

# RETINAL AND STRIATAL PROFILING OF THE R6/1 MOUSE MODEL OF HUNTINGTON'S DISEASE

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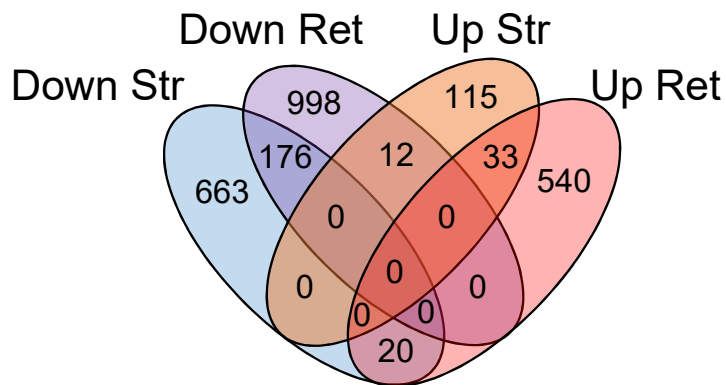
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## INTRODUCTION

Increasing evidence indicates compromised structural and functional integrity of visual pathways in Huntington's disease (HD) mouse models and patients. The retina can be an accessible window of the central nervous system through non-invasive methods to monitor the health status and response to potential treatments in patients. However, no correlation has been firmly established between the molecular events occurring in the retina and in the most affected brain area by the HTT polyQ expansion, the striatum. The aim of this study is to explore the molecular and cellular alterations linked to the disease in the retina of the R6/1 strain in order to define its suitability for biomarker screening.

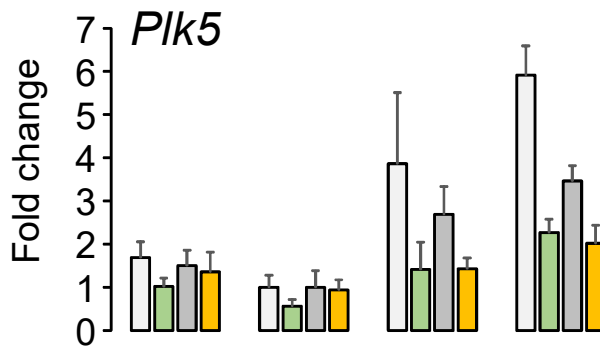
# 1. R6/1 RETINAS SHOW EXTENSIVE TRANSCRIPTIONAL DYSREGULATION



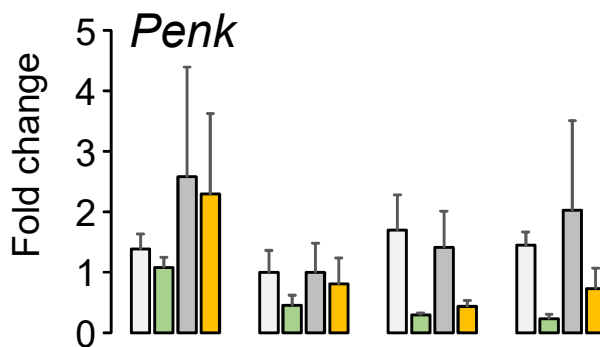
| Tissue   | Change | Top Tissue Expression / Biological Process term |
|----------|--------|---|
| Striatum | Down   | UP_TISSUE: Brain                                |
|          |        | GO_BP: Intracellular signal transduction        |
|          | Up     | UP_TISSUE: Brain                                |
|          |        | GO_BP: Cell adhesion-related                    |
| Retina   | Down   | UP_TISSUE: Retina                               |
|          |        | GO_BP: Visual perception                        |
|          | Up     | UP_TISSUE: Bone marrow                          |
|          |        | GO_BP: Immune system process                    |

RNA-seq analysis:

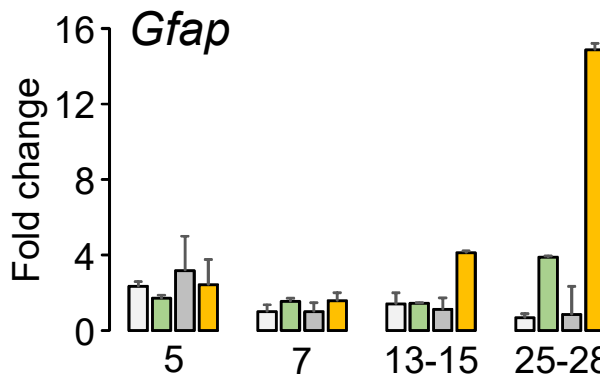
- Differentially expressed genes (DEG) between 15 weeks-old R6/1 and wt littermates in striatum (Str) and retina (Ret), adj  $P$ -value  $< 0.05$
- Top terms from DAVID Gene Enrichment tool



| Tissue | Variable | F, d.f. | $P$ -value |
|--------|----------|---------|------------|
| Striat | Genotype | 7.91, 1 | 0.007      |
|        | Age      | 5.14, 3 | 0.004      |
|        | Interact | 1.51, 3 | 0.226      |
| Retina | Genotype | 4.96, 1 | 0.032      |
|        | Age      | 5.89, 3 | 0.002      |
|        | Interact | 1.58, 3 | 0.218      |

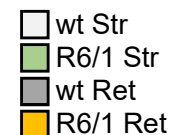


| Tissue | Variable | F, d.f.  | $P$ -value         |
|--------|----------|----------|--------------------|
| Striat | Genotype | 14.48, 1 | $4 \times 10^{-4}$ |
|        | Age      | 1.09, 3  | 0.362              |
|        | Interact | 1.52, 3  | 0.223              |
| Retina | Genotype | 0.92, 1  | 0.344              |
|        | Age      | 1.20, 3  | 0.324              |
|        | Interact | 0.13, 3  | 0.944              |



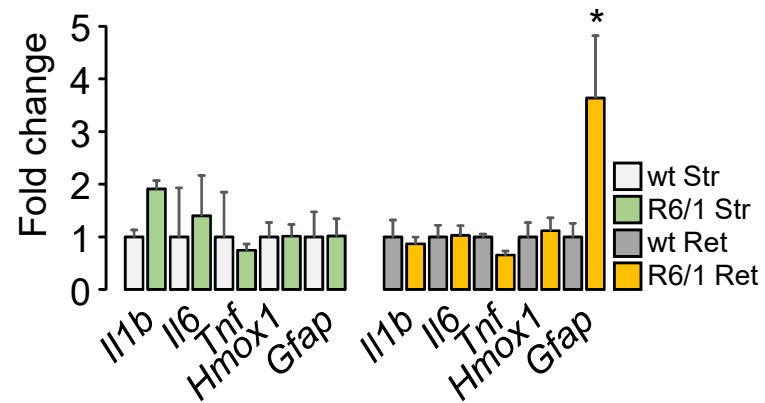
| Tissue | Variable | F, d.f.  | $P$ -value         |
|--------|----------|----------|--------------------|
| Striat | Genotype | 0.57, 1  | 0.455              |
|        | Age      | 0.73, 3  | 0.538              |
|        | Interact | 1.70, 3  | 0.183              |
| Retina | Genotype | 9.37, 1  | 0.004              |
|        | Age      | 8.57, 3  | $2 \times 10^{-3}$ |
|        | Interact | 10.88, 3 | $3 \times 10^{-5}$ |

Time-course analysis of selected DEG in RT-qPCR assays using independent samples. Two-way ANOVA results in the tables: **significant  $P$  values  $< 0.05$**

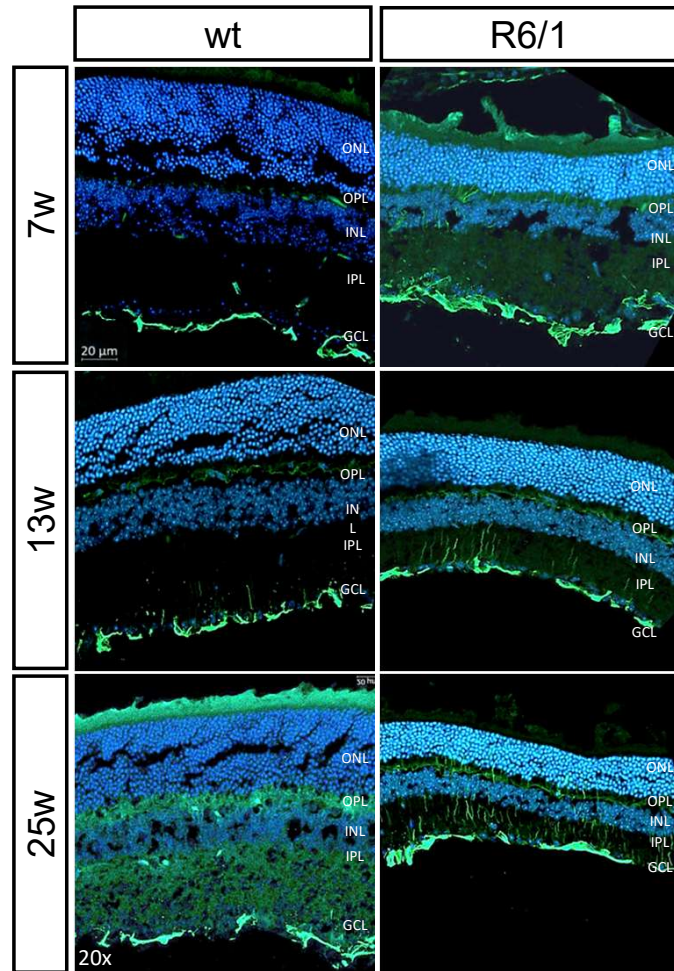
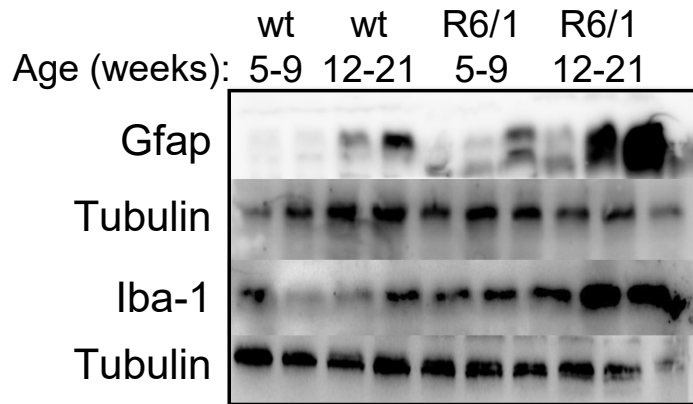


## 2. GFAP AND IBA-1 ARE INDUCED IN R6/1 RETINAS

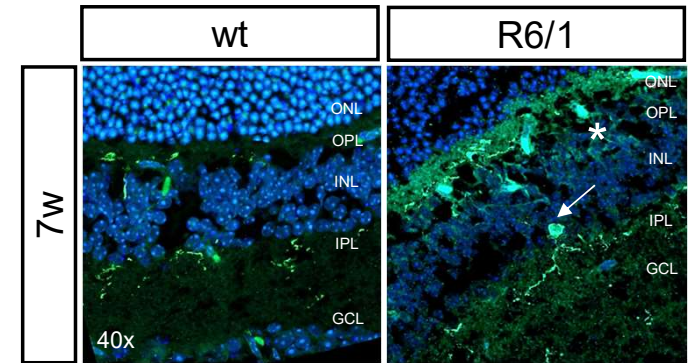
RT-qPCR assays in 13-15 weeks-old R6/1 and wt littermates: \*,  $P < 0.05$ , Mann-Whitney U-test. In agreement with RNA-seq results



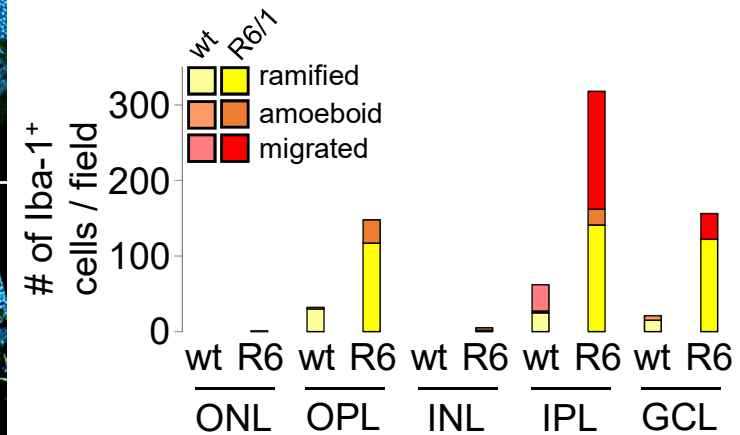
Western blotting assays in R6/1 and wt retinas



Gfap / DAPI: increasing staining of GCL and OPL, later comprising INL and IPL

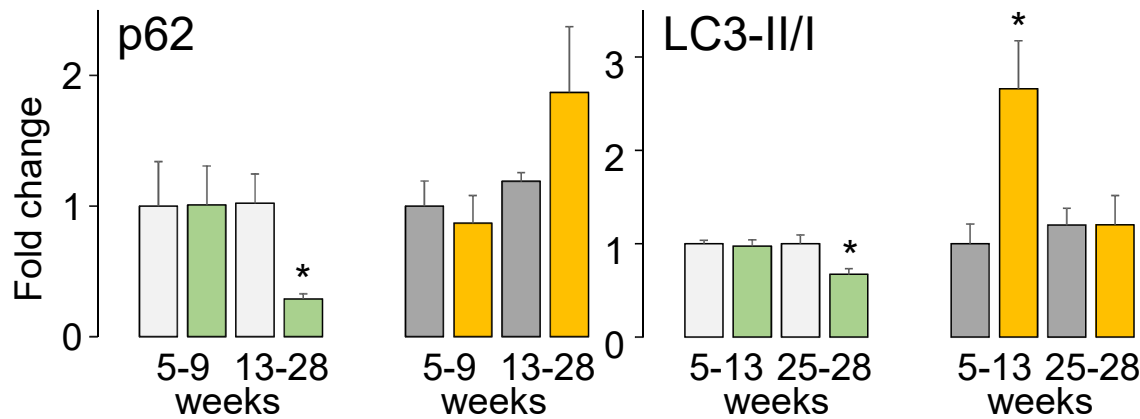
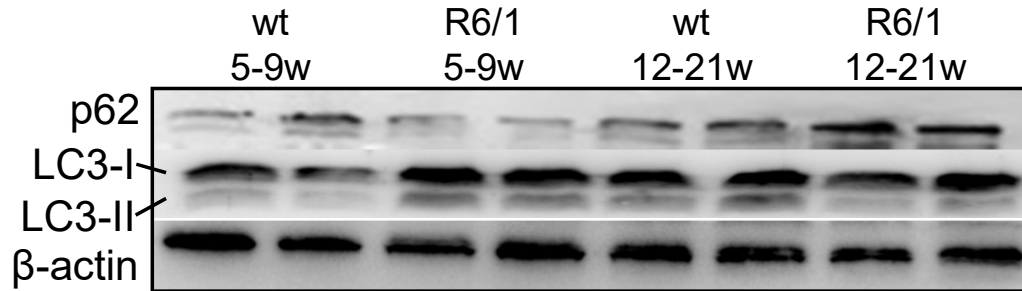
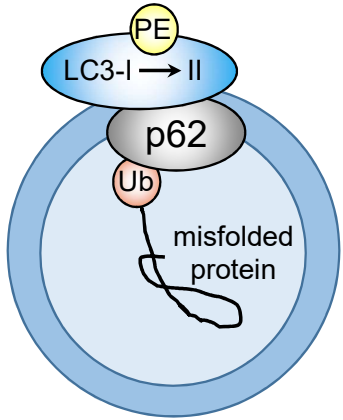


Iba-1 / DAPI: arrow, ramified resting microglia; \*, amoeboid microglia



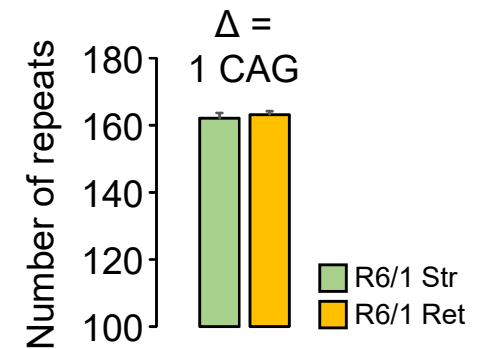
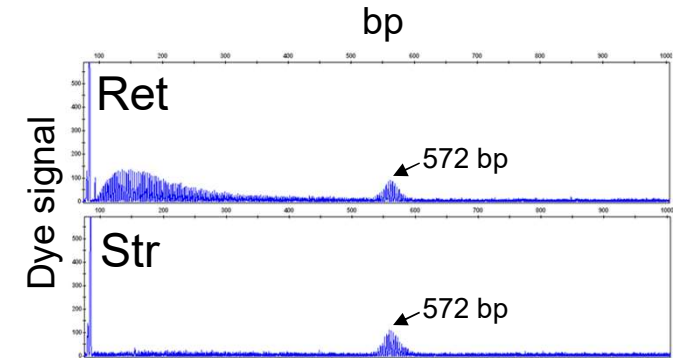
Morphology analysis indicates microglia activation. O, outer; I, inner; N, nuclear; P, plexiform; GCL, ganglion cells layer

### 3. LC3 AND p62 PROFILES ARE DIFFERENT IN THE RETINA AND STRIATUM OF R6/1 MICE



Western blotting analysis of R6/1 and wt. \*,  $P$ -value < 0.05 Mann-Whitney U-test. Representative blots from retina are shown. Autophagy may be altered in the retina (autophagic induction in early stages) and in the striatum (autophagic flux impairment in advance stages). p62 and LC3 may be compensated in retina by upregulation of *SQSTM1* and *MAP1LC3B* genes, respectively (RNA-seq results).

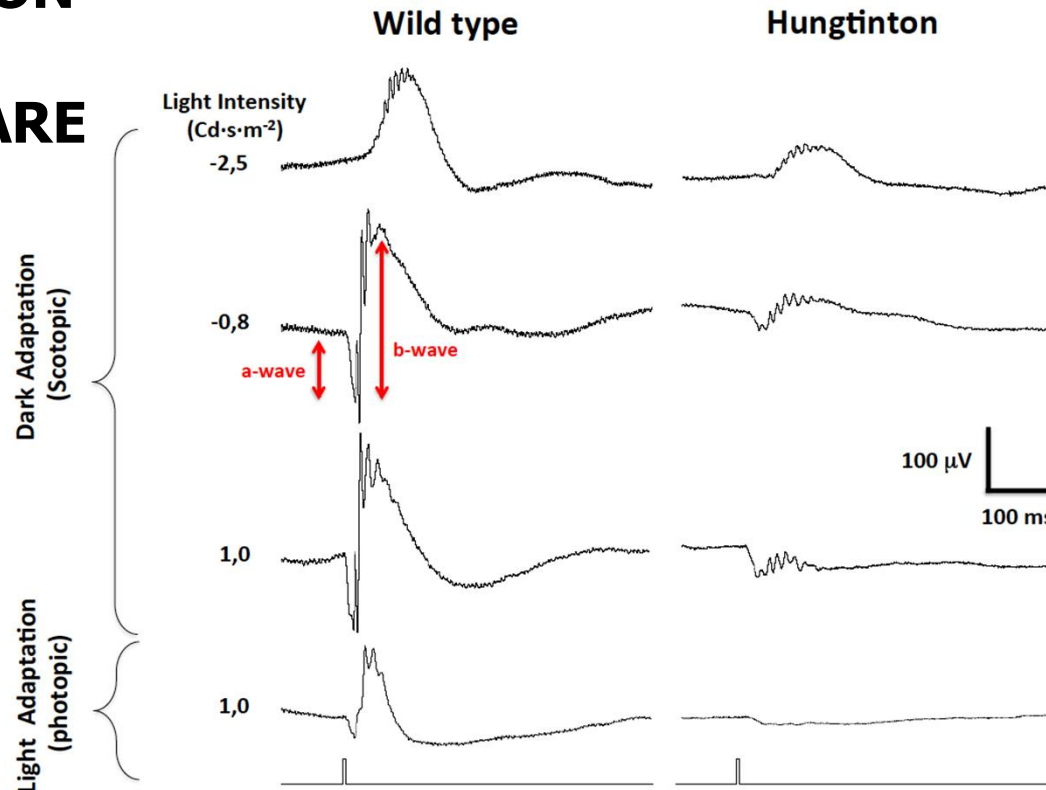
### 4. INSTABILITY OF CAG REPEATS DO NOT EXPLAIN TISSUE DIFFERENCES



CAG repeat analysis in retina and striatum of the same R6/1 animals

# 5. FUNCTION OF R6/1 RETINAS ARE IMPAIRED

Electroretinogram of 24 weeks-old R6/1 and wt littermates



# CONCLUSIONS

- \* R6/1 retinas show a profound transcriptional dysregulation containing an important retinal-specific component.
- \* There is a delimited inflammatory response characterized by gliosis and microglia activation but without the induction of other markers. The presence of degeneration-related markers in R6/1 retina justifies further exploration in other models. This is not clearly evident in the striatum.
- \* Currently, we are exploring in further detail autophagic processes as a potential explanation for retina malfunctioning in the R6/1 mice.

# FUNDING



MINISTERIO DE CIENCIA E INNOVACIÓN



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