

Gene identifier	Log2 fold change (HEPG2 vs. non-HEPG2)	Adjusted p-value	Product name
Smp_049850	1.01	5.22E-11	hypoxanthine guanine phosphoribosyltransferase
Smp_181510	0.96	1.00E-09	hypothetical protein
Smp_159810	0.94	2.75E-09	MEG-2 (ESP15) family
Smp_187410	0.9	1.06E-08	hypothetical protein
Smp_180620	0.9	1.84E-08	MEG 17
Smp_074560	0.89	3.05E-08	hypothetical protein
Smp_212180	0.88	1.43E-11	glucose dehydrogenase (acceptor)
Smp_180330	0.87	2.25E-10	MEG 2 (ESP15) family
Smp_126880	0.82	3.12E-07	hypothetical protein
Smp_136890	0.82	1.55E-07	universal stress protein family M13 unassigned peptidase (M13 family)
Smp_147740	0.82	1.33E-07	MEG-3 (Grail) family
Smp_138070	0.78	9.78E-07	Peptidase M8
Smp_167120	0.76	3.94E-06	Ferritin-2 heavy chain
Smp_047680	0.74	1.32E-06	Ferritin-2 heavy chain
Smp_047660	0.72	9.87E-06	Ferritin-2 heavy chain
Smp_138080	0.7	2.17E-05	MEG-3 (Grail) family
Smp_193400	0.7	1.19E-05	hypothetical protein
Smp_062900	0.7	5.39E-07	tryparedoxin peroxidase
Smp_074570	0.69	1.43E-05	hypothetical protein
Smp_148820	0.67	2.51E-07	hypoxanthine guanine phosphoribosyltransferase
Smp_138060	0.66	9.65E-06	MEG-3 (Grail) family
Smp_179860	0.66	8.97E-05	MEG-2 (ESP15) family
Smp_203400	0.65	2.56E-05	rhodopsin orphan GPCR
Smp_188780	0.65	4.59E-05	hypothetical protein
Smp_159800	0.65	1.18E-04	MEG-2 (ESP15) family
Smp_170610	0.64	9.72E-05	rhodopsin orphan GPCR
Smp_133840	0.64	1.18E-07	lipoygenase y domain containing
Smp_164170	0.62	1.24E-04	endoglycoceramidase
Smp_152590	0.62	2.19E-04	MEG-10
Smp_145920	0.61	1.84E-07	protein tyrosine sulfotransferase
Smp_124750	0.61	2.60E-04	hypothetical protein
Smp_155250	0.61	2.01E-04	nuclear receptor subfamily 2 group E member
Smp_063150	0.61	6.33E-06	hypothetical protein
Smp_182090	0.6	4.07E-04	hypothetical protein
Smp_170480	0.59	2.78E-04	family M13 unassigned peptidase (M13 family)
Smp_004710	0.59	6.31E-05	hypothetical protein
Smp_177380	0.59	3.23E-04	hypothetical protein

Smp_201260	0.58	4.02E-04	hypothetical protein
Smp_144790	0.58	2.81E-04	calmodulin
Smp_199230	0.58	6.93E-04	hypothetical protein
Smp_105370	0.57	4.64E-05	cathepsin B peptidase (C01 family)
Smp_194280	0.57	2.55E-05	hypothetical protein
Smp_186780	0.56	1.01E-05	hypothetical protein
Smp_153070	0.56	5.19E-05	hypothetical protein
Smp_169100	0.56	1.45E-04	monocarboxylate transporter
Smp_017720	0.56	1.34E-04	zinc finger protein
Smp_057710	0.55	7.46E-05	hypothetical protein
Smp_033010	0.55	6.16E-08	16 kda calcium binding protein
Smp_156600	0.55	3.83E-04	hypothetical protein
Smp_197150	0.54	1.63E-04	high affinity cGMP specific 3'
Smp_041700	0.54	1.56E-05	G protein coupled receptor fragment
Smp_132210	0.54	1.02E-10	f box only protein 32
Smp_154250	0.53	2.29E-05	ring finger protein unkempt
Smp_052880	0.53	1.98E-03	hypothetical protein
Smp_134500	0.53	1.51E-03	calcium:calmodulin dependent 3'
Smp_012920	0.52	4.23E-04	class A rhodopsin G protein coupled
Smp_071050	0.52	1.61E-05	hypothetical protein
Smp_157100	0.52	3.65E-05	sphingoid long chain base kinase
Smp_201700	0.52	2.78E-04	hypothetical protein
Smp_163740	0.52	6.45E-04	nesprin 1
Smp_159830	0.52	3.10E-03	MEG-2 (ESP15) family
Smp_179110	0.52	1.15E-03	purine nucleoside phosphorylase
Smp_022450	0.52	2.64E-04	hypothetical protein ras and EF hand domain containing
Smp_132500	0.51	8.92E-04	protein
Smp_125450	0.51	2.80E-03	retinal guanylyl cyclase 2
Smp_033000	0.51	5.08E-07	calcium-binding protein
Smp_144300	0.5	2.00E-04	tns1 protein