

term_ID	description	frequency	uniqueness	dispensabil	eliminated	n.genes	coeff	odds.ratio
GO:0006415	translation	0.18%	0.785	0	0	89	-0.18319	0.320309
GO:0006414	translation	0.72%	0.879	0.517	0	99	-0.17377	0.339617
GO:0006614	SRP-depen	0.10%	0.747	0.454	0	106	-0.161	0.367666
GO:0000184	nuclear-tra	0.00%	0.886	0.109	0	114	-0.15654	0.378006
GO:0006413	translation	0.27%	0.887	0.536	0	150	-0.14015	0.418542
GO:0019083	viral transc	0.06%	0.874	0.218	0	152	-0.13186	0.440678
GO:0006401	RNA catabo	0.27%	0.854	0.559	0	208	-0.11284	0.495959
GO:0044033	multi-orgar	0.23%	0.951	0.519	0	168	-0.11957	0.47566
GO:0006955	immune re	0.38%	0.66	0	0	921	0.059328	1.445854
GO:0006412	translation	4.70%	0.86	0.555	0	369	-0.08115	0.603937
GO:0019058	viral life cy	1.21%	0.922	0.677	0	283	-0.08231	0.599572
GO:0022411	cellular cor	0.29%	0.859	0.496	0	361	-0.06996	0.647404
GO:0016071	mRNA met	0.49%	0.907	0.271	0	492	-0.06107	0.684196
GO:0001775	cell activati	0.05%	0.928	0.072	0	521	0.055916	1.415516
GO:0006952	defense res	0.57%	0.876	0.412	0	911	0.044453	1.318185
GO:0006364	rRNA proce	0.76%	0.827	0.515	0	56	-0.1181	0.480026
GO:0016072	rRNA meta	0.76%	0.899	0.682	0	60	-0.11564	0.487418
GO:1902578	single-orga	12.73%	0.845	0.397	0	361	-0.06181	0.681065
GO:1902580	single-orga	0.30%	0.821	0.634	0	361	-0.06181	0.681065
GO:0006521	regulation	0.00%	0.796	0.056	0	54	0.106921	1.943469
GO:0050900	leukocyte r	0.01%	0.663	0.697	0	215	0.064356	1.491744
GO:0044270	cellular nitr	6.23%	0.862	0.644	0	938	-0.03485	0.805279
GO:0033238	regulation	0.00%	0.835	0.478	0	62	0.093335	1.786108
GO:0070661	leukocyte p	0.01%	0.913	0.037	0	109	0.077243	1.616127
GO:0046395	carboxylic a	1.14%	0.799	0.321	0	147	-0.07043	0.64553
GO:0034660	ncRNA met	3.21%	0.892	0.392	0	197	-0.06321	0.675139
GO:0042590	antigen prc	0.00%	0.785	0.574	0	75	0.086647	1.713392
GO:0010676	positive reg	0.00%	0.729	0.193	0	28	-0.11404	0.492264
GO:0044282	small mole	1.35%	0.853	0.537	0	181	-0.06483	0.66837
GO:0009064	glutamine f	1.29%	0.843	0.496	0	29	-0.10997	0.50489
GO:0061024	membrane	1.01%	0.92	0.582	0	653	-0.0369	0.795096
GO:0006396	RNA proces	2.78%	0.889	0.488	0	469	-0.04159	0.772244
GO:0045923	positive reg	0.00%	0.714	0.677	0	15	-0.12366	0.463702
GO:0015949	nucleobase	0.01%	0.906	0.114	0	18	0.122782	2.144789
GO:0031929	TOR signali	0.01%	0.772	0.297	0	52	-0.08945	0.573558
GO:0055114	oxidation-r	15.04%	0.908	0.21	0	431	-0.0405	0.777506
GO:0051340	regulation	0.00%	0.839	0.463	0	100	0.070503	1.549836
GO:0048208	COPII vesic	0.00%	0.779	0.65	0	13	0.130692	2.252864
GO:0051438	regulation	0.00%	0.754	0.258	0	98	0.070296	1.54784
GO:0042752	regulation	0.01%	0.867	0.154	0	55	-0.08582	0.586629
GO:0019884	antigen prc	0.00%	0.777	0.617	0	159	0.058341	1.437016
GO:0010885	regulation	0.00%	0.766	0.142	0	12	-0.12559	0.458184
GO:0055088	lipid homeo	0.01%	0.851	0.328	0	56	-0.08508	0.589343
GO:0015980	energy deri	2.41%	0.865	0.1	0	255	-0.04914	0.736819
GO:0030490	maturation	0.00%	0.884	0.592	0	10	-0.12977	0.446421
GO:0009607	response to	0.47%	0.889	0.42	0	332	0.042581	1.302943

GO:0044092	negative re	0.39%	0.805	0.583	0	622	0.0325	1.223819
GO:0000209	protein pol	0.01%	0.886	0.68	0	167	0.055127	1.408597
GO:0034097	response to	0.10%	0.866	0.355	0	418	0.03742	1.261813
GO:0042108	positive reg	0.00%	0.685	0.56	0	30	0.099707	1.858261
GO:0034369	plasma lipi	0.00%	0.77	0.508	0	17	-0.1129	0.495786
GO:0034367	macromole	0.00%	0.937	0.509	0	17	-0.1129	0.495786
GO:0070979	protein K11	0.00%	0.895	0.62	0	26	0.103917	1.907515
GO:0070328	triglyceride	0.00%	0.859	0.57	0	15	-0.11607	0.486107
GO:0010828	positive reg	0.00%	0.715	0.533	0	22	-0.10603	0.517394
GO:0032091	negative re	0.00%	0.84	0.595	0	36	0.09302	1.782619
GO:2000505	regulation i	0.00%	0.815	0.416	0	11	-0.12217	0.468021
GO:0043299	leukocyte c	0.00%	0.652	0.623	0	21	0.108268	1.959806
GO:0008089	anterograd	0.00%	0.812	0.589	0	22	0.106297	1.935945
GO:0002444	myeloid leu	0.00%	0.772	0.632	0	23	0.104055	1.909158
GO:0009250	glucan bios	0.15%	0.83	0.463	0	35	-0.09073	0.569026
GO:0016337	single orga	0.02%	0.949	0.038	0	200	0.04814	1.348738
GO:0086011	membrane	0.00%	0.796	0.292	0	10	0.130286	2.247178
GO:0001816	cytokine pr	0.13%	0.892	0.579	0	347	0.037933	1.265847
GO:0006629	lipid metab	3.09%	0.907	0.156	0	793	-0.0269	0.846037
GO:0032615	interleukin	0.00%	0.907	0.65	0	26	0.098524	1.844649
GO:0032655	regulation i	0.00%	0.811	0.652	0	26	0.098524	1.844649
GO:0007492	endoderm	0.00%	0.967	0	0	45	0.082528	1.670093
GO:1901605	alpha-amin	4.27%	0.829	0.684	0	127	-0.05768	0.698765
GO:0010743	regulation i	0.00%	0.82	0.39	0	25	-0.09666	0.548427
GO:0043648	dicarboxyli	1.12%	0.858	0.488	0	46	-0.08113	0.60401
GO:0006091	generation	3.22%	0.95	0.058	0	319	-0.03883	0.785584
GO:0098602	single orga	0.02%	0.95	0.608	0	232	0.043443	1.309943
GO:0001704	formation i	0.01%	0.892	0.594	0	61	0.073207	1.576095
GO:0045017	glycerolipic	0.21%	0.857	0.686	0	202	-0.04709	0.746299
GO:0006968	cellular def	0.00%	0.903	0.253	0	42	0.082986	1.674857
GO:0016458	gene silenc	0.03%	0.776	0.219	0	62	-0.07288	0.635751
GO:0032006	regulation i	0.01%	0.751	0.364	0	42	-0.08231	0.599564
GO:0043467	regulation i	0.01%	0.825	0.514	0	46	-0.08002	0.608183
GO:0044255	cellular lipi	2.58%	0.847	0.158	0	626	-0.02826	0.838922
GO:0030521	androgen r	0.00%	0.783	0.32	0	52	0.075635	1.60006
GO:0070672	response to	0.00%	0.895	0.61	0	11	0.121421	2.126722
GO:0050817	coagulation	0.02%	0.904	0.628	0	440	0.031495	1.216197
GO:0090322	regulation i	0.00%	0.854	0.194	0	10	0.123305	2.151773
GO:0090077	foam cell d	0.00%	0.915	0.513	0	28	-0.09038	0.57027
GO:0015804	neutral am	0.11%	0.869	0.153	0	17	-0.10304	0.527122
GO:0042060	wound hea	0.03%	0.896	0.454	0	490	0.029656	1.202374
GO:0016192	vesicle-mei	0.38%	0.915	0.234	0	714	0.02505	1.168447
GO:0010876	lipid localiz	0.10%	0.881	0.555	0	155	-0.04902	0.737395
GO:0070647	protein mo	0.13%	0.893	0.393	0	533	0.028076	1.19063
GO:0009066	aspartate fi	0.86%	0.849	0.663	0	28	-0.08851	0.576931
GO:0002886	regulation i	0.00%	0.698	0.669	0	10	0.120397	2.113239
GO:0031333	negative re	0.01%	0.792	0.596	0	47	0.074663	1.590422

GO:0051262	protein tet	0.03%	0.922	0.651	0	53	-0.07222	0.638372
GO:0071350	cellular res	0.00%	0.877	0.605	0	10	0.120063	2.108851
GO:0031330	negative re	0.07%	0.749	0.601	0	57	-0.07027	0.646162
GO:0050435	beta-amylc	0.00%	0.955	0.115	0	15	0.105949	1.931754
GO:0046903	secretion	0.63%	0.868	0.306	0	460	0.028797	1.195978
GO:0055081	anion hom	0.03%	0.841	0.658	0	16	-0.10063	0.535066
GO:0006082	organic aci	9.36%	0.854	0.353	0	716	-0.02388	0.862107
GO:0006887	exocytosis	0.03%	0.811	0.555	0	184	0.04222	1.300018
GO:0006536	glutamate i	0.27%	0.86	0.589	0	14	-0.10317	0.526665
GO:0051084	'de novo' p	0.01%	0.916	0.291	0	36	0.079165	1.635554
GO:0008625	extrinsic ap	0.00%	0.772	0.324	0	49	0.070393	1.548777
GO:0006954	inflammato	0.10%	0.873	0.546	0	252	0.036269	1.252819
GO:1901379	regulation i	0.01%	0.753	0.609	0	23	0.090694	1.75704
GO:0009057	macromole	1.64%	0.882	0.502	0	808	-0.02179	0.873365
GO:0061384	heart trabe	0.00%	0.94	0.408	0	10	-0.11013	0.50438
GO:0018196	peptidyl-as	0.00%	0.914	0.398	0	94	0.054164	1.400186
GO:1902600	hydrogen ic	1.80%	0.823	0.49	0	37	-0.07704	0.619538
GO:0006508	proteolysis	3.71%	0.905	0.578	0	705	0.022549	1.150428
GO:0010469	regulation i	0.01%	0.726	0.619	0	51	0.068331	1.529058
GO:0006112	energy rese	0.13%	0.889	0.642	0	125	-0.04914	0.736831
GO:0050878	regulation i	0.02%	0.786	0.642	0	474	0.026877	1.181792
GO:0042107	cytokine m	0.01%	0.937	0.231	0	54	0.066659	1.513249
GO:0034341	response to	0.00%	0.694	0.681	0	97	0.053056	1.39058
GO:1901685	glutathione	0.00%	0.952	0.027	0	23	-0.08826	0.577802
GO:1901687	glutathione	0.00%	0.941	0.086	0	23	-0.08826	0.577802
GO:1901264	carbohydr	0.23%	0.866	0.427	0	26	-0.08499	0.589682
GO:0071158	positive reg	0.00%	0.756	0.51	0	78	0.057485	1.429388
GO:0009611	response to	0.11%	0.889	0.511	0	618	0.023607	1.158013
GO:0006911	phagocytos	0.00%	0.848	0.577	0	13	0.105927	1.931494
GO:0034440	lipid oxidat	0.05%	0.872	0.602	0	61	-0.06418	0.671098
GO:0016042	lipid catab	0.18%	0.845	0.675	0	120	-0.04942	0.735575
GO:0071331	cellular res	0.00%	0.865	0.68	0	24	-0.08692	0.582646

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