

Figure 2-figure supplement 2

GREAT Analysis: shared enhancers



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Functional annotation of genes near putative enhancers present in both control cobblestone RPE and RPE exposed to TGFβ and TNFα.

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 log₁₀ of false discovery rate.