a)							[Predviđeno uređene MF-ključ	na rači i	z ovog rod	o (z ekor	< 0.2)		
10	9 MF-ključnih reči sa najznačajn	iiom pre	edviđenom	uređenoš	ću (org. 2	2007)	7	#	name	ne reci ii		a (z-skor avg_dis		р	
#			n families				اــ	0	Oxidoreductase	4126	472.25	0.28	-41.35	0	
0		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	k /4	5	Ligase	995	693.30	0.46	-18.05	0	
5	Glycosidase	1826	244	444.73	-13.98	0	1/0	6	Glycosyltransferase	1134	551.26	0.40	-17.04	0	
6		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	$\downarrow X$	9	Protease	1863	674.42	0.54	-13.20	0	
9	Kinase	7017	322	448.29	-10.22	0	$ X/ \setminus I $	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 Monooxygenase	1293 1668	63 73	345.26 444.87	-9.66 -9.26	0		12	Acyltransferase	867 195	531.58 488.21	0.42	-11.28 -10.70	0	
13	Metalloprotease	1100	109	553.73	-7.89	0	$\{ X \}$	14	Decarboxylase Aminotransferase	202	451.05	0.23	-10.70	0	
14	Aminopeptidase	452	39	509.17	-7.55	0		15	Aminopeptidase	130	668.72	0.24	-9.10	0	
15	Dioxygenase	360	66	433.20	-7.32	0	┧ / X 1	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		4423	380	549.70	-7.1	0	7 X 🔻	18	Methyltransferase	874	611.22	0.47	-8.33	0	
18	Aminotransferase	955	28	420.27	-6.02	0	7 \	19	Carboxypeptidase	116	631.16	0.37	-8.25	0	
						-	_ \	20	Threonine protease	138	246.88	0.18	-7.50	0	
							*	21	Dioxygenase	366	622.32	0.48	-7.39	0	
h	h)														
Ŋ	Predviđeno uređeni MF-termini dobijeni iz direktnog i izvedenog mapiranja														
	Predviđeno uređene MF-klju	čne reči	iz ovog ra	da (z-skoi	r < -0.2)		ļ	#	name	n	avg_len	avg_dis	z	р	
#	name	n	avg_len	avg_dis	S Z	p		0	catalytic activity	28179	569.86	0.48	-52.25	0	
0	Oxidoreductase	4126	5 472.25	0.28	-41.35	0	→	1	oxidoreductase activity	4986	450.49	0.30	-39.62	0	
1	Hydrolase	7564	614.81	0.51	-26.99	0	→	2	hydrolase activity	11097	623.94	0.53	-27.76	0	
2	Lyase	1431		0.30	-23.62	0	+	3	lyase activity	1624	480.74	0.31	-24.47	0	
3	Monooxygenase	555		0.20	-20.29	0	<u>+</u> _	4	monooxygenase activity	633	511.13	0.22	-21.40	0	
4	Transferase	8846		0.55	-19.72	0		5	transporter activity	5388	621.16	0.53	-19.15	0	
5	Ligase	995		0.46	-18.05	0	<u> </u>	6	transferase activity	10428	619.96	0.56	-18.80	0	
6	Glycosyltransferase	1134		0.40	-17.04	0	\vee	7	transferase activity, trans	1343	545.86	0.41	-18.49	0	
7 8	Glycosidase	697 931	570.50 422.72	0.37	-16.81 -13.60	0		8	ligase activity hydrolase activity, acting	1097 934	675.42 548.13	0.46	-16.68 -16.32	0	
9	Isomerase Protease	1863			-13.60	0		10	isomerase activity	1046	425.93	0.39	-16.32	0	
10	Transducer	1703		0.34	-13.20	0	\ I	11	transferase activity, trans	1235	540.97	0.33	-14.19	0	
11	G-protein coupled receptor	1385		0.39	-12.45	0	\ \ \ \	12	carboxy-lyase activity	292	494.47	0.43	-13.09	0	
12	Acyltransferase	867		0.42	-11.28	0	1 × × ! L	13	peptidase activity	2211	644.11	0.53	-13.05	0	
13	Decarboxylase	195		0.25	-10.70	0	\ \ \ \	14	G-protein coupled receptor	1473	465.64	0.40	-12.30	0	
14	Aminotransferase	202		0.24	-10.23	0	\ / 🖘	15	carboxylic ester hydrolase	628	482.21	0.37	-12.01	0	
15	Aminopeptidase	130	668.72	0.37	-9.10	0		16	serine-type peptidase activity	708	614.43	0.46	-10.79	0	
16	Serine protease	460	700.07	0.50	-8.87	0	+	17	aminopeptidase activity	177	633.80	0.35	-10.74	0	
17	Metalloprotease	507		0.56	-8.43	0	√ / ∟	18	transaminase activity	230	459.43	0.26	-10.46	0	
18	Methyltransferase	874		0.47	-8.33	0	\ N	19	methyltransferase activity	947	607.08	0.48	-8.77	0	
19	Carboxypeptidase	116		0.37	-8.25	0	1/′ /	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	// 47	22	ATP binding	7585	801.90	0.69	-8.22	0	
22	Serine esterase	141	423.09	0.28	-7.21 7.00	0	/ I N	23	dioxygenase activity	419	604.98	0.48	-7.83	0	
23	Receptor Nucleotidyltransferase	3424		0.59	-7.09 -6.18	0	11 \	24 25	peroxidase activity antioxidant activity	282 499	414.57 344.42	0.34	-7.11 -6.89	0	
25	Peroxidase	221	457.61	0.63	-5.50	0	\sim 1 \sim 1.	25 26	threonine-type endopeptidas	139	247.87	0.36	-6.79	0	
26	Serine protease inhibitor	182		0.40	-4.92	0	// ∕ ⊢	20 27	nucleotidyltransferase acti	892	805.20	0.19	-6.07	0	
27	Porin	63	318.84	0.38	-4.55	0	<u> </u>	28	porin activity	76	369.24	0.39	-5.22	0	
28	Dipeptidase	22	522.27	0.23	-4.43	0	1 1 \ \	29	serine-type endopeptidase i	218	485.65	0.40	-4.99	0	
29	Integrin	71	1038.25		-4.29	0	\	30	phospholipase A2 activity	93	377.39	0.30	-4.93	0	
30	Protease inhibitor	284		0.41	-4.15	0	X 1 -	31	phospholipase D activity	37	778.27	0.43	-4.87	0	
31	RNA-directed RNA polymerase	e 62	2506.11	0.87	-4.11	0		32	prenyltransferase activity	128	369.37	0.32	-4.70	0	
32	Neurotoxin	66	209.62	0.17	-3.80	0	-\+-	33	toxin activity	285	349.98	0.32	-4.59	0	
33	Hemagglutinin	23	281.22	0.13	-3.37	0		34	dipeptidase activity	41	557.73	0.34	-4.23	0	
34	Retinal protein	47	384.06	0.28	-3.37	0	`./ \	35	metalloendopeptidase activity	369	711.73	0.63	-4.22	0	
35	Myosin	185			-3.29	0	//\ /\' 	36	phospholipase A2 activity (70	365.39	0.30	-3.90	0	
36	Photoreceptor protein	77	540.68	0.47	-3.02	0.01	√ √ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \	37	endonuclease activity	773	658.33	0.59	-3.85	0	
37	Prenyltransferase	39	376.21	0.31	-2.59	0.01	// XX / L	38	antigen binding	277	345.87	0.38	-3.68	0	
38	Elongation factor	106		0.48	-2.59	0.01	1 A AN -	39	translation elongation fact	116	464.12	0.46	-3.35	0	
39	Endonuclease	443	_	0.60	-2.59	0.01	// \ / \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \	40	nuclease activity	1124	659.17	0.61	-3.20	0	
40	Toxin Rectariolytic enzyme	214		0.39	-2.56	0	// //	41	G-protein coupled photorece	105	390.23	0.30	-2.99	0.01	
41	Bacteriolytic enzyme Nuclease	35 703	434.66 668.54	0.31	-2.53 -2.49	0.01	/ \ \\\ \	42 43	chloride channel activity RNA-directed 5'-3' RNA poly	195 75	622.49	0.62	-2.67 -2.57	0.01	
43	Ion channel impairing toxin	60	74.92	0.00	-2.49	0.01	, \ L	43 44	tRNA binding	247	566.35	0.87	-2.37	0.02	
÷.7	ion chamici impaning toxin	00	74.92	0.13	-2.4/	U	, \ \\ <u>\</u>	τ∺τ	axiva omanig	∠+1	1 200.33	0.57	-4.39	U.UI	