19 MF-ključnih reči sa najznačajnijom predviđenom uređenošću (org. 2007)									
#	name	n prot	n families	avg_len	Z-score	P-value			
0	Oxidoreductase	14995	992	376.63	-29.54	0			
1	Transferase	26525	1606	445.17	-24.25	0			
2	Lyase	7262	347	377.92	-22.64	0			
3	Hydrolase	20464	1995	430.68	-21.75	0			
4	Isomerase	4487	220	383.98	-14.18	0			
5	Glycosidase	1826	244	444.73	-13.98	0			
6	Glycosyltransferase	2950	261	437.53	-12.51	0			
7	Acyltransferase	2239	179	402.83	-10.85	0			
8	Methyltransferase	3524	224	349.60	-10.53	0			
9	Kinase	7017	322	448.29	-10.22	0			
10	Ligase	8010	230	529.41	-10.06	0			
11	Decarboxylase	1293	63	345.26	-9.66	0			
12	Monooxygenase	1668	73	444.87	-9.26	0			
13	Metalloprotease	1100	109	553.73	-7.89	0			
14	Aminopeptidase	452	39	509.17	-7.55	0			
15	Dioxygenase	360	66	433.20	-7.32	0			
16	Aminoacyl-tRNA synthetase	3402	37	571.83	-7.15	0			
17	Protease	4423	380	549.70	-7.1	0			
18	Aminotransferase	955	28	420.27	-6.02	0			

	1	Predviđeno uređene MF-ključn	I			I	Γ	-
	#	name	n	avg_len	avg_dis	Z	p	-
—	0	Oxidoreductase	4126	472.25	0.28	-41.35	0	_
A	1	Hydrolase	7564	614.81	0.51	-26.99	0	_
	2	Lyase	1431	481.37	0.30	-23.62	0	_
	3	Monooxygenase	555	503.36	0.20	-20.29	0	_
7	4	Transferase	8846	631.95	0.55	-19.72	0	<u> </u>
//	5	Ligase	995	693.30	0.46	-18.05	0	
	6	Glycosyltransferase	1134	551.26	0.40	-17.04	0	<u> </u>
	7	Glycosidase	697	570.50	0.37	-16.81	0	_
	8	Isomerase	931	422.72	0.35	-13.60	0	
\times	9	Protease	1863	674.42	0.54	-13.20	0	
// /	10	Transducer	1703	482.28	0.41	-12.56	0	
XX	11	G-protein coupled receptor	1385	465.62	0.39	-12.45	0	
\mathcal{I}	12	Acyltransferase	867	531.58	0.42	-11.28	0	\mathbb{H}
	13	Decarboxylase	195	488.21	0.25	-10.70	0	1
	14	Aminotransferase	202	451.05	0.24	-10.23	0	\mathbb{I}
\times	15	Aminopeptidase	130	668.72	0.37	-9.10	0	1
/X	16	Serine protease	460	700.07	0.50	-8.87	0	1
	17	Metalloprotease	507	688.25	0.56	-8.43	0	
\times	18	Methyltransferase	874	611.22	0.47	-8.33	0	1
	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	X
	22	Serine esterase	141	423.09	0.28	-7.21	0	1
	23	Receptor	3424	647.92	0.59	-7.09	0	/
	24	Nucleotidyltransferase	600	969.68	0.63	-6.18	0	1
	25	Peroxidase	221	457.61	0.40	-5.50	0	\triangleright
	26		182	497.66	0.40	-4.92	0	ſ
		Serine protease inhibitor						
	27	Porin	63	318.84	0.21	-4.55	0	$\overline{}$
	28	Dipeptidase	22	522.27	0.23	-4.43	0	
	29	Integrin	71	1038.25	0.70	-4.29	0	\
	30	Protease inhibitor	284	446.29	0.41	-4.15	0	1
	31	RNA-directed RNA polymerase	62	2506.11	0.87	-4.11	0	1\\
	32	Neurotoxin	66	209.62	0.17	-3.80	0	1
	33	Hemagglutinin	23	281.22	0.13	-3.37	0	↓
	34	Retinal protein	47	384.06	0.28	-3.37	0	$\backslash \backslash$
	35	Myosin	185	1275.20	0.72	-3.29	0	<u>ا</u> ا
	36	Photoreceptor protein	77	540.68	0.47	-3.02	0.01	\mathcal{N}
	37	Prenyltransferase	39	376.21	0.31	-2.59	0.01	\mathbb{N}
	38	Elongation factor	106	467.68	0.48	-2.59	0.01	$H \lambda$
	39	Endonuclease	443	728.14	0.60	-2.59	0.01	\mathbb{M}
	40	Toxin	214	412.51	0.39	-2.56	0	Y
	41	Bacteriolytic enzyme	35	434.66	0.31	-2.53	0.01] /
	42	Nuclease	703	668.54	0.60	-2.49	0.01	
	43	Ion channel impairing toxin	60	74.92	0.15	-2.47	0] ,'
	44	Voltage-gated sodium channe	22	79.55	0.05	-2.42	0.01] /
	45	Aspartyl protease	120	942.30	0.64	-2.39	0.01],
	46	Hemostasis impairing toxin	55	297.55	0.35	-2.10	0.02	1 ′
	47	Thiol protease inhibitor	63	411.06	0.35	-2.04	0.03	1/
		_	ı	1	1			. /

	Predviđeno uređeni MF-termini dobijeni iz direktnog i izvedenog mapiranja									
	#	name	n	n avg_len avg_dis z			p			
	0	catalytic activity	28179	569.86	0.48	-52.25	0			
	1	oxidoreductase activity	4986	450.49	0.30	-39.62	0			
\parallel	2	hydrolase activity	11097	623.94	0.53	-27.76	0			
\dashv	3	lyase activity	1624	480.74	0.31	-24.47	0			
	4	monooxygenase activity	633	511.13	0.22	-21.40	0			
	5	transporter activity	5388	621.16	0.53	-19.15	0			
	6	transferase activity	10428	619.96	0.56	-18.80	0			
	7	transferase activity, trans	1343	545.86	0.41	-18.49	0			
	8	ligase activity	1097	675.42	0.46	-16.68	0			
	9	hydrolase activity, acting	934	548.13	0.39	-16.32	0			
	10	isomerase activity	1046	425.93	0.35	-14.19	0			
	11	transferase activity, trans	1235	540.97	0.43	-13.31	0			
	12	carboxy-lyase activity	292	494.47	0.26	-13.09	0			
	13	peptidase activity	2211	644.11	0.53	-13.05	0			
	14	G-protein coupled receptor	1473	465.64	0.40	-12.30	0			
	15	carboxylic ester hydrolase	628	482.21	0.37	-12.01	0			
	16	serine-type peptidase activity	708	614.43	0.46	-10.79	0			
	17	aminopeptidase activity	177	633.80	0.35	-10.74	0			
	18	transaminase activity	230	459.43	0.26	-10.46	0			
	19	methyltransferase activity	947	607.08	0.48	-8.77	0			
	20	carboxypeptidase activity	143	596.81	0.34	-8.54	0			
	21	metallopeptidase activity	632	663.90	0.56	-8.44	0			
	22	ATP binding	7585	801.90	0.69	-8.22	0			
	23	dioxygenase activity	419	604.98	0.48	-7.83	0			
	24	peroxidase activity	282	414.57	0.34	-7.11	0			
	25	antioxidant activity	499	344.42	0.36	-6.89	0			
	26	threonine-type endopeptidas	139	247.87	0.19	-6.79	0			
	27	nucleotidyltransferase acti	892	805.20	0.59	-6.07	0			
	28	porin activity	76	369.24	0.22	-5.22	0			
	29	serine-type endopeptidase i	218	485.65	0.40	-4.99	0			
	30	phospholipase A2 activity	93	377.39	0.30	-4.93	0			
	31	phospholipase D activity	37	778.27	0.43	-4.87	0			
A	32	prenyltransferase activity	128	369.37	0.32	-4.70	0			
	33	toxin activity	285	349.98	0.32	-4.59	0			
7	34	dipeptidase activity	41	557.73	0.34	-4.23	0			
	35	metalloendopeptidase activity	369	711.73	0.63	-4.22	0			
	36	phospholipase A2 activity (70	365.39	0.30	-3.90	0			
	37	endonuclease activity	773	658.33	0.59	-3.85	0			
	38	antigen binding	277	345.87	0.38	-3.68	0			
	39	translation elongation fact	116	464.12	0.46	-3.35	0			
	40	nuclease activity	1124	659.17	0.61	-3.20	0			
	41	G-protein coupled photorece	44	390.23	0.30	-2.99	0			
7	42	chloride channel activity	195	622.49	0.62	-2.67	0.01			
\	43	RNA-directed 5'-3' RNA poly	75	2341.31	0.87	-2.57	0.02			
M	44	tRNA binding	247	566.35	0.57	-2.39	0.01			
1	45	peptidase inhibitor activity	446	493.46	0.50	-2.38	0.01			
1	46	photoreceptor activity	91	574.22	0.54	-2.30	0.01			
1	47	glucan endo-1,3-beta-D-gluc	28	522.18	0.43	-2.29	0.02			
	48	signal transducer activity	4149	619.06	0.62	-2.17	0.02			
	49 50	aminoacyl-tRNA ligase activity	255 262	659.40 953.41	0.67	-1.93	0.04			
	50	cysteine-type endopeptidase	202	933.41	0.64	-1.65	0.05			