



Functional Annotation Chart

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Current Gene List: condition_specific_gene_list
Current Background: Homo sapiens
2052 DAVID IDs

Options

Rerun Using Options

Create Sublist

8 chart records [Download File](#)

Sublist	Category	Term	RT	Genes	Count	%	P-Value	Benjamini
<input type="checkbox"/>	CHROMOSOME	MT	RT		11	0.5	4.8E-12	1.2E-10
<input type="checkbox"/>	CHROMOSOME	17	RT		138	6.7	3.5E-4	4.6E-3
<input type="checkbox"/>	CHROMOSOME	19	RT		125	6.1	1.1E-3	9.6E-3
<input type="checkbox"/>	CHROMOSOME	11	RT		127	6.2	3.2E-2	1.7E-1
<input type="checkbox"/>	CHROMOSOME	3	RT		133	6.5	3.2E-2	1.7E-1
<input type="checkbox"/>	CHROMOSOME	12	RT		114	5.6	5.4E-2	2.3E-1
<input type="checkbox"/>	CHROMOSOME	16	RT		98	4.8	6.5E-2	2.3E-1
<input type="checkbox"/>	CHROMOSOME	X	RT		72	3.5	7.0E-2	2.3E-1

1234 gene(s)

 from your list are not in the output.

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