



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

8095 chart records

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◀◀◀ 1, 2, 3, 4, 5, 6, 7, 8 ▶▶▶

Sublist	Category	Term	RT	Genes	Count	%	P-Value	Benjamini
<input type="checkbox"/>	BIOGRID_INTERACTION	baculoviral IAP repeat containing 3(BIRC3)	RT	<div></div>	493	24.0	1.9E-164	3.8E-160
<input type="checkbox"/>	BIOGRID_INTERACTION	pleckstrin homology domain containing A4(PLEKHA4)	RT	<div></div>	750	36.5	1.7E-146	1.6E-142
<input type="checkbox"/>	BIOGRID_INTERACTION	elongation factor Tu GTP binding domain containing 2(EFTUD2)	RT	<div></div>	479	23.3	4.4E-145	2.9E-141
<input type="checkbox"/>	BIOGRID_INTERACTION	nuclear receptor subfamily 2 group C member 2(NR2C2)	RT	<div></div>	477	23.2	6.8E-145	3.4E-141
<input type="checkbox"/>	BIOGRID_INTERACTION	bromodomain containing 4(BRD4)	RT	<div></div>	475	23.1	3.3E-140	1.3E-136
<input type="checkbox"/>	BIOGRID_INTERACTION	estrogen receptor 1(ESR1)	RT	<div></div>	616	30.0	5.9E-134	2.0E-130
<input type="checkbox"/>	BIOGRID_INTERACTION	RNA binding motif protein 39(RBM39)	RT	<div></div>	365	17.8	2.8E-131	7.9E-128
<input type="checkbox"/>	BIOGRID_INTERACTION	Adenomatous polyposis coli 2(Apc2)	RT	<div></div>	335	16.3	1.7E-123	4.2E-120
<input type="checkbox"/>	BIOGRID_INTERACTION	UFM1 specific ligase 1(UFL1)	RT	<div></div>	369	18.0	9.0E-119	2.0E-115
<input type="checkbox"/>	BIOGRID_INTERACTION	ORF1a polyprotein;ORF1ab polyprotein(ORF1ab)	RT	<div></div>	751	36.6	6.9E-116	1.4E-112
<input type="checkbox"/>	BIOGRID_INTERACTION	MYC proto-oncogene, bHLH transcription factor(MYC)	RT	<div></div>	542	26.4	4.3E-114	7.7E-111
<input type="checkbox"/>	BIOGRID_INTERACTION	RecQ like helicase 4(RECQL4)	RT	<div></div>	441	21.5	5.0E-114	8.3E-111
<input type="checkbox"/>	BIOGRID_INTERACTION	zinc finger CCCH-type containing 18(ZC3H18)	RT	<div></div>	325	15.8	4.9E-110	7.5E-107
<input type="checkbox"/>	BIOGRID_INTERACTION	N-alpha-acetyltransferase 40, NatD catalytic subunit(NAA40)	RT	<div></div>	349	17.0	1.3E-108	1.8E-105
<input type="checkbox"/>	BIOGRID_INTERACTION	neurotrophic receptor tyrosine kinase 1(NTRK1)	RT	<div></div>	526	25.6	7.4E-107	9.8E-104
<input type="checkbox"/>	BIOGRID_INTERACTION	fizzy and cell division cycle 20 related 1(FZR1)	RT	<div></div>	375	18.3	6.4E-106	7.9E-103
<input type="checkbox"/>	BIOGRID_INTERACTION	DDR GK domain containing 1(DDR GK1)	RT	<div></div>	392	19.1	2.6E-105	3.1E-102
<input type="checkbox"/>	BIOGRID_INTERACTION	prohibitin 1(PHB1)	RT	<div></div>	353	17.2	1.9E-101	2.1E-98
<input type="checkbox"/>	BIOGRID_INTERACTION	kinesin family member 14(KIF14)	RT	<div></div>	485	23.6	4.6E-100	4.8E-97
<input type="checkbox"/>	BIOGRID_INTERACTION	fibronectin 1(FN1)	RT	<div></div>	299	14.6	6.5E-98	6.5E-95
<input type="checkbox"/>	BIOGRID_INTERACTION	H4 clustered histone 1(H4C1)	RT	<div></div>	453	22.1	5.1E-97	4.8E-94
<input type="checkbox"/>	BIOGRID_INTERACTION	anterior gradient 2, protein disulphide isomerase family member(AGR2)	RT	<div></div>	326	15.9	6.5E-97	5.9E-94
<input type="checkbox"/>	BIOGRID_INTERACTION	citron rho-interacting serine/threonine kinase(CIT)	RT	<div></div>	418	20.4	1.5E-96	1.3E-93
<input type="checkbox"/>	BIOGRID_INTERACTION	cullin 7(CUL7)	RT	<div></div>	302	14.7	3.8E-94	3.1E-91
<input type="checkbox"/>	BIOGRID_INTERACTION	AAR2 splicing factor(AAR2)	RT	<div></div>	261	12.7	5.1E-93	4.0E-90
<input type="checkbox"/>	BIOGRID_INTERACTION	charged multivesicular body protein 4C(CHMP4C)	RT	<div></div>	269	13.1	2.4E-92	1.8E-89
<input type="checkbox"/>	BIOGRID_INTERACTION	WD repeat domain 5(WDR5)	RT	<div></div>	322	15.7	1.3E-91	9.5E-89
<input type="checkbox"/>	BIOGRID_INTERACTION	small nuclear ribonucleoprotein U1 subunit 70(SNRNP70)	RT	<div></div>	309	15.1	4.3E-91	3.0E-88
<input type="checkbox"/>	BIOGRID_INTERACTION	WW domain containing E3 ubiquitin protein ligase 2(WWP2)	RT	<div></div>	288	14.0	4.5E-91	3.0E-88
<input type="checkbox"/>	BIOGRID_INTERACTION	basic transcription factor 3(BTF3)	RT	<div></div>	282	13.7	5.8E-91	3.8E-88
<input type="checkbox"/>	BIOGRID_INTERACTION	methylphosphate capping enzyme(MEPCE)	RT	<div></div>	296	14.4	2.2E-89	1.4E-86
<input type="checkbox"/>	BIOGRID_INTERACTION	cyclin dependent kinase 2(CDK2)	RT	<div></div>	273	13.3	2.0E-88	1.2E-85
<input type="checkbox"/>	BIOGRID_INTERACTION	ring finger and CCCH-type domains 2(RC3H2)	RT	<div></div>	258	12.6	9.4E-88	5.6E-85
<input type="checkbox"/>	BIOGRID_INTERACTION	estrogen receptor 2(ESR2)	RT	<div></div>	537	26.2	1.9E-85	1.1E-82
<input type="checkbox"/>	BIOGRID_INTERACTION	PIH1 domain containing 1(PIH1D1)	RT	<div></div>	264	12.9	3.0E-85	1.7E-82
<input type="checkbox"/>	BIOGRID_INTERACTION	vascular cell adhesion molecule 1(VCAM1)	RT	<div></div>	207	10.1	4.3E-82	2.4E-79
<input type="checkbox"/>	BIOGRID_INTERACTION	BRCA1 DNA repair associated(BRCA1)	RT	<div></div>	338	16.5	2.3E-81	1.2E-78
<input type="checkbox"/>	BIOGRID_INTERACTION	E1A binding protein p300(EP300)	RT	<div></div>	366	17.8	9.3E-81	4.9E-78
<input type="checkbox"/>	BIOGRID_INTERACTION	staufen double-stranded RNA binding protein 1(STAU1)	RT	<div></div>	301	14.7	1.5E-80	7.6E-78
<input type="checkbox"/>	BIOGRID_INTERACTION	cullin 3(CUL3)	RT	<div></div>	365	17.8	4.8E-80	2.4E-77
<input type="checkbox"/>	BIOGRID_INTERACTION	integrin alpha FG-GAP repeat containing 1(ITFG1)	RT	<div></div>	222	10.8	5.5E-80	2.6E-77
<input type="checkbox"/>	BIOGRID_INTERACTION	ring finger and CCCH-type domains 1(RC3H1)	RT	<div></div>	249	12.1	9.1E-80	4.3E-77
<input type="checkbox"/>	BIOGRID_INTERACTION	tumor protein p53(TP53)	RT	<div></div>				