

## **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 9 chart records March Download File Sublist Category Term RT Genes Count % P-Value Benjamini UP\_KW\_DOMAIN Coiled coil **RT** 361 17.6 8.5E-18 2.6E-16 UP\_KW\_DOMAIN <u>Transit peptide</u> <u>RT</u> 81 3.9 2.8E-5 4.3E-4 UP\_KW\_DOMAIN Redox-active center RT 15 0.7 1.8E-4 1.9E-3 UP\_KW\_DOMAIN <u>Zinc-finger</u> <u>RT</u> <u>205</u> 10.0 2.6E-4 2.0E-3 UP\_KW\_DOMAIN <u>WD repeat</u> RT 42 2.0 1.1E-3 7.1E-3 UP\_KW\_DOMAIN Bromodomain RT 10 0.5 7.0E-3 3.6E-2 
 UP\_KW\_DOMAIN
 SH2 domain
 RT
 19
 0.9
 8.2E-3
 3.7E-2
UP\_KW\_DOMAIN SH3 domain RT 30 1.5 2.9E-2 1.1E-1

1377 gene(s) from your list are not in the output.

UP\_KW\_DOMAIN <u>TPR repeat</u>

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**RT** 23 1.1 4.9E-2 1.7E-1