





















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	1.60e-016	1	1.10e-014	1	7	1.40	0.89	0.61	NFAT	AA, BP, MD, ME	2, 3, 4, 5	TR	
2	1.20e-011	7	7.00e-010	8	5	1.00	0.00	0.83	SRF_Q5	AA, ME	1, 2, 3, 4, 5		
3	3.70e-011	10	2.20e-009	12	5	1.00	0.00	0.70	MYOGEN	BP, MD	1, 2, 3, 4, 5	PAL	
4	1.20e-010	11	1.20e-009	9	6	1.20	1.10	0.61	TTF1	AA, BP, ME	1	PAL	
5	1.50e-010	12	8.90e-009	15	8	1.60	1.30	0.61	TANTIG	AA, BP	1, 2, 3, 4, 5		
6	4.00e-010	15	1.50e-009	11	3	0.60	0.55	0.75	SRF_Q5	AA, BP, MD	1, 3, 4, 5		
7	1.30e-009	17	1.30e-008	17	6	1.20	1.10	0.61	ZID	AA, BP, MD	1, 2, 3, 4, 5		
8	3.00e-009	19	1.60e-007	25	7	1.40	0.89	0.75	RSRFC4	AA	4	TR	
9	3.50e-009	20	1.30e-008	17	5	1.00	1.00	0.69	TAL1AL	AA, BP, MD, ME	1, 3, 4, 5		
10	1.00e-008	25	5.70e-007	27	8	1.60	0.89	0.70	CAP	BP	1, 2, 3, 4, 5		
11	3.50e-008	28	1.80e-006	34	7	1.40	0.89	0.63	GKLF	BP	2, 3, 5	TR	
12	9.30e-008	29	9.30e-007	30	6	1.20	1.10	0.68	TANTIG	BP, MD, ME	1, 2, 3		
13	2.80e-007	32	5.10e-007	26	2	0.40	0.55	0.66	MEF2_Q	AA	1, 2, 3, 5		
14	4.70e-007	36	4.50e-006	38	6	1.20	0.84	0.65	GRE	BP	2, 5		
15	8.40e-007	37	2.90e-006	36	5	1.00	1.20	0.62	GCM	AA, ME	2, 3, 4, 5		
16	1.20e-005	46	1.00e-004	46	7	1.40	0.89	0.65	OLF1	AA, ME	1		
17	1.60e-005	48	5.30e-004	55	11	2.20	2.20	0.65	GATA	AA	1, 2, 5		
18	2.60e-005	50	4.60e-005	45	4	0.80	1.10	0.65	GEN_IN	AA	2, 3, 4, 5		
19	8.20e-005	52	6.10e-004	56	7	1.40	1.10	0.67	SMAD3	BP	1, 3, 4		
20	1.10e-004	55	2.50e-003	60	9	1.80	0.84	0.64	MAZ	BP, ME	1, 2, 5		
21	1.20e-004	57	2.70e-003	62	9	1.80	1.10	0.64	LFA1	BP	2, 5		