



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

56 chart records

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Sublist	Category	Term	RT	Genes	Count	%	P-Value	Benjamini
<input type="checkbox"/>	GENE3D	-	RT		76	3.7	3.1E-16	2.9E-13
<input type="checkbox"/>	GENE3D	Glutaredoxin	RT		32	1.6	6.5E-6	3.0E-3
<input type="checkbox"/>	GENE3D	Zinc/RING finger domain, C3HC4 (zinc finger)	RT		79	3.8	3.8E-5	1.2E-2
<input type="checkbox"/>	GENE3D	Rel homology domain (RHD), DNA-binding domain	RT		7	0.3	1.6E-4	3.7E-2
<input type="checkbox"/>	GENE3D	-	RT		15	0.7	5.1E-4	7.8E-2
<input type="checkbox"/>	GENE3D	SAP domain	RT		10	0.5	5.1E-4	7.8E-2
<input type="checkbox"/>	GENE3D	-	RT		6	0.3	9.5E-4	1.3E-1
<input type="checkbox"/>	GENE3D	P-loop containing nucleotide triphosphate hydrolases	RT		110	5.4	1.1E-3	1.3E-1
<input type="checkbox"/>	GENE3D	-	RT		9	0.4	1.3E-3	1.4E-1
<input type="checkbox"/>	GENE3D	-	RT		7	0.3	1.6E-3	1.5E-1
<input type="checkbox"/>	GENE3D	ARID DNA-binding domain	RT		7	0.3	2.4E-3	1.8E-1
<input type="checkbox"/>	GENE3D	STAT transcription factor, all-alpha domain	RT		5	0.2	2.9E-3	1.8E-1
<input type="checkbox"/>	GENE3D	STAT transcription factor, DNA-binding domain	RT		5	0.2	2.9E-3	1.8E-1
<input type="checkbox"/>	GENE3D	DUSP-like	RT		5	0.2	2.9E-3	1.8E-1
<input type="checkbox"/>	GENE3D	STAT transcription factor, N-terminal domain	RT		5	0.2	2.9E-3	1.8E-1
<input type="checkbox"/>	GENE3D	AN1-like Zinc finger	RT		5	0.2	5.3E-3	3.0E-1
<input type="checkbox"/>	GENE3D	Nucleic acid-binding proteins	RT		18	0.9	7.6E-3	4.1E-1
<input type="checkbox"/>	GENE3D	Zinc finger, RanBP2-type	RT		6	0.3	9.8E-3	5.0E-1
<input type="checkbox"/>	GENE3D	Bromodomain-like	RT		11	0.5	1.1E-2	5.5E-1
<input type="checkbox"/>	GENE3D	Severin	RT		8	0.4	1.3E-2	5.9E-1
<input type="checkbox"/>	GENE3D	-	RT		6	0.3	1.4E-2	5.9E-1
<input type="checkbox"/>	GENE3D	-	RT		4	0.2	1.7E-2	6.4E-1
<input type="checkbox"/>	GENE3D	Nemo cc2-lz domain - 1d5 darpin complex	RT		4	0.2	1.7E-2	6.4E-1
<input type="checkbox"/>	GENE3D	Cysteine proteinases	RT		19	0.9	1.7E-2	6.4E-1
<input type="checkbox"/>	GENE3D	-	RT		8	0.4	1.9E-2	7.1E-1
<input type="checkbox"/>	GENE3D	SH2 domain	RT		20	1.0	2.1E-2	7.3E-1
<input type="checkbox"/>	GENE3D	MHC class I-like antigen recognition-like	RT		8	0.4	2.3E-2	7.4E-1
<input type="checkbox"/>	GENE3D	TCP-1-like chaperonin intermediate domain	RT		6	0.3	2.3E-2	7.4E-1
<input type="checkbox"/>	GENE3D	GroEL-like equatorial domain	RT		6	0.3	2.3E-2	7.4E-1
<input type="checkbox"/>	GENE3D	-	RT		8	0.4	2.8E-2	8.4E-1
<input type="checkbox"/>	GENE3D	-	RT		3	0.1	2.9E-2	8.4E-1
<input type="checkbox"/>	GENE3D	GroEL	RT		6	0.3	3.0E-2	8.4E-1
<input type="checkbox"/>	GENE3D	Death Domain, Fas	RT		18	0.9	3.0E-2	8.4E-1
<input type="checkbox"/>	GENE3D	Tetratricopeptide repeat domain	RT		32	1.6	3.8E-2	9.8E-1
<input type="checkbox"/>	GENE3D	-	RT		15	0.7	3.9E-2	9.8E-1
<input type="checkbox"/>	GENE3D	Mannose-6-phosphate receptor binding domain	RT		4	0.2	4.0E-2	9.8E-1
<input type="checkbox"/>	GENE3D	-	RT		4	0.2	4.0E-2	9.8E-1
<input type="checkbox"/>	GENE3D	-	RT		5	0.2	4.6E-2	1.0E0
<input type="checkbox"/>	GENE3D	Cyclin-like	RT		10	0.5	5.0E-2	1.0E0
<input type="checkbox"/>	GENE3D	F-actin capping protein, beta subunit	RT		3	0.1	5.4E-2	1.0E0
<input type="checkbox"/>	GENE3D	MRG domain	RT		3	0.1	5.4E-2	1.0E0
<input type="checkbox"/>	GENE3D	-	RT		4	0.2	5.5E-2	1.0E0
<input type="checkbox"/>	GENE3D	-	RT		10	0.5	5.6E-2	1.0E0
<input type="checkbox"/>	GENE3D	YVTN repeat-like/Quinoprotein amine dehydrogenase	RT		47	2.3	5.9E-2	1.0E0
<input type="checkbox"/>	GENE3D	Arf GTPase activating protein	RT		7	0.3	6.8E-2	1.0E0
<input type="checkbox"/>	GENE3D	SH3 Domains	RT		32	1.6	7.1E-2	1.0E0
<input type="checkbox"/>	GENE3D	-	RT		5	0.2	7.2E-2	1.0E0
<input type="checkbox"/>	GENE3D	-	RT		7	0.3	7.7E-2	1.0E0
<input type="checkbox"/>	GENE3D	SUI1-like domain	RT		3	0.1	8.4E-2	1.0E0
<input type="checkbox"/>	GENE3D	-	RT		5	0.2	8.6E-2	1.0E0
<input type="checkbox"/>	GENE3D	Phosphotransferase system, lactose/cellobiose-type IIA subunit	RT		5	0.2	8.6E-2	1.0E0
<input type="checkbox"/>	GENE3D	-	RT		7	0.3	8.8E-2	1.0E0
<input type="checkbox"/>	GENE3D	Cyclophilin-like	RT		6	0.3	8.9E-2	1.0E0
<input type="checkbox"/>	GENE3D	-	RT		14	0.7	9.3E-2	1.0E0
<input type="checkbox"/>	GENE3D	Cupin	RT		7	0.3	1.0E-1	1.0E0
<input type="checkbox"/>	GENE3D	Zinc finger, CCCH-type	RT		7	0.3	1.0E-1	1.0E0

1406 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**.