

## **Functional Annotation Chart**

Help and Manual

**Current Gene List: condition\_specific\_gene\_list Current Background: Homo sapiens** 2052 DAVID IDs

## **Options**

Rerun Using Options | Create Sublist 4 chart records March Download File Sublist Category RT Genes Count % P-Value Benjamini Term **RT** 306 14.9 3.9E-10 7.1E-9 GAD\_DISEASE\_CLASS INFECTION GAD\_DISEASE\_CLASS IMMUNE **RT** 364 17.7 1.2E-4 1.0E-3 **RT** 384 18.7 2.3E-4 1.4E-3 GAD\_DISEASE\_CLASS CANCER

GAD\_DISEASE\_CLASS PHARMACOGENOMIC RT 314 15.3 1.2E-2 5.6E-2

1210 gene(s) from your list are not in the output.

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