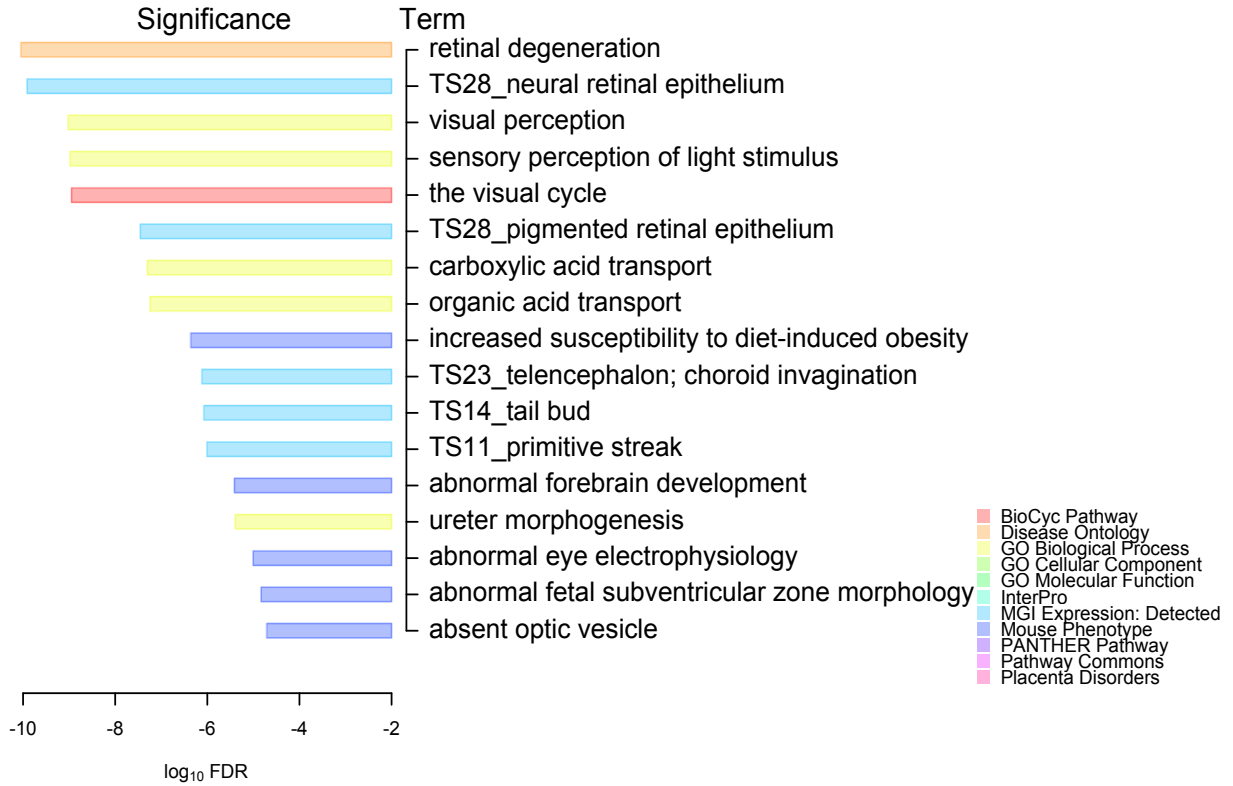1   **Figure 2-figure supplement 1**

GREAT Analysis: RPE-specific enhancers

2
3   **Figure 2-figure supplement 1.**
4   **Functional annotation of genes near putative enhancers present exclusively in control**
5   **cobblestone RPE and not in RPE exposed to TGFβ and TNFα.**
6   The top over-represented categories belonging to the different ontologies are shown in the
7   lower right legend. GO = Gene ontology, MGI = Mouse Genome Informatics. The *x* axes values
8   correspond to the $\log_{10}$ of false discovery rate.
9