a									Predviđeno uređene MF ključ	ne reči i	z ovog rad	a (z-skor	<u>-02)</u>		
10	9 MF ključnih reči sa najznačajn	niiom pr	edviđenom	uređenoš	ću (org. 2	2007)	1	#	name	ne reci ii		a (z-skor avg_dis		р	
#			n families					0	Oxidoreductase	4126	472.25	0.28	-41.35	0	
0	Oxidoreductase	14995	992	376.63	-29.54	0		1	Hydrolase	7564	614.81	0.51	-26.99	0	
1	Transferase	26525	1606	445.17	-24.25	0		2	Lyase	1431	481.37	0.30	-23.62	0	
2	Lyase	7262	347	377.92	-22.64	0		3	Monooxygenase	555	503.36	0.20	-20.29	0	
3	Hydrolase	20464	1995	430.68	-21.75	0		4	Transferase	8846	631.95	0.55	-19.72	0	
4	Isomerase	4487	220	383.98	-14.18	0	\ / 』	5	Ligase	995	693.30	0.46	-18.05	0	
5	Glycosidase	1826	244	444.73	-13.98	0		6	Glycosyltransferase	1134	551.26	0.40	-17.04	0	
6	Glycosyltransferase	2950	261	437.53	-12.51	0		7	Glycosidase	697	570.50	0.37	-16.81	0	
7	Acyltransferase	2239	179	402.83	-10.85	0		8	Isomerase	931	422.72	0.35	-13.60	0	
8	Methyltransferase	3524	224	349.60	-10.53	0	\	9	Protease	1863	674.42	0.54	-13.20	0	
9	Kinase	7017	322	448.29	-10.22	0	X/ \	10	Transducer	1703	482.28	0.41	-12.56	0	
10	Ligase	8010	230	529.41	-10.06	0		11	G-protein coupled receptor	1385	465.62	0.39	-12.45	0	
11 12	Decarboxylase	1293	63 73	345.26 444.87	-9.66 -9.26	0	1	12	Acyltransferase	867	531.58	0.42	-11.28	0	
13	Monooxygenase Metalloprotease	1668	109	553.73	-9.26 -7.89	0	ſ X ¯	13 14	Decarboxylase	195 202	488.21	0.25	-10.70 -10.23	0	
14	Aminopeptidase	1100 452	39	509.17	-7.55	0		15	Aminotransferase Aminopeptidase	130	451.05 668.72	0.24	-9.10	0	
15	Dioxygenase	360	66	433.20	-7.32	0		16	Serine protease	460	700.07	0.50	-8.87	0	
16	Aminoacyl-tRNA synthetase	3402	37	571.83	-7.15	0	\times	17	Metalloprotease	507	688.25	0.56	-8.43	0	
17	Protease	4423	380	549.70	-7.1	0	/X \	18	Methyltransferase	874	611.22	0.47	-8.33	0	
18	Aminotransferase	955	28	420.27	-6.02	0	/ \	19	Carboxypeptidase	116	631.16	0.37	-8.25	0	
			l				` \	20	Threonine protease	138	246.88	0.18	-7.50	0	
							×	21	Dioxygenase	366	622.32	0.48	-7.39	0	
L	1 \														
b)						Г	P	redviđeno uređeni MF termini dob	ijeni iz o	direktnog i	izvedeno	og mapira	anja	
	Predviđeno uređene MF klju	čne reči	iz ovog ra	da (z-skoi	r < -0.2)			#	name	n	avg_len	avg_dis	Z	р	
#	name	n	avg_len		_	p		0	catalytic activity	28179	569.86	0.48	-52.25	0	
0	Oxidoreductase	4120	5 472.25	0.28	-41.35	0		1	oxidoreductase activity	4986	450.49	0.30	-39.62	0	
1	Hydrolase	7564	4 614.81	0.51	-26.99	0	→	2	hydrolase activity	11097	623.94	0.53	-27.76	0	
2	Lyase	143	1 481.37	0.30	-23.62	0	→	3	lyase activity	1624	480.74	0.31	-24.47	0	
3	Monooxygenase	555	503.36	0.20	-20.29	0	→	4	monooxygenase activity	633	511.13	0.22	-21.40	0	
4	Transferase	8846	6 631.95	0.55	-19.72	0	[5	transporter activity	5388	621.16	0.53	-19.15	0	
5	Ligase	995	693.30	0.46	-18.05	0		6	transferase activity	10428	619.96	0.56	-18.80	0	
6	Glycosyltransferase	1134	4 551.26	0.40	-17.04	0	\sim	7	transferase activity, trans	1343	545.86	0.41	-18.49	0	
7	Glycosidase	697	570.50	0.37	-16.81	0		8	ligase activity	1097	675.42	0.46	-16.68	0	
8	Isomerase	931		0.35	-13.60	0		9	hydrolase activity, acting	934	548.13	0.39	-16.32	0	
9	Protease	1863			-13.20	0	\ I	10	isomerase activity	1046	425.93	0.35	-14.19	0	
10	Transducer	1703	_	0.41	-12.56	0		11	transferase activity, trans	1235	540.97	0.43	-13.31	0	
11	G-protein coupled receptor	1385		0.39	-12.45	0	1 × × L	12	carboxy-lyase activity	292	494.47	0.26	-13.09	0	
12 13	Acyltransferase	867 195		0.42	-11.28 -10.70	0	\ \ \ \	13 14	peptidase activity	2211 1473	644.11	0.53	-13.05 -12.30	0	
14	Decarboxylase Aminotransferase	202	_	0.23	-10.70	0	\ / 🤻	15	G-protein coupled receptor carboxylic ester hydrolase	628	465.64 482.21	0.40	-12.30	0	
15	Aminopeptidase	130		0.24	-9.10	0	\ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \	16	serine-type peptidase activity	708	614.43	0.37	-10.79	0	
16	Serine protease	460		0.50	-8.87	0		17	aminopeptidase activity	177	633.80	0.35	-10.74	0	
17	Metalloprotease	507		0.56	-8.43	0	1 1// -	18	transaminase activity	230	459.43	0.26	-10.46	0	
18	Methyltransferase	874	_	0.47	-8.33	0	√ / ∟	19	methyltransferase activity	947	607.08	0.48	-8.77	0	
19	Carboxypeptidase	116		0.37	-8.25	0	\ N	20	carboxypeptidase activity	143	596.81	0.34	-8.54	0	
20	Threonine protease	138		0.18	-7.50	0	1/′ / \ . ∟	21	metallopeptidase activity	632	663.90	0.56	-8.44	0	
21	Dioxygenase	366		0.48	-7.39	0	AL T	22	ATP binding	7585	801.90	0.69	-8.22	0	
22	Serine esterase	141	423.09	0.28	-7.21	0	// \	23	dioxygenase activity	419	604.98	0.48	-7.83	0	
23	Receptor	3424		0.59	-7.09	0	11	24	peroxidase activity	282	414.57	0.34	-7.11	0	
24	Nucleotidyltransferase	600		0.63	-6.18	0	√ 1 \.∟	25	antioxidant activity	499	344.42	0.36	-6.89	0	
25	Peroxidase	221		0.40	-5.50	0	// ∕ ⊢	26	threonine-type endopeptidas	139	247.87	0.19	-6.79	0	
26	Serine protease inhibitor	182		0.38	-4.92	0	<u> </u>	27	nucleotidyltransferase acti	892	805.20	0.59	-6.07	0	
27	Porin	63	318.84	0.21	-4.55	0	/ \ \ \	28	porin activity	76	369.24	0.22	-5.22	0	
28	Dipeptidase	22	522.27	0.23	-4.43	0	\	29	serine-type endopeptidase i	218	485.65	0.40	-4.99	0	
29 30	Integrin Protease inhibitor	71 284	1038.25	0.70	-4.29 -4.15	0	X 1 ⊢	30	phospholipase A2 activity	93	377.39 778.27	0.30	-4.93 -4.87	0	
31	RNA-directed RNA polymeras		2506.11		-4.15	0	\ / \	32	phospholipase D activity prenyltransferase activity	128	369.37	0.43	-4.87	0	
32	Neurotoxin	66	209.62	0.87	-3.80	0	\\	33	toxin activity	285	349.98	0.32	-4.70	0	
33	Hemagglutinin	23	281.22	0.17	-3.37	0	\\\ \ \ \ \ \	34	dipeptidase activity	41	557.73	0.34	-4.23	0	
34	Retinal protein	47	384.06	0.13	-3.37	0	\mathbb{N} X / \mathbb{N}	35	metalloendopeptidase activity	369	711.73	0.63	-4.22	0	
35	Myosin	185			-3.29	0	V / L	36	phospholipase A2 activity (70	365.39	0.30	-3.90	0	
36	Photoreceptor protein	77	540.68	0.47	-3.02	0.01	//\ /\' L	37	endonuclease activity (773	658.33	0.59	-3.85	0	
37	Prenyltransferase	39	376.21	0.31	-2.59	0.01	\/ \/\ \\\ \\\	38	antigen binding	277	345.87	0.38	-3.68	0	
38	Elongation factor	106		0.48	-2.59	0.01	// / X / L	39	translation elongation fact	116	464.12	0.46	-3.35	0	
39	Endonuclease	443	_	0.60	-2.59	0.01	1 A 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	40	nuclease activity	1124	659.17	0.61	-3.20	0	
40	Toxin	214	_	0.39	-2.56	0	// \	41	G-protein coupled photorece	44	390.23	0.30	-2.99	0	
41	Bacteriolytic enzyme	35	434.66	0.31	-2.53	0.01	/ // \\\	42	chloride channel activity	195	622.49	0.62	-2.67	0.01	
42	Nuclease	703	668.54	0.60	-2.49	0.01	, \ L	43	RNA-directed 5'-3' RNA poly	75	2341.31	0.87	-2.57	0.02	
'	Ion channel impairing toxin	60	74.92	0.15	-2.47	0	· / \ \ [44	tRNA binding	247	566.35	0.57	-2.39	0.01	