Motif	Group Specificity	Group Specificity Rank	Site Specificity	Site Specificy Rank	Testset Sites	Average Testset Sites	StDev Testset Sites	TRANSFAC Correlation	TRANSFAC ID	Motif Finders	Sets	Repeats	Logo
1	4.40e-011	1	1.60e-011	1	4	0.80	0.45			AA	1, 2, 3, 4, 5		
2	1.00e-010	2	2.70e-011	2	4	0.80	0.84			AA, BP	1, 2, 4, 5		
3	4.00e-010	3	9.00e-011	3	4	0.80	0.84			AA	1, 2, 3, 4, 5	PAL,TR	
4	6.60e-010	4	2.20e-010	4	4	0.80	0.45			BP	1, 3, 5	PAL,TR	
5	2.00e-009	5	5.50e-010	5	5	1.00	1.20	0.60	TANTIG	AA, BP, MD, ME	1, 2, 3, 4, 5	PAL,TR	
6	3.50e-009	6	7.90e-010	6	6	1.20	1.60			AA	1, 2, 4, 5	PAL,TR	
7	7.00e-009	7	2.60e-009	7	5	1.00	0.71	0.61	CETS16	AA, BP	1, 2, 3, 4		
8	1.10e-008	8	3.90e-009	8	5	1.00	0.71			BP	1, 2, 5		
9	2.50e-008	9	8.40e-009	9	4	0.80	0.45			BP	2		
10	6.80e-008	11	1.90e-008	11	4	0.80	0.84	0.61	AP2	AA, BP	1, 2, 3, 4, 5	PAL	
11	9.70e-008	12	2.80e-008	12	4	0.80	0.84			AA, BP, MD, ME	1, 2, 3, 4, 5		
12	1.10e-007	13	3.20e-008	13	4	0.80	0.84	0.61	ZF5	BP, MD, ME	2, 4	PAL	
13	1.80e-007	14	5.00e-008	14	4	0.80	0.84	0.65	MEF3	AA, BP, MD	1, 2, 3, 4, 5	PAL	
14	2.00e-007	15	5.50e-008	15	4	0.80	0.84	0.61	HEN1	AA, BP, MD, ME	1, 3, 4, 5	PAL	
15	2.60e-007	16	9.20e-008	17	5	1.00	0.71			BP	2, 3, 5		
16	2.80e-007	17	9.40e-008	18	2	0.40	0.55	0.63	TATA	AA	1, 2, 3, 5		
17	2.90e-007	18	6.90e-008	16	6	1.20	1.60			AA	1, 2, 3, 4	PAL,TR	
18	4.50e-007	19	1.70e-007	20	4	0.80	0.45	0.63	TATA	AA, ME	1, 2, 3, 4, 5		
19	4.50e-007	19	1.20e-007	19	5	1.00	1.00			AA, BP	1, 2, 3, 4		
20	8.50e-007	21	2.50e-007	21	4	0.80	1.30			AA	2	PAL,TR	
21	1.60e-006	22	4.40e-007	22	4	0.80	0.84			BP, MD	2	PAL,TR	
22	6.20e-006	24	1.40e-006	24	5	1.00	1.20			AA	1, 4	PAL,TR	
23	6.30e-006	25	2.20e-006	25	6	1.20	1.10			BP, ME	1, 2, 3, 4, 5	PAL	
24	9.80e-006	26	2.70e-006	26	4	0.80	0.84			BP, MD	3, 4	PAL	
25	2.50e-005	28	8.20e-006	28	5	1.00	0.71	0.63	GATA1	BP	1, 2, 3, 4, 5		
26	2.60e-005	29	8.20e-006	28	4	0.80	1.30			AA, BP, MD	1, 2, 3, 5	PAL,TR	
27	1.20e-004	30	3.90e-005	31	5	1.00	0.71			BP	2		
28	1.30e-004	31	3.50e-005	30	7	1.40	1.50			BP	1, 2	PAL,TR	
29	2.20e-004	32	6.30e-005	32	5	1.00	1.20			BP, MD	1, 2, 3, 4, 5		