

Motif	Group Specificity	Group Specificity Rank	Site Specificity	Site Specificity Rank	Testset Sites	Average Testset Sites	StdDev Testset Sites	TRANSFAC Correlation	TRANSFAC ID	Motif Finders	Sets	Repeats	Logo
1	1.20e-014	4	1.40e-015	4	9	0.38	0.58	0.82	MAZ	AA, ME	1, 2, 3, 4, 5		
2	2.50e-012	16	4.60e-013	18	5	0.21	0.41	0.75	GC	AA, BP, ME	1, 2, 3, 4, 5	PAL,TR	
3	2.50e-012	16	4.70e-013	19	6	0.25	0.53	0.69	NERF	AA, BP, MD, ME	1, 2, 3, 4, 5		
4	7.00e-012	26	8.40e-013	24	8	0.33	0.56	0.62	SP1_Q4	AA, BP	1, 2, 3, 4, 5	PAL,TR	
5	1.80e-009	40	2.10e-002	139	87	3.60	5.20	0.81	LMO2CO	AA, BP, ME	1, 2, 3, 4, 5		
6	3.10e-009	43	6.00e-010	43	6	0.25	0.53	0.64	ALPHAC	AA, ME	1, 2, 3, 5		
7	7.20e-009	47	1.10e-009	46	10	0.42	0.83	0.66	NFMUE1	AA, BP, ME	3, 4	TR	
8	1.20e-008	50	2.10e-009	51	7	0.29	0.55	0.67	SF1	AA, BP, MD, ME	1, 2, 4, 5		
9	1.30e-008	51	1.60e-009	48	11	0.46	0.78	0.74	E47	AA, BP, ME	1, 3, 4, 5		
10	1.90e-008	55	2.40e-009	53	11	0.46	0.66	0.64	NERF	AA, MD, ME	1, 2, 3, 4, 5		
11	1.60e-006	70	2.10e-002	139	34	1.40	2.60	0.68	TTF1	ME	1, 2, 3, 4, 5		
12	2.00e-006	72	2.40e-007	67	13	0.54	0.78	0.62	EGR3	BP	1, 2, 5	PAL	
13	2.40e-005	82	3.60e-006	77	9	0.38	0.65	0.63	AP2	AA, MD, ME	1, 2, 3, 4, 5		
14	2.80e-005	83	4.50e-006	78	8	0.33	0.48	0.73	TCF4	AA, ME	1, 2, 3, 4, 5		
15	2.90e-005	84	1.00e-001	183	94	3.90	5.40	0.66	IK3	BP	1, 3, 5		
16	3.30e-005	88	4.80e-006	79	12	0.50	0.66	0.62	NGFIC	BP, ME	1, 2, 3, 4, 5	PAL	
17	3.50e-005	89	6.80e-002	159	41	1.70	2.30	0.76	IK3	ME	1, 2, 3, 4, 5		
18	3.70e-005	90	6.60e-006	81	8	0.33	0.64	0.79	SP1_Q4	AA, BP, ME	1, 2, 3, 4, 5		
19	8.90e-005	95	1.30e-005	84	12	0.50	0.72	0.64	TITF1	ME	1, 2, 3, 4, 5	TR	
20	1.20e-004	96	1.70e-005	86	10	0.42	0.65	0.78	SP1	AA, BP, MD, ME	1, 3, 4, 5	PAL	
21	1.80e-004	97	3.30e-005	87	7	0.29	0.55	0.65	AP2GAM	AA, ME	2		
22	2.60e-004	100	1.10e-001	199	107	4.50	6.10	0.81	CREB	BP, MD, ME	1, 3, 4, 5		
23	2.90e-004	101	5.70e-005	91	9	0.38	0.88	0.61	MTF1	AA	3, 4		
24	2.90e-004	101	1.10e-001	199	121	5.00	5.20	0.73	CACBIN	BP	1, 2, 3, 4, 5		
25	3.20e-004	103	4.40e-005	88	10	0.42	0.58	0.62	AP4	ME	1, 2, 3, 4, 5		
26	3.50e-004	104	5.60e-005	90	8	0.33	0.56	0.67	SP1_Q2	MD, ME	1, 2, 3, 4, 5	TR	
27	3.50e-004	104	5.70e-005	91	10	0.42	0.88	0.60	MTF1	BP, ME	1, 2, 3, 5	PAL	
28	3.50e-004	104	7.10e-002	162	44	1.80	2.50	0.72	ACAAT	BP, ME	1, 2, 3, 4	PAL	
29	4.80e-004	112	7.60e-002	166	34	1.40	2.40	0.67	AP2GAM	BP, ME	2, 3, 4	PAL	
30	6.10e-004	117	1.10e-001	199	114	4.80	7.60	0.63	HIF1	BP	1, 2, 3		
31	6.60e-004	118	8.70e-002	173	34	1.40	2.00	0.66	GC	ME	1, 2, 3, 4, 5		