

ID	Description	GeneRatio	BgRatio	pvalue	p.adjust	qvalue	geneID	Count
hsa04520	Adherens j	17/406	71/7946	6.4E-08	1.77E-05	1.53E-05	TGFBR1/EF	17
hsa04115	p53 signali	13/406	72/7946	6.01E-05	0.00829	0.007177	CCNG1/CC	13
hsa04068	FoxO signa	18/406	131/7946	0.00011	0.010143	0.008782	RBL2/TGFB	18
hsa05203	Viral carcin	23/406	201/7946	0.000226	0.01118	0.00968	RBL2/NFKE	23
hsa04151	PI3K-Akt si	34/406	354/7946	0.00026	0.01118	0.00968	PRLR/RBL2	34
hsa05100	Bacterial ir	12/406	73/7946	0.00029	0.01118	0.00968	ARPC5L/IT	12
hsa04141	Protein prc	20/406	166/7946	0.00029	0.01118	0.00968	SAR1A/RAI	20
hsa04152	AMPK sign	16/406	120/7946	0.000373	0.01118	0.00968	SIRT1/PPP	16
hsa05215	Prostate ca	14/406	97/7946	0.000378	0.01118	0.00968	ERBB2/NF	14
hsa05131	Shigellosis	25/406	236/7946	0.000405	0.01118	0.00968	PFN1/ARPC	25
hsa04211	Longevity r	13/406	89/7946	0.000541	0.012453	0.010781	NFKB1/ATF	13
hsa04110	Cell cycle	16/406	124/7946	0.000541	0.012453	0.010781	RBL2/CDK7	16
hsa03015	mRNA surv	13/406	91/7946	0.000672	0.013815	0.011961	PPP2CA/HI	13
hsa05205	Proteoglyc	22/406	204/7946	0.000701	0.013815	0.011961	FZD7/CAM	22
hsa04213	Longevity r	10/406	62/7946	0.001071	0.019712	0.017066	SIRT1/IGF1	10
hsa04510	Focal adhe	21/406	199/7946	0.001218	0.019911	0.017238	ERBB2/ITG	21
hsa04066	HIF-1 signa	14/406	109/7946	0.001242	0.019911	0.017238	CAMK2D/E	14
hsa04218	Cellular sei	18/406	160/7946	0.001299	0.019911	0.017238	RBL2/TGFB	18
hsa03013	RNA trans	18/406	165/7946	0.001844	0.026784	0.023188	EIF3H/PAIF	18
hsa04114	Oocyte me	15/406	128/7946	0.00215	0.029675	0.025691	CAMK2D/C	15
hsa05132	Salmonella	11/406	80/7946	0.002344	0.030807	0.026672	PFN1/ARPC	11
hsa04922	Glucagon s	13/406	106/7946	0.002787	0.034334	0.029725	CAMK2D/A	13
hsa04350	TGF-beta s	12/406	94/7946	0.002861	0.034334	0.029725	TGFBR1/TF	12
hsa05130	Pathogenic	20/406	202/7946	0.003364	0.038684	0.03349	ARPC5L/NF	20
hsa04010	MAPK sign	26/406	295/7946	0.0045	0.048599	0.042075	TGFBR1/EF	26
hsa01230	Biosynthes	10/406	75/7946	0.004578	0.048599	0.042075	BCAT2/GO	10