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		-
	#	name	n	avg_len	avg_dis	Z 41.25	p	4
	0	Oxidoreductase	4126	472.25	0.28	-41.35	0	
1	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	
/1	5	Ligase	995	693.30	0.46	-18.05	0	
	6	Glycosyltransferase	1134	551.26	0.40	-17.04	0	
	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	
$\langle / \setminus /$	10	Transducer	1703	482.28	0.41	-12.56	0	
XX	11	G-protein coupled receptor	1385	465.62	0.39	-12.45	0	\downarrow
$\mathcal{I} \setminus \mathcal{I}$	12	Acyltransferase	867	531.58	0.42	-11.28	0	1
	13	Decarboxylase	195	488.21	0.25	-10.70	0	1
\wedge	14	Aminotransferase	202	451.05	0.24	-10.23	0	1
\times	15	Aminopeptidase	130	668.72	0.37	-9.10	0	1
/X	16	Serine protease	460	700.07	0.50	-8.87	0	14
\ / \	17	Metalloprotease	507	688.25	0.56	-8.43	0	
$X \rightarrow$	18	Methyltransferase	874	611.22	0.47	-8.33	0	
	19	Carboxypeptidase	116	631.16	0.37	-8.25	0	
	20	Threonine protease	138	246.88	0.18	-7.50	0	
1	21	Dioxygenase	366	622.32	0.48	-7.39	0	\rightarrow
	22	Serine esterase	141	423.09	0.48	-7.21	0	1/
	23		3424	647.92	0.28	-7.21	0	//
	24	Receptor		969.68	0.39	-6.18	0	-[
		Nucleotidyltransferase	600					$\langle \cdot \rangle$
	25	Peroxidase	221	457.61	0.40	-5.50	0	
	26	Serine protease inhibitor	182	497.66	0.38	-4.92	0	<u> </u>
	27	Porin	63	318.84	0.21	-4.55	0	┤ ̄
	28	Dipeptidase	22	522.27	0.23	-4.43	0	λ
	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	
	32	Neurotoxin	66	209.62	0.17	-3.80	0]-//
	33	Hemagglutinin	23	281.22	0.13	-3.37	0] \
	34	Retinal protein	47	384.06	0.28	-3.37	0	<u> </u>
	35	Myosin	185	1275.20	0.72	-3.29	0]
	36	Photoreceptor protein	77	540.68	0.47	-3.02	0.01	$\rfloor l$
	37	Prenyltransferase	39	376.21	0.31	-2.59	0.01	M
	38	Elongation factor	106	467.68	0.48	-2.59	0.01]
	39	Endonuclease	443	728.14	0.60	-2.59	0.01	111
	40	Toxin	214	412.51	0.39	-2.56	0	1
	41	Bacteriolytic enzyme	35	434.66	0.31	-2.53	0.01	1
	42	Nuclease	703	668.54	0.60	-2.49	0.01	/
	43	Ion channel impairing toxin	60	74.92	0.15	-2.47	0	1
	44	Voltage-gated sodium channe	22	79.55	0.15	-2.42	0.01	,
	45	Aspartyl protease	120	942.30	0.64	-2.42	0.01	/
	45	Hemostasis impairing toxin	55	297.55	0.04	-2.39	0.01	/
			<u> </u>					-
	47	Thiol protease inhibitor	63	411.06	0.35	-2.04	0.03	1/
	48	tRNA-binding	119	575.46	0.56	-2.01	0.03	ľ

	P	Predviđeno uređeni MF termini dobijeni iz direktnog i izvedenog mapiranja								
	#	name	n	avg_len	avg_dis	z	p			
	0	catalytic activity	28179	569.86	0.48	-52.25	0			
\blacksquare	1	oxidoreductase activity	4986	450.49	0.30	-39.62	0			
+	2	hydrolase activity	11097	623.94	0.53	-27.76	0			
+	3	lyase activity	1624	480.74	0.31	-24.47	0			
\vdash	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			
1	32	prenyltransferase activity	128	369.37	0.32	-4.70	0			
\langle	33	toxin activity	285	349.98	0.32	-4.59	0			
$\overline{}$	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			
	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			
	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	aminoacyl-tRNA ligase activity	255	659.40	0.67	-1.93	0.04			
	50	cysteine-type endopeptidase	262	953.41	0.64	-1.65	0.05			