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4 Afadin-deficient retinas exhibit severe neuronal lamination defects but preserve
5 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, 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 the first

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 defects in retinal

lamination (Inagaki et al., 2005). To clarify the effects of AJ molecule depletion

on retinal neural circuits and cell morphology while avoiding the issue of

functional redundancy, we focused on afadin, which is a scaffolding protein for

nectin and has no paralog in mice. Afadin regulates the localization of nectin,

which initiates cell–cell adhesion and promotes AJ formation by recruiting the

cadherin–catenin complex. (Ohama et al., 2018; Takai and Nakanishi, 2003). In

addition, afadin interacts with various cell adhesion and signaling molecules, as

well as the actin cytoskeleton, and contributes to the accumulation of β-catenin,

αE-catenin, and E-cadherin at AJs (Sakakibara et al., 2018; Sato et al., 2006).

Afadin KO mice exhibit severe disruption of AJs in the ectoderm, along with

other developmental defects, leading to embryonic lethality (Ikeda et al., 1999;

Zhadanov et al., 1999). Conditional deletion of afadin in RGCs leads to

disruption of dendrites in ON-OFF direction-selective RGCs (Duan et al., 2018).

However, the effect of afadin loss on retinal lamination, circuit formation, and

function is poorly understood.

In this study, using the *afadin* conditional KO (cKO) mouse, we revealed that the

afadin-deficient retinas exhibit severe pathological defects, such as outer retinal

lamination disruption, as well as decrease and mislocalization of photoreceptors

and their synapses. In contrast to severe disruption of the outer retina, the inner

plexiform layer (IPL) structure in the inner retina was relatively intact. Using the

retinas isolated from the *afadin* cKO mouse, we could record local field

potentials (micro electoretinograms: mERGs) and RGC firings. Based 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.

MATERIALS and METHODS

Animals

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

WT, *afadin* conditional heterozygous (cHet), and cKO mice on 129S6/SvEvTac background, and *R26R-H2B-EGFP* mice on C57BL/6J background were used in this study. The *afadin flox* mice (Majima et al., 2009), *Dkk3-Cre* mice (Sato et al., 2007; Yamamoto et al., 2020), and *R26R-H2B-EGFP* mice (Abe et al., 2011,CDB Accession number. CDB0203K), have been described previously. Exon 2 of the *afadin* gene is flanked by loxP sites in the *afadin flox* mice, and Cre activity is detected in the retina of the *Dkk3-Cre* mice from embryonic day 10.5 (E10.5). Mice of either sex were used for all animal experiments. The stages used for individual experiments are described in figure legends.

Immunohistochemistry

Immunohistochemical analysis was performed as described previously (Hori et al., 2019; Kubo et al., 2021). In brief, the retina used for MEA recordings and isolated mice eyes were fixed with 4% PFA in PBS for 30 min at room

temperature. After three-time washes, retinas were cryoprotected by 30%

sucrose in PBS overnight, embedded in an OCT compound (Sakura, Japan),

frozen on dry ice, and sectioned at 20 μm of thickness. Whole-mount

immunostaining was performed as previously described with some modifications

(Ueno et al., 2018). The retinas were gently peeled off from the sclera, fixed with

4%PFA in PBS for 30 min at room temperature, and washed three times. Retinal

sections and whole retinas were soaked in blocking buffer (5% NDS, 0.1% Triton

X-100, in 1x PBS) for 1-2 h at room temperature, and incubated with primary

antibodies in blocking buffer 1 or 2 overnight at 4°C. The sections were washed

with PBS three times and incubated with fluorescent dye-conjugated secondary

antibodies and DAPI (1:1000) for more than 2 h at room temperature or

overnight at 4°C under the light-shielded condition. The specimens were

observed using a laser confocal microscope (LSM900; Carl Zeiss, Germany).

For BrdU staining, mice were given an intraperitoneal injection of 20 mg/kg BrdU

and sacrificed 2 h later. Retinal sections were then pretreated with 2 N HCl at

37 °C for 30 min before blocking. The antibodies and dilution ratios were as

follows; anti-l-afadin (ab90809, Abcam, UK, 1:100), anti-nectin1 (D146-3, MBL,

Japan, 1:100), anti-nectin2 (D083-3, MBL, Japan, 1:100), anti-nectin3 (D084-3,

MBL, Japan, 1:100), anti-PKCα (P5704, Sigma, USA, 1:1000, P4334, Sigma,

1:10000), anti-SCGN (AF4878, R&D systems, USA, 1:2000), anti-Arr3

(AB15282, Millipore, USA, 1:1000), anti-Calbindin (PC253L-100, Millipore,

USA,1:200), anti-ChAT (AB144P, Millipore, USA, 1:50), anti-Bassoon

(SAP7F407, Enzo, USA, 1:1000), anti-mGluR6 (current study, 1:3000),

anti-PSD95 (#124 014, Synaptic Systems, Germany, 1:3000), anti-GluR5 (Grik1,

gift from Steve H DeVries, 1:2000), anti-PKARIIβ (#610625, BD biosciences,

USA, 1:500), anti-Calretinin (PC235L-100UCN, Millipore, USA, 1:5000),

anti-EAAT5 (HPA049124, Sigma, USA, 1:100), anti-vGlut1 (AB5905, Millipore,

USA, 1:6000), anti-HPC-1 (S0664, Sigma, USA, 1:10000), anti-active caspase3

(AF835, R&D Systems, USA, 1:300), anti-Chx10 (Hori et al., 2019, 1:100),

anti-Otx2 (AF1979, R&D systems, USA, 1:200), anti-Glutamine synthetase (GS,

MAB302, Millipore, USA, 1:500), anti-CD31 (#557355, BD Biosciences, USA,

1:100), anti-Lhx2 (sc-19344, Santa Cruz Biotechnology, USA, 1:100),

anti-RBPMS (GTX118619, GeneTex, USA, 1:500), anti-Ki67 (#556003, BD

Biosciences, USA, 1:100), anti-Tuj1 (#801201, BioLgend, USA, 1:500),

anti-AP2α (3B5-c, DSHB, USA, 1:1000), anti-Rom1 (gift from Robert Moldey,

1:10), anti-Rhodopsin (STJ95452, ST John's Laboratory, UK, 1:100),

anti-S-opsin (sc-14363, Santa Cruz Biotechnology, USA, 1:500), anti-M-opsin

(AB5405, Sigma, USA, 1:500), anti-GFP (GFP-1010, Aves labs, USA, 1:1000),

anti-β-catenin (#610153, BD Biosciences, USA, 1:1000), anti-N-cadherin

(#610920, BD Transduction Laboratories, USA, 1:500), anti-BrdU (#347580, BD

Biosciences, USA, 1:100), and anti-phospho-histone H3S10 (#06-570, Millipore,

USA, 1:2000) antibodies.

Counting of retinal cells and synapses

The cell number of each retinal cell type was counted around the area 500 μm

away from the optic nerve using immunostained retinal sections, and we

calculated the cell number per 100 μm width of retinal section. Each retinal cell

type was identified as follows: rod (Otx2 signal at the nuclear periphery), cone

(Arr3+, Otx2 signal in the soma), BC (Chx10+), horizontal cell (Calbindin+ and

AP2α-), AC (AP2α+), RGC (RBPMS+), and Müller glial cell (Lhx2+).

It was difficult to count the number of synapses between photoreceptor and BC

in 2D images of the cKO retina because the direction of the photoreceptor

synapse was aberrant. Thus, the number of synapses was counted using 3D

images reconstituted from immunostained vertical sections. The image of

vertical sections was acquired with a width (x-axis) of 126.8 μm, a length (y-axis)

of 126.8 μm, and a thickness (z-axis) of 12 μm. Then, the number of synapses

per 1 mm^2 of retinal surface was calculated. The regions distal to the IPL were

analyzed because photoreceptor synapses were not observed below the IPL.

Since the synapses between the rod photoreceptor and BC in the cHet retina

were highly dense, we counted them every 1.8 μm using 2D images and

integrated them. We confirmed that the number of synapses was not different

between 2D and 3D images.

Processing of tissues for electron microscopy

The eyeball was removed, cut in the cornea, and fixed with 2.5% glutaraldehyde

in PBS for 2.5 h on ice, and with 2% osmium tetroxide for 2 h. Retinas were

washed with 1x PBS, and dehydrated with a graded series of ethanol, followed

by propylene oxide, and embedded in Epon 812. Seventy-nm ultrathin sections

were cut by an ultramicrotome (Ultracut E; Reichert-Jung, Germany), mounted

on nickel grids, and stained with 2% uranyl acetate for 4 h and with nitrate for 5

min. Retinal sections were observed by transmission electron microscope

(H-7500; HITACHI Co, Japan).

Reverse transcription real-time quantitative PCR (RT-qPCR)

Total retinal RNA was isolated using Isogen II reagent (Nippon Gene, Japan)

following the manufacturer's instructions, and reverse transcribed into cDNA

using SuperScript II Reverse Transcriptase (Thermo Fisher Scientific, USA) with

random hexamers and oligo (dT) 12-18 primers. Diluted cDNA was used as a

template for qPCR using a TB Green® Premix Ex Taq™ II (Tli RNaseH Plus)

(Takara Bio, Japan) according to the manufacturer's instructions, on Thermal

Cycler Dice® Real Time System II (Takara Bio, Japan). The primer sequences

are listed in **Table S1**.

Western blot analysis

Retinal tissues were lysed in RIPA buffer (50 mM Tris-HCl pH7.6, 150 mM NaCl,

1mM EDTA, pH8.0, 1% Nonided-P40, 1% Sodium Deoxycholate, 0.1% SDS),

cooled for 10 min on ice, and centrifuged at 14,000 rpm for 10 min at 4^oC. Then,

a 3x sample buffer was added to the supernatant, and it was boiled for 10 min at

95^oC. Samples were separated by SDS-PAGE and transferred to PVDF

membranes (Millipore, USA) using a wet transfer cell (Bio-Rad, USA). The

membranes were soaked in blocking buffer (5% skim milk (W/V) and 0.1%

Tween 20 in TBS) and incubated with anti-l-afadin antibody (1:1000) overnight at

4°C or horseradish peroxidase-conjugated anti-GAPDH antibody (1:5000) for 2

h at room temperature. After washing with TBS/0.1% Tween-20 four times for 10

min each, the membranes were incubated with horseradish

peroxidase-conjugated anti-mouse IgG antibody for 2 h at room temperature.

Signals were detected using ImmunoStar LD (Fujifilm Wako, Japan).

Antibody production

cDNA fragments encoding a C-terminal portion of mouse mGluR6 (853-871

residues) were subcloned into *pGEX4T-1* plasmid (Amersham, UK). The

GST-mGluR6 fusion protein was expressed in *Escherichia coli* strain BL21 and

purified with glutathione Sepharose 4B (GE Healthcare, USA) according to the

manufacturer's instructions. The anti-mGluR6 antibody was obtained by

immunizing guinea pigs with the purified GST-mGluR6.

ERG recordings

Mice were dark-adapted for more than 1 h and then anesthetized deeply by

intraperitoneally injecting mixed anesthesia consisting of (in mg/kg) 0.3

medetomidine, 4 midazolam, and 5 butorphanol tartrate. We also administered

0.1% tropicamide and 0.1% phenylephrine to the eyes to dilate the pupil. ERGs

were recorded via contact lens electrodes (Mayo Corporation, Japan), and data

were sampled at 1,250 Hz by using the PuREC system (Mayo Corporation,

Japan). Reference and ground electrodes were placed in the mouth and on the

tail, respectively. Full-field stimulation was presented via an optical stimulator

equipped with white LED (RMG; Mayo Corporation, Japan), and its light intensity

and duration were controlled by an LED visual stimulator (LS-100; Mayo

Corporation, Japan). Scotopic ERGs were recorded by applying 5-ms flash (1.0

x 10^4 cd/m^2) three times in the dark. After 10 min light adaptation (31.6 cd/m^2),

photopic ERGs were recorded by superimposing 5-ms flash (1.0×10^4 cd/m^2)

sixteen times on the adapting background. After noise reduction by a program

installed in the PuREC system, ERGs were lowpass-filtered (<50 Hz) on

Python3 and then analyzed.

MEA recordings

The retinal preparation was made as described previously (Takeuchi et al.,

2018). In brief, a dark-adapted (>1 h) mouse was sacrificed by cervical

dislocation under a dim red light, and its eyes were enucleated. Under a

stereomicroscope equipped with an infrared (IR) image converter (V6833P,

Hamamatsu Photonics, Japan) and IR illumination (HVL-IRM, Sony, Japan), the

retina was isolated from the eye, and the dorsal retina was placed ganglion cell

layer side down onto the multielectrode array (MEA, Multichannel Systems,

Germany, 60pMEA200/30iR-Ti: 60 electrodes, electrode size 30x30 μm,

inter-electrode distance 200 μm) and attached to the MEA by suction using a

vacuum pump (Constant Vacuum Pump, Multichannel Systems, Germany).

When another type of MEA (Multi Channel Systems, Germany,

60MEA200/30iR-Ti-gr: 60 electrodes, electrode size 30×30 μm, inter-electrode

distance 200 μm) was used, a piece of anodisc (13 mm, 0.2 μm, 6809-7023,

Whatman, UK) and a weight with nylon fibers were placed on the retina to

improve the contact with MEA. Then, the MEA chamber was constantly

superfused with Ames' medium (A1372-25, United States Biological, USA)

bubbled with 95% O_2/5% CO_2 at the rate of 6.0 mL/min at 32°C. After >30 min

superfusion, we started recordings. Signals were amplified, sampled at 20 kHz,

and stored using MEA2100-Lite-System (Multichannel Systems, Germany). In

pharmacological experiment, L-(+)-2-amino-4-phosphonobutyric acid (10 μM;

L-AP4, 23052-81-5, Tocris, UK) dissolved in Ames' medium was bath-applied to

the retina.

Light stimulus generated by Psychtoolbox-3 on MATLAB was presented on a

monitor display (P2314H, Dell, Japan), and projected through optics on the

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 Uniform Manifold Approximation and Projection (UMAP). To check the

accuracy of spike sorting, we performed autocorrelation analysis to confirm the

presence of a refractory period (the number of spike events at ± 1 ms bin was

less than 1% of total spike events).

Classification of RGCs based on PSTH

Using the MEA system, we recorded firing responses of RGCs to light

stimulation (0.24 or 0.27 cd/m^2 in intensity, 2 s in duration) presented 7 times

from the monitor and calculated the peristimulus time histograms (PSTHs) with

20-ms bin. The PSTH was smoothed by a Gaussian filter (kernel size: 3 bins, σ =

1). The light-evoked response was defined when the firing rate of the PSTH

exceeded the threshold (increment >+4 SDs, decrement <-2 SDs) determined

by the spontaneous firing rate for 2 s before light stimulation. RGCs were

classified into "ON" (increment within 2 s after light onset), "ON-OFF" (increment

within 2 s after light onset and offset), "ON-OFF inhibition" (decrement within 2 s

after light onset and offset), "OFF" (decrement within 2 s after light onset and/or

increment within 2 s after light offset), and "None" (no clear change in firing rate)

types.

To determine the type of photoreceptor inputs, a 2-s flash of green LED (λ_{max} =

510 nm, 4.8×10^3 photons/s/μm^2) and 2-s flash of UV LED (λ_{max} = 360 nm,

3.3×10^4 photons/s/μm^2) were applied seven times at 8-s interval serially. We

classified RGCs into "Rod (+)" (responded to green LED), "Cone (+)-Rod (+)"

(responded to both green and UV LEDs), "Cone (+)" (responded to UV LED),

and "None" responsive types.

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, S4A-C, S4E, and 9B, 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

S3C, 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 Figures S4D, S4F,

and S6B–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.

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 (Figures 1B, C, S1A), 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 (Figures 1B, S1B). 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 S1C) (Koike et al., 2005).

Immunostaining with anti-afadin, anti-SCGN (Type 2-6 cone BC marker)

(Puthussery et al., 2010), and anti-PKCα (rod BC marker) revealed that afadin

localizes to the dendritic tips, cell bodies, and axons of ON- BCs and at least cell

bodies of OFF-BCs (Figure 1D).

Afadin is required for proper retinal lamination

To investigate the role of afadin in the retina, we generated retina-specific *afadin*

cKO mice by crossing *afadin flox* mice with *Dkk3-Cre* mice (Majima et al., 2009;

Sato et al., 2007; Yamamoto et al., 2020). Cre activity in *Dkk3-Cre* mice was

confirmed by crossing with *R26R-H2B-EGFP* reporter mice (Abe et al., 2011), in

which Cre-dependent excision of a *loxP*-flanked STOP cassette drives nuclear

EGFP expression from the *Rosa26* locus (Figure S2A, B). Afadin was not

detected at the protein level by Western blotting in the *afadin* cKO retina (Figure

S2C). The *afadin* cKO mouse was viable and showed no gross morphological

abnormalities. To examine retinal lamination and morphology, using the cKO

retina at 1M, we performed toluidine blue staining. We found that there were no

clear outer and inner segments of photoreceptors in the outer retina, no distinct

OPL, and no clear compartmentalization of the ONL and INL (Figure 2A). Rod

photoreceptors, identified by inverted nuclear architecture (Solovei et al., 2009)

(Figure 2A insets), were widely scattered in the cKO retina, whereas the IPL and

GCL appeared intact. Furthermore, rosette-like structures, which lacked cell

bodies centrally, were sometimes observed in the cKO outer retina (Figure 2A, B,

C: arrowhead).

To investigate the defects in the OS of the cKO retina in more detail, we stained

retinal sections with rhodamine-labeled peanut agglutinin (PNA, a marker of

cone photoreceptor outer segment and axon terminals) or antibodies against

Rom1 (rod OS marker), Rhodopsin (rod OS marker), S-opsin (S-cone OS

marker), and M-opsin (M-cone OS marker). Rhodopsin, Rom1, S-opsin, and

M-opsin play central roles in phototransduction in the OS (Clarke et al., 2000;

Greenwald et al., 2014; Tanimoto et al., 2015; Xu et al., 2020). Distribution of

Rom1, Rhodopsin, S-opsin, M-opsin, and PNA signals in the cKO retina differed

significantly from those of the a*fadin* cHet retina. In the cKO retina, Rom1,

Rhodopsin, S-opsin, M-opsin, and PNA signals were remarkably decreased in

number and mislocalized in the retina (Figure 2B, C). These markers were not

observed in the rosette-like structure of the cKO retina, suggesting that the

rosette-like structure is different from the photoreceptor rosette as reported

(Stuck et al., 2012). Electron microscopic analysis also demonstrated that only a

few OS structures were observed between retinal cells in the disrupted ONL and

INL regions of the cKO retina (Figure S2D). These findings indicate that rod and

cone phototransduction may be severely impaired in the cKO retina. To

investigate the distribution of cone and rod photoreceptors in the cKO retina, we

performed wholemount immunostaining with antibodies against Arr3, Rhodopsin,

and Rom1. Cell bodies of cone photoreceptors were arranged regularly in the

cHet retina, whereas cone photoreceptor arrangements and morphology were

significantly disordered in the cKO retina (Figure 2D). Rom1, Rhodopsin, and

Arr3 signals were decreased but relatively dense in some areas (Figure 2D).

These results suggest that photoreceptor distribution is disrupted, but

photoreceptors remain in some areas heterogeneously.

We further investigated AJs in the cKO retina by immunostaining with OLM

adherens junction markers β-catenin, N-cadherin, and nectin-1. We found that

these signals were scattered in the cKO retina (Figure S2E). This data suggests

that AJs in the OLM are disrupted in mature cKO retina.

As toluidine blue staining of the cKO retina suggested mislocalization of each

cell type, we investigated the position of each retinal cell body in the cKO retina

by immunostaining with retinal cell type-specific markers; Otx2 (a photoreceptor

and BC marker), Chx10 (a BC marker), Arr3, Calbindin (a horizontal cell and AC

marker), AP2α (an AC maker), RBPMS (a RGC marker), and Lhx2 (a Müller glial

cell marker) (Figure 2E). All retinal cell types were aligned in restricted and

orderly regions in the cHet retina. In the cKO retina, the spatial organization of

rod photoreceptors, BCs, cone photoreceptors, horizontal cells, and Müller glial

cells appeared particularly disrupted, whereas positions of ACs and RGCs

remained relatively intact, except for the immediately surrounding region of

rosette-like structures. To quantitatively assess the retinal cell positioning in the

cKO retina, the distance from the soma of each cell type to the outer or inner

retinal surface was measured (Figure 2F-L). The distance for each cell type in

the cKO retina was significantly different from that in the cHet retina. In the cKO

retina, BCs, horizontal cells, Müller glial cells were positioned close to the outer

retinal surface and ACs were close to the inner retinal surface, whereas

photoreceptors and RGCs were positioned away from the outer and inner retinal

surfaces, respectively.

Notably, BCs in the cKO retina were broadly distributed throughout the retina

(Figure 2H). Rod photoreceptors also tended to be dispersed throughout the

cKO retina. However, approximately 75% of them in the cKO retina were located

within 60 μm of the outer retinal surface, which corresponds to the region where

all rods are positioned in the cHet retina (Figure 2G). Cone photoreceptors,

horizontal cells, and Müller glial cells were similarly spread, but their distribution

was largely confined to the area distal to the IPL (Figure 2F, I, J). Although ACs

and RGCs were mainly located in the inner retina, a sparse distribution was also

observed throughout the retina, which may be attributed to the cells surrounding

the rosette-like structure (Figure 2K, L). The reduction in the distance from ACs

to the inner retinal surface in the cKO retina is conceivably due to a shortening of

the IPL (Figure 2K), though there was no significant difference in retinal

thickness between cHet and cKO retinas (Figure 2M). The cKO mice appeared

to show greater inter-individual variability in retinal layer structure and cell

positioning than the cHet mice.

Immunostaining with an anti-GS antibody (a Müller glial cell marker) showed that

Müller glial cell morphology was affected, especially at the distal part. However,

a distinct inner limiting membrane was retained in the cKO retina (Figure S2F). It

has been reported that defects in the distal processes of Müller glia are

associated with abnormal retinal vasculature (Shen et al., 2012). Thus, we

immunostained the cKO retina with anti-CD31, a blood vessel marker, but no

apparent vascular abnormalities were detected (Figure S2G). These results

indicate that afadin is essential for normal retinal lamination.

Ablation of afadin results in the remarkable decrease and mislocalization of photoreceptor-BC synapses

No distinct OPL structure in the cKO retina suggests that synapses between photoreceptor and BC may be disrupted. To assess their integrity, we immunostained retinal sections of the cKO mouse (1 M) with antibodies against Bassoon (a photoreceptor synaptic ribbon marker), PSD95 (a photoreceptor synaptic terminal marker), mGluR6 (an ON-BC dendritic tip marker), GluR5 (an OFF-BC dendritic tip marker), PKCα, and PKARIIβ (a type 3b OFF cone BC marker). In the cHet retina, photoreceptor synapses containing synaptic ribbons formed a row in the OPL (Figures 3A, B, S3A). On the other hand, in the cKO retina, the row of photoreceptor synapses was not observed, and photoreceptor-BC synapses were remarkably reduced in number. Bassoon and mGluR6 were aberrantly distributed, and the position of the soma and dendritic tips of rod BCs was noticeably disturbed (Figure 3A). Only a few ribbon synapses remained (Figure 3A arrowheads). The 3D reconstructions of the confocal image stacks showed more clearly that photoreceptor synapses decreased in number and mislocated in the cKO retina (Figure 3C, D). The contact between the axon terminal of photoreceptors and the dendrite of ON- and OFF-BCs was also significantly reduced in number, suggesting that photoreceptor-BC synapses were considerably lost in the cKO retina (Figures 3B, D, S3A, B). Along with these defects, GluR5 signals with a branch-like pattern were detected in the IPL of the cKO retina, though no GluR5 signals were found in the IPL of the cHet retina (Figure 3B, D). Photoreceptor-ON BC

ribbon synapses and photoreceptor-OFF BC synapses in the outer retinal region

beneath 1 mm^2 of the surface area of the cKO retina were significantly reduced

to about 5% and 15% of those in the cHet retina, respectively (Figures 3E, F,

S3C). These results indicate that afadin may regulate the formation and

localization of photoreceptor synapses.

Afadin is required for proper localization of GluR5 to OFF BC dendrites

GluR5, a kainate receptor subunit that forms functional homomeric and

heteromeric channels, is specifically expressed in OFF BCs and localizes to their

dendrites (Haverkamp et al., 2003). To investigate GluR5 localization in more

detail, we immunostained the cKO retina with anti-GluR5, anti-SCGN, and

anti-PKCα antibodies. We found that ectopic GluR5 signals in the IPL

overlapped not only with SCGN but also PKCα in the cKO retina (Figure 4A).

These data indicate that afadin may be required for specific localization of GluR5

in OFF BC dendrites. It seems likely that ectopically localized GluR5 may

receive glutamate released from surrounding cells in the IPL, resulting in

depolarization of ON- and OFF-BCs. It should be noted that there were no

remarkable differences between cKO and cHet retinas in the localization of

glutamate transporter EAAT5 and vesicular glutamate transporter vGlut1 to the

axon terminal of photoreceptors and BCs (Figures 4B, S3D, E). It has been

reported that mGluR6 localization to ON BC dendritic tips requires trans-synaptic

interaction with ELFN1 in the synaptic cleft and that ELFN1 ablation results in a

remarkable reduction of mGluR6 in ON BC dendritic tips (Cao et al., 2015). As

GluR5 localization may also be regulated by trans-synaptic interaction with

photoreceptor synaptic terminal proteins, it seems possible that the defect of

photoreceptor-OFF BC contacts prevents this trans-synaptic interaction,

resulting in GluR5 mislocalization in the cKO retina.

Then, we investigated the IPL structure in the cKO retina by immunostaining it

with anti-ChAT, Calbindin, and Calretinin antibodies. Distinct ON- and OFF ChAT

bands, Calbindin bands, and Calretinin bands were observed in the cKO retina

(Figure S3F). However, type 2-6 BCs immunostained with SCGN were slightly

disturbed in the axon terminal position (Figures 4A, B, S3D). These results

suggest that stratification patterns in the IPL may be slightly impaired but almost

intact in the cKO retina. Immunostaining analysis with anti-HPC-1 (also known

as syntaxin, an AC marker), anti-Calbindin, anti-RBPMS, and anti-Tuj1 (an RGC

marker) antibodies revealed that the rosette-like structure contains AC and RGC

processes inside (Figure 4C). These data indicate that the rosette-like structure

in the cKO may be an ectopic IPL, termed "acellular patches" (Nahar and Cho,

2022).

Retinal cell fate is altered in the cKO retina

To examine whether afadin loss affects retinal cell fate determination, we

counted the number of retinal cells immunostained with each retinal cell marker.

In the cKO retina at 1M, BCs significantly increased to about 150% of the cHet

retina, but rods, cones, and Müller glial cells were significantly decreased to

about 30%, 40%, and 80% of the cHet retina, respectively (Figure 4D).

PKCα-positive cells and SCGN-positive cells were significantly increased

(Figure S4A), indicating that both rod- and OFF-BCs were increased in the cKO

retina. In contrast, the number of rod BC processes in the IPL was significantly

decreased in the cKO retina (Figure S4B). These data suggest that ectopically

located BCs may increase in the cKO retina at 1M. A significant increase in BCs

and a significant decrease in photoreceptors were also observed in the cKO

retina at P14, a stage at which retinal differentiation is complete (Figure S4C).

Consistently, the expression of marker genes for photoreceptors and BCs was

significantly decreased and increased, respectively, in the cKO retina (Figure

S4D). In contrast, the number of Müller glial cells was unaltered in the cKO retina

at P14 (Figure S4C), suggesting that the reduction in Müller cells observed at

1M may result from cell death after P14 rather than a differentiation defect. To

assess whether retinal cell fate changes occur during differentiation, we

examined the number of BCs and the expression of rod marker genes in the

cKO and cHet retinas at P1, P3, and P5. No significant differences were

observed between genotypes at P1 and P3, whereas the number of BCs was

significantly increased, and the expression of rod marker genes was significantly

decreased in the cKO retina at P5 (Figure S4E, F). These results suggest that

defects in cell fate determination of BCs and rod photoreceptors begin to emerge

between P3 and P5. Based on these data and ectopic localization of GluR5 to

ON-BCs in the cKO retina (Figure 4A), we propose that afadin may be required

for normal retinal cell fate determination and gene expression.

Retinal structure is severely disrupted during development in *afadin* cKO

mice

To investigate whether the aberrant lamination in the cKO retina can be

attributed to a defect in development or maintenance, we immunostained the

developing cHet and cKO retinas using markers for individual retinal cell types,

outer segments, and photoreceptor synapses (Figures 4E, S5A-D). In the cKO

retina at P1, retinal cells appeared to be organized into distinct multicellular

compartments with apparent boundaries (Figures 4E, S5A, B). RGCs, ACs, and

their dendritic processes, which are components of the IPL, were localized along

the boundaries of these compartments (Figure S5A, B). Furthermore, RGCs and

ACs were observed in the outer part of the retina. At P3 and P5, the interspaces

between compartments appeared to be reduced, coinciding with an increase in

RGCs and ACs positioned on the inner side of the retina (Figure S5A, B). By

comparison, the positioning of BCs and rod photoreceptors appears to remain

disorganized throughout postnatal development (Figure 4E). Contrary to our

expectation that retinal structure would become increasingly disorganized as

development progresses, laminar disruption in the cKO retina appeared more

severe at earlier postnatal stages than at later ones, particularly in the IPL, ACs,

and RGCs (Figure S5A, B). Aberrant acellular regions were observed in the cKO

retina from P1 through P14 (Figures 4E, S5A-C). Some of these regions

contained AC processes and were surrounded by AC and RGC somata from P3

onward (Figure S5A, B). Correspondingly, clear signals of rod outer segment

markers were observed in the interior of some acellular regions at P8, P11, and

P14 (Figure S5C). At 1M, these structures were not observed in the cKO retina

(Figure 2B). These data suggest that both acellular patch and photoreceptor

rosette may be formed in the developing retina, but photoreceptor rosette may

degenerate after P14. The outer segment marker signals were disorganized in

the cKO retina at P8, P11, and P14 (Figure S5C). Moreover, the row of rod and

cone photoreceptor synapses was not observed in the cKO retina at P11 (Figure

S5D). These data indicate that proper development of retinal lamination and

synapse formation is impaired in the cKO retina.

Small local field potentials are recorded from the cKO retina

To examine whether the cKO mouse retina can respond to light stimulation, we

first recorded electoretinogram (ERG) from anaesthetized animals. Under the

scotopic condition, a flashlight was applied to eyes in the dark (Figure 5A *left*).

Scotopic ERG from the cHet mouse showed a negative a-wave and a following

positive b-wave, whereas both waves from the cKO mouse were almost flat.

Under the photopic condition, the mouse was adapted to the background light for

10 min, then a flashlight was superimposed on the adapting background (Figure

5A *right*). Photopic ERGs from the cHet mouse showed a small a-wave and a

prominent b-wave, whereas those from the cKO mouse were almost flat. These

results may be attributed to the scattered distribution of rod and cone

photoreceptors, a lack of a uniform photoreceptor layer, and the decrease of

outer segments and photoreceptor synapses in the cKO retina (see Figures 2

and 3).

Then, to examine whether local field potentials can be detected from the cKO

retina, we performed MEA recordings from the retinas isolated from the cKO and

cHet mice. Surprisingly, a flashlight could occasionally evoke micro-ERGs

(mERGs) from the cKO retina (SD of the amplitude during 300 ms after the

flashlight onset; >50 µV, 13/590 electrodes, n = 10 retinas from 6 mice) (Figure

5B *right*). However, the amplitude of mERGs from the cKO retina was smaller

than that from the cHet retina (SD of the amplitude during 200 ms after the

flashlight onset; >1,000 µV, 292/413 electrodes, n = 7 retinas from 4 mice)

(Figure 5B *left*). As the a-wave represents hyperpolarization of the

photoreceptors and the b-wave arises from the inner retina, predominantly from

ON BCs with a contribution from Müller glial cells (Perlman, 2020), the first

negative and following positive waves of mERGs may correspond to the a- and

b-waves, respectively. To confirm this, L-AP4, an agonist of mGluR6, was

bath-applied to suppress the response of ON BCs. In the presence of L-AP4, the

second positive wave disappeared, whereas the first negative wave was

enhanced in the cHet retina and remained in the cKO retina (Figure 5B),

indicating that the first negative and following positive waves are the a- and

b-waves, respectively. We compared the amplitude and implicit time

(time-to-peak latency) of the a- and b-waves and found that both a- and b-waves

were significantly smaller in amplitude (Figure 5C) and longer in implicit time in

the cKO retina than those in the cHet retina (Figure 5E). The ratio of b-wave

amplitude to a-wave amplitude (b/a) was significantly smaller in the cKO retina

than in the cHet retina (Figure 5D), suggesting that the efficiency of synaptic

transmission from photoreceptors to ON BCs may be decreased in the cKO

retina. However, the difference of the implicit time between a- and b-waves was

similar in the cKO and cHet retinas (Figure 5F), suggesting that signals from

photoreceptors to ON BCs may be transmitted properly. These results indicate

that despite severe disorganization of the cKO outer retina, photoreceptors can

still respond to light stimulation and that the synaptic transmission from

photoreceptors to ON-BCs remains functional.

RGCs in the cKO retina evoke firings to light stimulation

Using the MEA system, we recorded spike discharges from RGCs in retinas

isolated from the cKO and cHet mice. The frequency of spontaneous firings in

the dark was not different significantly between cKO and cHet retinas (cHet; 12.5

± 19.2 Hz, n = 434, cKO; 10.8 ± 14.6 Hz, n = 388, p = 0.79, Mann-Whitney U

test). We examined whether RGCs in the cKO retina respond to light stimulation.

The isolated retina was stimulated by a flash seven times every 8 s, and the

PSTH was calculated for each RGC after spike sorting. RGCs could be

classified into "ON", "ON-OFF", "ON-OFF inhibition", "OFF", and "None" types

(Figure 6A, B; see **MATERIALS and METHODS**). All types were found in both

cKO and cHet retinas, and the composition of RGC types was similar in both

retinas (Figure 6C). We also investigated whether RGCs received both rod and

cone inputs and found that > 60% of RGCs among successfully spike sorted

RGCs in both the cKO and cHet retinas received both inputs (Figure 6D).

Receptive fields are characterized in some RGCs

Using pseudorandom checkerboard pattern stimulation, we examined the

properties of spatiotemporal RF of RGCs by calculating the spike-triggered

average. Amazingly, we could identify RFs in the cKO retina. To check the

morphology of the retina where RFs were identified, the retina was removed

from the MEA after recording, and sliced retinal sections were immunostained

with anti-PKARIIβ, anti-PSD95, and anti-PKCα antibodies (Figure 7A). The

sectioned region clearly showed impairment of the outer retina, where it was

difficult to identify or locate photoreceptors. On the other hand, the inner retina

was apparently normal, and both the IPL and the proximal part of the INL could

be recognized. Even though the outer retina was severely impaired, we could

identify RF of RGCs (Figure 7B). However, the cells with clear RF were fewer in

the cKO retina (69/388 cells, n = 4 retinas) than those in the cHet retina (195/434

cells, n = 4 retinas). By fitting ellipse to each RF, we measured the short and long

axes and calculated the RF area (Figure 7C, D, E). The RF area in the cKO

retina was significantly smaller than that in the cHet retina. We tried to find an

antagonistic surround of RFs, but it was difficult to detect a clear surround in

both cKO and cHet retinas.

The temporal profile of RFs was calculated, and the principal component

analysis was performed. Then, using UMAP for clustering, we could recognize 5

clusters (Figure 7F, cKO; n = 4 retinas, cHet; n = 4 retinas). The temporal profile

belonged to each cluster was superimposed separately (Figure 7G). Among 5

clusters, clusters 1 and 2 did not include the temporal profiles calculated from

RGCs in the cKO retina. The temporal profiles of cluster 1 and cluster 2 were ON

and OFF types, respectively, each of which had a shorter time-to-peak latency

than the other temporal profiles obtained from RGCs in both cKO and cHet

retinas (Figure 7H). These results indicate that the light-evoked responses of

RGCs in the cKO retina are not as refined as cHet. Actually, the rising time of

transient responses (determined by an upward deflection soon after light

stimulation obtained from the cumulative curve of each PSTH) was significantly

longer in the cKO retina (58.5 ± 19.8 ms, n = 17) than in the cHet retina (43.0 ±

17.8 ms, n = 23) (***p < 0.001, Student's test).

It has been reported that degenerated retinas, such as the *rd1* retinas (Poria and

Dhingra, 2015), show oscillatory spontaneous firings in RGCs. Thus, we

checked whether RGCs in the cKO retina show RGC oscillations. RGCs in the

cKO retina showed oscillations (P30; 1.55% of 382 RGCs, 14.0 ± 2.4 Hz, n = 4

retinas, P40; 15.0% of 333 RGCs, 9.2 ± 3.8 Hz, n = 4 retinas). On the other hand,

RCGs in the cHet retina did not show obvious RGC oscillations (0% of 432

RGCs, n = 4 retinas).

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Afadin cKO mice retain partial visual function

As the MEA analyses suggested partial preservation of retinal functions in the

cKO retina, we further examined whether cKO mice retained visual function,

using visual behavior tests. We first conducted the optomotor response (OMR)

test, a robust method for assessing contrast-based visual function and

estimating visual acuity in mice (Thaung et al., 2002; Ye et al., 2021). The OMR

test measures head-tracking movements in response to a vertical square wave

grating stimulus (Figure 8A). Mice capable of detecting contrasting motion

passing through their visual field exhibit head-tracking movements in response

to stimuli (Kretschmer et al., 2015; Mitchiner et al., 1976; Schmucker et al.,

2005). Apparent head-tracking movements were observed in the cHet and cKO

mice during presentation of drifting grating stimuli at spatial frequencies of 0.056

and 0.1 cycles/degree, but not in those with optic nerve crush (ONC), in which

the optic nerve had been mechanically crushed, resulting in vision loss (Figure

8B, C). The number of head-tracking movements in response to these stimuli

was significantly higher in the cHet and cKO mice than in the ONC mice. At 0.27

cycles/degree, the cKO mice hardly exhibited head-tracking movements, in

contrast to the cHet mice, and no significant difference was observed between

the cKO and ONC mice under this condition. The number of movements was

significantly lower in the cKO mice than in the cHet mice at all three spatial

frequencies (Figure 8C). There was no significant difference in the number of

head-tracking movements across the three spatial frequencies in the cHet mice,

whereas in the cKO mice, the number tended to decrease with increasing spatial

frequency (Figure 8D). These data suggest that the cKO mice can recognize

moving visual stimuli, but their ability is inferior to that of the cHet mice.

We further conducted the visual cliff test to evaluate visual performance in a

perceptual task. This test exploits rodents' innate tendency to avoid the deep

side of the cliff—a plexiglass surface suspended 80 cm above a checkerboard

on the floor—and prefer the shallow side, where the plexiglass covered directly

atop the checkerboard (Figure 8E) (Fox, 1965). The cHet mice spent

approximately 80% of their time on the shallow side, significantly more time than

the cKO and ONC mice, while the cKO mice did not differ significantly from the

ONC mice in time spent on the shallow side (Figure 8F). Nonetheless, some

cKO mice exhibited a behavior in which they looked down toward the bottom of

the deep side and quickly backed away (Video 1). This behavior, observed in the

cHet mice but not in the ONC mice, may imply that they can visually recognize

the deep side (Video 2). Furthermore, the cKO mice, but not the ONC mice,

showed a significant preference for the shallow side over the deep side (57.35%,

p = 0.048, one-sample t-test, figure not shown). These results indicate that

visual information from the retina is transmitted to the brain and visual function is

partially maintained in *afadin* cKO mice.

Ectopic AJs and aberrant localization of progenitor cells are observed in

the developing *afadin* cKO retina

As RGC oscillations were observed in the cKO retina, like the *rd1* retinas with

photoreceptor degeneration (Choi et al., 2014), we examined whether RGC

oscillations were related to cell death in the cKO retina. Using anti-active

caspase3 (an apoptotic cell marker), we counted the number of dead cells

immunohistochemically. We found that dead cells were significantly increased in

the cKO retina at around P30 (Figure S6A, B). These findings suggest that the

progressive cell death observed in the cKO retina may lead to a further decrease

of photoreceptors, resulting in an increase of oscillatory RGCs at P40. Apoptotic

cells were significantly increased in the cKO retina in the developing stage at P1,

P3, P5, and P14 (Figure S6B). In the cKO retina at 1 M, the number of rod

photoreceptors and Müller glial cells was significantly reduced to approximately

60% and 70% of that at P14, respectively (P14; rod 98.3 ± 3.8, Müller glial cell

17.2 ± 3.3, n = 3, 1M; rod 57.3 ± 17.3, Müller glial cell 12.4 ± 1.5, n = 4. rod *p* =

0.011, Müller glial cell $p = 0.046$ by Student's t-test), in contrast to the cHet retina,

in which the number of rod photoreceptors remained unchanged (P14; rod

176.5± 14.3, Müller glial cell 17.3 ± 1.4, n = 4, 1M; rod 185.9 ± 17.5, Müller glial

cell 15.7 ± 2.3, n = 4, rod $p = 0.43$, Müller glial cell $p = 0.27$ by Student's t-test).

This reduction may result from ongoing rod cell death even after the completion

of differentiation.

Then we investigated the mechanism underlying lamination defects in the cKO

retina. We previously demonstrated that the photoreceptor-cell specific *aPKCλ*

KO retina shows disruption of AJs and mislocalization of mitotic progenitors in

developing stage, followed by severe lamination defects in adults (Koike et al.,

2005). In the developing retina, AJs are observed in the outer retinal surface,

where progenitor cells divide and migrate toward the inner retina. To investigate

the possibility that disruption of AJs may lead to abnormal cell migration and

lamination, we immunostained the cKO retina at P0 with anti-β-catenin,

N-cadherin, nectin-1, and phospho-histone H3S10 (pHH3, a marker of the

mitotic cell) (Figure 9A). N-cadherin, nectin-1, and β-catenin signals were

obviously detected, but these signals were localized in the inner retina with

aberrant patterns. These signals were slightly observed in the outer retina.

These data indicate that ectopic AJs are present in the cKO retina. Mitotic cells

immunostained by pHH3 were localized near the ectopic AJs. The number of

pHH3-positive cells was significantly increased to approximately 150% of that in

the cHet (Figure 9B). At P1, P3, and P5, mitotic cells were also mislocalized to

the middle retinal region in the cKO mice, and their distance from the outer

retinal surface was significantly larger —approximately 45 µm, 70µm, and 75 µm,

respectively— than that in the cHet (Figure 9C, D). Mitotic cells showed a

broader distribution in the cKO retina than in the cHet. We further examined the

distribution of retinal progenitor cells during development. In the cHet retina,

progenitor cells labeled with BrdU were scattered but relatively enriched in the

middle region at P1 (Figure 9E, F). As the development progressed, the number

of progenitor cells decreased, and their distribution was progressively confined

to the middle region of the retina. In the cKO retina, progenitor cells were more

broadly distributed (Figure 9E, F). At P5, no significant difference in the distance

was observed between the cHet and cKO retinas However, progenitor cells in

the cKO retina showed a trend toward a broader distribution compared to the

cHet retina. Retinal thickness and the number of progenitor cells did not differ

significantly between the cHet and cKO retinas at P1, P3, and P5 (Figure S6C,

D). These data suggest that progenitor cells may divide at ectopic regions and

migrate aberrantly into the cKO retina, resulting in lamination defects.

DISCUSSION

In the current study, we showed that the *afadin* cKO retina exhibits various

severe pathological defects, including photoreceptor morphological defects,

mislocalization, reduced number of photoreceptors and their synapses,

disruption of the outer retinal lamination, ectopic localization of GluR5, aberrant

rosette-like structure in the outer retina (Figures 2-4). These results suggest that

afadin is required for normal retinal lamination, photoreceptor cell morphology,

photoreceptor synapse formation, and proper GluR5 localization to OFF bipolar

cell dendrites. Conditional loss of AJ molecules in the retina has been reported

to cause partial or complete disruption of retinal lamination (Erdmann et al.,

2003; Fu et al., 2006; Masai et al., 2003). However, the effect of AJ molecule

loss on the retinal cell morphology and synapses was unclear. We revealed that

loss of afadin significantly but not completely reduces photoreceptor synapses.

In the *afadin* cKO mouse retina with severe outer retinal disorganization, we

detected the a- and b-waves of mERG and the RF of RGCs (Figures 2, 5, 7).

These results suggest that some visual information processing may be

performed by newly formed and preserved circuits even when the retinal

structure is severely impaired and the number of photoreceptors is remarkably

decreased. To our knowledge, this is the first report that the RF of RGCs could

be formed if several functional photoreceptors connect to BCs, even when the

outer retinal lamination is severely disrupted (Figure 7). The number and size of

RFs observed in the cKO retina were smaller than those in the cHet retina,

suggesting that the precise arrangement of retinal cells and lamination is

necessary for normal visual function. It is important to note that light stimulation

evoked firing responses in RGCs around the rosette-like structure, and that such

RGC firings could not be detected by a large distance between RGC soma and

MEA electrode.

We would like to speculate why the RF of RGCs was observed in the *afadin* cKO

retina. First, visual processing may be partially preserved without any

remodeling if only a small number of functional photoreceptor-BC-RGC

pathways remain intact. It has been shown that the RF of RGCs is detected even

when approximately half of cone photoreceptors are stimulated (Care et al.,

2019; Lee et al., 2022). Thus, in the *afadin* cKO retina, where cone

photoreceptors are reduced to about 40% of the cHet retina, such reduction *per*

se may not critically affect the RF organization. Furthermore, RFs of RGCs are

also detected in several mouse models of retinitis pigmentosa, in which rod

photoreceptors are degenerated and surviving cone photoreceptors lose their

OS discs and pedicles, instead forming abnormal processes resembling

synaptic dendrites (Barhoum et al., 2008; Ellis et al., 2023; Scalabrino et al.,

2022). A few photoreceptor-BC-RGC pathways (vertical pathways of the retina)

are inferred to be maintained in the cKO retina. In some regions, the density of

photoreceptors was high enough to make functional synapses between

photoreceptor and BC (Figure 2D), possibly resulting in the RF of RGCs. Second,

synaptic rewiring of interneurons such as BCs may recover retinal function in the

afadin cKO retina. Various studies show partial loss of photoreceptors or BCs, or

partial silencing of BCs causes rewiring in the developing and mature retinas

(Johnson and Kerschensteiner, 2014; Jones et al., 1995; Okawa et al., 2014;

Shen et al., 2020; Strettoi et al., 2022). Also, in the *rd1* retina, neural connections

between inner retinal neurons via gap junctions may be enhanced, potentially

resulting in the widespread distribution of electrically evoked RGC responses

(Ahn et al., 2022). These reports suggest that reduced input from photoreceptors

and BCs may lead to synaptic reorganization among surviving cells. It is possible

that synaptic reorganization triggered by reduced photoreceptor and BC inputs

may occur in the *afadin* cKO retina with severe defects of the photoreceptor

synapses and outer segments in the developing stages, and that this might

contribute to RF formation. Synaptic rewiring may occur between BC dendrite

and their closely located photoreceptor terminal in the cKO retina. It may be

necessary to count the number of BC dendritic connections to each cone pedicle

to verify the synaptic rewiring between photoreceptor and BC, but this was quite

difficult because of the remarkably aberrant cone morphology in the cKO retina.

As the IPL was relatively intact and BC axon terminals were surrounded by many

RGC dendrites, synaptic reconnections between BC axon terminal and RGC

dendrite are presumed to be more easily established than those between

photoreceptor terminal and BC dendrite in the cKO retina. Thus, several BCs

connected with multiple functional photoreceptors may rewire to RGC dendrites

and transmit signals efficiently. Our findings imply that a very few BCs receiving

input from functional photoreceptors responded to light stimulation, and that the

output signal can be amplified downstream. The above two possibilities could

both contribute to the formation of RGC RFs. Furthermore, we cannot rule out

the possibility that gene expression may be changed in the *afadin* cKO retina,

resulting in facilitation of RF formation.

The ratio of responding to non-responding RGCs to light stimulation in the *afadin*

cKO retina was comparable to that in the cHet retina despite notable reduction of

photoreceptors (Figures 4D, 5, 6). It seems likely that the number of

photoreceptors connected to spreading dendrite tips of one BC may be

increased, and that glutamate spillover from BC terminals may contribute to

increasing the responsiveness of RGCs in the cKO retina because glutamate

transporter EAAT5 and vesicular glutamate transporter vGlut1 were similar in

both cKO and cHet retinas (Figures 4B, S3D, E). Glutamate released from BC

terminals may spread and bind GluR5, which is ectopically localized in BC

processes in the IPL, enhancing the response and expanding the responding

area (Figure 4A, B). These factors may contribute to the emergence of RF, but

the RF size in the cKO retina was still smaller than that in the cHet retina (Figure

7). Photoreceptors in the cKO retina were reduced to 30 to 40% of the cHet

retina and scattered at 1M (Figures 2, 4D). The random checkerboard pattern

used for STA was projected onto the assumed photoreceptor layer of the normal

retina. However, as the morphology and position of photoreceptors were

disturbed in the cKO retina (Figures 2, S2D), the remaining photoreceptors

would be stimulated by an unfocused image with decreased intensity.

Accordingly, photoreceptors possibly responded weakly to unfocused images,

and thresholding of noise (increment >+4 SDs, decrement <-2 SDs, determined

by the spontaneous firing rate, see **MATERIALS and METHODS**) may result in

reduction of the RF size. Rod-driven responses were observed in RGCs in the

cKO retina (Figures 5, 6). However, it is unlikely that the rod photoreceptors

transmit signals to cone photoreceptors through gap junctions in the disrupted

cKO outer retina, and thus, rod signals may not be transmitted to RGCs via the

traditional AII amacrine-ON cone BC pathway.

In the cKO retina, the temporal profile of RFs with short time-to-peak latency was

not included (Clusters 1 and 2; Figure 7), indicating that RGCs in the cKO retina

cannot respond to rapid changes in luminance. There are several ways to

explain this result. For example, disruption of the outer retinal lamination may

make it difficult to synchronize photoreceptor responses without gap junctions,

resulting in dull responses. BCs with fast response kinetics may not be present

or not functional in the cKO retina (Kuo et al., 2024). Furthermore, glutamate

spillover from BC terminals may amplify the signals, but at the same time, the

time-to-peak of RGC responses may be delayed. These possibilities should be

investigated in the future.

Ectopic continuous AJs formed in the developing *afadin* cKO retina (Figure 9A)

are possibly associated with partial disruption of retinal lamination. The *β-catenin*

cKO retina also shows ectopic continuous AJs in developing stage and the

disruption of the outer retinal lamination with relatively intact IPL and GCL

structure in adult (Fu et al., 2006), suggesting that afadin and β-catenin are

required for normal continuous AJ formation at the outer retinal surface but

dispensable for maintaining AJs. The photoreceptor-specific *aPKCλ* cKO retina

in which AJs are dispersed in the developing stage exhibits disruption of whole

retinal lamination despite aPKCλ remaining in cells excluding photoreceptors.

Individual cells contain AJs, but the AJs are not continuous in the *aPKCλ* cKO

retina (Koike et al., 2005). These findings, based on the studies using each

mutant mouse, imply that continuous AJs are important for the formation of the

outer retinal structure. The ONL was severely disrupted in the cKO retina,

despite the fact that afadin is not localized in the layer. The disruption is

presumed to be caused by defects in AJs in the outer retina during development.

In the developing *afadin* cKO retina, ectopic AJs and mislocalized progenitor

cells were observed (Figure 9). These defects may lead to abnormal cell

migration and positioning, ultimately resulting in ONL disorganization. In contrast,

the IPL structure appeared relatively intact in the cKO retina, possibly due to

functional redundancy between afadin and other cell adhesion molecules.

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In the *afadin* cKO retina, significant increase in BCs and significant decrease in

photoreceptors and Müller glial cells were observed at 1M (Figure 4D). The

changes in the number and/or marker gene expression of photoreceptors and

BCs were also observed in the developing cKO retina (Figure S4C-F). These

results suggest that retinal cell fate was affected in the *afadin* cKO mice. As

retinal cell fate is not altered in the *β-catenin* cKO retina (Fu et al., 2006), the

phenotypic difference between *β-catenin* cKO and *afadin* cKO retinas may imply

distinct functional properties of the in AJ molecules.

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One of the major concerns of retinal regeneration therapy is that while

transplanted photoreceptors can form synapses with donor BCs, the

transplanted tissue often shows aberrant structures, such as rosettes (Yamasaki

et al., 2022). Our study demonstrated that RGCs show RFs and visual behaviors

are partially preserved even when the outer retinal lamination is severely

disrupted, provided that synaptic connectivity is maintained to a certain extent.

Therefore, mice with photoreceptor degeneration that have undergone retinal

transplantation might recognize stimulus patterns to some extent, even if

aberrant structures are formed after retinal transplantation. Moreover, retinal

regeneration therapy could be more effective in restoring vision than previously

anticipated. In recent studies, distinct RFs of RGCs were detected in some

mouse lines with progressive photoreceptor degeneration or substantial loss of

cone photoreceptors or BCs, indicating the robustness of retinal circuitry (Care

et al., 2019; Ellis et al., 2023; Johnson and Kerschensteiner, 2014; Lee et al.,

2022; Scalabrino et al., 2022). However, whether visual function is retained in

these models remains unclear. Behavioral visual responses are partially

preserved in the *afadin* cKO mice, in which RFs of RGCs are reduced in number

and size but clearly detectable. Given that RGCs are intact in the mouse lines

used in the aforementioned studies, it is possible that visual functions are

preserved to a substantial extent in those models. Although the *afadin* cKO mice

responded to certain visual stimuli, the extent of what visual patterns or stimuli

they can perceive remains unclear. Further detailed analyses of visual functions

are required in the future.

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Author Contributions

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C.K., M.T., and A.U. designed the project. Y.T. and J.M. generated *afadin flox*

mice. M.H generated the ONC mice. A.U. and H.T. carried out

immunohistochemical analysis. A.U. performed molecular biochemical analyses.

T.K. performed electron microscopic analysis. H.O. and S.M. carried out ERG

experiments. H.O. and M.W. performed multi-electrode recordings. K.S., T.N.,

and M.T. analyzed data of the multi-electrode recordings. A.H. carried out visual

behavioral tests. A.H. and Y.E. analyzed the data of the visual behavioral tests.

A.U., M.T., and C.K. wrote the manuscript. C.K. supervised the project.

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 FIGURE LEGENDS

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

 A. Immunostaining of the wild-type (WT) mouse retina (1M) with anti-afadin

 antibody. The afadin signals disappeared in the pre-absorbed (peptide+) sample.

 The signal observed in the peptide+ image represents the background and

 non-specific staining. Nuclei were stained with DAPI (blue).

 B. Immunostaining of the WT retinal section (1M) using anti-afadin (green)

 combined with anti-Nectin-1 (red) antibody. Necin-1 was partially co-localized

 with afadin in the OLM, OPL, and IPL.

 C. Afadin was colocalized with nectin-1, nectin-2, and nectin-3 at the OLM. The

 1M WT retinal sections were immunostained with anti-afadin (green),

 anti-nectin-1 (red, upper panels), anti-nectin-2 (red, middle panels), and

 anti-nectin-3 (red, lower panels) antibodies.

 D. Immunostaining of 1M WT retinal sections using anti-afadin (green),

 anti-SCGN (red), and anti-PKCα(white) antibodies. Afadin signals overlapped

 with SCGN and PKCα in the OPL and INL, and PKCα in the IPL.

 OS, the outer segment; IS, the inner segment; OLM, the outer limiting

 membrane; ONL, the outer nuclear layer; OPL, the outer plexiform layer; INL,

 the inner nuclear layer; IPL, the inner plexiform layer; GCL, the ganglion cell

 layer.

 Figure 2 Outer retinal lamination is severely disrupted in the afadin cKO

retina.

A. Toluidine blue staining of the cHet and cKO mouse retinal sections at 1M. The

retinal layer structure was disrupted in the cKO retina. Rod photoreceptor nuclei

(insets) were deeply stained. The red asterisk indicates a rod photoreceptor. The

arrowhead indicates a rosette-like structure. Scale bar in the inset, 2.5 µm.

B. Representative images of the cHet and cKO retinas (1M) stained with

anti-Rom1 (green) and anti-Rhodopsin (Rho, red) antibodies and

PNA-rhodamine (white). Rom1, Rhodopsin, and PNA signals were remarkably

decreased and scattered in the cKO mice. The arrowhead indicates the

rosette-like structure.

C. Representative images of the cHet and cKO retinas (1M) stained with

anti-S-opsin (green) and anti-M-opsin (white) antibodies and PNA-rhodamine

(red). Cone OSs were aberrantly located in the cKO retina. The arrowhead

indicates the rosette-like structure.

D. Retinal flat-mount immunostaining of the cHet and cKO retinas

immunostained with anti-Arr3 (green, left panels), anti-Rom1 (green, right

panels), and anti-Rhodopsin (red, right panels) antibodies. The inset, a

magnified view of the area enclosed by the dashed box, shows an area with a

relatively high density of photoreceptors in the cKO retina.

E. Immunofluorescent analysis of the cHet and cKO retinas (1M) using anti-Arr3

(cone marker, green in the left panels), anti-Otx2 (photoreceptor and BC marker,

red in the left panels), anti-RNA binding protein with multiple splicing (RBPMS,

RGC marker, green in second left panels), anti-Chx10 (BC marker, red in second

left panels), anti-Lhx2 (Müller glia marker, green in third left and third right

panels), anti-AP2α (AC marker, red in third left and second right panels), and

anti-Calbindin (horizontal cell and AC marker, white in third left and right panels)

antibodies. Arrowheads indicate RGCs and ACs near the rosette-like structure.

F-L. Distribution of each retinal cell type based on their distance from the retinal

surface in the cHet and cKO mice. The shortest distance between each cell body

and either the outer or inner retinal surface was measured using ImageJ. For

cones (**F**), rods (**G**), bipolar cells (**H**), horizontal cells (**I**), and Müller glia (**J**), the

distance was measured from the outer retinal surface. For amacrine cells (**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).

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Figure 3 Photoreceptor-BC synapses were severely impaired and

mislocalized in the afadin cKO retina.

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

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

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

views of the area enclosed by the dashed box.

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

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

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

of the area enclosed by the dashed box.

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

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

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

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

indicates the ribbon synapse between rod and rod-BC.

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

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

retina (1M).

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

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

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

n = 3, cKO; 29.1 ± 12.75, n = 4, 10 images from each mouse. Error bars, mean ±

SD. ***$p < 0.001$ by a GLMM). The number of synapses decreased to about 5 %

of the cHet in the cKO retina.

F. Quantification of the number of synapses between photoreceptor cells and

type 3 OFF-BC under 1mm^2 of the retinal surface in the cHet and cKO mice

immunostained with anti-GluR5, anti-PSD95, and anti-PKARIIβ antibodies

(cHet; 24.0 ± 2.8, n = 3, cKO; 4.1 ± 1.3, n = 3, 10 images from each mouse. Error

bars, mean ± SD. ***$p < 0.001$ by a GLMM). The number of synapses between

the photoreceptor and Type 3 OFF-BC in the cKO retina decreased to about

15% of the cHet retina.

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

A. Immunostaining of the cHet and cKO retinal sections at 1M with anti-GluR5

(green), anti-SCGN (red), and anti-PKCα antibodies. GluR5 signal was

obviously observed in the IPL of the cKO retina. Yellow and white arrowheads

indicate overlap of GluR5 and PKCα, and GluR5 and SCGN, respectively. Small

panels are magnified views of the area enclosed by the dashed box.

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

C. Immunofluorescent analysis of the cHet and cKO retinas (1M) with anti-HPC-1 (AC marker, green in the left panels), anti-Calbindin (red in the left panels), anti-Tuj1 (RGC marker, green in the right panels), and anti-RBPMS (red in the right panels) antibodies. Arrowhead indicates acellular patches.

D. The number of each retinal cell type per 100 μm width of retinal section at 1M (cHet; Rod 185.9 ± 17.5, Cone 8.4 ± 1.4, Bipolar cell 45.5 ± 5.2, Ganglion cell 7.4 ± 1.5, Horizontal cell 2.1 ± 0.4, Amacrine cell 30.1 ± 3.2, Müller glial cell 15.7 ± 2.3, n = 4, cKO; Rod 57.3 ± 17.3, Cone 3.5 ± 1.3, Bipolar cell 64.6 ± 7.8, Horizontal cell 1.9 ± 0.7, Amacrine cell 27.6 ± 6.3, Ganglion cell 5.7 ± 0.9, Müller glial cell 12.4 ± 1.5, n = 4. Error bars, mean ± SD. Horizontal cell p = 0.58, AC p = 0.52, RGC p = 0.10. ***p < 0.001, **p < 0.01, *p < 0.05 by Student's t test. In the cKO retina, rod and cone photoreceptors and Müller glial cells significantly decreased to about 30, 40, and 80% of the cHet retina, respectively, and BCs significantly increased to about 150% of the cHet retina.

E. Immunofluorescent analysis of the cHet and cKO retinas at P1, P3, P5, P8, P11, and P14 using anti-Chx10 (green), anti-Otx2 (red), and anti-Ki67 (progenitor cell marker, white) antibodies. NBL, the neuroblastic layer.

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 x 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 x 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).

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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.

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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.

F. Clustering of RGCs based on the temporal profile of RFs. Five clusters were

visualized (cHet; circle, n = 4 retinas, cKO; x, n = 4 retinas).

G. Temporal profiles from each cluster were superimposed separately (cHet;

black, cKO; magenta, mean; solid line, SD: pale line).

H. Time-to-peak latency of the temporal profile (cHet; filled bar, cKO; open bar).

Mean ± SD. (cHet; cluster 1; -113 ± 15 msec, n = 31, cluster 2; -111 ± 13 msec, n

= 36, cluster 3; -200 ± 31 msec, n = 51, cluster 4; -171 ± 19 msec, n = 37, cluster

5; -178 ± 28 msec, n = 40, cKO; cluster 3; -163 ± 21 msec, n = 10, cluster 4; -238

± 34 msec, n = 36, cluster 5; -217 ± 41 msec, n = 23).

Figure. 8 Visual function is partially retained in the *afadin* cKO mice.

A. Schematic diagram of the experimental setup for the OMR test. Drifting

square-wave gratings were displayed on monitors surrounding the mouse, and

head-tracking movements were recorded under photopic conditions.

B. Representative traces of head-tracking movements in the cHet, cKO, and

ONC mice. Blue lines indicate head movement trajectories, and red segments

denote the periods when the head movement aligned with the direction and

timing of the visual stimulus.

C. Summary of head-tracking movements in response to drifting gratings with

spatial frequencies of 0.056, 0.10, and 0.27 cycles/degree, as well as under

no-stimulus conditions, in the cHet, cKO, and ONC mice (cHet; 0.056

cycles/degree 7.30 ± 1.94, 0.10 cycles/degree 7.89 ± 2.00, 0.27 cycles/degree

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

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

A. Immunostaining of the cHet and cKO retinas at postnatal day 0 (P0) with

anti-phospho-histone H3S10 (pHH3, mitotic cell marker, green), anti-N-Cadherin

(red in left, second left, and second right panels), anti-Nectin-1 (white in left,

second left, and third right panels), anti-β-catenin (red in the right panels)

antibodies. Ectopic N-cadherin, nectin-1, and β-catenin signals were observed in

the cKO retinas, and pHH3-positive cells were localized near these ectopic

signals.

B. Quantification of pHH3-positive cells per 100 μm width of retinal sections in

the cHet and cKO retinas at P0 (cHet; 4.40 ± 0.937, n = 3, cKO; 6.69 ± 0.507, n =

4). pHH3 positive cells significantly increased in the cKO retina. Error bars,

mean ± SD. **$p < 0.01$ by Student's t-test).

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

D. Distribution of mitotic cells based on their distance from the outer retinal surface in the cHet and cKO mice at P1 (top), P3 (middle), and P5 (bottom). The shortest distance between mitotic cell body and the outer retinal surface was measured using ImageJ. Bin width was 5 μm (0–5 up to 245–250 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; P1 3.76 ± 17.5 μm, n = 300 cells from 5 mice, P3 6.73 ± 28.8 μm, n = 469 cells from 3 mice, P5 16.7 ± 44.7 μm, n = 180 cells from 4 mice, cKO; P1 48.4 ± 24.1 μm, n = 326 cells from 4 mice, P3 76.0 ± 41.9 μm, n = 902 cells from 4 mice, P5 94.0 ± 44.8 μm, n = 163 cells from 4 mice). ***$p < 0.001$ by log1p-transformation followed by a LMM.

E. Retinal sections from the cHet and cKO mice immunostained with anti-BrdU antibody (green) at P1 (upper panels), P3 (middle panels), and P5 (lower panels).

F. Distribution of progenitor cells based on their distance from the outer retinal surface in the cHet and cKO mice at P1 (top), P3 (middle), and P5 (bottom). Bin width was 5 μm (0–5 up to 245–250 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; P1 58.0 ± 22.1 μm, n = 2527 cells from 4 mice, P3 83.5 ± 31.7 μm, n = 1904 cells from 4 mice, P5 95.3 ± 32.4 μm, n = 620 cells from 5 mice, cKO; P1 62.7 ± 35.2 μm, n = 1713 cells from 4 mice, P3 96.5 ± 48.5 μm, n = 2025 cells from 3 mice, P5 90.7 ± 40.5 μm, n = 630 cells from 3 mice). P5 $p = 0.49$, ***$p < 0.001$, **$p < 0.01$ by log1p-transformation followed by a LMM.

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Figure S1 Afadin is localized to AJs in developing and mature retinas.

A. Immunostaining of the WT retinal section (1 M) with anti-afadin (green)

antibody combined with nectin-2 (red, upper and middle panels) and nectin-3

(red, lower panels) antibodies. Nectin-2 and nectin-3 were localized in the OLM.

The nectin-2 signal in the IPL was insufficient for reliable assessment of its

localization and colocalization.

B. Immunostaining with anti-afadin (green), anti-nectin-1 (red), and anti-Arr3 (a

cone marker, white) antibodies. Afadin was co-localized with nectin-1 in cone

synapses.

C. Immunostaining of the WT retinal section (P0) with anti-afadin (green) and

anti-nectin-1 (red) antibodies.

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Figure S2 AJs in the OLM are disrupted in the afadin cKO retina.

A, B. Immunostaining of *Dkk3*-Cre; *R26R-H2B-EGFP* and *R26R-H2B-EGFP*

retinas at P0 (**A**) and 1M (**B**) using an anti-GFP antibody (green). The Cre

recombinase activity was detected in the *Dkk3*-Cre; *R26R-H2B-EGFP* retina at

both stages.

C. Western blot analysis of the cHet and cKO retinas using anti-afadin (upper

panel) and anti-GAPDH (lower panel) antibodies. No significant afadin band was

detected in the cKO retina.

D. Electron microscopic analysis of the cHet and cKO retinas (1M). A few ectopic

disc structures were observed in the cKO retina. The inset, a magnified view of

the area enclosed by the dashed box, shows the ectopic disc structure at high

magnification. The arrowhead indicates the outer segment disc structure. RPE,

the retinal pigment epithelium.

E. Immunostaining of the cHet and cKO retinal sections at 1M using

anti-β-catenin (green, left panels), anti-nectin-1 (red, left panels), and

anti-N-cadherin (green, right panels). Signals of these markers were not

observed at the outer retinal surface in the cKO mice.

F. Immunostaining of the cHet and cKO retinal sections at 1M using

anti-glutamine synthetase (GS, green) antibody. Obvious GS signals were

observed in the inner side of the retina but not at the outer retinal surface in the

cKO mice.

G. Retinal sections from the cHet and cKO mice at 1M were stained with an

anti-CD31 antibody (red). No notable vascular differences were observed in the

cKO retina.

**Figure S3 The OS and photoreceptor-BC synapses are affected in the
developing afadin cKO retina.**

A. Immunostaining of the cHet and cKO retinas at 1M with anti-PKCα (green),

anti-PSD95 (red) antibodies. Small panels are magnified views of the area

enclosed by the dashed box.

B. 3D projection of confocal image immunostaining with anti-PKCα (green),

anti-PSD95 (red) antibodies.

C. Quantification of the number of contacts between photoreceptor and rod

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

with anti-GluR5, anti-PSD95, and anti-PKARIIβ antibodies (cHet; 550.5 ± 29.1, n

= 3, cKO; 65.2 ± 7.9, n = 3, 10 images from each mouse. Error bars, mean ± SD

***p < 0.001 by a LMM). The number of synapses between the rod and ON BC

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

D, E. The cHet and cKO retinas (1M) stained with anti-EAAT5 (green),

anti-SCGN (red in **D**), and anti-PKCα (white in **D**), anti-PSD95 (red in **E**)

antibodies and PNA-rhodamine (white in **E**). Glutamate transporter EAAT5

localization to ON and OFF BC axon terminals and photoreceptor axon terminals

were unaffected in the cKO retina. Small panels are magnified views of the area

enclosed by the dashed box.

F. Immunostaining of the cHet and cKO retinal sections at 1M with anti-ChAT

(green), anti-Calretinin (red), and anti-Calbindin (white) antibodies. Obvious

ChAT bands, Calretinin bands, and Calbindin bands were observed in the IPL of

the cKO mice.

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Figure S4 Retinal cell fate determination is altered during development in

the *afadin* cKO mice.

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

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

5.2, SCGN+ 37.4 ± 7.7, n = 4. Error bars, mean ± SD. *p < 0.05 by Student's

t-test). PKCα+ cells and SCGN+ cells significantly increased to 150% of the

cHet.

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

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;

P1 4.71 ± 1.75, n = 4, P3 18.7 ± 1.74, n = 4, P5 96.4 ± 6.91, n = 3). BCs

(Chx10-positive and Ki67-negative) were significantly increased in the cKO

retina at P5, but not at P1 or P3. Error bars, mean ± SD. P1 $p = 0.77$, P3 $p = 0.56$,

**$p < 0.01$ by Student's t-test.

F. Expression of rod photoreceptor cell marker genes (Rho and Nrl) in the cHet

and cKO at P1, P3, and P5. Each gene expression was normalized to GAPDH, a

housekeeping gene. The expression of Nrl and Rhodopsin significantly

decreased in the cKO retina at P5. No significant difference was observed at P1

and P3. Error bars, mean ± SD. P1: Nrl $p = 0.50$, Rho $p = 0.81$, P3: Nrl $p = 0.97$,

Rho $p = 0.81$, *$p < 0.05$. Statistical tests: P1 Rho, Mann–Whitney U test; P1 Nrl,

P3 Rho, P3 Nrl, P5 Nrl, and P5 Rho, Student's t-test.

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Figure S5 Retinal lamination is disputed in the *afadin* cKO retina during

development.

A. Immunostaining of the cHet and cKO retinal sections at P1, P3, P5, P8, P11,

and P14 with anti-HPC-1 (green), anti-RBPMS (red) antibodies. Yellow and

white arrowheads indicate acellular patches and aberrant acellular regions not

containing IPL components, respectively.

B. Immunostaining of the cHet and cKO retinal sections at P1, P3, P5, P8, P11,

and P14 with anti-AP2α (green), anti-Calbindin (red) antibodies. Yellow and

white arrowheads indicate acellular patches and aberrant acellular regions not

containing IPL components, respectively.

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.

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Figure. S6 The number of apoptotic cells decreased in the developing cKO

retina.

A. Immunostaining of the cHet and cKO retinas (1M) with anti-active caspase 3

(AC3, green) antibody. Arrowhead indicates AC3 positive cell.

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.

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A GluR5 SCGN PKCα   **B** GluR5 vGlut1 SCGN

cHet / cKO

ONL / OPL / INL / IPL / GCL

IPL

20 μm / 10 μm / 20 μm

C HPC-1 Calbindin   Tuj1 RBPMS

cHet / cKO

ONL / OPL / INL / IPL / GCL

40 μm

D

Number of cells /100 μm

□ cHet
■ cKO

*** Rod   ** Cone   ** Bipolar cell   n.s. Ganglion cell   n.s. Horizontal cell   n.s. Amacrine cell   * Müller glia

E Chx10 Otx2 Ki67   P1   P3   P5   P8   P11   P14

cHet / cKO

NBL / ONL / OPL / INL / IPL / GCL

20 μm

A

pHH3 N-Cadherin Nectin-1   pHH3 β-Catenin

cHet

NBL
GCL

50 μm   20 μm

cKO

B

○ cHet
● cKO

Number of cells /100 μm

**

cHet   cKO

C   cHet   cKO

P1   NBL   GCL   50 μm

P3   NBL   GCL   50 μm   GCL

P5   NBL   GCL   50 μm   GCL

D

cHet   cKO

Percentage (%)

***

Distance from the outer retinal surface (μm)

***

***

E   cHet   cKO

P1   NBL   GCL   50 μm

P3   NBL   GCL   50 μm   GCL

P5   NBL   GCL   50 μm   GCL

F

Percentage (%)

**

cHet   cKO

Distance from the outer retinal surface (μm)

***

n.s.

1   **Table S1 The primer sequences used for RT-qPCR**

Primer name	Sequence (5' to 3')
Opnsw Fw	CAGCATCCGCTTCAACTCCAA
Opnsw Rv	GCAGATGAGGGAAAGAGGAATGA
Opnmw Fw	CTCTGCTACCTCCAAGTGTGG
Opnmw Rv	AAGTATAGGGTCCCCAGCAGA
Nrl Fw	GCTGTGCCTTTCTGGTTCTGA
Nrl Rv	GCTCCCGCTTTATTTCGAACT
Rho Fw	GACTCTGCCAGCTTTCTTTGCT
Rho Rv	GCGTCGTCATCTCCCAGTGGA
Trpm1 Fw	ATGCGCCCATTGTCAAGTTC
Trpm1 Rv	TTCTCCAATGCAAGGCTCACA
Grm6 Fw	GTCCATCATGGTCGCCAATGT
Grm6 Rv	AGTCATAGCGTGTGGAGTCAC
Chx10 Fw	GGCGACACAGGACAATCTTTA
Chx10 Rv	TTCCGGCAGCTCCGTTTTC

A

B

AC3+ cell number

C

BrdU+ cell number

D