

1 **Figure 2-figure supplement 3**

GREAT Analysis: TnT-specific enhancers



2
3 **Figure 2-figure supplement 3.**
4 **Functional annotation of genes near putative enhancers present exclusively in RPE exposed**
5 **to TGFβ and TNFα and not in control cobblestone RPE.**

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 log₁₀ of false discovery rate.

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