Disorder related MF keywords (z-skor > 0.2)														
#	name	n	avg_len	avg_dis	Z	p								
0	DNA-binding	6518	546.53	0.87	46.90	1		Disorder related MF terms (obtained from KW 2 GO mapping)						
1	Developmental protein	3897	655.21	0.86	31.10	1	#	пате	n	avg_len	avg_dis	Z	p	
2	Activator	2574	600.51	0.88	28.12	1	0	DNA binding	8254	574.89	0.86	46.13	1	
3	Repressor	1988	589.29	0.85	22.63	1	1	sequence-specific DNA binding	3836	536.95	0.92	41.77	1	
4	RNA-binding	2728	575.76	0.76	16.62	1	2	chromatin binding	1538	713.64	0.93	23.54	1	
5	Chromatin regulator	1038	847.06	0.90	13.91	1	3	RNA binding	5797	565.80	0.73	19.79	1	
6	Ribonucleoprotein	1886	272.29	0.60	13.39	1	4	receptor binding	4109	529.68	0.68	14.04	1	
7	Serine/threonine-protein ki	1782	802.24	0.84	11.56	1	5	protein serine/threonine ki	1942	784.87	0.84	12.88	1	
8	Chaperone	937	430.43	0.71	10.02	1	6	structural molecule activity	2884	558.88	0.64	10.84	1	
9	Ribosomal protein	1408	186.38	0.53	9.34	1	-i → 7	structural constituent of r	1350	182.95	0.53	9.25	1	
10	Growth factor	398	299.63	0.70	8.98	1	8	calmodulin binding	710	1079.79	0.88	9.22	1	
11	Protein kinase inhibitor	49	337.20	0.96	8.34	1	9	growth factor activity	458	312.38	0.70	8.99	1	
12	Calmodulin-binding	520	1229.00	0.90	7.57	1	10	protein kinase inhibitor ac	237	401.60	0.82	8.94	1	
13	Hormone	338	221.13	0.59	7.24	1	11	hormone activity	452	218.09	0.61	8.84	1	
14	Cyclin	133	422.71	0.87	7.18	1	1 12	cell adhesion molecule binding	1090	805.74	0.80	8.60	1	
15	Signal transduction inhibitor	115	408.43	0.84	6.76	1	1 1 13	GTPase activator activity	633	820.60	0.87	8.15	1	
16	Guanine-nucleotide releasin	319	1144.39	0.96	6.40	1	14	actin binding	1228	1000.47	0.80	7.73	1	
17	GTPase activation	424	867.35	0.88	6.28	1	15	cyclin-dependent protein se	142	357.65	0.77	6.58	1	
18	Growth factor binding	50	593.98	1	6.09	1	16	protein tyrosine kinase act	504	933.02	0.86	6.26	1	
19	Neuropeptide	105	234.96	0.68	5.99	1	\ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \	guanyl-nucleotide exchange	575	990.55	0.86	6.22	1	
20	Potassium channel	191	621.52	0.85	5.10	1	18	protein phosphatase inhibit	89	425.83	0.83	6.19	1	
21	Calcium channel	193	1397.77	0.93	5.09	1	19	heparin binding	403	652.76	0.76	6.01	1	
22	Protein phosphatase inhibitor	64	352.86	0.81	5.07	1	20	neuropeptide hormone activity	117	158.64	0.65	5.88	1	
23	Tyrosine-protein kinase	376	863.00	0.89	5.04	1	21	microtubule motor activity	225	1394.78	0.97	5.65	1	
24	Mitogen	137	286.03	0.68	4.80	1	22	enzyme inhibitor activity	1059	462.66	0.63	5.60	1	
25	Vasoactive	46	267.00	0.76	4.29	1	23	potassium channel activity	339	612.39	0.79	4.88	1	
26	Heparin-binding	221	650.97	0.73	3.88	1	// \ 24	damaged DNA binding	284	680.80	0.80	4.31	1	
27	Muscle protein	193	920.01	0.73	3.81	1	25	growth factor binding	351	737.85	0.79	4.20	1	
28	Actin-binding	837	974.92	0.77	3.80	1	/\ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \	chloride channel inhibitor	25	827.64	0.96	3.96	1	
29	Amphibian defense peptide	49	85.80	0.53	3.34	1	27	calcium channel activity	327	1205.13	0.86	3.54	1	
30	Helicase	739	1086.05	0.87	3.29	1	28	morphogen activity	23	331.57	0.83	3.20	1	
31	Prion	22	497.05	0.91	3.06	1	29	helicase activity	766	1068.02	0.86	3.15	1	
32	Ion channel	1027	861.88	0.76	2.89	1	30	motor activity	447	1275.57	0.86	3.10	1	
33	Voltage-gated channel	386	816.98	0.78	2.87	1	31	voltage-gated ion channel a	501	767.73	0.76	3.06	1	
34	Viral nucleoprotein	39	1202.79	0.90	2.46	1	32	rRNA binding	541	266.55	0.52	3.00	1	
35	Tumor antigen	26	428.81	0.77	2.25	0.99	33	calcium ion binding	1878	859.00	0.70	2.80	1	
36	Exonuclease	239	725.99	0.75	2.19	0.99	, 34	RNA-directed DNA polymerase	75	1427.97	0.97	2.72	1	
37	Segmentation polarity protein	24	712.75	0.92	2.17	0.99	35	kinase activity	3593	726.02	0.71	2.70	1	
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							37	exonuclease activity	371	749.41	0.75	2.40	0.99	
							38	translation initiation fact	268	482.02	0.65	1.99	0.97	
							39	acetylcholine-gated cation	60	504.17	0.75	1.99	0.99	
							40	metalloendopeptidase inhibi	40	323.35	0.65	1.86	0.98	
							41	cysteine-type endopeptidase	153	424.63	0.58	1.48	0.95	
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