

ID	Description	setSize	enrichment	NES	pvalue	p.adjust	qvalues	rank
hsa01240	Biosynthesis of amino acids	142	0.618439	2.889374	0.001433	0.009559	0.004224	2731
hsa01200	Carbon metabolism	110	0.613343	2.755941	0.001484	0.009559	0.004224	2369
hsa00190	Oxidative phosphorylation	98	0.582297	2.564267	0.001515	0.009559	0.004224	4286
hsa04146	Peroxisome	77	0.660703	2.769133	0.00155	0.009559	0.004224	2614
hsa00983	Drug metabolism - cytochrome P450	69	0.614971	2.517995	0.001558	0.009559	0.004224	1053
hsa05204	Chemical carcinogenesis	69	0.62515	2.559675	0.001558	0.009559	0.004224	1974
hsa04976	Bile secretion	71	0.603031	2.484144	0.001563	0.009559	0.004224	1608
hsa00071	Fatty acid metabolism	42	0.690356	2.554009	0.001577	0.009559	0.004224	1458
hsa03320	PPAR signaling	68	0.486715	1.982446	0.001577	0.009559	0.004224	1953
hsa00310	Lysine degradation	59	0.442005	1.762923	0.001585	0.009559	0.004224	1454
hsa00270	Cysteine and methionine metabolism	46	0.552752	2.079402	0.00159	0.009559	0.004224	4306
hsa00280	Valine, leucine and isoleucine degradation	46	0.777012	2.923052	0.00159	0.009559	0.004224	1811
hsa00980	Metabolism of drugs and xenobiotics	60	0.68006	2.714795	0.00159	0.009559	0.004224	2091
hsa00140	Steroid hormone biosynthesis	50	0.589899	2.255535	0.001592	0.009559	0.004224	1895
hsa00830	Retinol metabolism	50	0.680946	2.603663	0.001592	0.009559	0.004224	2555
hsa04978	Mineral absorption	50	0.643913	2.462063	0.001592	0.009559	0.004224	2227
hsa00010	Glycolysis / Gluconeogenesis	57	0.530474	2.092994	0.001595	0.009559	0.004224	1803
hsa00982	Drug metabolism - cytochrome P450	57	0.727868	2.871817	0.001595	0.009559	0.004224	764
hsa00380	Tryptophan metabolism	40	0.719565	2.618659	0.001597	0.009559	0.004224	1255
hsa00640	Propanoate metabolism	30	0.7555	2.590059	0.0016	0.009559	0.004224	1271
hsa01230	Biosynthesis of nucleosides and nucleotides	65	0.546763	2.207659	0.001603	0.009559	0.004224	2173
hsa00330	Arginine and proline metabolism	48	0.665939	2.512357	0.001605	0.009559	0.004224	1693
hsa00760	Nicotinate and nicotinamide metabolism	31	0.543599	1.88529	0.001605	0.009559	0.004224	1039
hsa00020	Citrate cycle (TCA cycle)	29	0.689358	2.350209	0.001608	0.009559	0.004224	3338
hsa00062	Fatty acid metabolism	25	0.579764	1.901915	0.001608	0.009559	0.004224	3760
hsa00630	Glyoxylate and glycolate metabolism	29	0.79552	2.712144	0.001608	0.009559	0.004224	1255
hsa00620	Pyruvate metabolism	36	0.65353	2.323807	0.00161	0.009559	0.004224	2615
hsa00053	Ascorbate and aldehyde metabolism	23	0.899049	2.911575	0.001618	0.009559	0.004224	1361
hsa00790	Folate biosynthesis	23	0.662317	2.144918	0.001618	0.009559	0.004224	1965
hsa00410	beta-Alanine metabolism	28	0.738414	2.502502	0.001621	0.009559	0.004224	2358
hsa00650	Butanoate metabolism	24	0.783167	2.533822	0.001621	0.009559	0.004224	2358
hsa00260	Glycine, serine and alanine metabolism	35	0.820255	2.88183	0.001623	0.009559	0.004224	1478
hsa00480	Glutathione metabolism	53	0.560926	2.16545	0.001623	0.009559	0.004224	2203
hsa00561	Glycerolipid metabolism	55	0.442077	1.721743	0.001626	0.009559	0.004224	878
hsa01212	Fatty acid metabolism	55	0.540029	2.103234	0.001626	0.009559	0.004224	3509
hsa00350	Tyrosine metabolism	33	0.650796	2.261502	0.001629	0.009559	0.004224	1223
hsa00250	Alanine, aspartate and glutamate metabolism	34	0.648693	2.262751	0.001631	0.009559	0.004224	1223
hsa00860	Porphyria and heme biosynthesis	34	0.671249	2.341432	0.001631	0.009559	0.004224	1002
hsa00040	Pentose and glucuronate interconversions	26	0.818333	2.71063	0.001634	0.009559	0.004224	1763
hsa04966	Collecting duct acid secretion	27	0.668635	2.23543	0.001642	0.009559	0.004224	4181
hsa00220	Arginine biosynthesis	20	0.687544	2.138307	0.001664	0.009559	0.004224	2111
hsa00360	Phenylalanine metabolism	17	0.725374	2.173129	0.001664	0.009559	0.004224	1223
hsa00770	Pantothenate and CoA biosynthesis	20	0.733571	2.281452	0.001664	0.009559	0.004224	825
hsa04964	Proximal tubule acid secretion	20	0.734192	2.283384	0.001664	0.009559	0.004224	2227
hsa04977	Vitamin D metabolism	20	0.6336	1.970538	0.001664	0.009559	0.004224	2312
hsa00340	Histidine metabolism	21	0.803042	2.537354	0.001667	0.009559	0.004224	1575
hsa00450	Selenocompound metabolism	16	0.66971	1.967554	0.001672	0.009559	0.004224	1691
hsa00900	Terpenoid biosynthesis	22	0.626749	1.98498	0.001675	0.009559	0.004224	4843
hsa01210	2-Oxocarboxylate metabolism	18	0.635577	1.911435	0.001704	0.009559	0.004224	2111
hsa00670	One carbon folate metabolism	19	0.619562	1.887175	0.001706	0.009559	0.004224	2669
hsa00120	Primary bile acid metabolism	14	0.714484	2.001	0.001751	0.009559	0.004224	1978
hsa05340	Primary immunodeficiency	34	-0.55304	-2.11676	0.002571	0.009559	0.004224	5043
hsa05330	Allograft rejection	33	-0.58581	-2.22693	0.002577	0.009559	0.004224	5783
hsa05321	Inflammatory bowel disease	55	-0.48496	-2.09143	0.002584	0.009559	0.004224	4402
hsa05310	Asthma	23	-0.55362	-1.88739	0.002604	0.009559	0.004224	5526
hsa05322	Systemic lupus erythematosus	48	-0.50527	-2.09606	0.002639	0.009559	0.004224	5596
hsa04664	Fc epsilon receptor signaling pathway	63	-0.40925	-1.81471	0.002646	0.009559	0.004224	2994

hsa05144	Malaria	47	-0.43865	-1.80844	0.002646	0.009559	0.004224	4156
hsa05221	Acute mye	65	-0.39679	-1.76725	0.002646	0.009559	0.004224	2655
hsa05416	Viral myoc	58	-0.54261	-2.34663	0.00266	0.009559	0.004224	4689
hsa05320	Autoimmu	44	-0.47876	-1.94365	0.002674	0.009559	0.004224	4689
hsa05012	Parkinson	203	0.323691	1.590482	0.002706	0.009559	0.004224	3989
hsa05220	Chronic my	76	-0.38207	-1.75465	0.002755	0.009559	0.004224	4719
hsa04714	Thermoge	189	0.331101	1.609794	0.002762	0.009559	0.004224	5346
hsa05133	Pertussis	71	-0.43975	-1.98296	0.002762	0.009559	0.004224	3454
hsa04662	B cell rece	79	-0.44683	-2.06859	0.00277	0.009559	0.004224	4335
hsa05150	Staphylocc	73	-0.41362	-1.872	0.002778	0.009559	0.004224	5366
hsa05332	Graft-vers	41	-0.47119	-1.86042	0.002778	0.009559	0.004224	5854
hsa04512	ECM-rece	82	-0.4384	-2.04002	0.002817	0.009559	0.004224	4226
hsa04540	Gap juncti	81	-0.37561	-1.74159	0.002825	0.009559	0.004224	3325
hsa05140	Leishmania	74	-0.53913	-2.44314	0.002849	0.009559	0.004224	3524
hsa04640	Hematopo	91	-0.39136	-1.86004	0.002882	0.009559	0.004224	5064
hsa05222	Small cell	90	-0.4008	-1.89676	0.00289	0.009559	0.004224	4226
hsa01522	Endocrine	94	-0.3676	-1.75845	0.002907	0.009559	0.004224	3186
hsa04670	Leukocyte	105	-0.41794	-2.04186	0.002907	0.009559	0.004224	2922
hsa04064	NF-kappa	98	-0.43621	-2.10538	0.002924	0.009559	0.004224	5413
hsa04620	Toll-like	98	-0.40706	-1.96465	0.002924	0.009559	0.004224	4495
hsa04625	C-type lec	98	-0.36949	-1.78334	0.002924	0.009559	0.004224	2669
hsa05146	Amoebiasi	98	-0.42892	-2.07019	0.002924	0.009559	0.004224	4764
hsa04666	Fc gamma	95	-0.40976	-1.96714	0.002959	0.009559	0.004224	2994
hsa04668	TNF signal	108	-0.48309	-2.37199	0.002959	0.009559	0.004224	4499
hsa05145	Toxoplas	108	-0.41522	-2.03878	0.002959	0.009559	0.004224	4226
hsa04061	Viral prote	87	-0.39878	-1.86413	0.002967	0.009559	0.004224	5192
hsa04658	Th1 and Th	86	-0.47652	-2.21727	0.002967	0.009559	0.004224	4719
hsa05235	PD-L1 exp	87	-0.51222	-2.3944	0.002967	0.009559	0.004224	4335
hsa04933	AGE-RAGE	99	-0.36187	-1.74364	0.002976	0.009559	0.004224	4719
hsa05142	Chagas dis	97	-0.39904	-1.91605	0.002985	0.009559	0.004224	4023
hsa04659	Th17 cell	96	-0.48955	-2.3467	0.003003	0.009559	0.004224	4746
hsa04660	T cell rece	96	-0.43507	-2.08554	0.003003	0.009559	0.004224	4335
hsa04650	Natural kill	119	-0.38376	-1.9137	0.003058	0.009559	0.004224	4394
hsa04926	Relaxin sig	119	-0.37375	-1.86381	0.003058	0.009559	0.004224	4715
hsa04611	Platelet act	114	-0.38577	-1.90359	0.003077	0.009559	0.004224	3240
hsa04110	Cell cycle	117	-0.33202	-1.64492	0.003145	0.009559	0.004224	4052
hsa04380	Osteoclast	122	-0.47133	-2.35293	0.003175	0.009559	0.004224	3598
hsa01040	Biosynthes	25	0.562857	1.846451	0.003215	0.009559	0.004224	3509
hsa04550	Signaling p	130	-0.33503	-1.68834	0.003215	0.009559	0.004224	5298
hsa05135	Yersinia inf	130	-0.38045	-1.91722	0.003215	0.009559	0.004224	5386
hsa04072	Phospholip	138	-0.32883	-1.67781	0.003236	0.009559	0.004224	3919
hsa04921	Oxytocin s	138	-0.32357	-1.65097	0.003236	0.009559	0.004224	5021
hsa00970	Aminoacyl	24	0.585357	1.893836	0.003241	0.009559	0.004224	2995
hsa04210	Apoptosis	129	-0.31414	-1.58153	0.003247	0.009559	0.004224	4335
hsa04630	JAK-STAT	139	-0.32916	-1.67552	0.003289	0.009559	0.004224	4844
hsa05162	Measles	132	-0.40495	-2.04568	0.003289	0.009559	0.004224	4862
hsa04614	Renin-ang	20	0.606565	1.886458	0.003328	0.009559	0.004224	1549
hsa00910	Nitrogen n	16	0.614975	1.806746	0.003344	0.009559	0.004224	3962
hsa04621	NOD-like	161	-0.37896	-1.96941	0.003344	0.009559	0.004224	2669
hsa05203	Viral carcin	161	-0.38903	-2.02171	0.003344	0.009559	0.004224	4428
hsa05152	Tuberculos	171	-0.29821	-1.56147	0.003378	0.009559	0.004224	4463
hsa04218	Cellular sei	148	-0.35381	-1.81505	0.00339	0.009559	0.004224	4335
hsa05161	Hepatitis B	157	-0.40255	-2.0834	0.003401	0.009559	0.004224	4444
hsa05202	Transcripti	158	-0.35666	-1.84562	0.003413	0.009559	0.004224	3920
hsa05164	Influenza A	162	-0.38974	-2.0239	0.003425	0.009559	0.004224	5366
hsa05206	MicroRNA	163	-0.40061	-2.07319	0.00346	0.009559	0.004224	4056
hsa05167	Kaposi sarc	175	-0.37164	-1.94696	0.003484	0.009559	0.004224	4339
hsa04360	Axon guid	173	-0.29931	-1.56011	0.003497	0.009559	0.004224	4061

hsa04062	Chemokine	177	-0.37992	-1.996	0.003509	0.009559	0.004224	4719
hsa05130	Pathogenic	184	-0.33676	-1.78799	0.003571	0.009559	0.004224	4023
hsa05170	Human im	191	-0.30514	-1.62752	0.003636	0.009559	0.004224	4335
hsa05163	Human cyt	210	-0.36087	-1.9546	0.00365	0.009559	0.004224	3977
hsa05166	Human T-c	211	-0.42656	-2.31591	0.003676	0.009559	0.004224	4402
hsa04810	Regulation	204	-0.29813	-1.60735	0.00369	0.009559	0.004224	3104
hsa05131	Shigellosis	209	-0.31013	-1.67141	0.00369	0.009559	0.004224	4148
hsa05171	Coronavir	195	-0.36922	-1.97132	0.00369	0.009559	0.004224	4873
hsa04015	Rap1 signa	194	-0.32454	-1.73189	0.003704	0.009559	0.004224	3306
hsa05169	Epstein-Ba	194	-0.38973	-2.07974	0.003704	0.009559	0.004224	5102
hsa04510	Focal adhe	196	-0.35993	-1.92476	0.003717	0.009559	0.004224	2820
hsa05205	Proteoglyc	199	-0.34676	-1.85134	0.003802	0.0097	0.004286	2859
hsa04060	Cytokine-c	249	-0.28026	-1.54778	0.003861	0.009773	0.004318	5202
hsa05132	Salmonella	244	-0.34749	-1.90562	0.003953	0.009927	0.004386	4210
hsa04010	MAPK sign	274	-0.30031	-1.68003	0.004	0.009969	0.004405	2729
hsa04151	PI3K-Akt s	321	-0.267	-1.52504	0.004032	0.009973	0.004406	4231
hsa05165	Human pa	316	-0.29025	-1.64967	0.004065	0.009978	0.004409	4226
hsa05168	Herpes sim	427	-0.3596	-2.10685	0.004405	0.010732	0.004742	5681
hsa05200	Pathways i	492	-0.27596	-1.64596	0.004739	0.011459	0.005063	4232
hsa01523	Antifolate	29	0.529322	1.804602	0.004823	0.011576	0.005115	1295
hsa04975	Fat digesti	35	0.488832	1.717429	0.00487	0.011602	0.005126	6657
hsa00532	Glycosamir	20	-0.5818	-1.90679	0.004988	0.011795	0.005212	4953
hsa05134	Legionello	57	-0.38369	-1.65378	0.005333	0.012522	0.005533	4026
hsa04672	Intestinal ir	42	-0.45387	-1.80451	0.005435	0.012668	0.005597	5526
hsa04115	p53 signali	72	-0.35136	-1.5849	0.00551	0.012751	0.005634	5441
hsa04940	Type I diak	41	-0.46799	-1.8478	0.005556	0.012766	0.005641	4689
hsa05210	Colorectal	85	-0.35278	-1.62856	0.00597	0.013622	0.006019	2655
hsa04721	Synaptic ve	68	0.39329	1.601915	0.006309	0.014295	0.006316	4336
hsa04514	Cell adhesi	136	-0.31576	-1.6039	0.006645	0.014897	0.006582	3620
hsa04145	Phagosome	144	-0.29641	-1.51608	0.006667	0.014897	0.006582	3865
hsa04657	IL-17 signa	86	-0.33076	-1.53904	0.008902	0.019755	0.008729	2729
hsa04919	Thyroid ho	114	-0.30141	-1.48731	0.009231	0.020345	0.00899	2979
hsa05224	Breast can	135	-0.30027	-1.5254	0.009868	0.021604	0.009546	3061
hsa04932	Non-alcoh	137	0.323095	1.503688	0.01	0.021745	0.009608	4759
hsa04330	Notch sign	52	-0.36985	-1.56628	0.010582	0.022766	0.010059	2979
hsa05100	Bacterial in	67	-0.3583	-1.60531	0.01061	0.022766	0.010059	4509
hsa05016	Huntingtor	254	0.287287	1.454768	0.010724	0.022859	0.0101	4189
hsa05414	Dilated car	87	-0.31749	-1.48412	0.011869	0.025135	0.011106	3958
hsa04935	Growth ho	117	-0.29478	-1.46043	0.012579	0.026464	0.011693	3311
hsa05412	Arrhythmo	68	-0.33784	-1.51647	0.013587	0.028354	0.012528	2392
hsa04950	Maturity of	19	0.537327	1.636688	0.013652	0.028354	0.012528	1114
hsa04066	HIF-1 sign	100	-0.30689	-1.47887	0.014925	0.030801	0.013609	2662
hsa00730	Thiamine r	14	0.595226	1.667005	0.015762	0.032322	0.014281	1739
hsa00030	Pentose ph	28	0.478261	1.620836	0.016207	0.033027	0.014593	2062
hsa00601	Glycosphir	27	0.473136	1.581823	0.01642	0.033251	0.014692	4687
hsa05160	Hepatitis C	144	-0.28114	-1.43799	0.016667	0.03354	0.01482	4339
hsa05231	Choline me	94	-0.31383	-1.50124	0.017442	0.034884	0.015413	3728
hsa04350	TGF-beta s	87	-0.31007	-1.44944	0.017804	0.03539	0.015637	4053
hsa04974	Protein dig	87	0.355001	1.536533	0.018045	0.03565	0.015752	977
hsa05211	Renal cell c	65	-0.33181	-1.47786	0.018519	0.036364	0.016067	4335
hsa04912	GnRH sign	89	-0.30943	-1.46117	0.020408	0.039833	0.0176	3311
hsa05215	Prostate ca	97	-0.29376	-1.41053	0.020896	0.04054	0.017912	4150
hsa05110	Vibrio chol	49	0.396331	1.503971	0.024116	0.046509	0.02055	1751
hsa00061	Fatty acid l	16	0.533991	1.568821	0.025084	0.048089	0.021248	3118
hsa05323	Rheumatoi	90	-0.29922	-1.41604	0.026012	0.049575	0.021904	4023

leading_edcore_enrichment

tags=51%, 210/8564/8836/23475/54490/10327/29968/3242/224/219/217/8942/55163/54575/205/
tags=54%, 8802/414328/2653/51179/1962/229/2203/29968/4329/26275/48/5096/5313/6470/1892/
tags=63%, 127124/4706/529/155066/29796/525/1337/245972/27089/9167/528/64077/4695/5061/
tags=60%, 1610/8309/3155/51268/51179/1962/30/5827/11001/10901/5264/26061/8528/6647/104/
tags=36%, 2938/2940/7172/1807/54490/1576/54575/54658/54657/54600/54576/54659/54579/545/
tags=43%, 2938/2940/130/54490/1576/1551/54575/54658/54657/54600/54576/54659/54579/545/
tags=41%, 123264/1244/6555/200931/54490/1576/9376/54575/54658/54657/54600/54576/54659/
tags=55%, 130/36/1962/224/30/219/217/501/1892/10455/223/128/3033/34/38/51/10449/2639/33/
tags=34%, 8309/5106/2168/1962/30/11001/345/1582/2710/34/51/364/5105/1593/33/81616/1376/
tags=29%, 51268/1962/224/219/217/8424/501/1892/223/123688/3033/55526/38/2639/51166/39/
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