

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	9.20e-023	1	4.00e-019	1	14	0.58	0.58	0.63	ERR1	AA, ME	1, 2, 3, 4, 5	PAL,TR	
2	4.00e-022	2	1.10e-016	6	20	0.83	0.64	0.82	RSRFC4	AA, ME	1, 2, 3, 4, 5		
3	4.50e-021	3	1.30e-018	2	14	0.58	0.83	0.89	srf	AA, ME	1, 2, 3, 4, 5	PAL	
4	1.50e-019	6	1.80e-016	7	16	0.67	0.82	0.81	myf	AA, ME	1, 2, 3, 4, 5		
5	3.80e-018	8	1.70e-015	8	13	0.54	0.66	0.61	MUSCLE	AA, ME	1, 2, 3, 4, 5		
6	7.30e-017	11	1.50e-014	11	13	0.54	0.78	0.66	TGIF	AA, ME	1, 3, 4		
7	1.40e-016	13	1.80e-013	14	15	0.62	0.88	0.67	RSRFC4	AA, ME	1, 2, 3, 5		
8	5.70e-016	16	2.60e-014	12	8	0.33	0.48	0.67	CP2	AA, ME	1, 2, 3, 4		
9	9.60e-016	17	2.00e-012	22	17	0.71	0.75	0.66	AHRARN	AA, ME	1, 2, 3, 4, 5		
10	9.80e-016	18	7.10e-012	25	18	0.75	0.74	0.72	MAZR	AA, BP, MD, ME	1, 2, 4	TR	
11	2.70e-015	19	1.20e-012	21	14	0.58	0.72	0.68	AP2GAM	AA, MD, ME	1, 2, 3, 4, 5	TR	
12	9.30e-015	22	8.60e-013	20	11	0.46	0.66	0.62	LFA1	AA, ME	1, 2, 3, 4, 5	PAL	
13	1.10e-014	23	2.20e-012	23	13	0.54	0.78	0.79	SRF_Q5	AA, ME	1, 3, 4, 5	PAL,TR	
14	1.70e-014	25	1.90e-011	29	21	0.88	1.20	0.78	SRF_Q5	AA, ME	1, 2, 3, 4, 5		
15	3.20e-014	26	2.80e-011	31	15	0.62	0.77	0.66	BEL1	AA, ME	2, 5	PAL	
16	1.10e-013	30	1.90e-010	39	14	0.58	0.58	0.79	MAZ	AA, ME	3		
17	1.20e-013	31	2.00e-010	40	18	0.75	0.79	0.75	SP1	AA, MD, ME	2, 3, 4, 5		
18	1.50e-013	32	1.60e-009	49	21	0.88	0.99	0.67	MAZR	AA, BP, MD, ME	3	TR	
19	2.40e-013	35	1.10e-011	28	10	0.42	0.72	0.69	CACBIN	AA	1, 2, 3, 4, 5		
20	5.50e-013	36	1.10e-010	35	13	0.54	0.78	0.64	HMEF2	AA	1, 3, 4, 5		
21	1.10e-012	37	2.70e-011	30	7	0.29	0.46	0.66	LFA1	AA	3, 4		
22	1.70e-012	41	6.10e-010	44	14	0.58	0.78	0.70	srf	AA, ME	2, 3	PAL	
23	2.20e-012	43	8.90e-010	47	14	0.58	0.72	0.62	PTF1BE	AA	2, 4, 5	TR	
24	5.70e-012	46	4.60e-010	43	13	0.54	0.83	0.65	HEB	AA, MD, ME	1, 2, 3, 4, 5		
25	2.60e-010	56	4.10e-005	76	39	1.60	1.80	0.81	MEIS1	AA, BP, MD, ME	1, 2, 4, 5		
26	1.10e-009	61	3.10e-006	70	22	0.92	0.88	0.77	MAZ	AA, ME	1, 2, 3, 4, 5	PAL	
27	2.80e-009	63	9.60e-008	58	12	0.50	0.78	0.78	tef	AA, ME	1, 2, 3, 4, 5		
28	1.20e-008	67	1.10e-006	66	10	0.42	0.50	0.93	tef	AA, ME	1, 2, 3, 5		
29	1.60e-008	68	1.40e-005	73	20	0.83	0.76	0.90	mef2	AA, ME	2, 3, 4		
30	2.30e-008	69	1.40e-006	68	17	0.71	1.10	0.72	MEF2_Q	AA, ME	2, 5		
31	4.80e-008	71	6.10e-007	61	10	0.42	0.93	0.63	BARBIE	AA, ME	1, 2, 4, 5	TR	
32	1.00e-007	74	3.20e-006	71	9	0.38	0.58	0.62	SP3	ME	1, 2, 3, 4, 5	PAL	
33	3.60e-006	78	1.40e-004	82	15	0.62	1.10	0.66	BARBIE	AA	5		
34	7.00e-006	79	4.80e-004	88	20	0.83	1.10	0.74	MYOGEN	AA, BP, ME	1, 2, 3, 4, 5		
35	1.80e-005	84	2.40e-004	86	12	0.50	0.88	0.65	NERF	ME	1, 2, 3, 4, 5		
36	4.20e-005	87	5.60e-003	96	26	1.10	1.10	0.63	SP1_Q2	BP, ME	1, 2, 3, 4, 5	TR	
37	5.00e-005	89	4.30e-004	87	8	0.33	0.70	0.62	TCF4	ME	1, 2, 3, 4, 5		
38	1.10e-004	91	8.90e-004	91	8	0.33	0.64	0.75	SMAD	ME	1, 2, 3, 4, 5		
39	2.00e-004	93	2.80e-003	93	15	0.62	0.92	0.81	TBP	AA	5		