



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_CELLULAR_COMPONENT	<a href="#">Cytoplasm</a>	RT	<div></div>	938	45.7	3.5E-63	2.0E-61
<input type="checkbox"/>	UP_KW_CELLULAR_COMPONENT	<a href="#">Nucleus</a>	RT	<div></div>	909	44.3	1.2E-50	3.3E-49
<input type="checkbox"/>	UP_KW_CELLULAR_COMPONENT	<a href="#">Endoplasmic reticulum</a>	RT	<div></div>	225	11.0	3.0E-12	5.6E-11
<input type="checkbox"/>	UP_KW_CELLULAR_COMPONENT	<a href="#">Endosome</a>	RT	<div></div>	128	6.2	9.6E-11	1.3E-9
<input type="checkbox"/>	UP_KW_CELLULAR_COMPONENT	<a href="#">Spliceosome</a>	RT	<div></div>	45	2.2	2.9E-10	3.2E-9
<input type="checkbox"/>	UP_KW_CELLULAR_COMPONENT	<a href="#">Lysosome</a>	RT	<div></div>	87	4.2	5.2E-10	4.9E-9
<input type="checkbox"/>	UP_KW_CELLULAR_COMPONENT	<a href="#">Mitochondrion</a>	RT	<div></div>	197	9.6	1.9E-6	1.5E-5
<input type="checkbox"/>	UP_KW_CELLULAR_COMPONENT	<a href="#">Golgi apparatus</a>	RT	<div></div>	140	6.8	2.1E-4	1.5E-3
<input type="checkbox"/>	UP_KW_CELLULAR_COMPONENT	<a href="#">Proteasome</a>	RT	<div></div>	16	0.8	4.0E-4	2.5E-3
<input type="checkbox"/>	UP_KW_CELLULAR_COMPONENT	<a href="#">MHC II</a>	RT	<div></div>	9	0.4	5.8E-4	3.2E-3
<input type="checkbox"/>	UP_KW_CELLULAR_COMPONENT	<a href="#">Cytoskeleton</a>	RT	<div></div>	185	9.0	9.8E-4	5.0E-3
<input type="checkbox"/>	UP_KW_CELLULAR_COMPONENT	<a href="#">MHC I</a>	RT	<div></div>	6	0.3	1.1E-3	5.2E-3
<input type="checkbox"/>	UP_KW_CELLULAR_COMPONENT	<a href="#">Chromosome</a>	RT	<div></div>	90	4.4	1.3E-3	5.5E-3
<input type="checkbox"/>	UP_KW_CELLULAR_COMPONENT	<a href="#">Cytoplasmic vesicle</a>	RT	<div></div>	103	5.0	2.2E-3	8.9E-3
<input type="checkbox"/>	UP_KW_CELLULAR_COMPONENT	<a href="#">Telomere</a>	RT	<div></div>	12	0.6	1.3E-2	4.9E-2
<input type="checkbox"/>	UP_KW_CELLULAR_COMPONENT	<a href="#">Centromere</a>	RT	<div></div>	24	1.2	3.4E-2	1.2E-1
<input type="checkbox"/>	UP_KW_CELLULAR_COMPONENT	<a href="#">Mitochondrion inner membrane</a>	RT	<div></div>	48	2.3	4.0E-2	1.3E-1

344 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**.