



Functional Annotation Chart

[Help and Manual](#)

Current Gene List: condition\_specific\_gene\_list  
Current Background: Homo sapiens  
2052 DAVID IDs




Options

Rerun Using Options

Create Sublist

3 chart records

 [Download File](#)

Sublist	Category	Term	RT	Genes	Count	%	P-Value	Benjamini
<input type="checkbox"/>	ENTREZ_GENE_SUMMARY	Enables RNA binding activity. [provided by Alliance of Genome Resources, Apr 2022]	<a href="#">RT</a>		3	0.1	2.4E-2	1.0E0
<input type="checkbox"/>	ENTREZ_GENE_SUMMARY	Involved in constitutive secretory pathway. Located in Golgi apparatus. [provided by Alliance of Genome Resources, Apr 2022]	<a href="#">RT</a>		2	0.1	2.7E-2	1.0E0
<input type="checkbox"/>	ENTREZ_GENE_SUMMARY	Predicted to enable nucleic acid binding activity. Predicted to be located in nucleus. [provided by Alliance of Genome Resources, Apr 2022]	<a href="#">RT</a>		2	0.1	2.7E-2	1.0E0

2045 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**.