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]

9 chart records   [Download File](#)

Sublist	Category	Term	RT	Genes	Count	%	P-Value	Benjamini
☐	UP_KW_DISEASE	Diamond-Blackfan anemia	RT		18	0.9	1.5E-13	1.5E-11
☐	UP_KW_DISEASE	Intellectual disability	RT		121	5.9	1.5E-4	5.4E-3
☐	UP_KW_DISEASE	Leber hereditary optic neuropathy	RT		8	0.4	1.6E-4	5.4E-3
☐	UP_KW_DISEASE	Parkinsonism	RT		13	0.6	3.0E-3	7.5E-2
☐	UP_KW_DISEASE	Proto-oncogene	RT		43	2.1	4.7E-3	9.4E-2
☐	UP_KW_DISEASE	Neurodegeneration	RT		69	3.4	5.8E-3	9.7E-2
☐	UP_KW_DISEASE	Primary mitochondrial disease	RT		36	1.8	1.6E-2	2.3E-1
☐	UP_KW_DISEASE	Parkinson disease	RT		8	0.4	1.9E-2	2.4E-1
☐	UP_KW_DISEASE	Amyotrophic lateral sclerosis	RT		10	0.5	3.8E-2	4.3E-1

[1782 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**.