National Institutes of Health
DAVID Bioinformatics

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 |

14 chart records

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Sublist	Category	Term	RT	Genes	Count	%	P-Value	Benjamini
☐	GOTERM_BP_1	metabolic process	RT		1514	73.8	5.2E-37	1.1E-35
☐	GOTERM_BP_1	cellular component organization or biogenesis	RT		968	47.2	1.4E-35	1.6E-34
☐	GOTERM_BP_1	immune system process	RT		464	22.6	2.0E-20	1.4E-19
☐	GOTERM_BP_1	cellular process	RT		1867	91.0	3.9E-18	2.1E-17
☐	GOTERM_BP_1	localization	RT		815	39.7	1.6E-14	7.2E-14
☐	GOTERM_BP_1	multi-organism process	RT		363	17.7	2.9E-11	1.1E-10
☐	GOTERM_BP_1	response to stimulus	RT		1095	53.4	2.9E-10	9.1E-10
☐	GOTERM_BP_1	biological regulation	RT		1422	69.3	6.7E-7	1.8E-6
☐	GOTERM_BP_1	signaling	RT		735	35.8	1.3E-3	3.2E-3
☐	GOTERM_BP_1	cell killing	RT		38	1.9	4.2E-3	9.3E-3
☐	GOTERM_BP_1	developmental process	RT		724	35.3	9.3E-3	1.9E-2
☐	GOTERM_BP_1	single-organism process	RT		1520	74.1	1.4E-2	2.5E-2
☐	GOTERM_BP_1	rhythmic process	RT		48	2.3	1.5E-2	2.5E-2
☐	GOTERM_BP_1	biological adhesion	RT		181	8.8	5.4E-2	8.5E-2

| 108 gene(s) | from your list are not in the output.

Please cite DAVID within any publication that makes use of any methods inspired by **DAVID**.