

Functional Annotation Chart

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Current Gene List: Topic 135

Current Background: Homo sapiens

10 DAVID IDs

Options

Rerun Using Options

Create Sublist

60 chart records

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Sublist	Category	Term	RT	Genes	Count	%	P-Value	Benjamini
<input type="checkbox"/>	UNIGENE_EST_QUARTILE	stomach_normal_3rd	RT	<div></div>	9	90.0	2.4E-6	1.3E-4
<input type="checkbox"/>	GNF_U133A_QUARTILE	Trigeminal Ganglion_3rd	RT	<div></div>	8	80.0	4.1E-6	2.5E-4
<input type="checkbox"/>	CGAP_SAGE_QUARTILE	stomach adenocarcinoma_3rd	RT	<div></div>	6	60.0	2.3E-4	1.7E-2
<input type="checkbox"/>	UP_KEYWORDS	Secreted	RT	<div></div>	6	60.0	7.2E-4	2.5E-2
<input type="checkbox"/>	UP_SEQ_FEATURE	signal peptide	RT	<div></div>	7	70.0	1.1E-3	6.2E-2
<input type="checkbox"/>	UNIGENE_EST_QUARTILE	gastrointestinal tumor_disease_3rd	RT	<div></div>	7	70.0	1.6E-3	4.2E-2
<input type="checkbox"/>	CGAP_SAGE_QUARTILE	colon normal epithelium_3rd	RT	<div></div>	5	50.0	2.6E-3	9.5E-2
<input type="checkbox"/>	GOTERM_CC_DIRECT	extracellular space	RT	<div></div>	5	50.0	2.8E-3	5.7E-2
<input type="checkbox"/>	UP_KEYWORDS	Signal	RT	<div></div>	7	70.0	3.2E-3	5.7E-2
<input type="checkbox"/>	CGAP_SAGE_QUARTILE	gastrointestinal tract_null_3rd	RT	<div></div>	4	40.0	4.2E-3	9.9E-2