



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

39 chart records

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Sublist	Category	Term	RT	Genes	Count	%	P-Value	Benjamini
<input type="checkbox"/>	DIP	receptor for activated C kinase 1(RACK1)	RT		68	3.3	4.4E-33	7.6E-30
<input type="checkbox"/>	DIP	proliferation-associated 2G4(PA2G4)	RT		37	1.8	1.6E-20	1.3E-17
<input type="checkbox"/>	DIP	eukaryotic translation initiation factor 3 subunit B(EIF3B)	RT		10	0.5	5.4E-5	3.1E-2
<input type="checkbox"/>	DIP	eukaryotic translation initiation factor 3 subunit C(EIF3C)	RT		11	0.5	1.4E-4	6.0E-2
<input type="checkbox"/>	DIP	eukaryotic translation initiation factor 3 subunit A(EIF3A)	RT		10	0.5	1.0E-3	3.1E-1
<input type="checkbox"/>	DIP	innate immunity activator(INAVA)	RT		15	0.7	1.4E-3	3.1E-1
<input type="checkbox"/>	DIP	eukaryotic translation initiation factor 3 subunit K(EIF3K)	RT		8	0.4	1.9E-3	3.1E-1
<input type="checkbox"/>	DIP	eukaryotic translation initiation factor 3 subunit H(EIF3H)	RT		8	0.4	1.9E-3	3.1E-1
<input type="checkbox"/>	DIP	eukaryotic translation initiation factor 3 subunit M(EIF3M)	RT		8	0.4	1.9E-3	3.1E-1
<input type="checkbox"/>	DIP	eukaryotic translation initiation factor 3 subunit L(EIF3L)	RT		8	0.4	1.9E-3	3.1E-1
<input type="checkbox"/>	DIP	cereblon(CRBN)	RT		10	0.5	2.0E-3	3.1E-1
<input type="checkbox"/>	DIP	eukaryotic translation initiation factor 3 subunit J(EIF3J)	RT		7	0.3	2.5E-3	3.5E-1
<input type="checkbox"/>	DIP	p21 (RAC1) activated kinase 1(PAK1)	RT		9	0.4	2.9E-3	3.9E-1
<input type="checkbox"/>	DIP	eukaryotic translation initiation factor 3 subunit F(EIF3F)	RT		8	0.4	4.2E-3	4.8E-1
<input type="checkbox"/>	DIP	eukaryotic translation initiation factor 3 subunit E(EIF3E)	RT		8	0.4	4.2E-3	4.8E-1
<input type="checkbox"/>	DIP	eukaryotic translation initiation factor 3 subunit D(EIF3D)	RT		7	0.3	6.0E-3	6.4E-1
<input type="checkbox"/>	DIP	protein phosphatase 2 regulatory subunit B'gamma(PPP2R5C)	RT		5	0.2	1.1E-2	1.0E0
<input type="checkbox"/>	DIP	nuclear protein 1, transcriptional regulator(NUPR1)	RT		9	0.4	1.5E-2	1.0E0
<input type="checkbox"/>	DIP	G protein pathway suppressor 1(GPS1)	RT		9	0.4	1.5E-2	1.0E0
<input type="checkbox"/>	DIP	reticulophagy regulator 1(RETREG1)	RT		8	0.4	2.3E-2	1.0E0
<input type="checkbox"/>	DIP	MDS1 and EVI1 complex locus(MECOM)	RT		5	0.2	2.7E-2	1.0E0
<input type="checkbox"/>	DIP	mitochondrial ribosomal protein L18(MRPL18)	RT		6	0.3	3.3E-2	1.0E0
<input type="checkbox"/>	DIP	reticulophagy regulator family member 3(RETREG3)	RT		4	0.2	3.9E-2	1.0E0
<input type="checkbox"/>	DIP	TCL1 family AKT coactivator A(TCL1A)	RT		4	0.2	3.9E-2	1.0E0
<input type="checkbox"/>	DIP	ORF129 ankyrin repeat protein(ORFVgORF129)	RT		4	0.2	3.9E-2	1.0E0
<input type="checkbox"/>	DIP	reticulophagy regulator family member 2(RETREG2)	RT		4	0.2	3.9E-2	1.0E0
<input type="checkbox"/>	DIP	F-box and WD repeat domain containing 5(FBXW5)	RT		5	0.2	5.1E-2	1.0E0
<input type="checkbox"/>	DIP	aquarius intron-binding spliceosomal factor(AQR)	RT		15	0.7	5.2E-2	1.0E0
<input type="checkbox"/>	DIP	UPF1 RNA helicase and ATPase(UPF1)	RT		7	0.3	5.3E-2	1.0E0
<input type="checkbox"/>	DIP	defender against cell death 1(DAD1)	RT		6	0.3	5.4E-2	1.0E0
<input type="checkbox"/>	DIP	lamin A/C(LMNA)	RT		4	0.2	8.1E-2	1.0E0
<input type="checkbox"/>	DIP	S-phase kinase associated protein 2(SKP2)	RT		4	0.2	8.1E-2	1.0E0
<input type="checkbox"/>	DIP	cyclin K(CCNK)	RT		4	0.2	8.1E-2	1.0E0
<input type="checkbox"/>	DIP	caspase 8(CASP8)	RT		6	0.3	8.1E-2	1.0E0
<input type="checkbox"/>	DIP	cyclin dependent kinase 9(CDK9)	RT		6	0.3	8.1E-2	1.0E0
<input type="checkbox"/>	DIP	inhibitor of nuclear factor kappa B kinase subunit beta(IKBKB)	RT		6	0.3	8.1E-2	1.0E0
<input type="checkbox"/>	DIP	damage specific DNA binding protein 1(DDB1)	RT		9	0.4	8.2E-2	1.0E0
<input type="checkbox"/>	DIP	COP9 signalosome subunit 4(COPS4)	RT		5	0.2	8.4E-2	1.0E0
<input type="checkbox"/>	DIP	F-box and WD repeat domain containing 8(FBXW8)	RT		8	0.4	9.2E-2	1.0E0

1857 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**.