

Table S8. KEGG pathway enrichment analysis results for FBXO32-associated genes in neuroblastoma.

ID	Description	setSize	enrichmentScore	NES	pvalue	p.adjust	qvalue	rank	leading_ed
hsa03030	"DNA replication - Homo sapiens (human)"	36	-0.831938921310906	-3.09795679610531	1				
hsa04110	"Cell cycle - Homo sapiens (human)"	154	-0.608606276102003	-2.99488088918019	1e-10				
hsa03013	"Nucleocytoplasmic transport - Homo sapiens (human)"	105	-0.624113660463082	-2.8830					
hsa03040	"Spliceosome - Homo sapiens (human)"	130	-0.601675166118941	-2.87680305166459	1e-				
hsa04610	"Complement and coagulation cascades - Homo sapiens (human)"	86	0.617274430236405						
hsa03460	"Fanconi anemia pathway - Homo sapiens (human)"	49	-0.685600884014663	-2.75994426					
hsa05144	"Malaria - Homo sapiens (human)"	50	0.674876980788658	2.73007518482318	1e-10	1.66			
hsa05323	"Rheumatoid arthritis - Homo sapiens (human)"	89	0.5966505156652	2.69898907740366					
hsa04060	"Cytokine-cytokine receptor interaction - Homo sapiens (human)"	290	0.50318026155930						
hsa04512	"ECM-receptor interaction - Homo sapiens (human)"	88	0.582300910102746	2.615060467					
hsa05146	"Amoebiasis - Homo sapiens (human)"	100	0.568672289675254	2.609048508954	1e-10	1.			
hsa04061	"Viral protein interaction with cytokine and cytokine receptor - Homo sapiens (human)"	98							
hsa04510	"Focal adhesion - Homo sapiens (human)"	200	0.496603069692248	2.51726891766059	1e-				
hsa04974	"Protein digestion and absorption - Homo sapiens (human)"	102	0.539522968154023	2.48					
hsa04380	"Osteoclast differentiation - Homo sapiens (human)"	125	0.516061560004748	2.45529122					
hsa04145	"Phagosome - Homo sapiens (human)"	148	0.502863538676107	2.43853808162657	1e-10				
hsa04020	"Calcium signaling pathway - Homo sapiens (human)"	238	0.465543504439207	2.4199579					
hsa05205	"Proteoglycans in cancer - Homo sapiens (human)"	202	0.444250036502874	2.252647786					
hsa04151	"PI3K-Akt signaling pathway - Homo sapiens (human)"	350	0.398605811143346	2.176002					
hsa04080	"Neuroactive ligand-receptor interaction - Homo sapiens (human)"	363	0.3683390559339						
hsa05165	"Human papillomavirus infection - Homo sapiens (human)"	328	0.37264518661785	2.0010					
hsa04621	"NOD-like receptor signaling pathway - Homo sapiens (human)"	178	0.44639532324243	2.					
hsa04142	"Lysosome - Homo sapiens (human)"	132	0.485530980754089	2.32175426315528	3.8571				
hsa04010	"MAPK signaling pathway - Homo sapiens (human)"	299	0.379130753085801	2.02733973					
hsa05150	"Staphylococcus aureus infection - Homo sapiens (human)"	87	0.540245217114455	2.4378					
hsa05145	"Toxoplasmosis - Homo sapiens (human)"	110	0.513356494441278	2.37993889897885	1e-				
hsa04514	"Cell adhesion molecules - Homo sapiens (human)"	146	0.463767959363541	2.241699346					
hsa03410	"Base excision repair - Homo sapiens (human)"	43	-0.656037092280527	-2.547086777292					
hsa05140	"Leishmaniasis - Homo sapiens (human)"	73	0.566559269603776	2.44806290498633	2.85				
hsa03015	"mRNA surveillance pathway - Homo sapiens (human)"	95	-0.50773399428915	-2.273216					
hsa05321	"Inflammatory bowel disease - Homo sapiens (human)"	63	0.58363679014708	2.4548084					
hsa05171	"Coronavirus disease - COVID-19 - Homo sapiens (human)"	229	0.386865627005928	2.000					
hsa05322	"Systemic lupus erythematosus - Homo sapiens (human)"	51	0.622188385839754	2.52439					
hsa05152	"Tuberculosis - Homo sapiens (human)"	176	0.42384278758546	2.10944646801028	1.624				
hsa04640	"Hematopoietic cell lineage - Homo sapiens (human)"	93	0.506902592851754	2.30469674					
hsa04810	"Regulation of actin cytoskeleton - Homo sapiens (human)"	227	0.390590100640825	2.020					
hsa03440	"Homologous recombination - Homo sapiens (human)"	37	-0.656542390340179	-2.46700					
hsa05330	"Allograft rejection - Homo sapiens (human)"	35	0.682698197461391	2.53947724325307					
hsa05418	"Fluid shear stress and atherosclerosis - Homo sapiens (human)"	136	0.443777584583708						
hsa04668	"TNF signaling pathway - Homo sapiens (human)"	113	0.47001187127474	2.18861345083					
hsa05202	"Transcriptional misregulation in cancer - Homo sapiens (human)"	172	0.41380759085435						
hsa05332	"Graft-versus-host disease - Homo sapiens (human)"	38	0.660119548902952	2.508246576					
hsa04670	"Leukocyte transendothelial migration - Homo sapiens (human)"	110	0.473430733876953						
hsa05320	"Autoimmune thyroid disease - Homo sapiens (human)"	50	0.583776504509674	2.361547					
hsa03430	"Mismatch repair - Homo sapiens (human)"	23	-0.72918843751938	-2.39834810457865	2.				
hsa05133	"Pertussis - Homo sapiens (human)"	76	0.513501828104333	2.24685871913068	3.470598				
hsa05417	"Lipid and atherosclerosis - Homo sapiens (human)"	212	0.373483854361316	1.91845621					
hsa04630	"JAK-STAT signaling pathway - Homo sapiens (human)"	165	0.403722494448279	1.989402					

hsa04940,"Type I diabetes mellitus - Homo sapiens (human)",41,0.610216342076157,2.3912787481
hsa04611,"Platelet activation - Homo sapiens (human)",121,0.43699978561216,2.06853674853014,
hsa04933,"AGE-RAGE signaling pathway in diabetic complications - Homo sapiens (human)",99,0.468
hsa04659,"Th17 cell differentiation - Homo sapiens (human)",106,0.448767316617907,2.068913983
hsa04350,"TGF-beta signaling pathway - Homo sapiens (human)",96,0.456263493912495,2.0954134
hsa05167,"Kaposi sarcoma-associated herpesvirus infection - Homo sapiens (human)",193,0.367666
hsa05132,"Salmonella infection - Homo sapiens (human)",249,0.343102575818557,1.785220272320
hsa04120,"Ubiquitin mediated proteolysis - Homo sapiens (human)",137,-0.402344460833043,-1.94
hsa05014,"Amyotrophic lateral sclerosis - Homo sapiens (human)",332,-0.309369233057186,-1.7055
hsa05410,"Hypertrophic cardiomyopathy - Homo sapiens (human)",90,0.462800473894637,2.09150
hsa04620,"Toll-like receptor signaling pathway - Homo sapiens (human)",102,0.448969238290123,2
hsa04672,"Intestinal immune network for IgA production - Homo sapiens (human)",45,0.572215415
hsa04979,"Cholesterol metabolism - Homo sapiens (human)",50,0.544823183849519,2.2039694591
hsa05016,"Huntington disease - Homo sapiens (human)",278,-0.326922086158876,-1.75923953551
hsa05164,"Influenza A - Homo sapiens (human)",168,0.378693778774119,1.86590398496602,4.103
hsa04022,"cGMP-PKG signaling pathway - Homo sapiens (human)",163,0.377657658499755,1.85831
hsa04658,"Th1 and Th2 cell differentiation - Homo sapiens (human)",90,0.447566775732072,2.0226
hsa05414,"Dilated cardiomyopathy - Homo sapiens (human)",95,0.449233163659201,2.0620290906
hsa04612,"Antigen processing and presentation - Homo sapiens (human)",68,0.492863068007087,2
hsa04064,"NF-kappa B signaling pathway - Homo sapiens (human)",102,0.434677843936516,1.9996
hsa04014,"Ras signaling pathway - Homo sapiens (human)",230,0.335385737362587,1.7364906611
hsa04015,"Rap1 signaling pathway - Homo sapiens (human)",207,0.345966017935254,1.755861516
hsa05131,"Shigellosis - Homo sapiens (human)",217,0.337088763092467,1.7370441300824,1.32514
hsa05217,"Basal cell carcinoma - Homo sapiens (human)",63,0.49642879215965,2.08800677672676
hsa03008,"Ribosome biogenesis in eukaryotes - Homo sapiens (human)",72,-0.454398955384836,-1.
hsa04217,"Necroptosis - Homo sapiens (human)",133,0.384552286098988,1.83861620750705,2.017
hsa04390,"Hippo signaling pathway - Homo sapiens (human)",153,0.380416648204746,1.857854146
hsa03420,"Nucleotide excision repair - Homo sapiens (human)",62,-0.483992454968487,-2.0268486
hsa05134,"Legionellosis - Homo sapiens (human)",57,0.497944487163977,2.07352254947369,3.324
hsa04928,"Parathyroid hormone synthesis, secretion and action - Homo sapiens (human)",106,0.407
hsa04114,"Oocyte meiosis - Homo sapiens (human)",125,-0.381198182167406,-1.80256850542085,
hsa05142,"Chagas disease - Homo sapiens (human)",101,0.417525218912002,1.91961361386009,4.
hsa05412,"Arrhythmogenic right ventricular cardiomyopathy - Homo sapiens (human)",77,0.449997
hsa05310,"Asthma - Homo sapiens (human)",28,0.613782920799301,2.16127149470639,5.2167779
hsa00511,"Other glycan degradation - Homo sapiens (human)",17,0.700235835334243,2.152656549
hsa04914,"Progesterone-mediated oocyte maturation - Homo sapiens (human)",97,-0.40489536275
hsa04270,"Vascular smooth muscle contraction - Homo sapiens (human)",131,0.380975614797928,1.
hsa04714,"Thermogenesis - Homo sapiens (human)",202,-0.322022130440085,-1.66451342279121,
hsa04625,"C-type lectin receptor signaling pathway - Homo sapiens (human)",104,0.3952188589954
hsa05224,"Breast cancer - Homo sapiens (human)",147,0.351187678481808,1.70031564731413,0.0
hsa05130,"Pathogenic Escherichia coli infection - Homo sapiens (human)",194,0.333392605261795,
hsa04392,"Hippo signaling pathway - multiple species - Homo sapiens (human)",28,0.595079461772
hsa04072,"Phospholipase D signaling pathway - Homo sapiens (human)",144,0.355264143843934,1.
hsa04024,"cAMP signaling pathway - Homo sapiens (human)",224,0.310939186169001,1.602661924
hsa05163,"Human cytomegalovirus infection - Homo sapiens (human)",221,0.314844310007967,1.6
hsa04978,"Mineral absorption - Homo sapiens (human)",60,0.451774089194036,1.89951334081888
hsa04062,"Chemokine signaling pathway - Homo sapiens (human)",188,0.323721158639536,1.6288
hsa00120,"Primary bile acid biosynthesis - Homo sapiens (human)",17,0.647193047936524,1.98959
hsa05416,"Viral myocarditis - Homo sapiens (human)",57,0.454729166115205,1.89356686166654,0
hsa00900,"Terpenoid backbone biosynthesis - Homo sapiens (human)",23,-0.585587749739906,-1.9

hsa04650,"Natural killer cell mediated cytotoxicity - Homo sapiens (human)",124,0.35987012467717
hsa04936,"Alcoholic liver disease - Homo sapiens (human)",141,0.341138540853335,1.63528895861
hsa05012,"Parkinson disease - Homo sapiens (human)",236,-0.289962934394904,-1.5304502128281
hsa00531,"Glycosaminoglycan degradation - Homo sapiens (human)",19,0.635822574325157,2.0191
hsa00970,"Aminoacyl-tRNA biosynthesis - Homo sapiens (human)",25,-0.579297015883886,-1.94817
hsa00532,"Glycosaminoglycan biosynthesis - chondroitin sulfate / dermatan sulfate - Homo sapiens (human)",134,-0.341124134410178,-1.63868399201571,0.000
hsa03010,"Ribosome - Homo sapiens (human)",134,-0.341124134410178,-1.63868399201571,0.000
hsa04210,"Apoptosis - Homo sapiens (human)",134,0.339345374186727,1.62222059574501,0.0009
hsa04310,"Wnt signaling pathway - Homo sapiens (human)",168,0.321649564080029,1.5848351280
hsa04924,"Renin secretion - Homo sapiens (human)",67,0.41933259890528,1.77401251022577,0.00
hsa05143,"African trypanosomiasis - Homo sapiens (human)",36,0.503722306589215,1.8879800647
hsa05100,"Bacterial invasion of epithelial cells - Homo sapiens (human)",69,0.418867371409717,1.7
hsa00830,"Retinol metabolism - Homo sapiens (human)",67,0.415601409051981,1.75822748064529
hsa04520,"Adherens junction - Homo sapiens (human)",88,0.374162961937652,1.68033185820373,
hsa04927,"Cortisol synthesis and secretion - Homo sapiens (human)",65,0.413792147681207,1.7421
hsa04657,"IL-17 signaling pathway - Homo sapiens (human)",92,0.370129560536447,1.68268758521
hsa03018,"RNA degradation - Homo sapiens (human)",75,-0.38769119014366,-1.68609278295943,0
hsa05204,"Chemical carcinogenesis - DNA adducts - Homo sapiens (human)",66,0.401331030160948
hsa05135,"Yersinia infection - Homo sapiens (human)",134,0.328071832636654,1.56832809364933,
hsa04623,"Cytosolic DNA-sensing pathway - Homo sapiens (human)",70,0.404606335293685,1.7310
hsa04371,"Apelin signaling pathway - Homo sapiens (human)",136,0.323680232754053,1.54991941
hsa00982,"Drug metabolism - cytochrome P450 - Homo sapiens (human)",67,0.401409920567086,1.
hsa03022,"Basal transcription factors - Homo sapiens (human)",44,-0.435914078450349,-1.7090968
hsa00514,"Other types of O-glycan biosynthesis - Homo sapiens (human)",44,0.45198503047337,1.7
hsa04216,"Ferroptosis - Homo sapiens (human)",41,0.453737565384912,1.77807921962891,0.0031
hsa04750,"Inflammatory mediator regulation of TRP channels - Homo sapiens (human)",98,0.339422
hsa00980,"Metabolism of xenobiotics by cytochrome P450 - Homo sapiens (human)",72,0.38247367
hsa05226,"Gastric cancer - Homo sapiens (human)",147,0.305018982784044,1.47678458252748,0.0
hsa04926,"Relaxin signaling pathway - Homo sapiens (human)",128,0.328632357371299,1.56684125
hsa03320,"PPAR signaling pathway - Homo sapiens (human)",74,0.363261031771478,1.5759437994
hsa04916,"Melanogenesis - Homo sapiens (human)",101,0.341609442782512,1.57058330200332,0.
hsa04970,"Salivary secretion - Homo sapiens (human)",91,0.358587136460195,1.62532107202099,0
hsa00910,"Nitrogen metabolism - Homo sapiens (human)",17,0.593383090214692,1.824171130805
hsa05166,"Human T-cell leukemia virus 1 infection - Homo sapiens (human)",220,0.27760449468038
hsa04925,"Aldosterone synthesis and secretion - Homo sapiens (human)",98,0.334253520634183,1.
hsa05169,"Epstein-Barr virus infection - Homo sapiens (human)",198,0.287944450295629,1.459709
hsa03450,"Non-homologous end-joining - Homo sapiens (human)",12,-0.65342845635325,-1.815050
hsa04913,"Ovarian steroidogenesis - Homo sapiens (human)",51,0.41881381518758,1.69924408996
hsa05120,"Epithelial cell signaling in Helicobacter pylori infection - Homo sapiens (human)",70,0.389
hsa03020,"RNA polymerase - Homo sapiens (human)",31,-0.481001127706934,-1.73484828785901,
hsa04662,"B cell receptor signaling pathway - Homo sapiens (human)",81,0.358548688457795,1.593
hsa01240,"Biosynthesis of cofactors - Homo sapiens (human)",147,-0.284682792007118,-1.3895241
hsa05162,"Measles - Homo sapiens (human)",138,0.305010706793263,1.45910517223444,0.007511
hsa00512,"Mucin type O-glycan biosynthesis - Homo sapiens (human)",34,0.453448698979534,1.67
hsa00590,"Arachidonic acid metabolism - Homo sapiens (human)",57,0.382475628549113,1.592691
hsa04972,"Pancreatic secretion - Homo sapiens (human)",100,0.337151709508411,1.546840211761
hsa00513,"Various types of N-glycan biosynthesis - Homo sapiens (human)",38,0.444865546853433,
hsa04614,"Renin-angiotensin system - Homo sapiens (human)",23,0.506764760446514,1.722532436
hsa05222,"Small cell lung cancer - Homo sapiens (human)",92,0.335885696208062,1.527007705773
hsa04915,"Estrogen signaling pathway - Homo sapiens (human)",137,0.297277255465559,1.423526

hsa00190,"Oxidative phosphorylation - Homo sapiens (human)",105,-0.32345277939589,-1.4941433
hsa01521,"EGFR tyrosine kinase inhibitor resistance - Homo sapiens (human)",79,0.34780877363364
hsa05215,"Prostate cancer - Homo sapiens (human)",97,0.326903683969871,1.49679029393995,0.0
hsa04613,"Neutrophil extracellular trap formation - Homo sapiens (human)",108,0.31584320088663
hsa04261,"Adrenergic signaling in cardiomyocytes - Homo sapiens (human)",152,0.28659901495607
hsa04934,"Cushing syndrome - Homo sapiens (human)",155,0.298697179048267,1.4601249363076,
hsa00600,"Sphingolipid metabolism - Homo sapiens (human)",50,0.392733379792171,1.588721626,
hsa04976,"Bile secretion - Homo sapiens (human)",89,0.337360048827947,1.52607106343293,0.01
hsa00670,"One carbon pool by folate - Homo sapiens (human)",19,-0.534778454077316,-1.6860616
hsa04666,"Fc gamma R-mediated phagocytosis - Homo sapiens (human)",93,0.321660468890832,1.4
hsa04550,"Signaling pathways regulating pluripotency of stem cells - Homo sapiens (human)",143,0.0
hsa04911,"Insulin secretion - Homo sapiens (human)",86,0.321630522363957,1.44281565514137,0.
hsa05218,"Melanoma - Homo sapiens (human)",72,0.346407511643207,1.49267643473663,0.02283
hsa04068,"FoxO signaling pathway - Homo sapiens (human)",129,0.292141764854321,1.396555875
hsa05207,"Chemical carcinogenesis - receptor activation - Homo sapiens (human)",195,0.265143714

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389e-06,2776,"tags=42%, list=15%, signal=36%", "2152/2308/7040/4313/183/2335/7043/7056/5054
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