



## DAVID Bioinformatics Resources 6.8

Laboratory of Human Retrovirology and Immunoinformatics (LHRI)

\*\*\* Welcome to DAVID 6.8 \*\*\*  
 \*\*\* If you are looking for [DAVID 6.7](#), please visit our [development site](#). \*\*\*

### Functional Annotation Chart

[Help and Manual](#)

Current Gene List: Topic 39

Current Background: Homo sapiens

16 DAVID IDs













Options

Rerun Using Options

Create Sublist

12 chart records

 [Download File](#)

Sublist	Category	Term	RT	Genes	Count	%	P-Value	Benjamini
<input type="checkbox"/>	UNIGENE_EST_QUARTILE	eye_normal_3rd	<a href="#">RT</a>		9	56.2	1.7E-3	8.3E-2
<input type="checkbox"/>	UNIGENE_EST_QUARTILE	normal_disease_3rd	<a href="#">RT</a>		10	62.5	2.9E-3	7.3E-2
<input type="checkbox"/>	UNIGENE_EST_QUARTILE	brain_normal_3rd	<a href="#">RT</a>		9	56.2	3.9E-3	6.5E-2
<input type="checkbox"/>	UP_TISSUE	Amygdala	<a href="#">RT</a>		4	25.0	9.5E-3	1.5E-1
<input type="checkbox"/>	UP_TISSUE	Brain	<a href="#">RT</a>		11	68.8	3.4E-2	2.5E-1
<input type="checkbox"/>	UP_KEYWORDS	<a href="#">Membrane</a>	<a href="#">RT</a>		10	62.5	5.4E-2	9.8E-1
<input type="checkbox"/>	UP_KEYWORDS	<a href="#">Alternative splicing</a>	<a href="#">RT</a>		12	75.0	7.4E-2	9.3E-1
<input type="checkbox"/>	UP_KEYWORDS	<a href="#">Transmembrane helix</a>	<a href="#">RT</a>		8	50.0	8.7E-2	8.8E-1
<input type="checkbox"/>	UP_KEYWORDS	<a href="#">Transmembrane</a>	<a href="#">RT</a>		8	50.0	8.8E-2	8.0E-1
<input type="checkbox"/>	UNIGENE_EST_QUARTILE	glioma_disease_3rd	<a href="#">RT</a>		6	37.5	9.2E-2	7.1E-1
<input type="checkbox"/>	UP_TISSUE	Cerebellum	<a href="#">RT</a>		3	18.8	9.3E-2	4.3E-1
<input type="checkbox"/>	UNIGENE_EST_QUARTILE	embryo_development_3rd	<a href="#">RT</a>		8	50.0	9.5E-2	6.5E-1