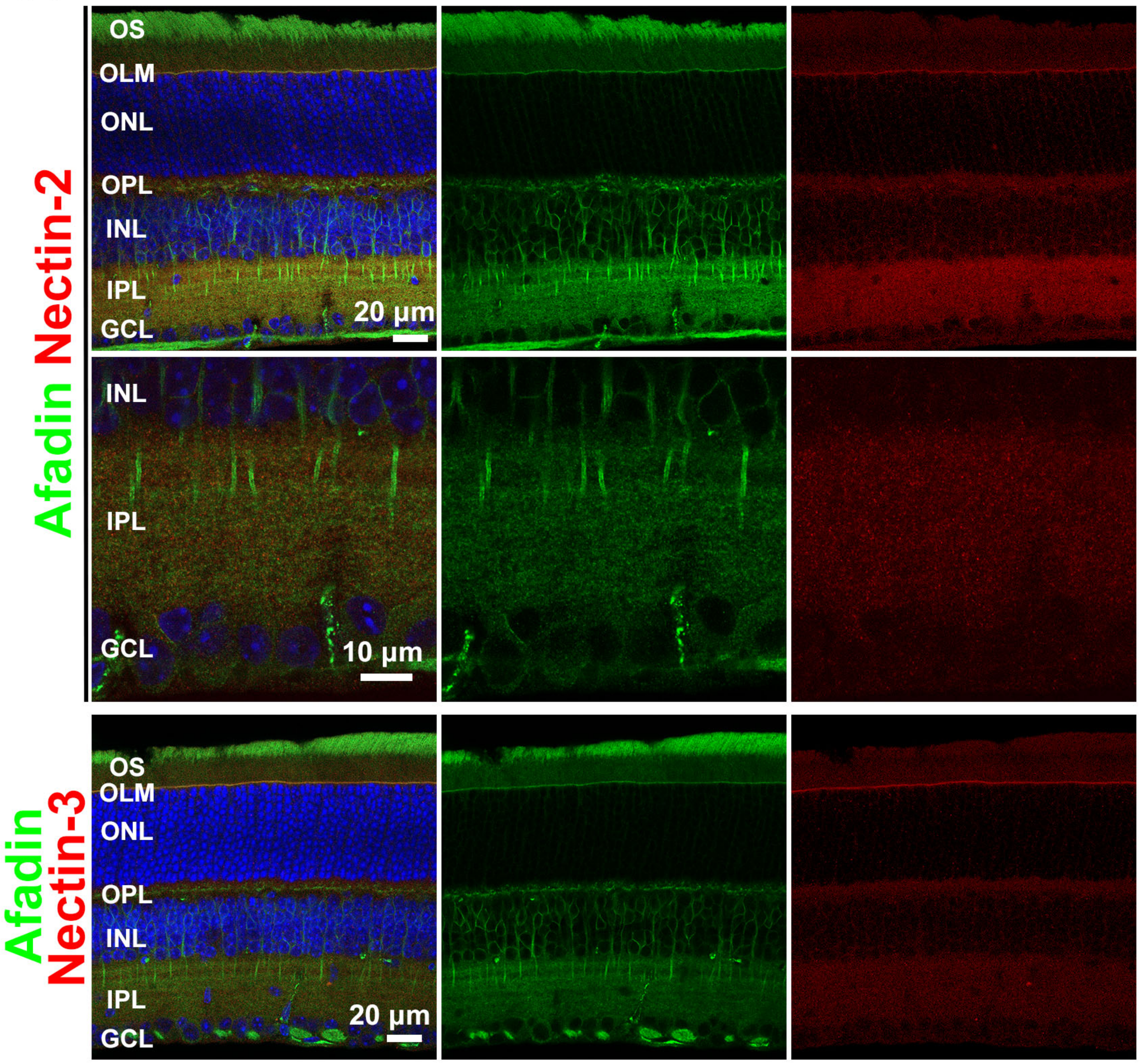

D
Number of cells (%)
Rod (+)
Rod (+) & Cone (+)
Cone (+)
None
cHet
(n = 434)
cKO
(n = 388)


A
200 µm
PKARIIβ PSD95 PKCα
Outer retina
IPL
Inner retina
B
Clear RF
Vague RF
Number of cells
400
200
0
cHet
cKO
C
ON type
OFF type
High
Low
cHet
cKO
D
***
RF area (mm²)
0.02
0.01
0.00
cHet
cKO
E
Long axis (µm)
600
400
200
0
Short axis (µm)
cKO
cHet
F
Component 2
Component 1
Clusters
cHet cKO
Cluster 1
Cluster 2
Cluster 3
Cluster 4
Cluster 5
G
Cluster 1
Cluster 2
Cluster 3
Cluster 4
Cluster 5
Tau (ms)
H
cHet
cKO
Time-to-peak latency (ms)
Cluster 1
Cluster 2
Cluster 3
Cluster 4
Cluster 5

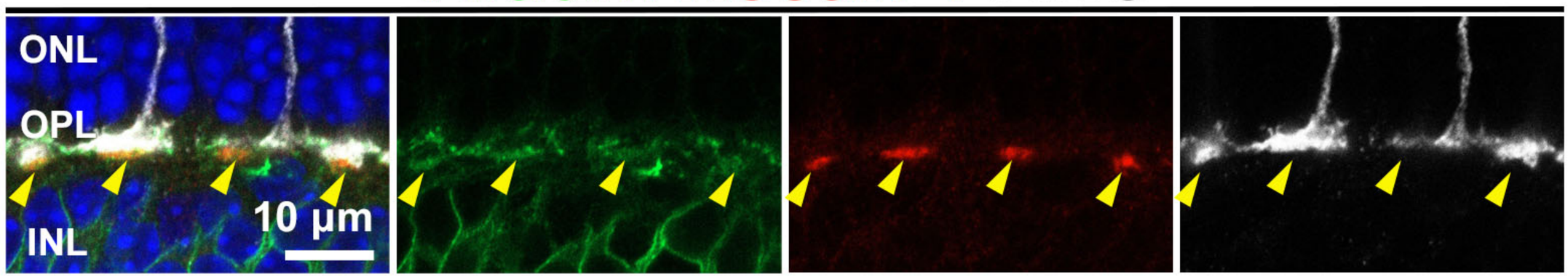

A
Left
Right
B
cHet
cKO
ONC
10 degree
10 s
C
Number of head movements / min
cHet
cKO
ONC
n.s.
***
No stimulus
0.056
0.10
0.27
Spatial frequency (cycles/degree)
D
n.s.
***
Number of head movements / min
- cHet
- cKO
- ONC
Spatial frequency (cycles/degree)
E
Shallow side
Deep side
F
Time spent in shallow side (%)
***
n.s.
cHet
cKO
ONC


A
pHH3 N-Cadherin Nectin-1
pHH3
β-Catenin
cHet
cKO
NBL
GCL
50 μm
20 μm
B
Number of cells /100 μm
cHet
cKO
**
C
cHet
cKO
P1
P3
P5
D
Percentage (%)
Distance from the outer retinal surface (μm)
***
E
F
n.s.

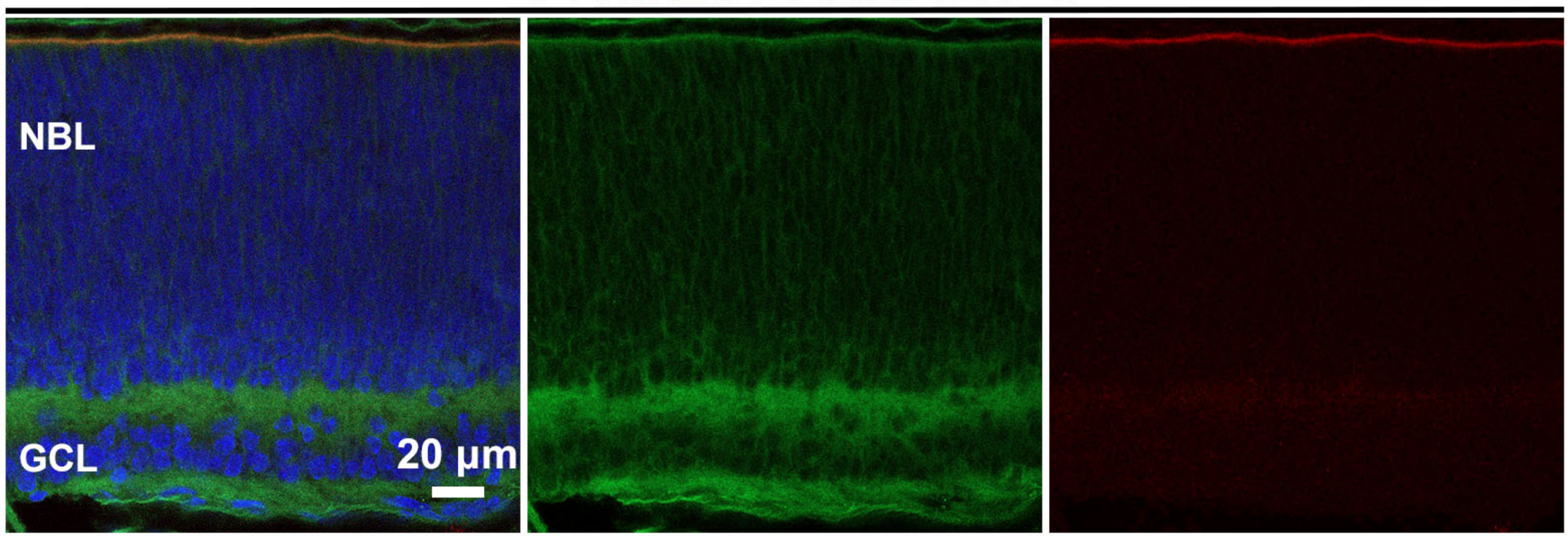
A

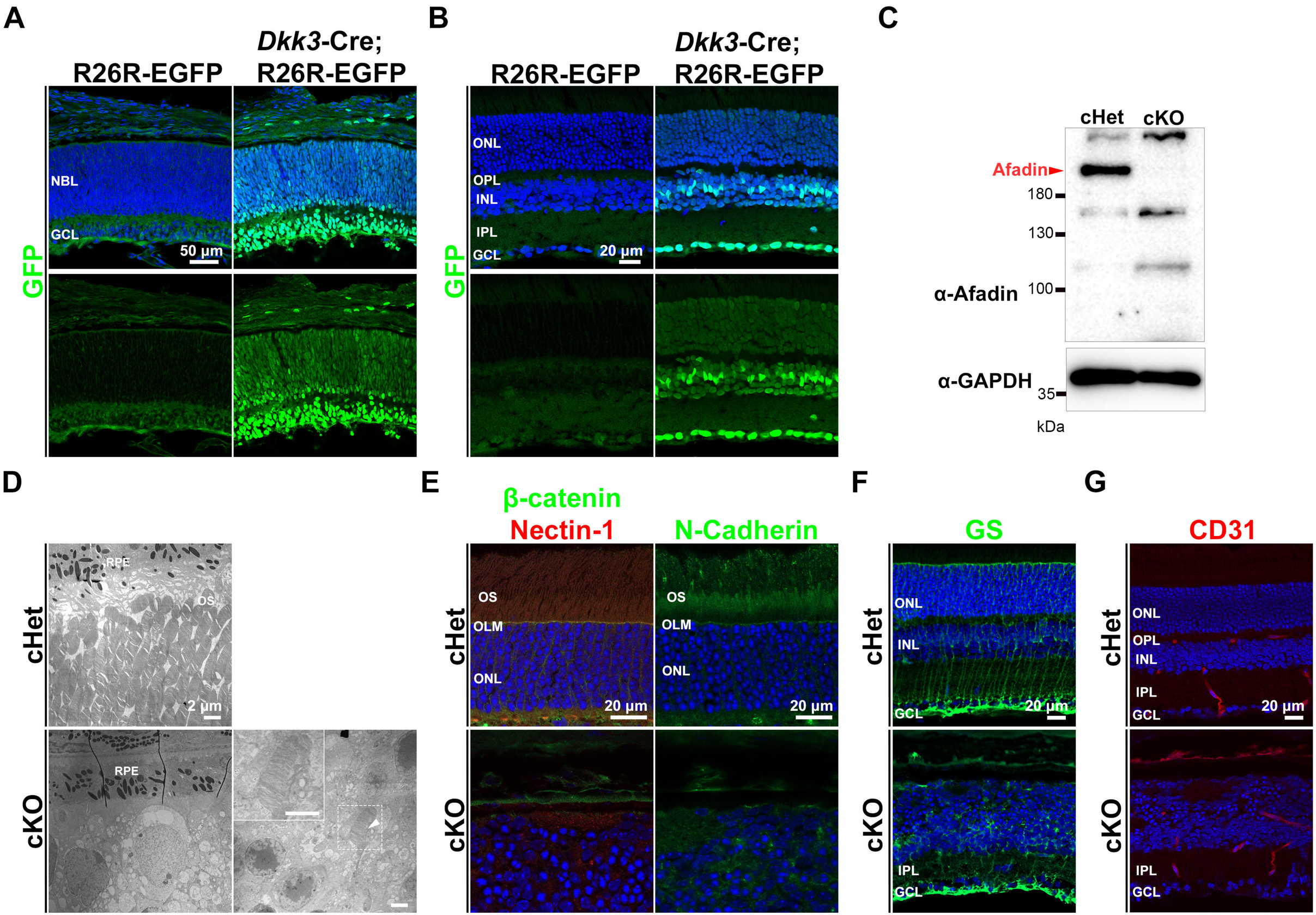

Afadin Nectin-2
OS
OLM
ONL
OPL
INL
IPL
GCL
20 µm
INL
IPL
GCL
10 µm
Afadin Nectin-3
OS
OLM
ONL
OPL
INL
IPL
GCL
20 µm

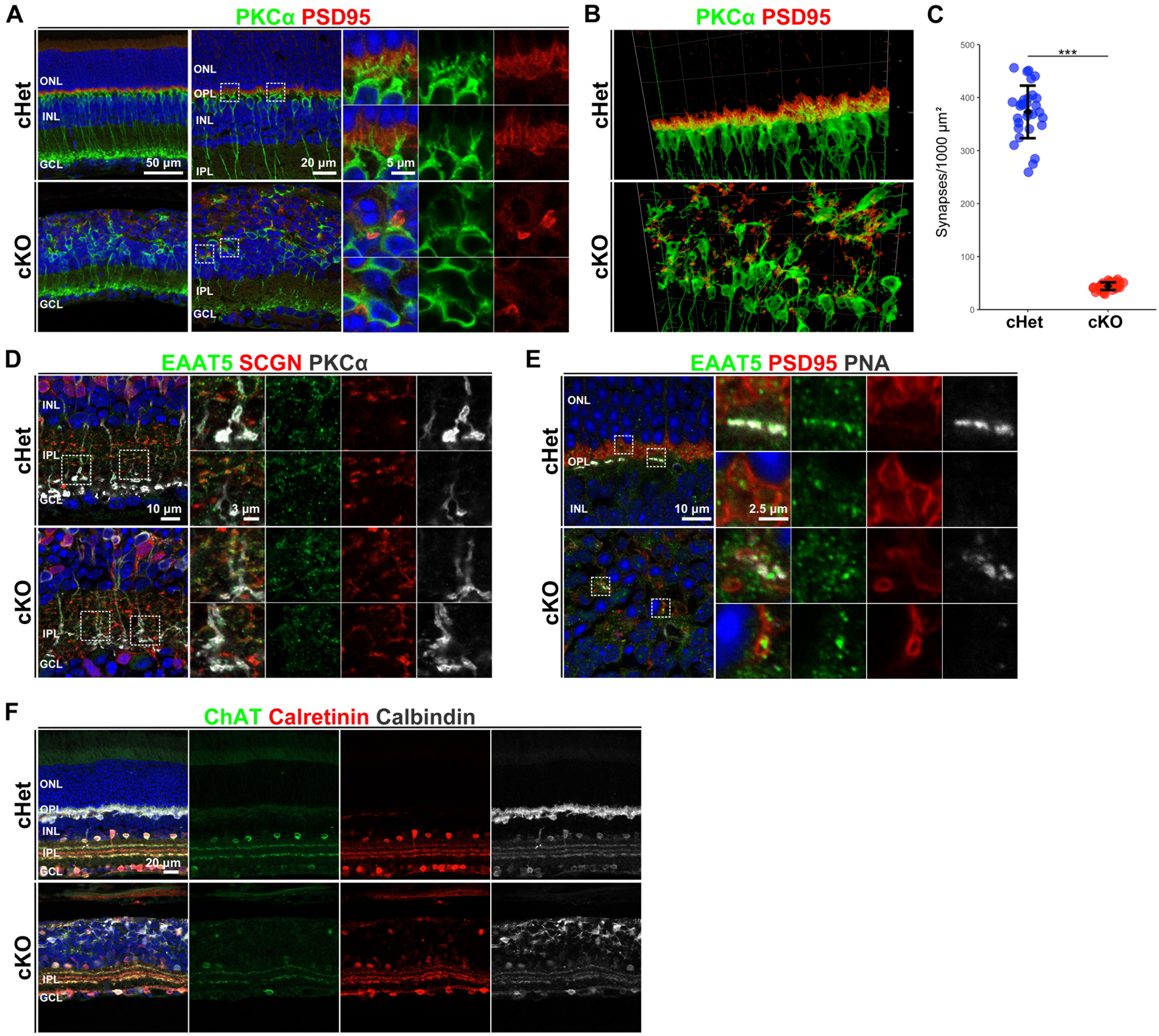
B

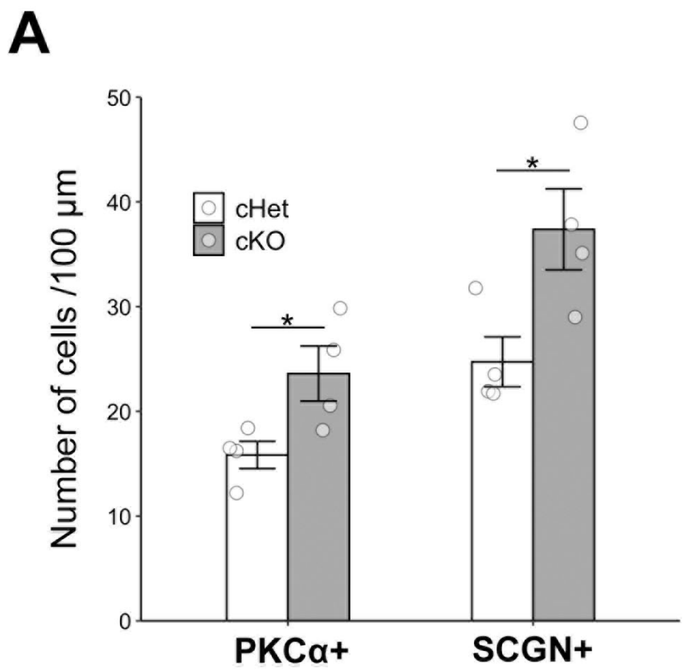

Afadin Nectin-1 Arr3
ONL
OPL
INL
10 µm

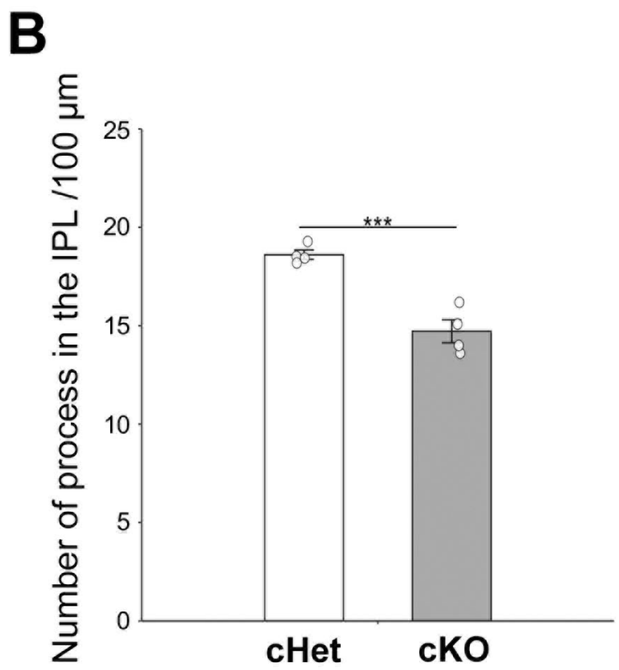
C

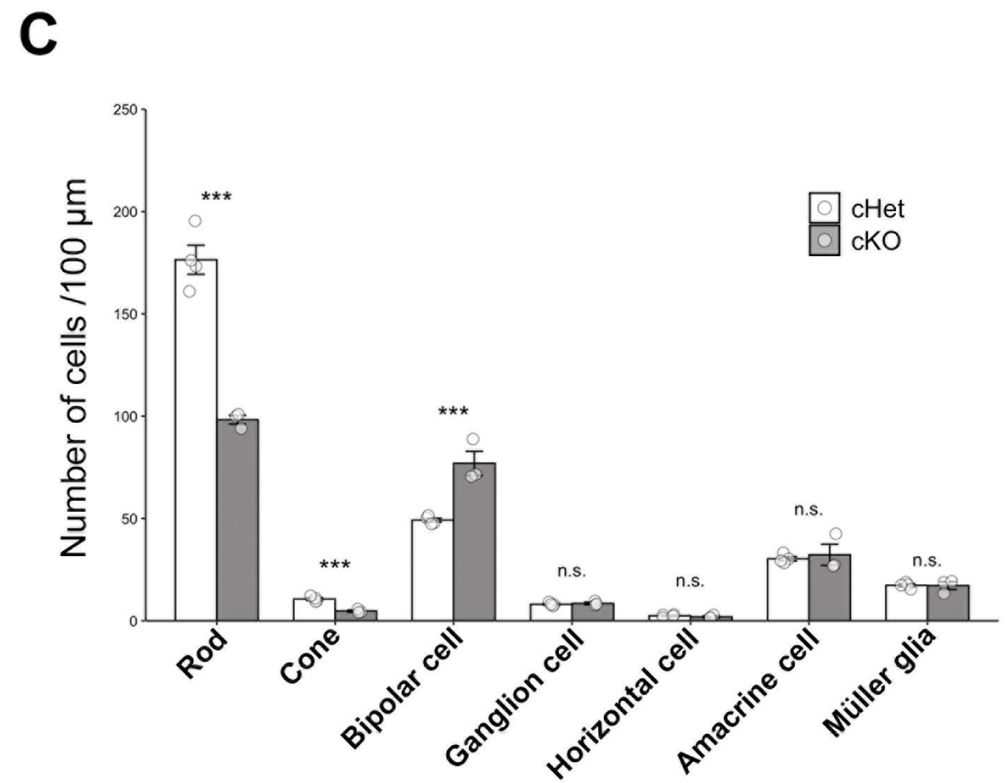

Afadin Nectin-1
NBL
GCL
20 µm

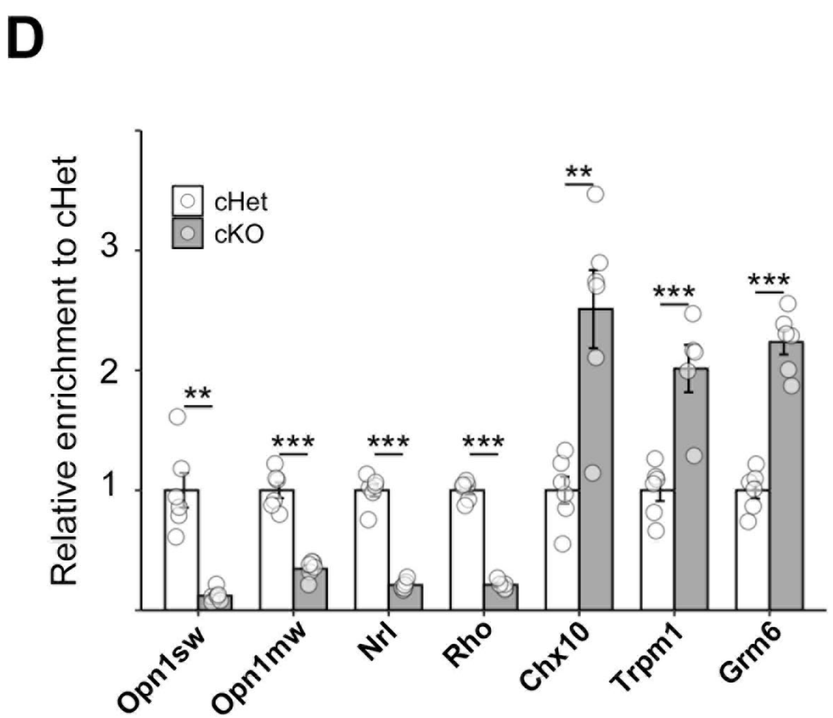

A
R26R-EGFP
Dkk3-Cre; R26R-EGFP
GFP
NBL
GCL
50 μm
B
R26R-EGFP
Dkk3-Cre; R26R-EGFP
GFP
ONL
OPL
INL
IPL
GCL
20 μm
C
cHet
cKO
Afadin
180
130
100
α-Afadin
α-GAPDH
35
kDa
D
cHet
cKO
RPE
OS
2 μm
RPE
E
β-catenin
Nectin-1
N-Cadherin
cHet
cKO
OS
OLM
ONL
20 μm
F
GS
cHet
cKO
ONL
INL
GCL
IPL
20 μm
G
CD31
cHet
cKO
ONL
OPL
INL
IPL
GCL
20 μm

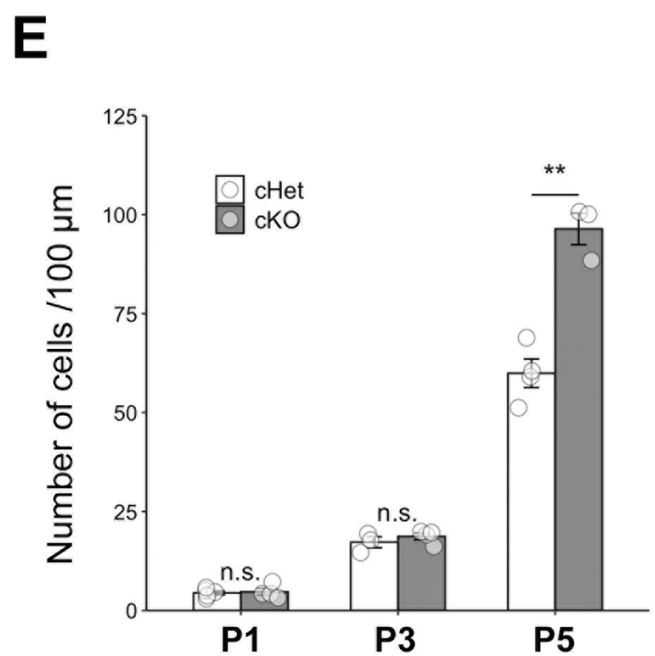

A
PKCα PSD95
cHet
cKO
ONL
INL
GCL
OPL
IPL
50 µm
20 µm
5 µm
B
PKCα PSD95
C
Synapses/1000 µm²
***
cHet
cKO
D
EAAT5 SCGN PKCα
10 µm
3 µm
E
EAAT5 PSD95 PNA
2.5 µm
F
ChAT Calretinin Calbindin
20 µm

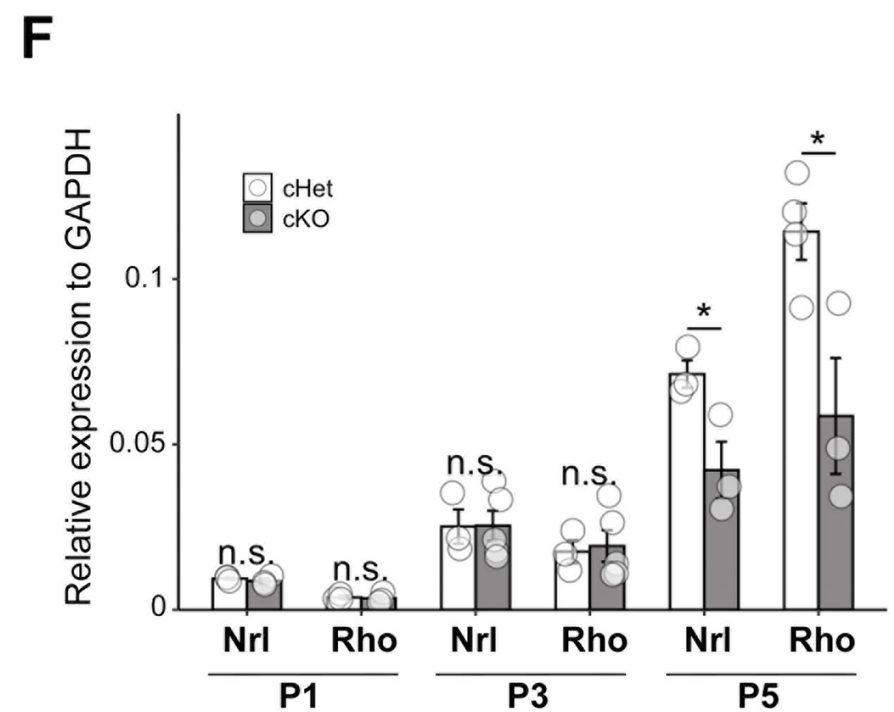

A
Number of cells /100 µm
cHet
cKO
PKCα+
SCGN+

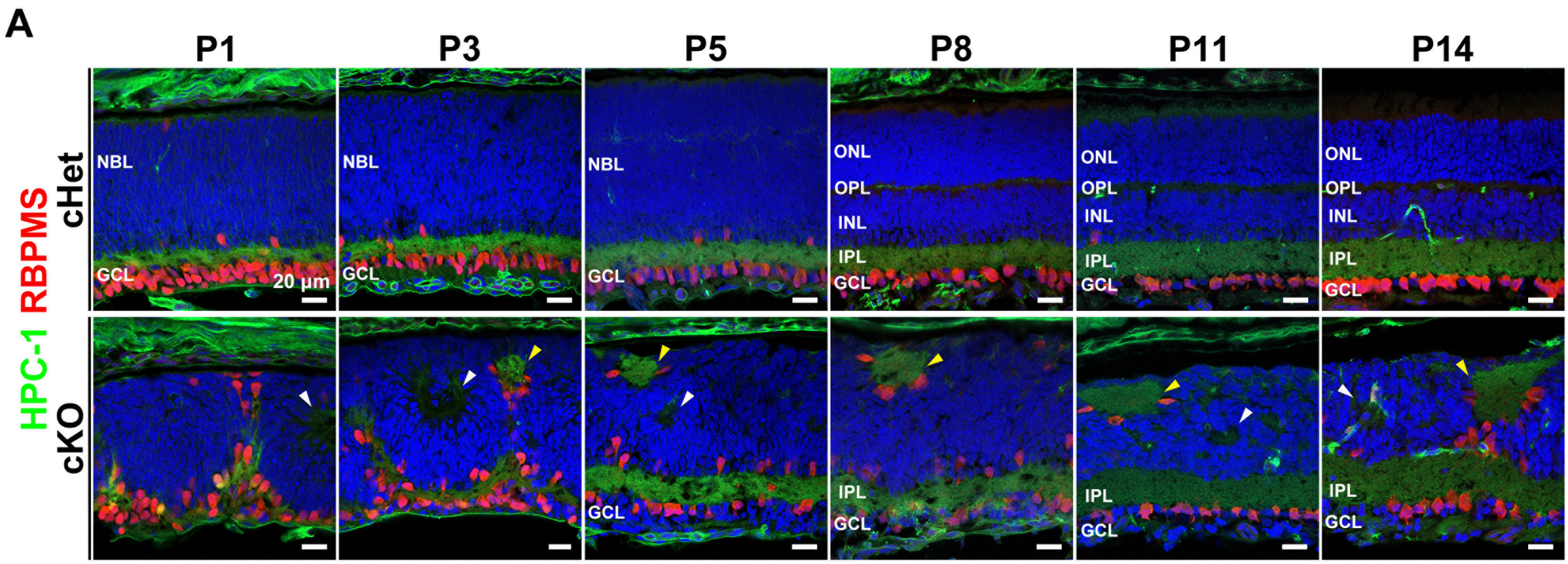

B
Number of process in the IPL /100 µm
cHet
cKO

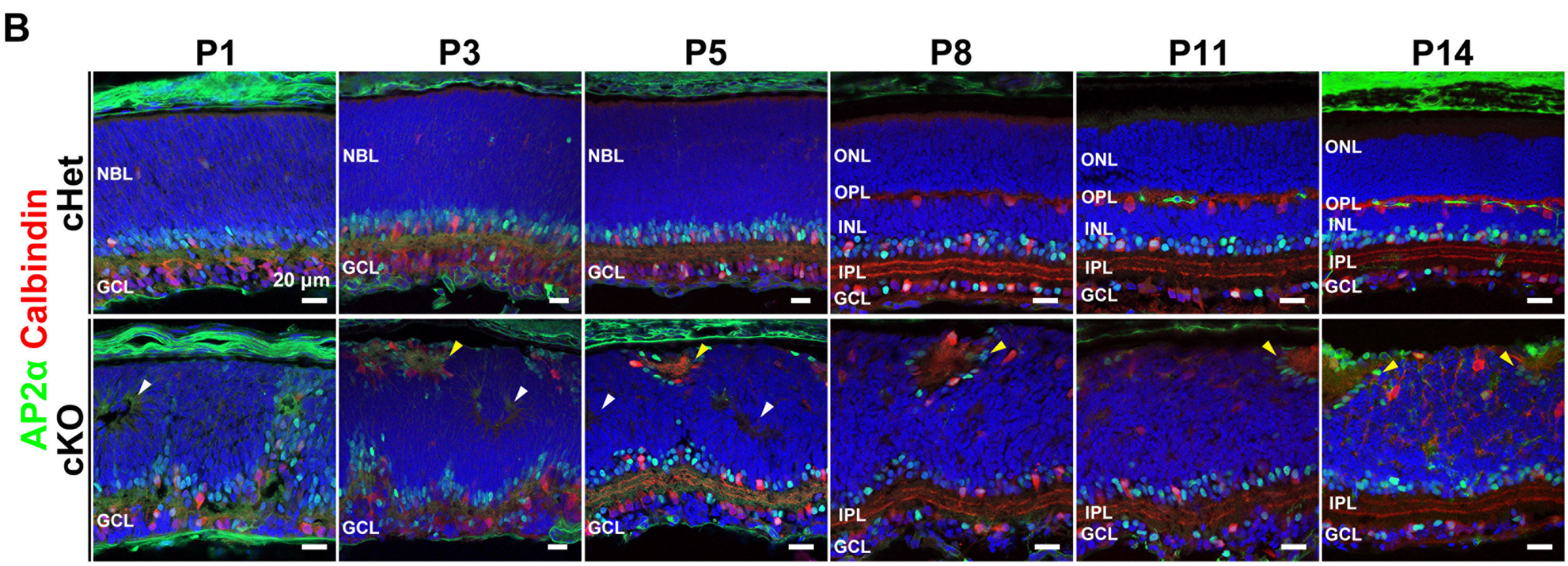

C
Number of cells /100 µm
cHet
cKO
Rod
Cone
Bipolar cell
Ganglion cell
Horizontal cell
Amacrine cell
Müller glia

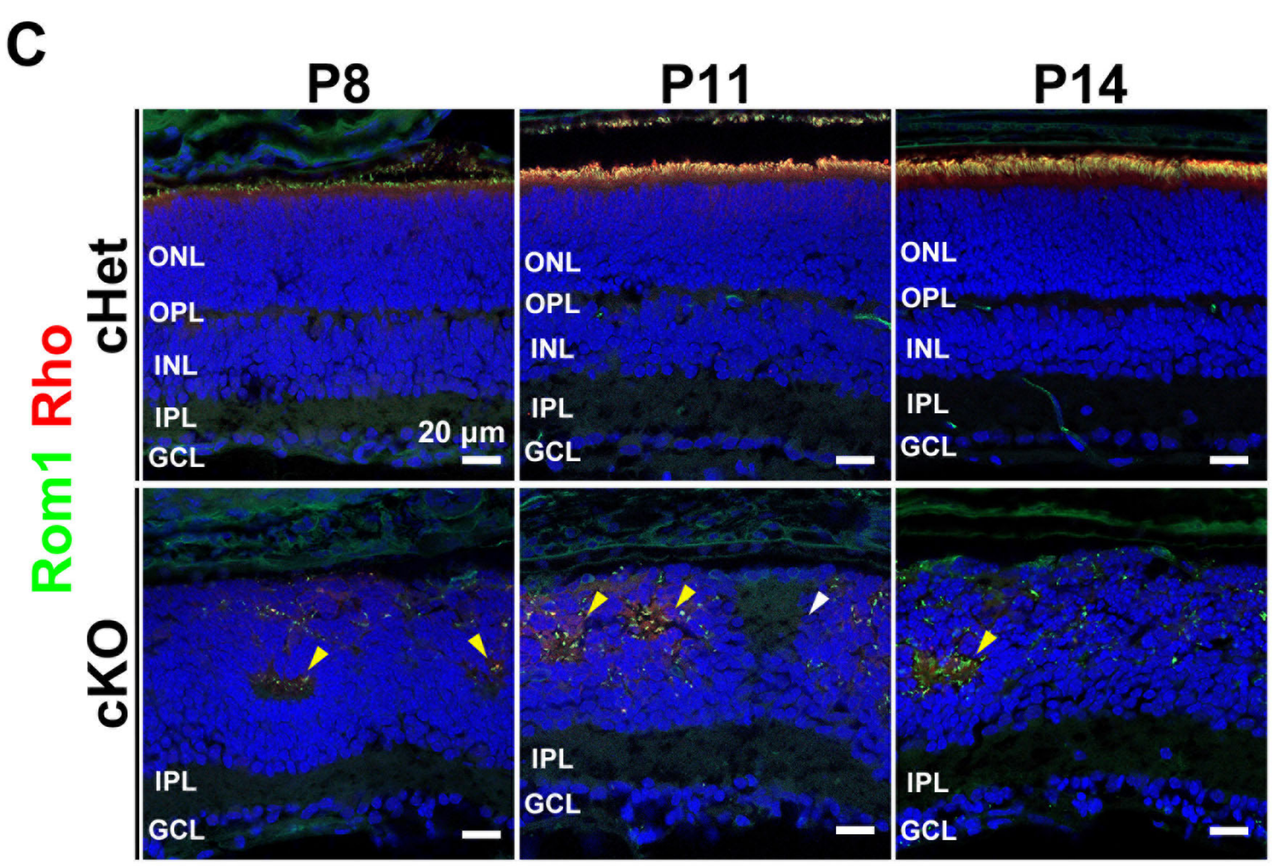

D
Relative enrichment to cHet
cHet
cKO
Opn1sw
Opn1mw
Nrl
Rho
Chx10
Trpm1
Grm6

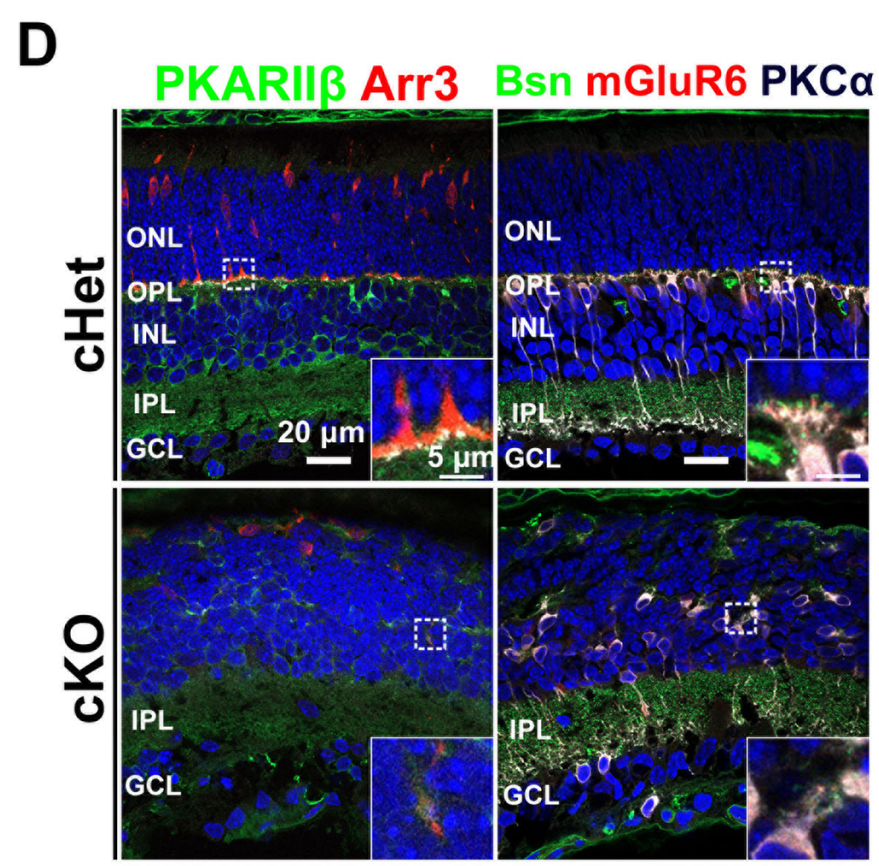

E
Number of cells /100 µm
cHet
cKO
P1
P3
P5

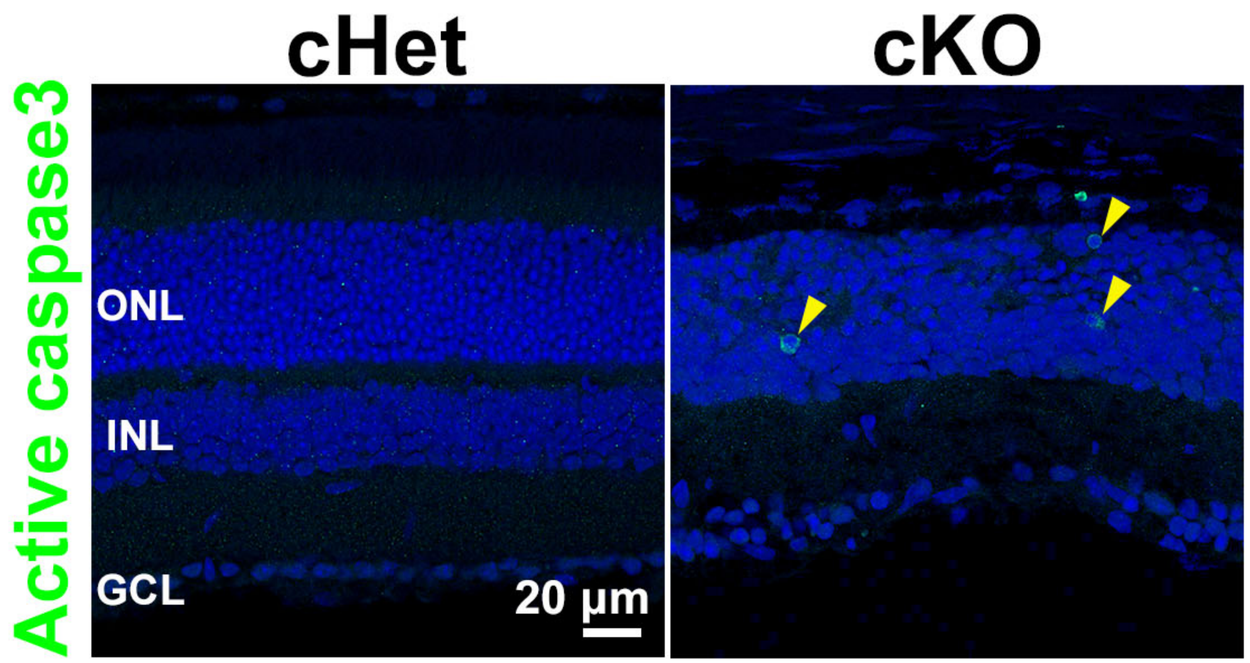

F
Relative expression to GAPDH
cHet
cKO
Nrl
Rho
P1
P3
P5

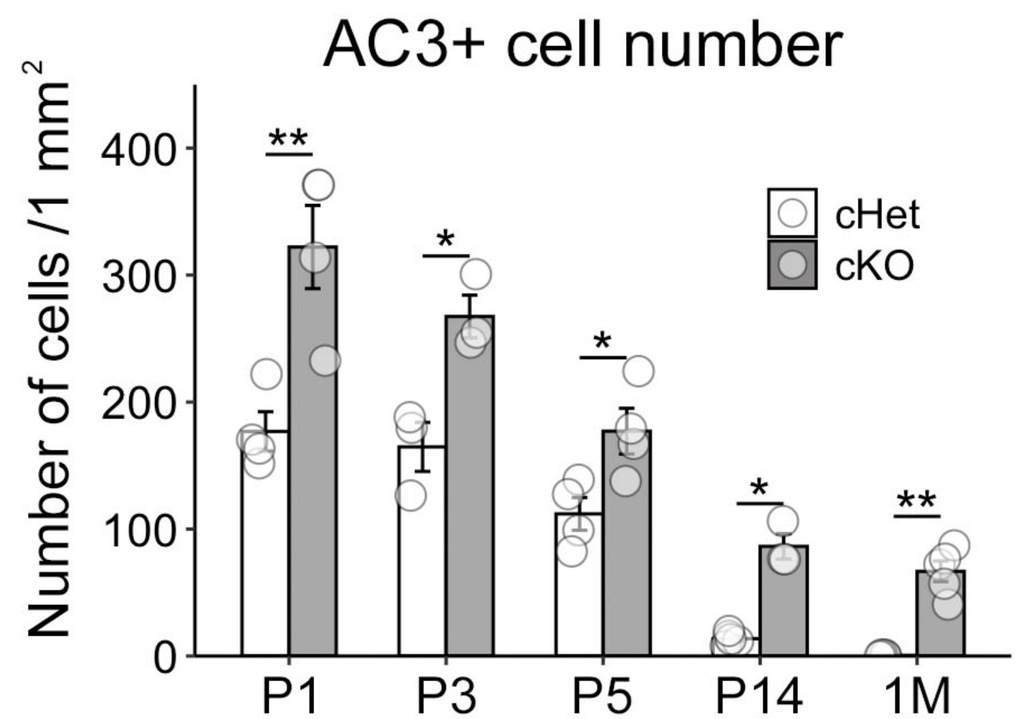

A
P1
P3
P5
P8
P11
P14
HPC-1 RBPMS
cHet
cKO
NBL
GCL
ONL
OPL
INL
IPL
20 µm

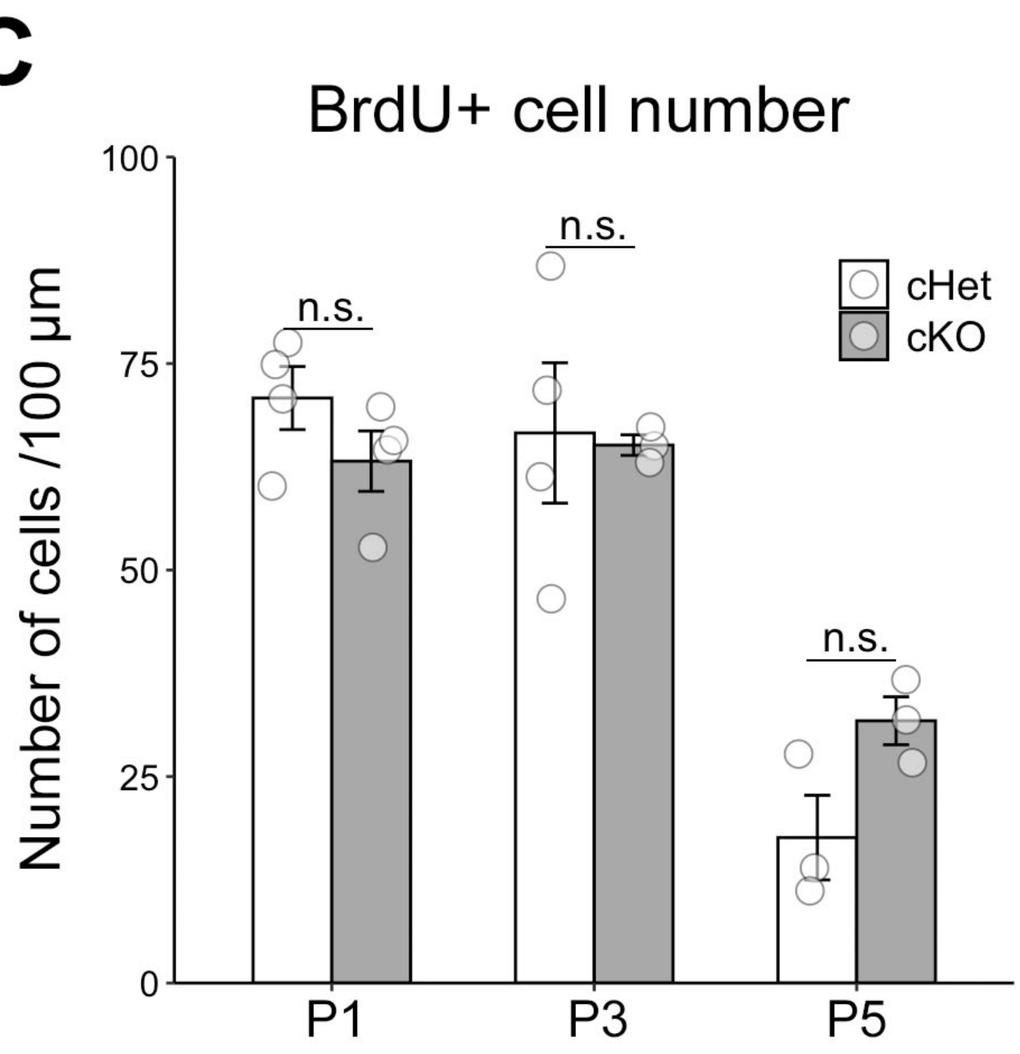

B
P1
P3
P5
P8
P11
P14
AP2α Calbindin
cHet
cKO
NBL
GCL
ONL
OPL
INL
IPL
20 µm

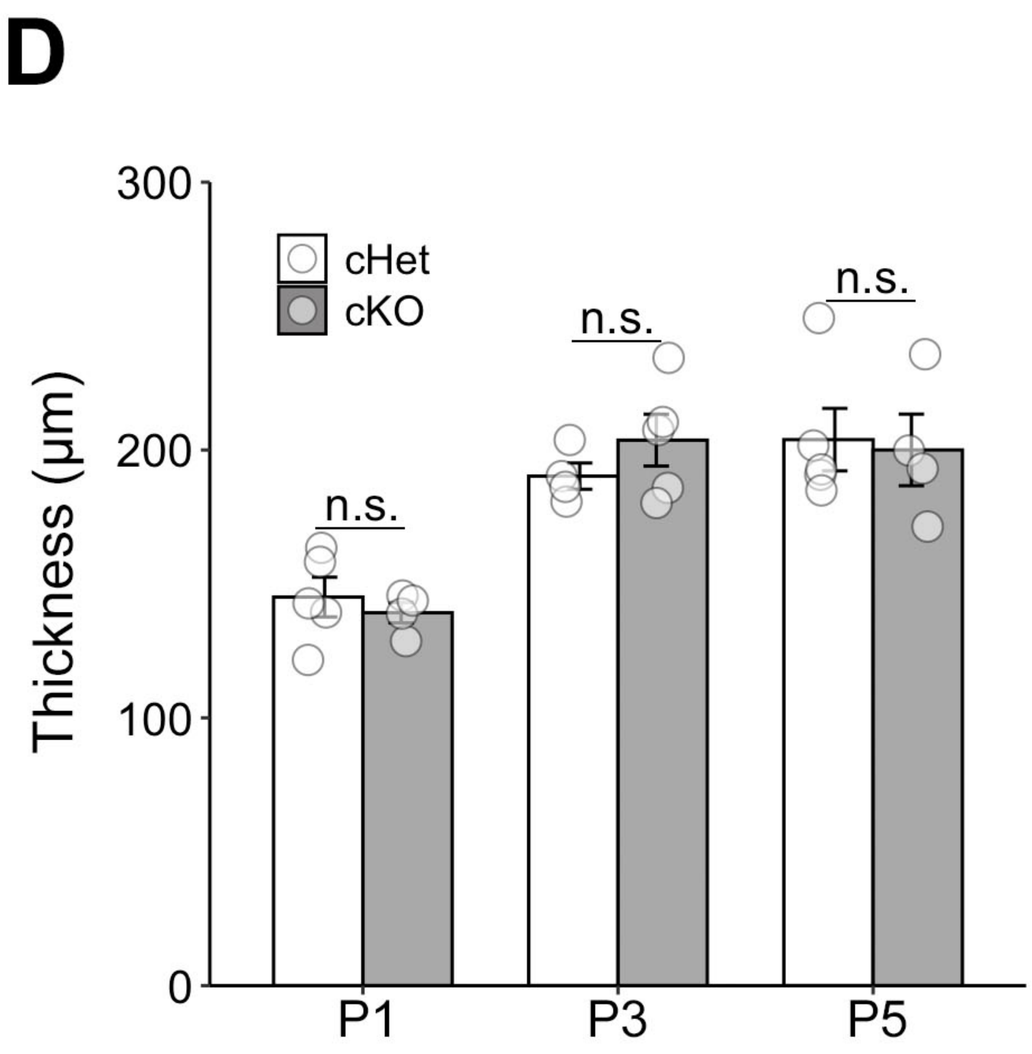

C
P8
P11
P14
Rom1 Rho
cHet
cKO
ONL
OPL
INL
IPL
GCL
20 µm


D
PKARIIβ Arr3
Bsn mGluR6 PKCα
cHet
cKO
ONL
OPL
INL
IPL
GCL
20 µm
5 µm


A
cHet
cKO
Active caspase3
ONL
INL
GCL
20 µm


B
AC3+ cell number
Number of cells /1 mm2
400
300
200
100
0
**
*
*
*
**
cHet
cKO
P1
P3
P5
P14
1M


C
BrdU+ cell number
Number of cells /100 µm
100
75
50
25
0
n.s.
n.s.
n.s.
cHet
cKO
P1
P3
P5


D
Thickness (µm)
300
200
100
0
cHet
cKO
n.s.
n.s.
n.s.
P1
P3
P5