



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

73 chart records [Download File](#)

Sublist	Category	Term	RT	Genes	Count	%	P-Value	Benjamini
<input type="checkbox"/>	CYTOGENETIC_LOCATION	6p21.32	<a href="#">RT</a>		20	1.0	3.9E-11	2.4E-8
<input type="checkbox"/>	CYTOGENETIC_LOCATION	12q24.31	<a href="#">RT</a>		18	0.9	6.2E-6	1.9E-3
<input type="checkbox"/>	CYTOGENETIC_LOCATION	19p13.3	<a href="#">RT</a>		26	1.3	1.8E-5	3.6E-3
<input type="checkbox"/>	CYTOGENETIC_LOCATION	11q13.1	<a href="#">RT</a>		16	0.8	9.8E-5	1.5E-2
<input type="checkbox"/>	CYTOGENETIC_LOCATION	19q13.2	<a href="#">RT</a>		19	0.9	1.5E-4	1.8E-2
<input type="checkbox"/>	CYTOGENETIC_LOCATION	3p21.31	<a href="#">RT</a>		20	1.0	2.1E-4	1.9E-2
<input type="checkbox"/>	CYTOGENETIC_LOCATION	17p13.1	<a href="#">RT</a>		18	0.9	2.2E-4	1.9E-2
<input type="checkbox"/>	CYTOGENETIC_LOCATION	1q21.3	<a href="#">RT</a>		19	0.9	4.7E-4	3.6E-2
<input type="checkbox"/>	CYTOGENETIC_LOCATION	1q22	<a href="#">RT</a>		10	0.5	9.8E-4	6.5E-2
<input type="checkbox"/>	CYTOGENETIC_LOCATION	22q13.1	<a href="#">RT</a>		12	0.6	1.2E-3	6.5E-2
<input type="checkbox"/>	CYTOGENETIC_LOCATION	22q13.2	<a href="#">RT</a>		12	0.6	1.2E-3	6.5E-2
<input type="checkbox"/>	CYTOGENETIC_LOCATION	2q11.2	<a href="#">RT</a>		13	0.6	1.4E-3	7.0E-2
<input type="checkbox"/>	CYTOGENETIC_LOCATION	1p34.2	<a href="#">RT</a>		12	0.6	1.6E-3	7.6E-2
<input type="checkbox"/>	CYTOGENETIC_LOCATION	6p21.33	<a href="#">RT</a>		13	0.6	1.8E-3	7.9E-2
<input type="checkbox"/>	CYTOGENETIC_LOCATION	1p35.3	<a href="#">RT</a>		8	0.4	2.5E-3	1.0E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	1p35.2	<a href="#">RT</a>		8	0.4	3.0E-3	1.2E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	Xq28	<a href="#">RT</a>		17	0.8	3.5E-3	1.3E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	19p13.13	<a href="#">RT</a>		7	0.3	5.2E-3	1.8E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	11p15.5	<a href="#">RT</a>		12	0.6	5.6E-3	1.8E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	11q12.2	<a href="#">RT</a>		8	0.4	5.7E-3	1.8E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	16p11.2	<a href="#">RT</a>		19	0.9	7.0E-3	2.0E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	12p13.31	<a href="#">RT</a>		14	0.7	7.4E-3	2.1E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	17q21.33	<a href="#">RT</a>		8	0.4	8.2E-3	2.2E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	19p13.11	<a href="#">RT</a>		12	0.6	8.4E-3	2.2E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	2p23.3	<a href="#">RT</a>		10	0.5	1.0E-2	2.4E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	10q22.1	<a href="#">RT</a>		9	0.4	1.1E-2	2.6E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	16p13.3	<a href="#">RT</a>		20	1.0	1.4E-2	3.2E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	11q13.2	<a href="#">RT</a>		10	0.5	1.5E-2	3.2E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	2q33.1	<a href="#">RT</a>		10	0.5	1.6E-2	3.3E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	15q22.31	<a href="#">RT</a>		8	0.4	1.6E-2	3.3E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	17q25.3	<a href="#">RT</a>		14	0.7	1.7E-2	3.3E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	14q21.3	<a href="#">RT</a>		6	0.3	2.0E-2	3.9E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	19q13.33	<a href="#">RT</a>		14	0.7	2.2E-2	4.0E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	5q31.1	<a href="#">RT</a>		10	0.5	2.4E-2	4.3E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	20q11.22	<a href="#">RT</a>		7	0.3	2.5E-2	4.3E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	2q23.3	<a href="#">RT</a>		5	0.2	3.6E-2	6.1E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	4p16.3	<a href="#">RT</a>		10	0.5	4.1E-2	6.7E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	12q13.2	<a href="#">RT</a>		7	0.3	4.1E-2	6.7E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	3p25.3	<a href="#">RT</a>		7	0.3	4.3E-2	6.7E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	2p13.3	<a href="#">RT</a>		7	0.3	4.5E-2	6.7E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	2q14.3	<a href="#">RT</a>		7	0.3	4.5E-2	6.7E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	10q26.11	<a href="#">RT</a>		5	0.2	4.7E-2	6.7E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	15q21.2	<a href="#">RT</a>		6	0.3	4.8E-2	6.7E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	20p13	<a href="#">RT</a>		10	0.5	4.8E-2	6.7E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	7p13	<a href="#">RT</a>		6	0.3	5.0E-2	6.8E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	7q21.2	<a href="#">RT</a>		4	0.2	5.1E-2	6.8E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	17q12	<a href="#">RT</a>		13	0.6	5.5E-2	7.1E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	17q23.3	<a href="#">RT</a>		5	0.2	5.7E-2	7.3E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	Xq24	<a href="#">RT</a>		7	0.3	6.1E-2	7.5E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	1p22.1	<a href="#">RT</a>		6	0.3	6.1E-2	7.5E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	3p22.2	<a href="#">RT</a>		6	0.3	6.4E-2	7.7E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	17p13.2	<a href="#">RT</a>		8	0.4	6.6E-2	7.8E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	16q24.2	<a href="#">RT</a>		5	0.2	6.8E-2	7.9E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	6p21.1	<a href="#">RT</a>		10	0.5	7.5E-2	8.0E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	1p36.33	<a href="#">RT</a>		9	0.4	7.7E-2	8.0E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	15q22.2	<a href="#">RT</a>		6	0.3	7.7E-2	8.0E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	9p13.3	<a href="#">RT</a>		9	0.4	7.9E-2	8.0E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	15q24.1	<a href="#">RT</a>		6	0.3	8.0E-2	8.0E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	16q24.3	<a href="#">RT</a>		6	0.3	8.0E-2	8.0E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	5q33.3	<a href="#">RT</a>		6	0.3	8.0E-2	8.0E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	22q12.3	<a href="#">RT</a>		8	0.4	8.1E-2	8.0E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	11q23.3	<a href="#">RT</a>		10	0.5	8.1E-2	8.0E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	15q21.1	<a href="#">RT</a>		7	0.3	8.3E-2	8.1E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	17q25.1	<a href="#">RT</a>		9	0.4	8.6E-2	8.1E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	14q13.2	<a href="#">RT</a>		4	0.2	8.6E-2	8.1E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	2p15	<a href="#">RT</a>		5	0.2	8.8E-2	8.1E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	21q22.11	<a href="#">RT</a>		8	0.4	8.8E-2	8.1E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	8q24.3	<a href="#">RT</a>		11	0.5	9.2E-2	8.1E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	6p21.31	<a href="#">RT</a>		7	0.3	9.2E-2	8.1E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	8q24.22	<a href="#">RT</a>		5	0.2	9.3E-2	8.1E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	2q37.1	<a href="#">RT</a>		8	0.4	9.4E-2	8.1E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	3p22.3	<a href="#">RT</a>		5	0.2	9.7E-2	8.1E-1
<input type="checkbox"/>	CYTOGENETIC_LOCATION	14q24.1	<a href="#">RT</a>		5	0.2	9.7E-2	8.1E-1

1325 gene(s) from your list are not in the output.

