

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	3.70e-011	2	2.40e-010	2	7	0.78	0.67			BP	1, 2, 3, 4, 5		AG I...AAAG...CAI
2	2.00e-010	3	1.30e-009	3	8	0.89	0.78	0.70	AREB6	AA, BP	1, 2, 4, 5		CAAGGIR...I...I...I...
3	1.40e-008	4	8.20e-008	4	9	1.00	1.20	0.60	SMAD3	BP, MD	1, 4, 5		AI...I...GAG...G...G...
4	3.30e-008	5	1.40e-006	13	16	1.80	1.80	0.62	HOX13	BP	2, 3, 4		A...AAI...AG...A...I...
5	5.70e-008	7	7.60e-007	11	10	1.10	0.93			AA, BP	1, 2, 3, 4, 5		AI...G...I...G...I...G...
6	6.00e-008	8	1.10e-007	5	5	0.56	0.73			BP	4		AG...G...I...A...I...I...I...
7	1.40e-007	10	4.30e-007	9	12	1.30	1.70			AA	2, 3, 4, 5	TR	I...G...A...AA...A...GA...A
8	1.90e-007	11	5.40e-007	10	8	0.89	1.30	0.61	RP58	ME	1, 2, 3, 4, 5	PAL	...G...A...G...A...A...IG
9	2.40e-007	12	3.10e-007	8	6	0.67	1.10	0.67	TITF1	BP	3, 5	PAL,TR	IAG...GAG...I...G...A...
10	4.90e-007	14	1.40e-006	13	7	0.78	0.83			BP	2, 3		I...I...I...A...G...I...
11	6.00e-007	16	1.10e-006	12	9	1.00	1.50			AA, BP	1, 5	PAL,TR	I...I...I...I...G...I...I...I...I...
12	7.00e-007	17	3.70e-006	20	10	1.10	1.20			BP	2, 3, 4, 5	PAL	A...I...I...I...I...I...I...
13	7.20e-007	18	8.40e-005	49	16	1.80	1.10	0.81	AP4_Q6	AA, BP, MD, ME	1, 2, 3, 4, 5		A...AG...I...G...
14	8.30e-007	19	1.50e-006	15	11	1.20	2.00			AA	2, 3	PAL	I...I...I...I...I...I...AAA...G...C...
15	1.10e-006	22	3.30e-006	19	13	1.40	1.90			AA	4, 5	PAL,TR	I...I...I...A...A...A...I...I...
16	2.30e-006	24	2.40e-005	32	24	2.70	3.30			AA	1, 3		A...I...A...A...I...A...I...
17	2.70e-006	25	6.80e-005	45	19	2.10	2.00			AA, BP	1, 2, 3, 4		I...I...I...I...I...I...I...
18	2.90e-006	26	2.90e-005	34	16	1.80	1.70	0.60	AREB6	AA, ME	2, 4		I...G...A...G...A...I...
19	4.10e-006	27	5.20e-006	22	5	0.56	1.00			AA, BP	1, 2, 3, 4	PAL	I...G...A...A...I...I...I...A
20	4.40e-006	29	2.10e-005	29	10	1.10	1.20	0.68	NKX61	AA, BP	1, 2, 4	PAL	A...G...I...A...I...A...
21	4.90e-006	30	1.30e-005	25	8	0.89	1.10			BP	3		A...I...G...A...I...A...
22	7.50e-006	34	3.60e-005	39	20	2.20	2.90			AA	1, 2, 4, 5	PAL,TR	I...I...A...A...A...A...A...
23	9.00e-006	35	2.30e-005	31	6	0.67	0.71	0.92	tef	AA, BP, MD, ME	1, 2, 3, 4, 5		IGCAATG
24	9.30e-006	36	2.50e-005	33	7	0.78	0.97	0.60	NRSF	BP, ME	1, 2, 3, 4, 5		I...I...A...G...I...A...A...
25	9.60e-006	37	1.60e-005	27	10	1.10	1.80			AA, BP	5		G...A...A...A...A...A...
26	9.80e-006	38	2.00e-004	63	18	2.00	1.60	0.68	TAL1BE	BP	1, 2, 5		A...A...A...A...A...A...
27	1.20e-005	39	3.20e-005	37	8	0.89	0.93	0.65	MAF	ME	1, 2, 3, 4, 5		A...G...G...G...G...G...I...
28	1.20e-005	39	1.10e-004	55	15	1.70	1.70	0.61	CLOX	BP	1, 5		A...G...I...I...A...I...
29	1.20e-005	39	2.10e-005	29	12	1.30	1.80			AA	2, 3, 4	PAL,TR	A...A...A...I...I...A...A...
30	1.20e-005	39	1.00e-004	54	13	1.40	1.80	0.66	MEIS1	AA, BP, MD, ME	1, 2, 3, 4, 5		T...G...A...A...G...
31	1.20e-005	39	3.00e-005	35	7	0.78	0.97			AA, BP, ME	1, 3, 4		A...A...G...A...A...I...
32	1.50e-005	45	6.40e-005	43	10	1.10	1.20	0.80	SF1	BP, ME	1, 2, 3, 4, 5		I...A...A...I...A...I...
33	1.60e-005	46	4.00e-005	40	11	1.20	1.50	0.75	LHX3	BP, ME	1, 3, 4, 5	PAL	C...A...I...T...A...I...
34	2.50e-005	50	4.30e-004	74	15	1.70	1.70			BP	2, 3		A...I...I...A...G...I...A...
35	2.60e-005	51	6.60e-005	44	11	1.20	1.60	0.62	FREAC7	BP	1, 2, 4, 5		A...A...A...A...A...I...G...
36	2.60e-005	51	4.50e-005	41	10	1.10	1.50			AA	3, 4, 5	PAL	I...I...I...I...A...A...A...
37	2.70e-005	53	2.20e-004	66	23	2.60	3.30			AA, BP	1, 2		I...A...I...I...A...A...A...
38	2.70e-005	53	1.10e-004	55	12	1.30	1.70	0.61	MEF2	BP	1, 4, 5		A...I...G...A...A...I...A...
39	2.90e-005	56	2.20e-004	66	18	2.00	2.00	0.65	CHX10	BP	1, 3, 4, 5		I...A...I...A...I...I...I...
40	2.90e-005	56	7.20e-005	46	13	1.40	1.70	0.60	OCT1_Q	AA, BP, ME	1, 2, 3, 4, 5		I...I...I...I...I...A...I...
41	3.00e-005	58	1.40e-003	100	28	3.10	3.00			BP	4, 5		I...I...A...I...A...A...
42	3.10e-005	59	1.30e-004	60	12	1.30	1.30	0.65	CEBPGA	BP	1, 3	PAL	I...G...A...I...I...A...
43	3.40e-005	61	8.50e-005	50	13	1.40	2.00	0.72	BRN2	BP, ME	2, 3, 4		I...G...A...I...A...I...
44	4.40e-005	62	6.70e-004	78	16	1.80	1.50	0.76	OCT1_Q	BP	1, 2, 3, 4, 5		I...G...A...I...I...A...
45	5.20e-005	63	8.70e-005	51	7	0.78	1.10			BP	3, 5		A...I...G...I...I...I...I...
46	5.80e-005	64	8.30e-004	83	22	2.40	2.30	0.74	HNF6	BP	1, 4		I...G...A...I...I...I...
47	8.50e-005	69	1.40e-004	62	14	1.60	2.50			AA, BP	1, 2, 5		AA...A...A...A...A...A...I...
48	9.10e-005	71	2.10e-004	64	10	1.10	1.30	0.64	BRN2	BP	2, 3, 5		A...I...A...A...I...I...
49	9.70e-005	72	1.20e-004	57	11	1.20	1.90			AA	2, 3	PAL,TR	I...I...I...A...A...A...I...
50	1.10e-004	75	1.40e-003	100	17	1.90	1.50	0.68	SREBP1	AA, BP, MD, ME	1, 5		A...I...G...A...I...G...
51	1.30e-004	78	4.90e-004	75	14	1.60	1.80			BP	1, 5	PAL	I...I...I...A...I...I...I...
52	1.40e-004	80	8.70e-004	85	27	3.00	3.80			AA	1, 2, 3, 5	PAL,TR	I...I...I...I...I...I...A...
53	1.50e-004	82	4.10e-003	123	31	3.40	3.70	0.64	BRACH	BP, ME	1, 2, 3, 4, 5		I...I...I...I...I...I...I...
54	1.50e-004	82	2.40e-004	68	9	1.00	1.50			BP	2, 4		I...I...A...I...I...I...I...
55	2.50e-004	86	2.70e-003	115	39	4.30	5.50			AA	1, 2, 3, 4, 5	PAL	I...I...I...I...I...I...A...
56	2.70e-004	87	2.70e-003	115	40	4.40	5.70			AA	4	PAL	I...I...I...I...I...I...A...