Figure 2-figure supplement 2

**GREAT Analysis: shared enhancers** 

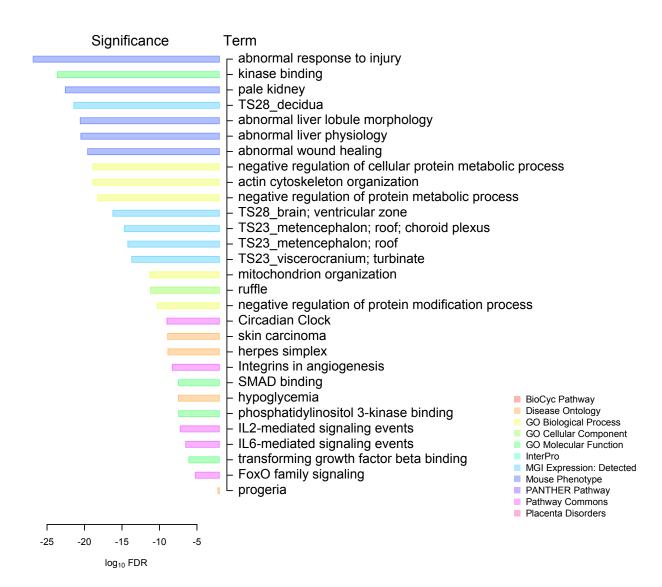


Figure 2-figure supplement 2. Functional annotation of genes near putative enhancers present in both control cobblestone RPE and RPE exposed to TGF $\beta$  and TNF $\alpha$ .

The top over-represented categories belonging to the different ontologies are shown in the lower right legend. GO = Gene ontology, MGI = Mouse Genome Informatics. The x axes values correspond to the  $\log_{10}$  of false discovery rate.