Morphological and functional alteration of light perception circuits in AD patients

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Liste of abbreviations

p-tau hyperphosphorylated tau

Aβ amyloid beta

NFT neurofibrillary tangles

SP senile plaques

POA preoptic area

SCN suprachiasmatic nucleus

ACC accumbens nucleus

AMY amygdala central nucleus

LGN lateral geniculate nucleus

LHB/pLHB lateral habenula/peri lateral habenula

SC superior colliculus

EWN Edinger-Westphal nucleus

LDT laterodorsal tegmental nucleus

PPT pedunculopontine tegmental nucleus

LC locus coeruleus

PBN parabrachial nucleus

VC primary visual cortex

Abstract

Disruption of sleep and circadian rhythms is one of the earliest symptoms of Alzheimer's disease

(AD). Circadian entrainment and the modulation of alertness are non-visual responses to light

driven by intrinsically photosensitive retinal ganglion cells (ipRGCs). However, little is known

about ipRGCs in the context of AD. To explore structural and functional changes of ipRGCs and

ipRGC circuits, we analyzed the retinas and brains of 13 elderly patients (9 males, 4 females, aged

69-103) ranging from normal to neuropathologically defined AD.

We performed extracellular electrophysiological recordings on freshly harvested retinas using

multi-electrode arrays. While rod and cone responses were moderately affected, we found a

decreased density of ipRGCs in donors with advanced AD pathology, which was confirmed by

immunostaining. The remaining ipRGCs displayed morphological alterations as well as abnormal

responses to light, including hyperexcitability. These changes appeared to be subtype-specific and

correlated with the advancement of the disease.

Altered ipRGCs circuits and function could contribute to the disruption of sleep and circadian

rhythms reported in AD patients. The measurement of ipRGC-dependent responses to light could

constitute a promising tool to predict or monitor pathological changes in the brain.

Introduction

Alzheimer's disease (AD) is a devastating neurodegenerative disorder and the most common cause of dementia. characterized by progressive cognitive decline, memory loss, and impaired daily functioning(*1*). AD neuropathological features, such as the presence of amyloid-beta plaques and tau protein containing tangles, play a central role in the disease progression. A definitive and unambiguous diagnosis is difficult to establish, and by the time it is obtained, the disease has usually progressed over decades, causing significant neurodegeneration. Dementia is estimated to affect more than 55 million patients, however its prevalence continues to rise with the aging of the population, and it's projected to double by 2050(*2*).Understanding the underlying disease mechanisms and identifying potential early biomarkers is urgently needed to allow timely diagnosis and intervention.

One of the earliest symptoms in AD is a disruption of sleep and circadian rhythms. Individuals with Alzheimer's often experience increasingly fragmented sleep and altered core body temperature rhythm(*3–6*). There is now abundant evidence that disruption of the daily rhythms can occur years before the emergence of classical symptoms, such as memory loss in AD(*7, 8*). Excessive daytime sleepiness and sleep behavior disorders are associated with an increased risk of developing AD and dementia suggesting that dysregulation of sleep may be important in the early pathogenesis of neurodegeneration(*9–12*). These behavioral disruptions not only compromise the quality of life of the patient, but they cause physical and psychological fatigue in the caregiver and are often the reason for institutional placement.

Regulation of non-visual responses to light including synchronizing the circadian clock to the day-night cycle and regulating sleep, alertness, mood, and the pupillary light reflex (PLR) is mainly mediated by a class of intrinsically photosensitive retinal ganglion cells (ipRGCs) that express the light-sensing protein melanopsin. Accordingly, when ipRGCs are selectively ablated or when signaling properties of melanopsin are changed, photoentrainment of the clock, acute light-modulation of sleep, and PLR to light are abolished or maladapted(*13–16*). Interestingly, a recent histological study has shown a specific decline in ipRGCs number in the retina of AD patients(*17*). Beta amyloid accumulation was also observed inside and around ipRGCs suggesting that the response properties of ipRGCs from AD patients may be altered(*17*). In mouse models of AD, ipRGC anomalies have also been reported(*18, 19*). While these observations suggest that light

perception circuits may be damaged in Alzheimer's disease, this has never been investigated
further in patients.

The aim of this study was to evaluate the alteration of light perception in AD patients
compared to normal individuals, using a comprehensive exploration in eye-brain donors.
Specifically, we investigated the histological and functional changes in ipRGC circuits in the retina
and brain of patients at different stages of Alzheimer's disease.

Results

To assess the impact of AD on light perception, we conducted an exploratory, multi-angle
study on 13 eye and brain donors, deceased at the Geneva Stroke Center (Geneva, Switzerland,
FigS1A). The demographic characteristics of the cohort are presented in **Table 1**. Age ranged from
69 to 103 years (81.31 ± 3.011). There were 9 males (mean age 79.44 ± 3.17 years; 69 to 97 years)
and 4 females (mean age 85.50 ± 7.01 y ears; 70 to 103 years, p=0.49, Mann Whitney test). The
cause of death was brain infarct and/or brain hemorrhage (**Table1**). Among the included donors,
two showed ante-mortem signs of dementia (donors 3 and 5), which together with their post-
mortem neuropathology support a diagnosis of Alzheimer's disease and are therefore referred to in
this manuscript as AD donors, as opposed to donors without known cognitive impairments (ND,
non-demented, donors). The neuropathological severity of tau lesions and amyloid deposition were
assessed in the anterior hippocampus, entorhinal, transentorhinal, and inferior temporal cortices
using the Braak staging for neurofibrillary tangles (NFT) and the Thal phases for senile plaques
(SP) (**FigS1B-D**)([20], [21]). Thal phases ranged from 0 to 4 and Braak scores ranged from I to VI.
The two AD patients scored 3 and 4 and V and VI respectively **(Table1)**. Neither Thal phases nor
Braak stages were correlated with the age of the donors (**FigS1E**).

To quantify AD hallmarks along some of the brain circuits implicated in the regulation of
visual and non-visual responses to light, we also examined tau pathology and amyloid deposition
in 13 brain regions implicated in vision (LGN, SC, Visual cortex), circadian rhythms (SCN),
alertness and sleep regulation (POA, pons nuclei), mood (AMY, LHB/pLHB, ACC), and pupillary
reflex to light (EWN) in 13 donors (**Fig1A**). Uniform brain sampling was performed by a
neuropathologist and SP and NFT were scored on a 0 to 3 scale by trained pathologists (0=none,
1=sparse, 2=moderate, 3=abundant) (**Fig1B**). We found aggregates in the majority of the regions
examined. While deposits of Amyloid beta (Aβ) were more abundant as the amyloid pathology

progresses, hyperphosphorylated tau (p-tau) burden did not follow this pattern and its presence in light perception circuits appeared uncorrelated with Braak stages (**Fig1C, FigS2A**). Aβ burden in the VC and Acc as well as in all retino-recipient nuclei at the exception of the LGN correlated with the Thal phase of the respective donor (**Fig1C**).

To evaluate the correlation between the brain and retina pathologies, we performed immunostaining of Aβ and Tau in the retina of the donors. We manually segmented the outer retina (OR: photoreceptor (PR) and outer plexiform layers (OPL)) and inner retina (IR: inner nuclear (INL), inner plexiform (IPL), and retinal ganglion cell (RGC) layers) (**Fig1D**) and determined the layer distribution of p-tau and Aβ burdens in retinal cross sections. Representative microscopic images show an accumulation of p-tau in patients that correlates with the Braak stage (**Fig1E**). Immunostaining of p-tau revealed diffuse staining in the OPL and intense staining in the INL and IPL including probable intracellular staining in the INL. Aβ deposits were identified in the inner retina, in particular in the innermost RGC and retinal nerve fiber layers. Aβ42 burden (**Fig1E, FigS2B**) also increases, although not significantly, with the Thal phase of the donor. Finally, we also measured the thickness of the segmented IR and OR as a proxy of neurodegeneration. The thickness of both layers decreased with AD and was negatively correlated with the increase of Thal phases and Braak stages (**Fig1F**).

The presence in the retina of both Aβ and Tau aggregates and degeneration (**Fig1**) suggested a possible alteration of the light information sent to the brain. To evaluate the function of the rods, cones, and ipRGCs, we recorded responses to light from a donor's retina preparation using multi-electrode array (MEA)(*22, 23*). Briefly, the eyes of the donors were collected within 2 hour postmortem (**Table1**), immediately preserved in oxygenated Ames 'medium placed in a light-proof container, transported to the lab where circular (3mm diameter) patches of retina were dissected under dim red light, placed on MEA, and allowed to recover in darkness for at least 1hour.

First, to evaluate photoreceptors' function, we recorded micro-electroretinogram (mERG)(*24*) on retinal patches from 5 donors (**Table 1**), including the two AD donors in response to sequences of 10ms light flashes of increasing irradiance (470nm, 6.10^9 to 2.10^{13} photons/cm^2/s) (**FigS3**). mERG did not display major differences in response profiles between AD and age-matched control donors (**FigS3A**). A-waves were detected in all donors at all light irradiances used while b-waves were only clearly detected at the highest light stimulation but in all donors too(*23*).

While the amplitude of the a-wave from the AD donors was the smallest, it was not different from

that of the other donors. Similarly, the b wave amplitude and the latencies of both waves did not

differ between donors (**FigS3B, C**).

Then, to evaluate ipRGCs'function, we performed extracellular electrophysiological

recordings of freshly harvested retina from 6 donors including the two AD donors (**Table 1**). For

this purpose, the recording medium was supplemented with synaptic blockers to block rod and

cone input to RGCs and capture ipRGCs intrinsic, melanopsin-initiated signal. In response to 30-

s pulses of monochromatic blue light (470 nm), photoresponsive cells were found in retinas from

each of the six donors (**Fig2A**). However, the number of ipRGCs varied from one donor to the

other. For each retina sample, we identified photoresponsive cells at an average density ranging

from <1 to ~8 cells/mm^2 (**Fig2B**). The number of light responsive cells was negatively correlated

with the Braak stage but not with the Thal phase, the age of the donor, or the delay of collection

of the eye (**Fig2B and FigS4A**). Grouped analysis reveals a two-time decrease in ipRGCs density

in retina pieces from AD donors (**FigS4B**).

To confirm the ipRGC loss in AD patients, we performed immunostaining against

melanopsin as previously described(*25*) and we counted melanopsin positive (mel+) cells in 3mm

diameter retina patches (**Fig2C**). We found about two time-decreased density of mel+ in the retina

of AD donors (**FigS4B**). When examined individually, mel+ density correlated with the patient

Braak stage and ipRGCs density but not with the Thal phase (**Fig2D, E and FigS4C**). Melanopsin

staining revealed that not only the number but also the morphology of mel+ cells is affected by the

disease. In each retina sample, a blind investigator used semi-automated tracing to reconstruct the

dendritic arborization of 4 mel+ cells per retina sample (**Fig2F**). Scholl analysis revealed that while

the extend radius of the ipRGCs dendritic tree was not different between donors, its number of

intersections midway between the soma and the extremities was decreased in AD donors (**Fig2G,**

FigS4D). Accordingly, both total neurites length and the number of terminal branches were lower

in the AD patients (**Fig2H-I, FigS4D**). These results suggest a decrease in dendritic tree

complexity in AD. Interestingly, AD ipRGCs had larger soma (**Fig2J, FigS4D**). Altogether, these

major morphological alterations strongly suggest that the function of the surviving ipRGCs is also

altered.

To evaluate potential functional alteration in the remaining ipRGCs from AD donors, we

characterized their response properties. We stimulated the retinas for 30 s at increasing irradiances

of monochromatic blue light (470 nm, 2.10^{12} to 1.10^{15} photons/cm^2/s). Overall, responses to light
were slow, sustained, and outlasted the stimulus by several seconds after the light was switched
off, the sensitivity was low as previously described (*22*) (**Fig2A**). Response properties followed
similar dynamics with increasing stimulation irradiance; latency decreased while the duration and
number of spikes of the response increased (**Fig3A**). Interestingly, responses to the lowest intensity
of light were correlated with the Thal and Braak scores reflecting an increased sensitivity or
excitability with the progression of the disease (**Fig3B,C and FigS4E**). Conversely, responses to
the highest intensity of light were inversely correlated with the Thal and Braak scores suggesting
a failure to sustain responses at advanced stages of AD (**Fig3B,C**).

Several morphological and functional ipRGC subtypes have been described in the human
retina (*22, 25–28*). To assess whether the alterations reported above reflect alteration and/or
degeneration of specific subtype(s), we performed principal components analysis (PCA) on the
response parameters of each ipRGCs recorded. Data separated in at least three main clusters
(**Fig3D**) whose response to light properties and time course closely resemble those from previously
described ipRGC subtypes (**Fig3D,E**) (*22*). The three subtypes were found in all donors (**Fig3E**)
but their distribution was changed in the AD donors. Type1 accounted for half of the ipRGCs in
AD donors while type 2 and 3 ipRGCs fraction was strongly decreased (**Fig3F**). When examined
individually, type1 fraction correlates with the Braak stage of the donor (**FigS4F**). This result
suggests that type 1 cells may be more resilient than type 2 and 3. As type1 ipRGCs are the subtype
that is most sensitive to light, the increased excitability observed could be due to this change of
distribution. To test this hypothesis, we examined the sensitivity to light of each subtype. We found
that type1 ipRGC sensitivity was increased in AD donors (**Fig3G**). Overall, all ipRGC subtypes
are affected and the increased excitability observed in AD may rely on both a shift in subtypes
distribution toward M1, the most sensitive ipRGCs, and an increase in their sensitivity.

Discussion

In this prospective study, we examined several circuits of light perception in a group of
elderly adult donors at various stages of AD pathological changes. We employed a comprehensive
suite of measurements to compose the first picture of how AD may impact ipRGCs abundance,
function, and brain targets and found that ipRGC circuits are injured both morphologically and
functionally in the brain and the retina.

In the retina, our results are in agreement with the presence and distribution of AD
hallmarks and their correlation with the neuropathological changes observed in the brain, reported
early and in extensive postmortem retinal histological studies ((*29–33*)(for review see(*34*)). We
also confirmed the loss of ipRGCs observed in AD patients(*17*). We extended these findings by
measuring *ex vivo* the function of the different photoreceptors. Both outer and inner retina
photoreceptors seem to malfunction with AD but, while rods/cones response alteration may be
limited to small decrease in amplitude at specific light intensity levels, ipRGCs injury is massive.

AD has been associated with various forms of visual dysfunction including abnormal visual
fields, reduced color and contrast vision which can be, at least in part, attributed to retinal and
optic nerve damage(*35–38*). Retinal abnormalities can be identified with electroretinogram (ERG)
in AD (*39–43*). Here, while we found a decrease in IR thickness, the mERG a and b waves did not
show significant differences between AD and ND patients in response to our stimulation protocol.
Because IR photoreceptor function declines rapidly with hypoxia (*23*), some differences between
donors may have been masked by the postmortem functional decline in all donors.

ipRGCs are the main contributors and in some cases the exclusive conduits of light
regulation to non-visual light responses (*13, 14*). Accordingly, they project to more than a dozen
of brain regions implicated in visual and non-visual responses to light including the nuclei
investigated here, SCN, POA, HB/pHb, AMY, LGN, and SC(*44, 45*). We found twice less ipRGCs
in AD donors, and the surviving one displayed abnormal morphologies and responses to light that
may precede degeneration. In particular, the increase in soma size resembles the phenomenon
called neuronal ballooning that is thought to result from deficient axonal trafficking and
subsequent accumulation of cargos at the initial segment (*46–48*). Then, the increased excitability
is reminiscent of cerebral neurons in which an increased excitability is observable at the earlier
stages of the disease before there is pronounced cell loss and hypoactivity(*49–51*). The initial loss
of ipRGCs seems biased toward types 2 and 3, with type 1 ipRGC appear more resilient. M1
ipRGCs, the morphological equivalent of functional type1 ipRGCs, had already been shown to
demonstrate increased survival to mechanical and metabolic stresses(*52*). However, the
morphological and functional alterations observed in AD donor remaining ipRGCs suggest that
type1 ipRGCs probably also degenerate. Overall, these alterations are consistent with and could
contribute to the alterations of ipRGC-regulated functions, such as circadian rhythm,
sleep/alertness, and mood, experienced by AD patients. These results also suggest that some

ipRGC-dependent responses to light may be impacted earlier than others and could constitute stage

specific markers.

As PLR is thought to rely on both M1 and M2 ipRGCs(*53*), our results predict that pupillary

responses to light should also be affected. Pupillometry is a scalable, cost effective, non-invasive

assay that a few groups have started to explore as a biomarker of AD in pre-symptomatic cases

(*54, 55*) and early AD cases (*56, 57*). However, the results have been mixed. This may reflect the

diversity of light stimuli and protocols (*58, 59*) and emphasizes the need to determine the spectral,

intensity, and/or temporal domains where ipRGCs malfunction and PLR is impacted. For example,

our results suggest that injured ipRGCs will fail to sustain response to long stimulations at high

irradiance.

The retina, an extension of the central nervous system, offers a unique opportunity to study

AD pathology outside of the brain. Over the past decades, accumulating evidence has shed light

on the presence of AD-related histological changes in the retina, suggesting the potential of

different retinal imaging modalities as a non-invasive method to detect and study the disease (for

review see(*34*)). Our results suggest that the retinal functions (light perception and transmission to

the brain) are also altered and add ipRGC-dependent responses to light as potential biomarkers of

AD. In AD donors, surviving ipRGCs seems to recapitulate several phenotypes observed in brain

neurons: hyperexcitability and ballooning. This, together with the fact that they are more accessible

and have clear cellular and behavioral readouts, makes ipRGC a promising model to study the

mechanisms of neuronal degeneration directly in human. Elucidating the mechanisms underlying

ipRGC vulnerability to AD could lead to the discovery of therapeutic targets to improve patients'

quality of life and potentially improve the overall disease trajectory.

Limitations

Our study has a number of limitations. First, our sample size is small because of the

challenges presented to obtain and collect postmortem eye and brain tissues from well-

characterized participants meeting the criteria of inclusion. However, we partially compensated

for this limitation by using in-depth objective physiopathological and functional measurements

from each donor. Second, our cohort has a greater proportion of male subjects, leading to potential

sex-related bias. Our study population was composed primarily of self-reported White and thus

may not be representative of other populations. Third, because our focus was on physiology, the

pieces of retina were collected in the same central region irrespective of the orientation. While this
strategy may have caused some variability and may leave apart more extreme phenotypes in the
peripheral retina(*17*, *29*, *60*, *61*), ipRGC density and distribution is relatively homogenous in this
region(*25*). Finally, our study is primarily correlational and, therefore, caution must be taken
before making cause-and-effect inferences.

Despite these limitations, the present study gives a glimpse of pathophysiological
mechanisms of neurodegeneration at work in humans.

Data availability
Code and data used to generate the figures described in this paper will be made available via
Zenodo (Zenodo.org) at the time of publication, with a DOI to be included here.

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Contributions
NS, DM, and LSM conceptualized the study. NS and LSM designed it. NS coordinated tissue
collections and harvests. NS, EK, YT, TAZ, KLE, and LSM performed the experiments. JH
provided the melanopsin antibodies. EK, JH, JAL, EC, GT, SK, and AA provided scientific
expertise and critical reading of the manuscript. LSM analyzed the data. LSM wrote the first draft,
all authors edited and approved the manuscript.

Competing interests
DM is an advisory board member for Optomed, Finland. All other authors declare that they have
no competing interest.

Methods

Donors

Patients were prospectively included in the study after obtention of informed consent from the

family or healthcare proxys. At a second phase, following the confirmation of death, the patient

was promptly transported to the hospital morgue where enucleation was immediately performed,

day or night, by a trained physician (NS). Only patients in whom enucleation was performed within

2 hours after death were included in the study.

Eye sampling procedure

Bilateral enucleation was performed in all eligible patients following a standard surgical

procedure. After insertion of a lid speculum, a conjunctival peritomy was performed using a #15

blade, followed by dissection of Tenon's capsule. The extraocular muscles were identified,

isolated, and transected, facilitating identification and isolation of the optic nerve from the

surrounding structures. Using microscissors, the optic nerve was then sharply transsected, as far

as possible from its bulbar insertion to the ocular globe, closest possible to the orbital apex. After

being freed from its orbital attachments, the eyeballs were removed from the orbital cavity. A 360-

degree concentric limbal incision allowed the removal of cornea, iris, and the lens. Following this

procedure, eyeballs were transported to the laboratory in a custom-designed container under

continuous oxygenation as described previously (*22*).

Exclusion criteria were the presence of significant posterior pole ocular pathology such as age-

related macular degeneration, glaucoma, or other retinopathies or optic neuropathies.

Brain histopathology

The tissue blocks were embedded in paraffin and 12 μm-thick adjacent sections were stained with

hematoxylin-eosin or with antibodies against tau (AT8; Pierce Biotechnology, Rockford, IL, USA,

1:1,000, that recognizes the Ser202 and Thr205 epitopes) and Aβ (4G8, Signet Laboratories,

Dedham, MA, USA, 1:2,000, that recognizes amino acids 18–22 of the Aβ protein). The

neuropathological assessment of Alzheimer disease pathological changes was performed

following NIA-AA guidelines(*62*). The severity of tau--immunoreactive lesions, amyloid

deposition and neuritic plaques was determined using the Braak stages, Thal phases, and CERAD

classification(*20*, *21*, *63*).

Retina sampling

The vitreous was removed from the eyecup under dim red light. The macula was identified

according to its pigmentation and the pattern of blood vessels in the retina. Circular patches (3 mm

diameter) of retina were punched out in an annular region comprised within ~2 to 10 mm from the

fovea (except in the blind spot) and from both left and right eyes. Given the number of challenges,

including punching out adequate patches, vitreous removal, patches adhesion on the MEA, length

of dark adaptation periods and protocol, sampling was done indifferently of the quadrant

orientation. While orientation difference may be a source of variability, it seemed a reasonable

compromise as this region is relatively homogenous regarding ipRGC density.

Samples used for histological experiments were taken in the same region at the end of the recording

session and immediately drop in 4% paraformaldehyde (PFA).

Retina histopathology

Retinas were fixed in 4% PFA at 4°C for 24 hours, followed by paraffin embedding. Sagittal-

oriented sections of the posterior segment were sliced at 10μm thickness. These paraffin sections

were treated with 70% formic acid (Roth, 5355.1) for 10 minutes at room temperature, followed

by antigen retrieval in citrate buffer (pH 6.0, Merck, 100244) at 100 °C for 1 hour. Sections were

then incubated in a blocking solution containing 5% normal goat serum (Sigma, D9663) and 0.05%

Triton X-100 (Sigma, T9284) in Tris-buffered saline (TBS; Sigma, 94158) at room temperature

for 1 hour. Then, sections were incubated overnight at 4 °C with primary antibodies: anti-Amyloid

Beta (ABeta x-42, clone 12F4, mouse, Merck, 05-831-I, 1:500) or anti-Tau (AT8, p-tau

Ser202/Thr205, mouse, ThermoFisher, MN1020, 1:250), diluted in 2% blocking buffer. Sections

were washed with TBS containing 0.05% Triton X-100, followed by incubation with secondary

antibodies (anti-mouse Alexa Fluor® 488, ThermoFisher, A11029, 1:1000) and counterstaining

of nuclei with 4′,6-diamidino-2-phenylindole (DAPI; Sigma, D9542) for 3 hours at room

temperature. Finally, sections were mounted with Inova Mounting Medium (Ruwag Handels AG,

508005). Images were acquired at 40x magnification using a scanning laser microscope (Nikon,

Eclipse 80i). Three to six distinct sections imaged per donor. Controls were processed using

identical protocols while omitting primary antibody to assess non-specific labelling. The outer and

inner retinas, including respectively the photoreceptor and outer plexiform layers and the inner

nuclear layer, inner plexiform layer and RGCs layer, were manually segmented using Fiji software

(NIH). The average intensity of the aggregates fluorescent signal and the thickness of these regions

were measured.

Melanopsin Immunofluorescence

Immunohistochemistry was performed as described in detail previously(*25, 64*). The flat-mount

pieces of retinas were incubated in 5% normal donkey serum for 20 min to avoid nonspecific

staining. The flat-mount was incubated for 3–4 days with melanopsin primary antibody at 4°C.

The patches were then washed and incubated with secondary antibodies for 2 to 3hours at room

temperature. Control experiments were performed by eliminating the primary antibody, which

abolished all specific staining. Experimenters blinded for the donor ID and pathology were asked

to count all cells in the pieces of retina and to trace four representative cells in each patches using

semi-automated tracing (pluggin SNT in Fiji).

Multi-electrode array recordings

Patches of retina were dark incubated in oxygenated Ames solution until recording (at least

60min). Retinal patches were mounted on MEA (256 electrodes arranged in a 16×16 square grid,

Multichannel Systems, Reutlingen, Germany), ganglion cell side contacting the electrodes, and

perfused with oxygenated Ames' medium (Sigma-Aldrich, A1420-10X1L) at 34°C. The

MEA2100 amplifier was enclosed in a lightproof box located in a room kept in dark with light

emitted only from the controller computer screen with red filter on the screen. ipRGC recordings

were done in presence of synaptic blockers (CNQX 20µM, L-AP4 20µM, and D-AP5 50µM) and

9-cis retinaldehyde. Stock solutions of 9-cis-Retinal (R5754; Sigma–Aldrich) in ethanol (100 mM)

were prepared in darkness and stored at −80°C. On the day of the experiment, an aliquot was

thawed and diluted to a final concentration of 100 µM in Ames' medium.

Data collection, and analysis were performed as previously described(*16, 22, 65*). Briefly, the

signal was acquired at 10 kHz and filtered using a low pass (<500Hz) and a high pass (>300Hz)

filters for the mERGs and for the RGC multiunit spike responses respectively. Thresholds for spike

detection were set at 5 times the standard deviation of the noise on each channel. Spike cutouts,

consisting of 1 msec preceding and 2 msec after a suprathreshold event, along were sorted into

trains of a single cell after recording using Offline Sorter (Plexon, Denton, TX). Data analysis and

display were performed using Neuroexplorer (Plexon) and custom software written in MATLAB.

Light stimulations

To elicit the mERGs, we used sequences of five 10ms monochromatic light pulses (470nm) separated by 10s of darkness at four increasing irradiances (6.10^9, 7.10^{10}, 2.10^{12}, and 2.10^{13} photons/cm^2/s). To generate ipRGC responses, we applied a protocol consisting in a sequence of four 30s light stimulations of increasing irradiance of light (7.10^{10}, 2.10^{12}, 2.10^{13}, 1.10^{14}, and 1.10^{15} photons/cm^2/s at 470nm,) separated by 2min of darkness. All light stimuli were generated by high brightness LED, LuxeonStar 5 located ~20cm at the vertical above the MEA. Stimuli were controlled by a homemade C+ program and synchronized to the electrophysiological recordings. The light intensities were measured with a powermeter (PM120D, Thor labs). The ENLIGHT Checklist form will be filled out at the time of publication to show the characteristics of our study and the light used (Supplementary Table XX), as recommended (*66*).

ipRGC response parameters analysis

The response discharge rate was calculated as the average of the discharge rate measured during the stimulation (30s) minus the average discharge rate measure during the 30s prior to the stimulation (baseline). The response latency was defined as the delay between the start of the stimulation and the response crossing a threshold set at baseline +2*SD. The response was considered over after firing rate dropped below the baseline level for more than 3 seconds. The dose-response curves (DRCs) were obtained by plotting the response (Y, discharge rate) as a function of the irradiance of the stimulation (X, photon/cm^2/s). DRCs were fitted by a Michaelis-Menten function of the form; Y=Rmax*X/(EC50+X) where Rmax is the maximum response and EC50 is the irradiance required to drive a 50 percent response. A few inhibitory responses were manually discarded.

ipRGCs clustering

To distinguish functional types of ipRGCs in the mouse, we performed principal component analysis (PCA) on sensitivity, duration, and latency of the response as previously described(*22, 67, 68*). PCA allowed reducing the dimensionality of the data set; the first two principal components accounted for almost 65.5% of the variance (PC1=47.1% and PC2=18.4%). Thus, we next performed k-means clustering analysis on the first two principal components. This model

produced clusters that corresponded to the initial analysis. To determine the number of ipRGC

clusters, the first two principal components were fitted with Gaussian mixture models (GMMs).

We produced separate GMMs for 1 to 5 ipRGCs clusters and then used the Akaike information

criterion (AIC) to determine which GMM reported best the ipRGCs distribution. When we

compared the 1 to 5 cluster models, the models with the smallest AIC value (best model) were the

3, 4, and 5 clusters model. We represented the 3 clusters model as the minimum number of

functional subtypes.

Statistical analysis

All data are expressed as means ± SEM unless stated otherwise. GraphPad Prism was used for all

statistical analyses. Statistical tests used are stated in the figure's legends. Differences between

groups were considered statistically significant for $p < 0.05$. Bivariate correlation analysis between

histological (ipRGC/mRGC density, morphological parameters, Thal/Braak scores, severity of

brain pathology) and clinical data (age, collection delay) was performed using Pearson coefficient.

Matlab (2019a) was used for k-means and hierarchical clustering, PCA, GMM, and AIC analysis

(Statistics and Machine Learning Toolbox).

454 **Table1: Donor's demographic and pathology.** Amyloid and tau pathologies were evaluated

455 with Thal phases and Braak stages respectively.

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	Age (years)	Sex	Delay to Brain collection (hours)	Delay to Eyes collection (hours)	Cause of death	Thal βAmyloid (A)	Braak NFT (B)	CERAD (C)	CAA	Brain histology	Retina histology	Retina function	Figure symbols
D1	71	M	6	2	Brain infarct, hemorrhage	0 (0)	II (1)	0	+	X	x	x	●
D2	80	F	25	2	Brain infarct	2 (1)	III (2)	2	-	X	x	x	■
D3	77	M	13	2	Brain hemorrhage	4 (3)	VI (3)	3	+	X	x	x	▲
D4	85	M	34	2	Brain hemorrhage	2 (3)	I (1)	0	+	X			
D5	89	F	114	1.5	Brain infarct	3 (2)	V (3)	2	+	X	x	x	▼
D6	97	M	27	9	Brain infarct	4 (3)	II (1)	0	+	X			
D7	103	F	3	1	Brain infarct	1 (1)	II (1)	0	-	X			
D8	74	M	11	0.5	Brain infarct	1 (1)	II (1)	0	-	X	x	x	○
D9	74	M	46	2	Brain infarct	0 (0)	I (1)	0	-	X			
D10	70	F	82	82	Brain hemorrhage	0 (0)	II (1)	0	-	X			
D11	77	M	41	2	Brain hemorrhage	2(1)	II (1)	0	-	X	x		◆
D12	69	M	20	2	Brain infarct	0 (0)	I (1)	0	-	X			
D13	91	M	18	2	Brain hemorrhage	2 (1)	0 (0)	0	-	X	x	x	□

Figure 1: AD hallmarks are present along the circuit of light perception. (**A**) Several brain nuclei contributing to the main light-regulated function (vision, entrainment of the circadian clock, PLR, sleep/alertness, mood) were collected. (**B**) Hypothalamic region including POA from 4 donors exhibiting increasing levels of Aβ deposits (scored 1 to 4). *Left, middle, and right columns,* H&E staining, Aβ staining, and related enlargement respectively (*dotted lines*, fornix; scale bars, 1mm). (**C**) Aβ deposits scores for each brain nuclei in each donor ranked as a function of their

Thal phase. (**D**) Fluorescence micrographs of retinal cross-sections from donor 7 (ND, left column)

and donor 3 (AD, right column) with Thal phases 2 and 5 and Braak scores II and VI respectively.

On the upper left panel, dotted lines materialized the definition of the Outer and Inner retinas (OR

and IR). (*Upper row*) Tissues were immunolabeled for p-tau (AT8, green), and DAPI+-nuclei

(blue;). Scale bar: 50 μm. (*Lower row*) Greyscale micrographs the same individuals

immunolabeled with p-tau (Scale bar: 50 μm). Scatter plots of the thickness of the donors' IR (**E**)

and OR (**F**) as a function of their Braak score (n= 3 to 9 sections measured, Pearson correlation

coefficient and p value). Scatter plots of the Aβ (**G**) and p-tau (**H**) burdens measured in each donors

as a function of the Thal phase and Braak scores of the donors respectively (n=3 to 5 for Aβ and

n=3 for Tau per donor, Pearson correlation coefficient and p value).

Figure 2: ipRGCs decline in AD. (A) Representative ipRGC spike trains to a 30-s light pulses of

increasing irradiance (~1010 to 1015 photons/cm2/s, 470 nm) from AD and age-matched ND

donors. (**B**) ipRGCs density in pieces of retina from each donor as a function of their Braak score

(n=1 to 5 retina pieces per donor). (**C**, *Upper row*) Microphotographs of patches of retina

immunostained against melanopsin (scale bar 500μm, *upper right corner*, Braak stage and Thal

phase of the donors). (*Lower row*) Enlargements from the first (*) and fourth (#) patch

microphotographs (scale bar 100μm). The mel+ cells density in each donor correlates with their

Braak score (**D**, n=1 to 2 retina pieces per donor) and with the density of ipRGCs reported in the

same donors (**E**). (**F**) Examples of dendritic shrinkage, mel+ cells from AD (Donor3, *purple*) and

age-matched (Donor1, *grey*) donors (scale bar 500μm). Scholl analysis (**G**), neurites length (**H**),

number of terminal branches (**I**), and soma diameter (**J**) from AD (n=8 cells per donor, 2 donors,

purple) and age-matched (n= 4 to 8 cells per donor, 4 donors, *grey*) donors were different.

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Figure 3: ipRGC function is altered in AD. (**A**) Average ipRGCs'response latency, duration,

number of spikes, and normalized response in each donor (n=28, 15, 18, 19, 77, and 13 for donors

1, 2, 3, 5, 8, and 13 respectively, where n is the number of cells) to a sequence of 30s light pulses

of increasing irradiance (~10^{10} to 10^{15} photons/cm^2/s, 470 nm). (**B**) Responses to low and high

light stimulations as a function of donors' Thal phase ($2x10^{12}$ and $1.5x10^{15}$ photons/cm^2/s,

respectively).   (**D**) Average ipRGC sensitivity to light as a function of Thal phase. (**E**)

Representative responses from types 1, 2, and 3 ipRGCs to increasing irradiance light pulses (30

s, 470 nm, irradiances (irr) 1, 2, 3, and 4: $2x10^{12}$, $2x10^{13}$, $1.5x10^{14}$, and $1.5x10^{15}$ photons/cm^2/s,

respectively). Blue bars indicate light pulses. (**F**) Principal components of the ipRGCs' response

parameters (sensitivity, latency, and duration; n =133 and n=37, four control and two AD donors

(*purple*) respectively). (**G**) Distribution of ipRGC subtypes in ND and AD donors. (**H**) ipRGC

sensitivity by subtype in NC and AD donors (n=14, 77, and 45, type1, 2, and 3 cells and n=7, 12,

and 18 type1, 2, and 3 cells respectively).

Supplemental Figure 1: Donor's inclusion and neuropathology. (**A**) Donor inclusion flow
chart. 13 brain-eye donors were included in our protocol in 2021 and 2022 of which two had to be
excluded; one due to AMD diagnostic and a second following an error in eye conditioning. 7 pair
of eyes were collected within 2h and could be recorded on MEA. (**B**) NFT and Aβ deposition in
the hippocampal formation and the temporal neocortex (CA1 sector of the hippocampus; EC =
entorhinal cortex; TC = temporal cortex). Left column: anti-tau (AT8); right column: anti amyloid

(4G8) immunostaining. High magnification of neurofibrillary tangle (**C**) and cored senile plaque

(**D**) (C: AT8 tau immunohistochemistry; D: 4G8 amyloid immunohistochemistry. (**E**) Neither the

Thal phase nor the Braak stage were correlated with the age of the donors.

Supplemental Figure 2: AD hallmarks in the brain and in the retina. (**A**) Heatmap of the tau
deposits scores for each brain nuclei in each donors ranked as a function of their Braak score. (**B**)
Fluorescence micrographs of retinal cross-sections from donors 7 (left column) and 3 (right
column) with Thal phases 2 and 5 and Braak stages II and VI respectively. On the upper left panel,
dotted lines materialized the definition of the Outer and Inner retinas (OR and IR). (*Upper row*)
Tissues were immunolabeled for 12F4+-Aβ42 (green), and DAPI+-nuclei (blue;). Scale bar: 50
μm. (*Lower row*) Greyscale micrographs the same individuals immunolabeled with 12F4+-Aβ42.
(**C**) Correlation of outer and inner retina thickness with the Thal phase of the donors.

Supplemental Figure 3: ipRGC abundancy and morphology decline in AD. (**A**) ipRGC density do not correlate with either the Thal phase, age of the donor, or delay of eye collection. (**B**) Comparison of density of mel+ cells and ipRGCs measured in patches of retina from ND and AD donors. (**C**) Correlation of each mel+ cell measurement (neurites length, number of nodes, and soma diameter) with Thal phases and Braak stages of the donors (*upper* and *lower* rows respectively).

Supplemental Figure 4: *ex vivo* micro Electroretinogram (mERG). (**A**) Representative mERG

traces from AD and age-matched donor in response to 10ms light pulses of increasing irradiance

$(6.10^9, 7.10^{10}, 2.10^{12},$ and 2.10^{13} photons/cm^2/s, 470nm) (average of n=247 and n=244 mERG

respectively). Dose response curves reporting the average amplitude (**B**) and latency (**C**) of the a

and b waves in donors 2, 3, 5, 8, and 13 in response to 10ms light pulses of increasing irradiance

$(6.10^9, 7.10^{10}, 2.10^{12},$ and 2.10^{13} photons/cm^2/s, 470nm) (n=2, 3, 3, 2, and 3 retina pieces in donors

2, 3, 5, 8, and 13 respectively). (**D**) ipRGC subtypes distribution as a function of Braak stage.

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