



**Purpose**

- Diabetic retinopathy (DR) is a microvascular disease associated with diabetes mellitus (DM) complications

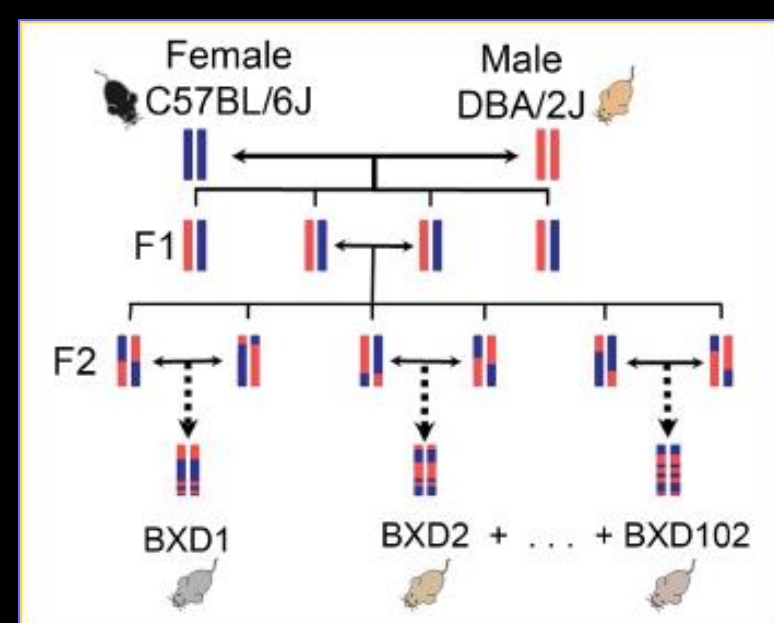
- Alterations in blood flow, vasodilation, & blood sugar can negatively affect the vasculature & neurons of the retina, resulting in blindness

- Animal models elicit a better understanding of the development, genetics, & mechanisms of multiple retinal degenerative diseases seen in humans

- Despite the complex etiology making DR difficult to model in animals, a spontaneous model of DR--the BXD79 mouse--has been uncovered

- This model displays naturally occurring phenotypes of hyperglycemia, obesity, vascular leakage, cataracts, and retinal degeneration

- This ongoing study seeks to investigate the etiology and pathology of DR in the BXD79 model.



- The BXD79 strain was found from an AMD genetic study; however, observed to be different in phenotype

**Methods**

- Single nucleotide polymorphism analysis (SNPs), reveals 9 predicted deleterious SNPs linked to many AMD & DR genes. More specifically the genes involved in angiogenesis, *Vegf*, *Kdr*, & *Flt1*.

- BXD79 & C57B/6J (wildtype) mice underwent functional examinations--electroretinograms (ERG), optokinetic nystagmography (OKN), optical coherence tomography (OCT), & fluorescein angiography (FA)--every 3 months beginning at 6 months of age until 18 months with tissues being observed using fluorescent immunohistochemistry (fIHC).

**Conclusions**

- Excessive sugar along with other DR-related phenotypes loosen tight junctions of the inner retinal vascular endothelial leading to vascular leakage & edema, respectively

- Functional test results (Fig. A & B) suggest a progressive reduction in retinal function, & disruption of retinal & vascular integrity (Fig. C & D) from 9 to 18 months

- fIHC reveal minimal mislocalization of inner/outer segment proteins, thus appearing normal (Fig. E). However, a loss of ganglion cell bodies & dendrites (Fig. F), thinning of the inner plexiform layer (Fig. H), and upregulation of KDR (fig. I & J)

- The combination of genetic mutations contributing to neovascularization & ganglion cell layer death can likely be linked to *Vegfa* and VEGF receptors *Flt1* and *Kdr*.

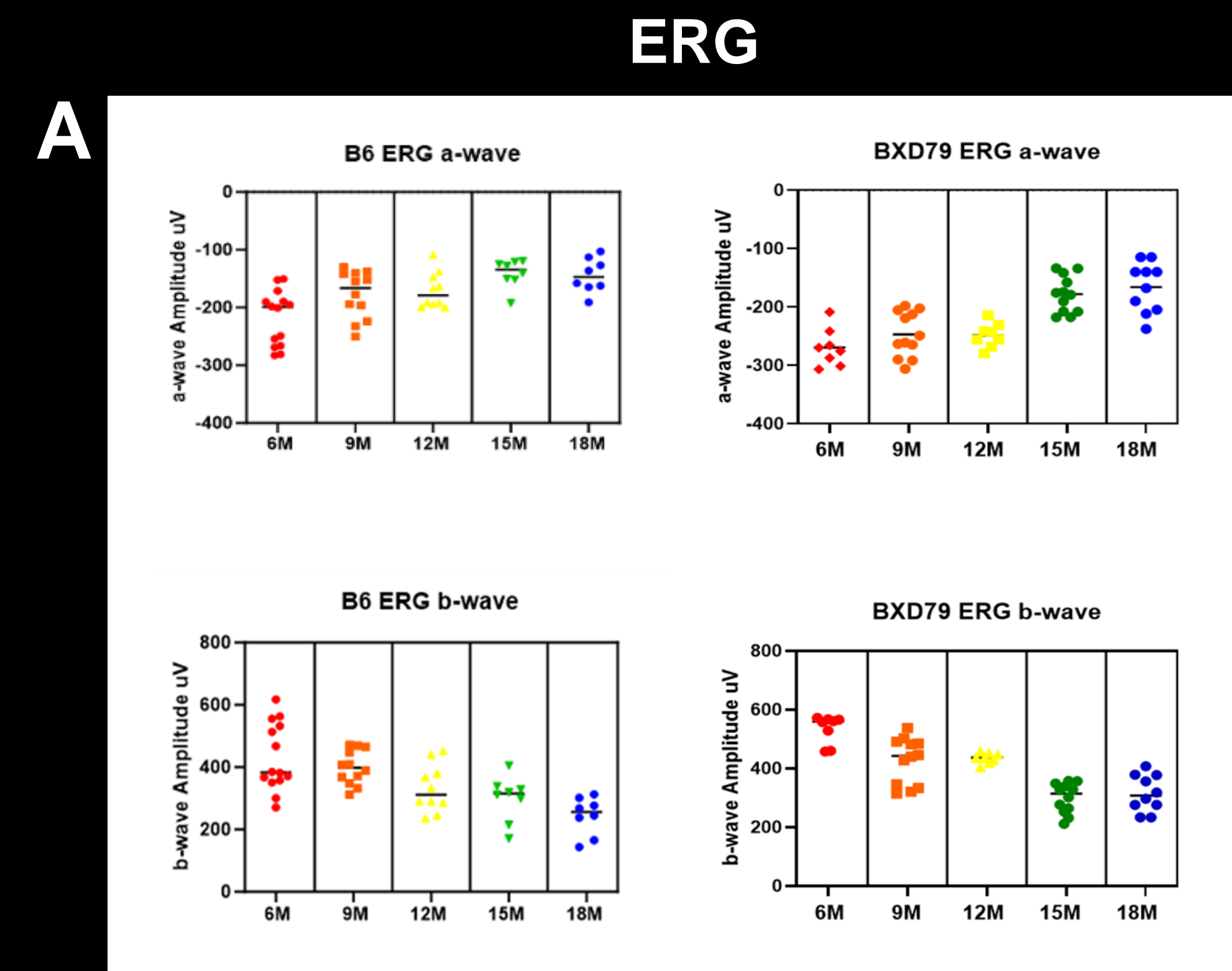


Fig A. Electroretinograms (ERGs) of wildtype (WT) & BXD79 from ages 6-18 months accessing retinal function. Data reveals dysfunction starting around 9 months in BXD79.

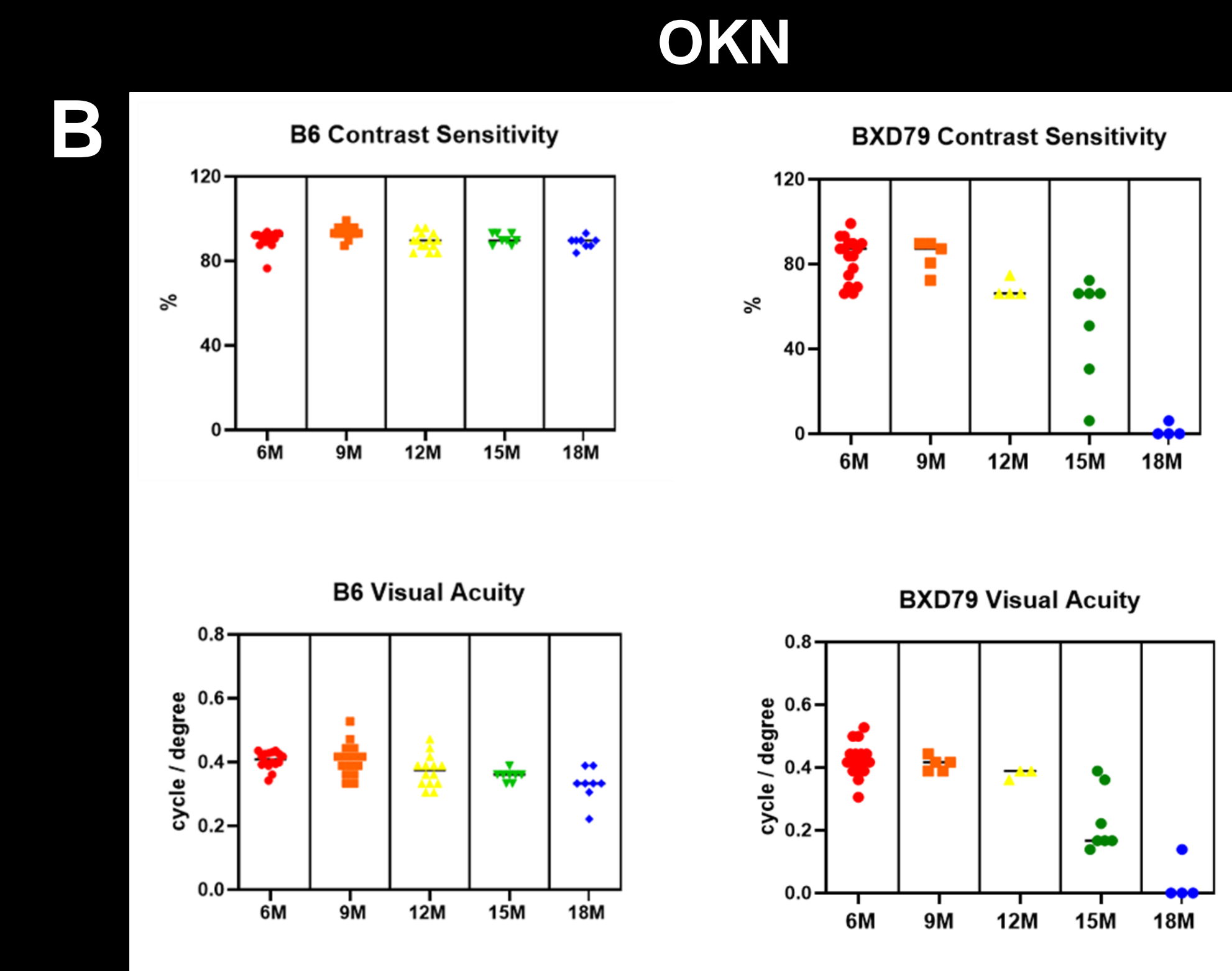


Fig B. Optokinetic nystagmography (OKN) of WT & BXD79 ages 6-18 months accessing visual acuity & contrast sensitivity. Data reveals loss of function starting at 9 months, persisting to 18 months in BXD79.

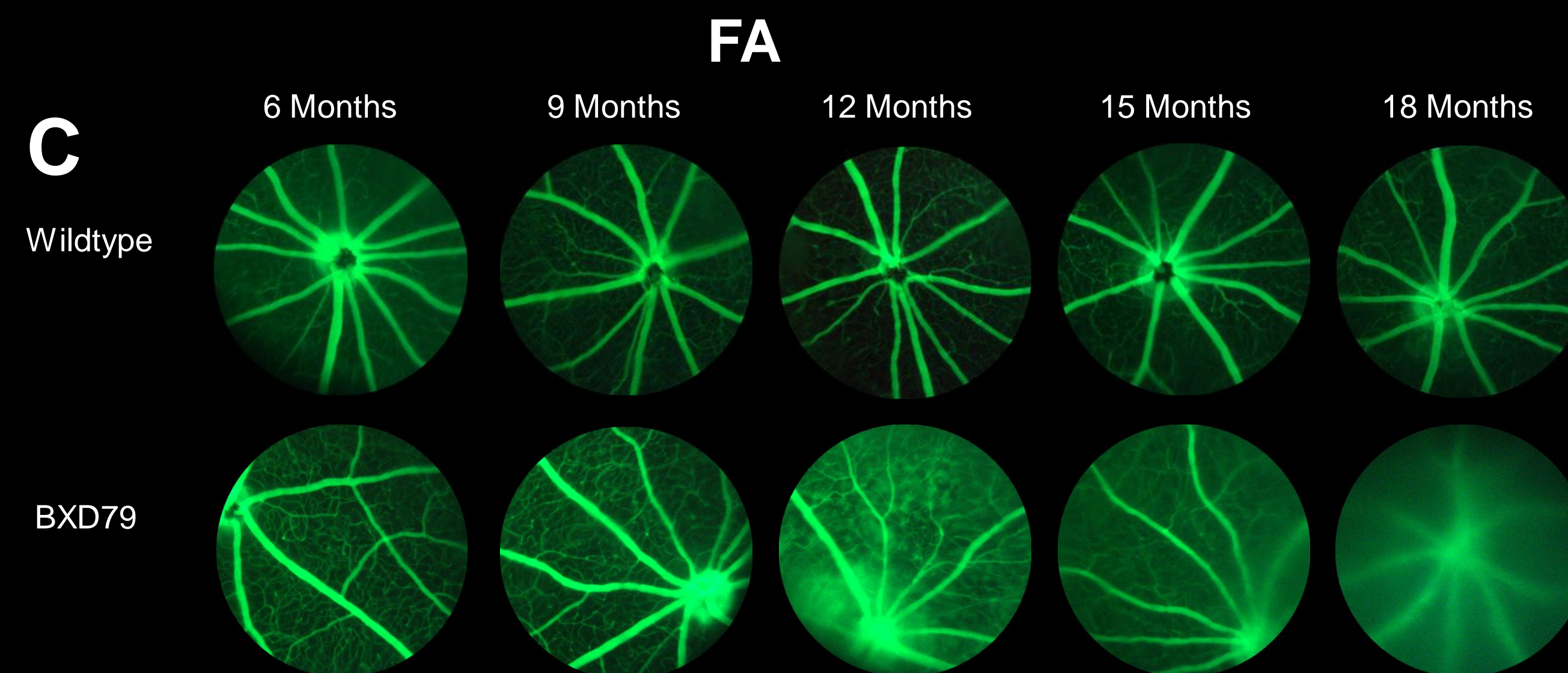


Fig C-D. C) Fluorescent angiography (FA) of WT & BXD79 (6-18 months) examining the vasculature of the retina. Vascular abnormalities & leakage are seen in BXD79 around 9 months. D) Optical coherence tomography (OCT) of WT & BXD79 (6-18 months) accessing the retinal layers reveals retinal thinning around 12 months.

## fIHC

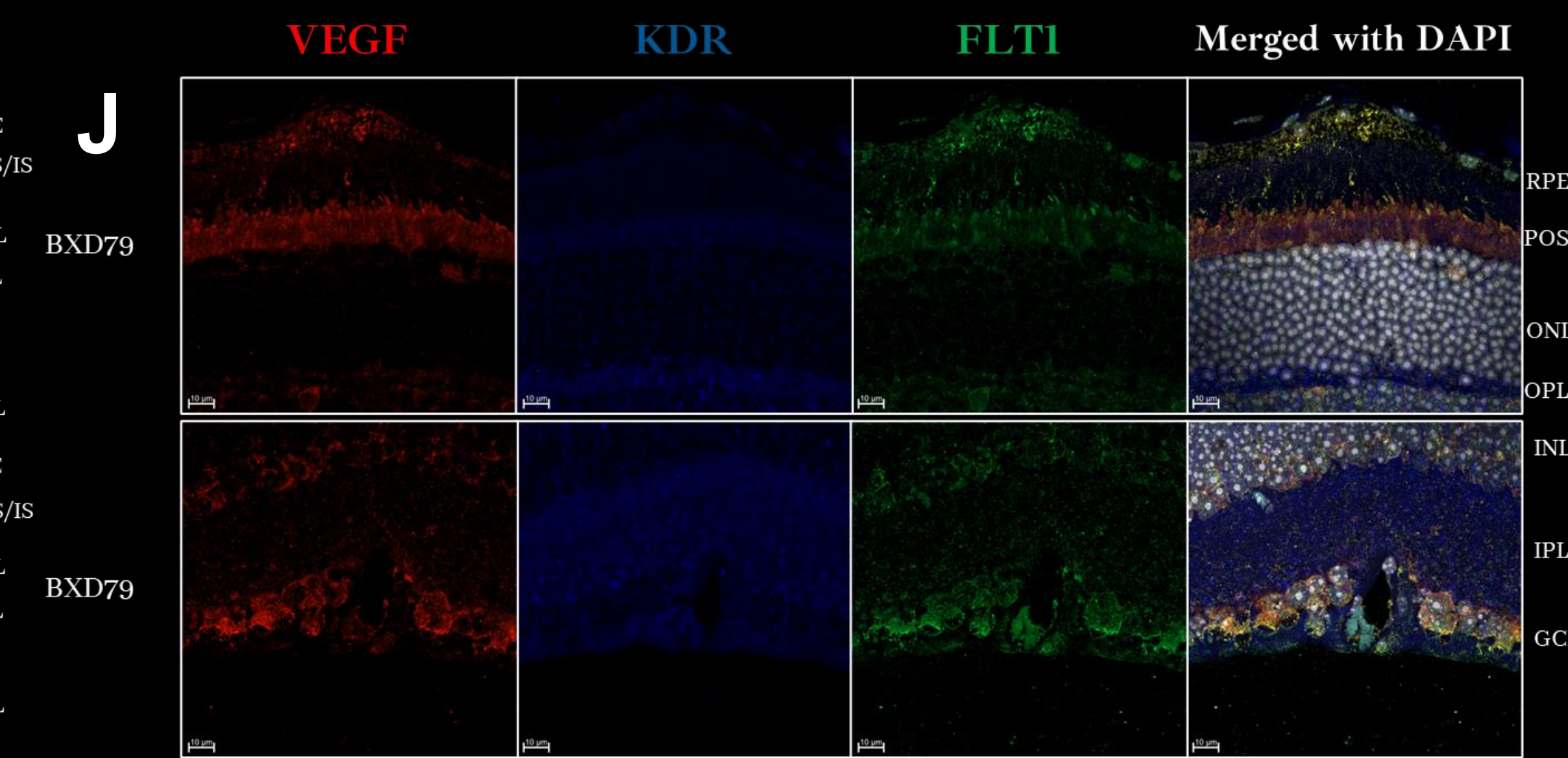
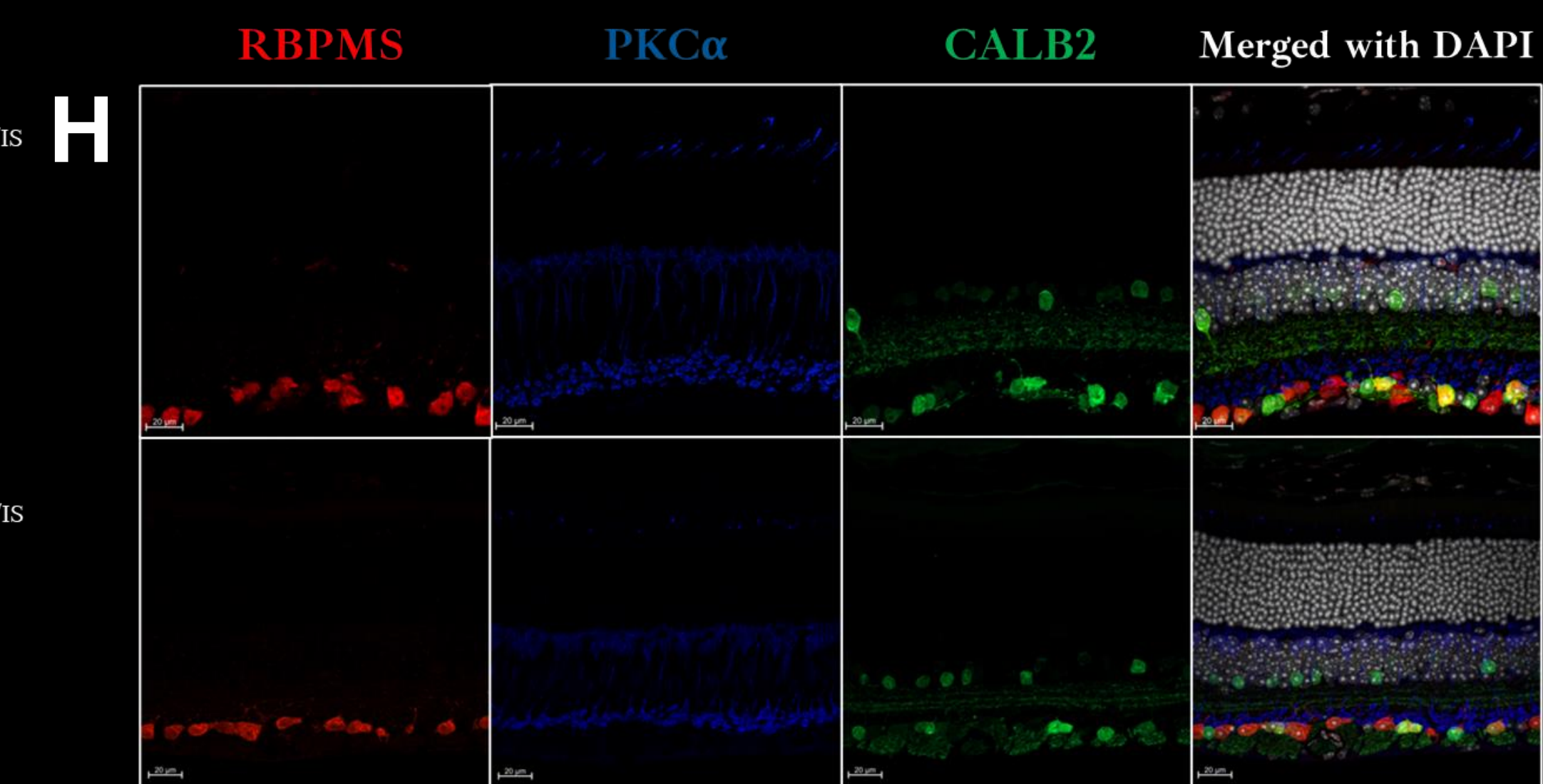
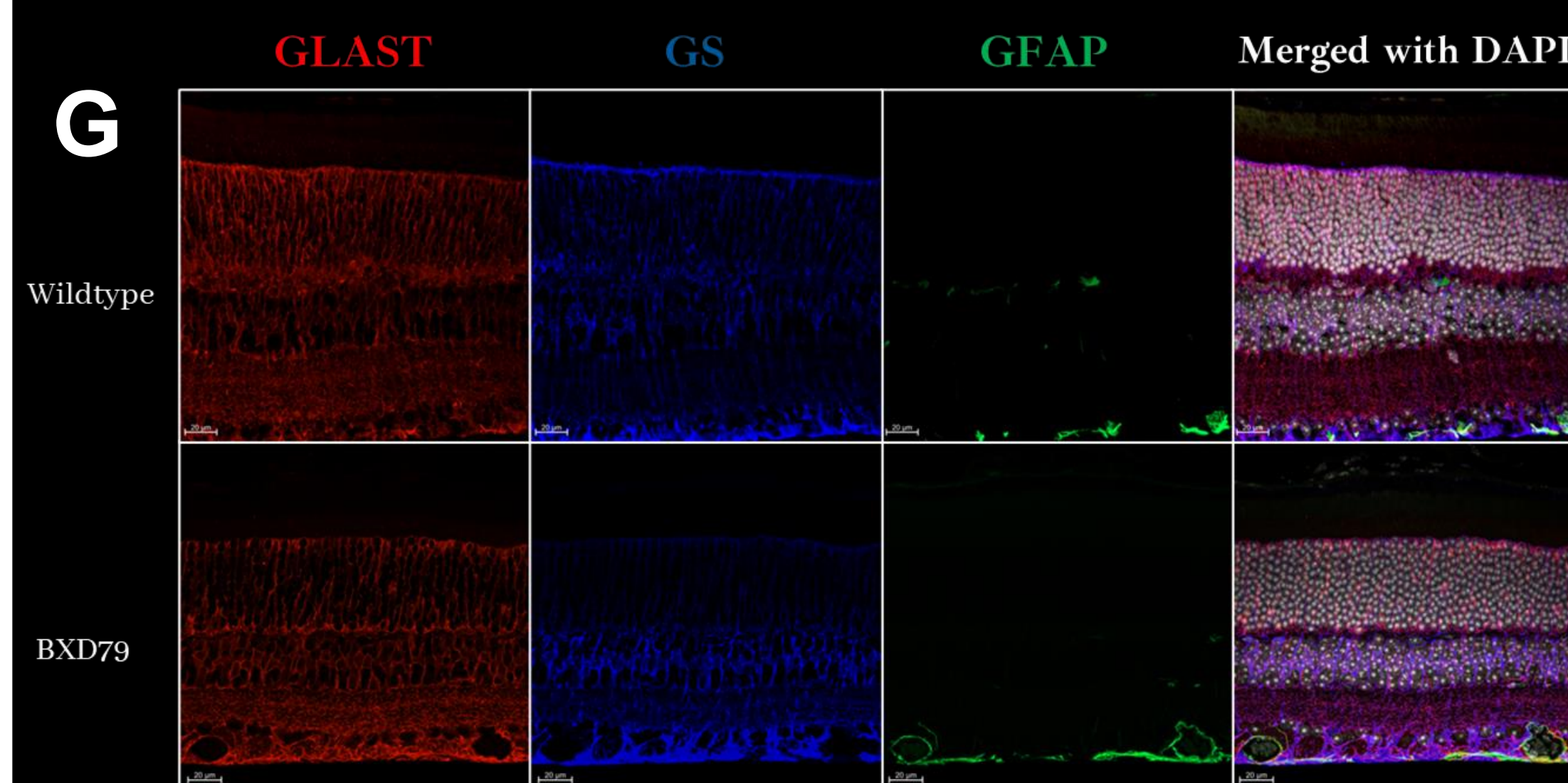
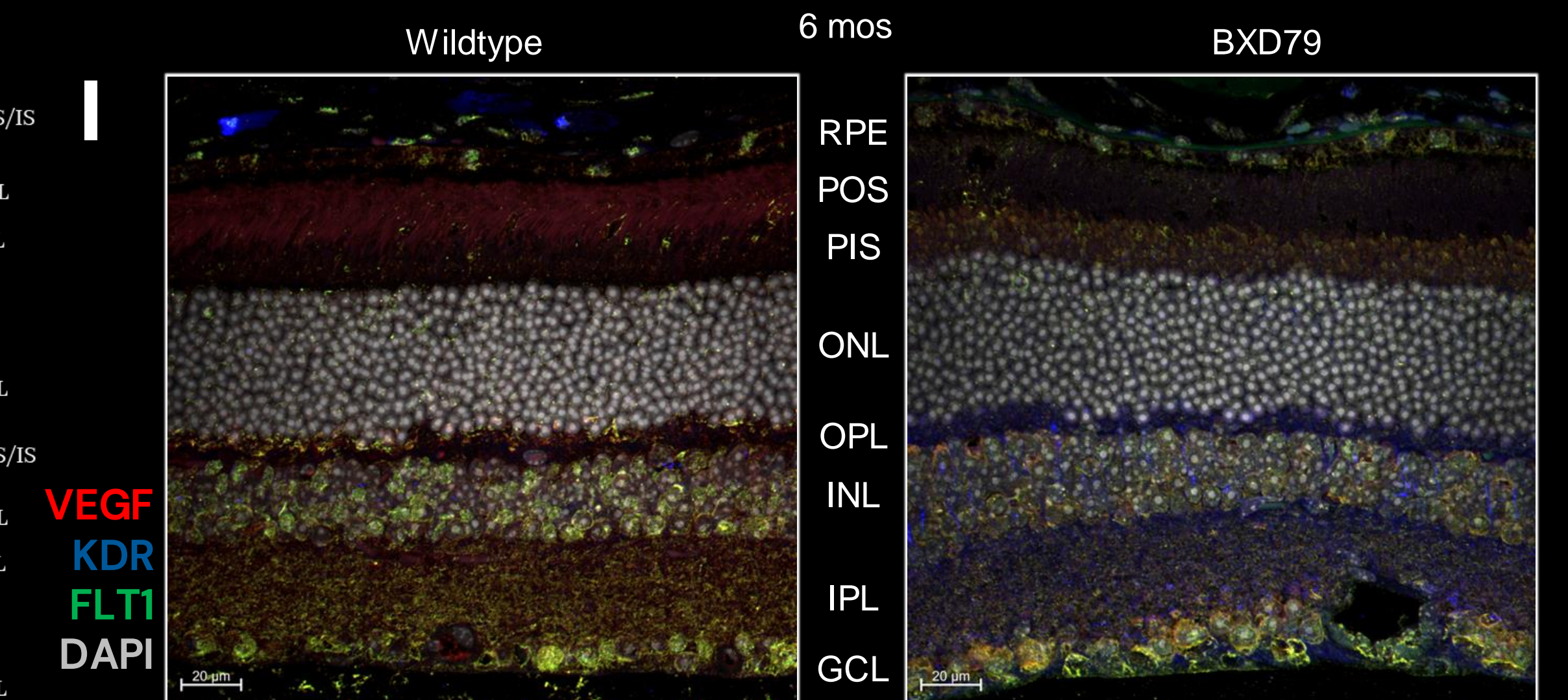
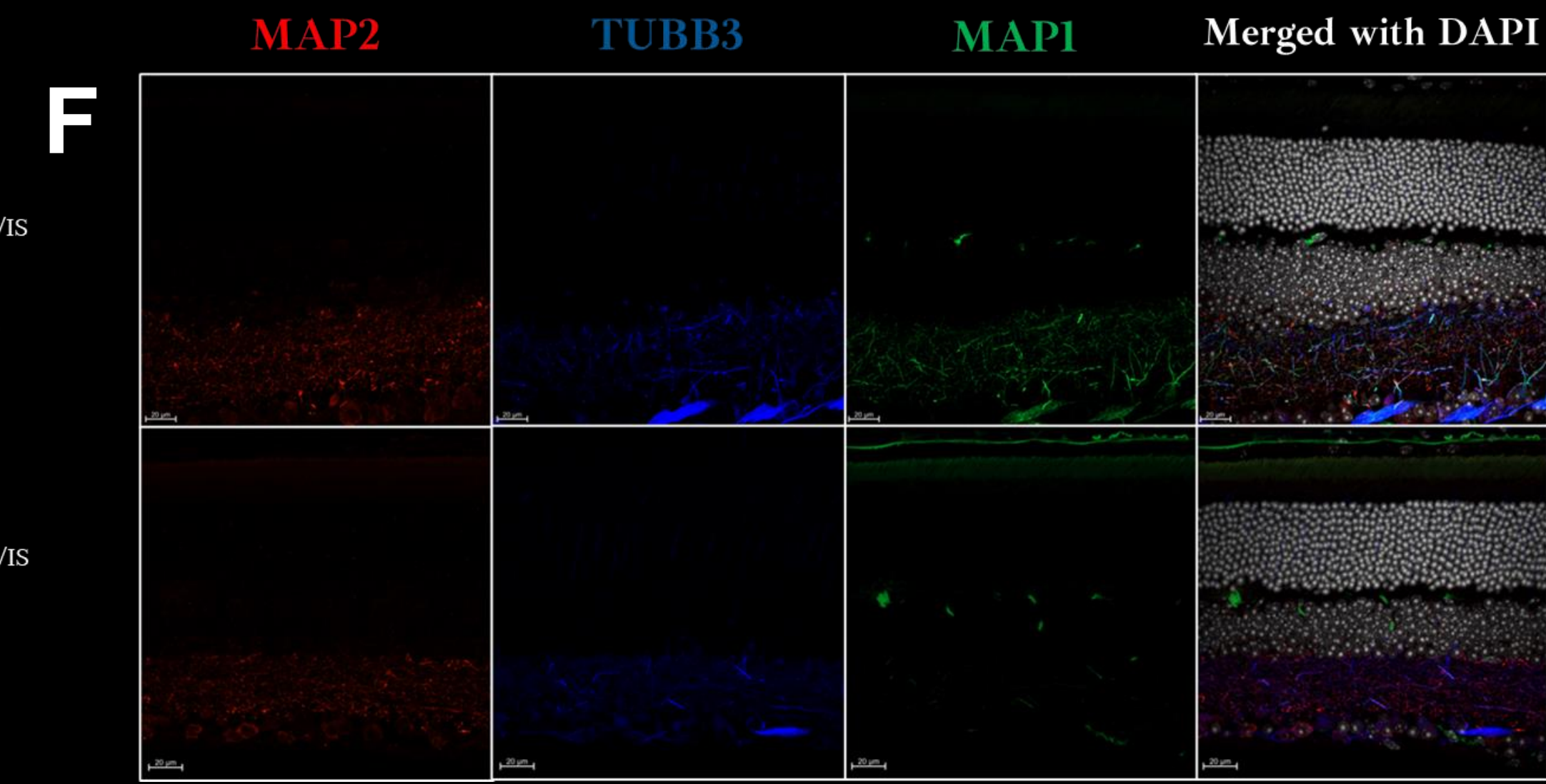
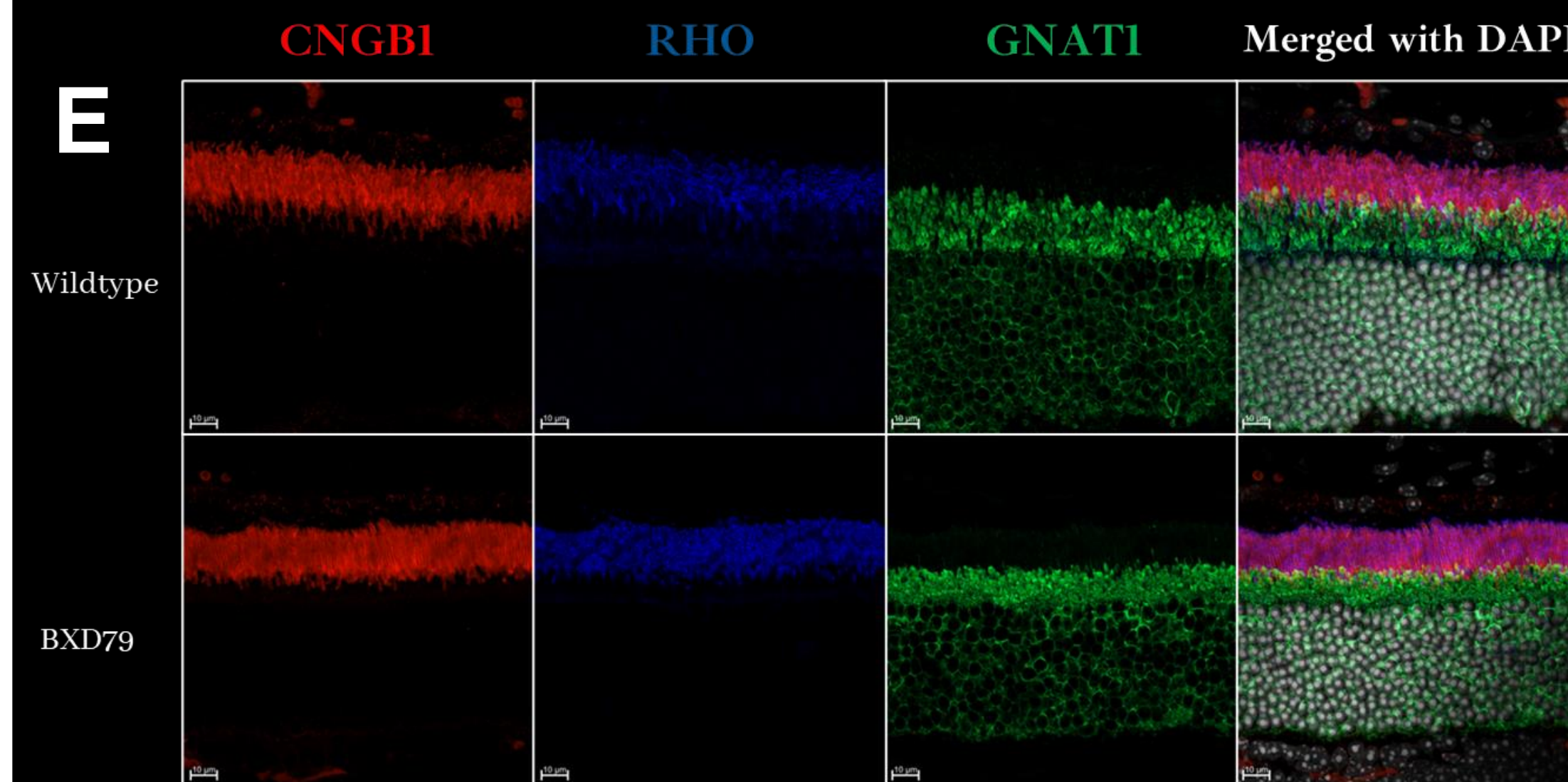


Fig. E-J. Fluorescent immunohistochemistry (fIHC) of WT & BXD79 at 18 months examining the localization of proteins in the retina. Data depicts fairly normal photoreceptors (E); however, ganglion cell body & dendrite loss (F), increased retinal stress with the upregulation of GFAP (G), retinal thinning (H), & upregulation of KDR (I-J) can also be observed. Scale bars = 20  $\mu$ m.

## Results/Discussion

## Purpose

## Methods

## Conclusions

## Support

