

Functional Annotation Chart

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Current Gene List: Topic 39

Current Background: Homo sapiens

16 DAVID IDs

Options

Rerun Using Options

Create Sublist

12 chart records

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