

Motif	Group Specificity	Group Specificity Rank	Site Specificity	Site Specificity Rank	Testset Sites	Average Testset Sites	StDev Testset Sites	TRANSFAC Correlation	TRANSFAC ID	Motif Finders	Sets	Repeats	Logo
1	4.40e-011	1	3.90e-011	1	7	1.40	1.10			AA, BP, ME	1, 2, 3, 5	PAL	
2	4.40e-011	1	3.90e-011	1	6	1.20	1.10			AA	1, 2, 3, 4, 5		
3	1.00e-010	3	5.30e-011	3	4	0.80	0.84			AA	1, 3, 4	PAL	
4	1.00e-010	3	5.80e-011	5	3	0.60	0.55			AA	2, 4	PAL	
5	1.00e-010	3	5.60e-011	4	4	0.80	0.84			AA	5	PAL	
6	1.00e-009	6	5.90e-010	7	3	0.60	0.55			AA	1, 2, 3, 4, 5	PAL	
7	1.00e-009	6	5.40e-010	6	5	1.00	1.00			AA	1, 2, 3	TR	
8	1.00e-009	6	6.00e-010	8	5	1.00	1.00	0.61	FOXJ2	AA	1, 3, 4, 5	PAL	
9	2.00e-009	9	1.10e-009	9	5	1.00	1.00			AA	1, 2, 3	TR	
10	5.60e-009	10	3.10e-009	10	6	1.20	1.30			AA	1, 3, 4	TR	
11	8.50e-009	11	4.70e-009	11	5	1.00	1.00			AA	1, 2, 4	PAL	
12	1.10e-008	13	1.10e-008	16	6	1.20	0.84			AA, BP, MD, ME	1, 2, 3, 4, 5		
13	1.20e-008	14	6.60e-009	12	5	1.00	1.00			AA	3	PAL	
14	1.20e-008	14	6.90e-009	13	4	0.80	0.84			AA	1, 2, 3, 4, 5		
15	1.70e-008	16	9.90e-009	15	6	1.20	1.30			AA, BP, MD, ME	5		
16	2.90e-008	17	1.70e-008	17	4	0.80	0.84	0.61	ATATA	BP, MD, ME	1, 2, 3, 4, 5		
17	3.20e-008	18	2.60e-008	19	10	2.00	1.90	0.64	HNF4AL	AA	1, 2, 3		
18	3.70e-008	19	2.20e-008	18	5	1.00	1.20	0.67	PIT1	BP	1, 2, 3, 4, 5	TR	
19	3.80e-008	20	3.70e-008	21	6	1.20	1.10			AA, BP	1, 2, 3, 4, 5		
20	4.60e-008	21	2.80e-008	20	5	1.00	1.00			AA	2, 3, 5	PAL	
21	6.80e-008	23	3.70e-008	21	4	0.80	0.84			AA	1, 3, 5	PAL,TR	
22	8.20e-008	24	4.80e-008	24	6	1.20	1.10			AA	2, 3, 4, 5		
23	8.20e-008	24	7.90e-008	26	6	1.20	0.84	0.72	CIZ	AA, BP, MD	1, 2, 3, 4, 5		
24	1.10e-007	27	6.90e-008	25	4	0.80	0.84			AA, BP, MD, ME	1, 2, 3, 4		
25	1.80e-007	28	1.00e-007	28	7	1.40	1.50			AA	2, 4	PAL	
26	2.00e-007	29	1.20e-007	29	5	1.00	1.20			AA	5	PAL	
27	2.30e-007	30	2.10e-007	32	8	1.60	1.30			AA	3, 4	PAL	
28	2.90e-007	31	1.80e-007	30	5	1.00	1.20			AA, BP	1		
29	2.90e-007	31	2.80e-007	33	6	1.20	1.10			BP	1, 2, 3, 4, 5		
30	3.30e-007	33	1.80e-007	30	6	1.20	1.10			AA	3		
31	4.30e-007	34	4.10e-007	34	5	1.00	0.71			AA, BP	1, 2, 3, 4, 5		
32	4.80e-007	35	4.60e-007	36	6	1.20	1.10	0.62	ATATA	BP	1, 2, 3, 5	PAL,TR	
33	7.20e-007	37	4.10e-007	34	8	1.60	1.70			AA	2, 3, 4, 5		
34	8.40e-007	38	4.80e-007	37	4	0.80	0.84			AA	3	PAL,TR	
35	9.10e-007	39	8.90e-007	39	5	1.00	0.71			AA, BP, ME	1, 2, 3, 4, 5		
36	1.80e-006	40	1.70e-006	40	6	1.20	0.84	0.63	PU1	BP, MD, ME	2, 3, 4		
37	2.70e-006	41	2.60e-006	44	8	1.60	1.50	0.67	TAL1AL	AA, ME	2, 3		
38	2.70e-006	41	2.40e-006	42	10	2.00	1.40			AA	4, 5	PAL	
39	3.20e-006	43	2.00e-006	41	5	1.00	1.00	0.65	HNF6	AA	1, 2, 3, 4, 5		
40	4.60e-006	44	2.40e-006	42	7	1.40	1.30			AA	3		
41	5.70e-006	45	3.30e-006	45	6	1.20	1.30			AA, BP	1, 5		
42	6.20e-006	46	5.70e-006	46	6	1.20	0.84			BP	4	PAL	
43	1.70e-005	48	1.50e-005	50	6	1.20	1.10	0.65	IRF7	BP	1, 2, 4		
44	1.90e-005	49	1.00e-005	48	4	0.80	1.10			AA	3		
45	2.80e-005	51	2.50e-005	53	9	1.80	1.50			AA, BP	2, 3, 5	PAL,TR	
46	2.90e-005	52	2.60e-005	54	6	1.20	0.84	0.68	TCF11M	BP	1, 4		
47	3.60e-005	53	2.10e-005	51	5	1.00	1.20			BP	3	TR	