



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

17 chart records

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Sublist	Category	Term	RT	Genes	Count	%	P-Value	Benjamini
<input type="checkbox"/>	UP_KW_MOLECULAR_FUNCTION	Ribosomal protein	RT	<div></div>	91	4.4	5.6E-37	3.2E-35
<input type="checkbox"/>	UP_KW_MOLECULAR_FUNCTION	Ribonucleoprotein	RT	<div></div>	118	5.8	6.5E-37	3.2E-35
<input type="checkbox"/>	UP_KW_MOLECULAR_FUNCTION	RNA-binding	RT	<div></div>	183	8.9	1.5E-28	4.9E-27
<input type="checkbox"/>	UP_KW_MOLECULAR_FUNCTION	Chaperone	RT	<div></div>	67	3.3	1.2E-13	2.8E-12
<input type="checkbox"/>	UP_KW_MOLECULAR_FUNCTION	Repressor	RT	<div></div>	120	5.8	2.4E-10	4.7E-9
<input type="checkbox"/>	UP_KW_MOLECULAR_FUNCTION	Chromatin regulator	RT	<div></div>	68	3.3	1.9E-8	3.1E-7
<input type="checkbox"/>	UP_KW_MOLECULAR_FUNCTION	Activator	RT	<div></div>	123	6.0	1.0E-7	1.4E-6
<input type="checkbox"/>	UP_KW_MOLECULAR_FUNCTION	Initiation factor	RT	<div></div>	20	1.0	4.6E-6	5.5E-5
<input type="checkbox"/>	UP_KW_MOLECULAR_FUNCTION	rRNA-binding	RT	<div></div>	11	0.5	5.3E-4	5.8E-3
<input type="checkbox"/>	UP_KW_MOLECULAR_FUNCTION	Helicase	RT	<div></div>	29	1.4	1.5E-3	1.5E-2
<input type="checkbox"/>	UP_KW_MOLECULAR_FUNCTION	Thiol protease	RT	<div></div>	28	1.4	3.9E-3	3.4E-2
<input type="checkbox"/>	UP_KW_MOLECULAR_FUNCTION	Rotamase	RT	<div></div>	11	0.5	1.5E-2	1.2E-1
<input type="checkbox"/>	UP_KW_MOLECULAR_FUNCTION	Transferase	RT	<div></div>	247	12.0	2.4E-2	1.8E-1
<input type="checkbox"/>	UP_KW_MOLECULAR_FUNCTION	Hydrolase	RT	<div></div>	216	10.5	2.8E-2	1.8E-1
<input type="checkbox"/>	UP_KW_MOLECULAR_FUNCTION	Elongation factor	RT	<div></div>	8	0.4	2.9E-2	1.8E-1
<input type="checkbox"/>	UP_KW_MOLECULAR_FUNCTION	Protein phosphatase	RT	<div></div>	24	1.2	3.0E-2	1.8E-1
<input type="checkbox"/>	UP_KW_MOLECULAR_FUNCTION	Kinase	RT	<div></div>	93	4.5	8.0E-2	4.6E-1

1134 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**.