

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	7.10e-021	1	2.20e-019	1	9	1.00	0.50	0.64	EGR3	AA, BP, MD, ME	1, 2, 3, 4, 5		
2	3.30e-014	5	2.70e-013	4	9	1.00	0.71	0.68	ER_Q6	AA, BP	1, 2, 3, 4, 5		
3	3.90e-013	7	1.30e-012	9	6	0.67	0.50	0.62	ALX4	AA, BP	2, 3, 4, 5	PAL	
4	1.90e-012	10	3.00e-010	28	11	1.20	0.67	0.67	MTF1	AA, BP, ME	1, 2, 3, 4, 5	TR	
5	2.20e-012	11	3.80e-012	10	6	0.67	0.71	0.60	TITF1	AA, MD, ME	1, 2, 3, 4, 5	PAL	
6	6.10e-012	16	1.90e-011	17	7	0.78	0.67	0.62	CIZ	AA, MD, ME	1, 2, 3, 4, 5		
7	9.10e-012	17	1.00e-011	14	4	0.44	0.53	0.66	RP58	AA, BP, ME	1, 2, 4, 5	PAL	
8	9.30e-012	19	2.90e-011	20	6	0.67	0.50	0.62	ATF3	AA, BP	1, 4		
9	1.20e-011	20	3.00e-010	28	11	1.20	0.67	0.60	HEB	AA	3, 4, 5		
10	2.10e-011	23	2.40e-011	18	4	0.44	0.53	0.72	PR	AA, BP, ME	1, 2, 3, 4, 5		
11	1.60e-010	30	5.10e-010	32	9	1.00	0.87	0.64	AP2	AA, BP, ME	1, 2, 3, 4, 5		
12	2.00e-010	32	3.40e-010	31	6	0.67	0.71	0.62	DR4	AA, BP	1, 3, 4, 5	PAL	
13	3.50e-009	42	7.00e-008	51	9	1.00	0.50	0.71	BRACH	AA, BP	2, 3, 5	PAL	
14	3.60e-009	43	6.10e-009	43	6	0.67	0.71	0.66	MEIS1	AA, MD, ME	1, 2, 3, 4, 5		
15	5.30e-009	44	5.90e-009	41	5	0.56	0.73	0.63	MAF	AA	1, 2		
16	1.80e-007	58	2.90e-007	59	5	0.56	0.53	0.61	E12	AA, ME	5		
17	3.40e-007	62	1.90e-006	66	10	1.10	0.78	0.66	SMAD4	BP	1, 2, 5	TR	
18	4.40e-007	64	6.10e-006	71	11	1.20	0.67	0.62	HAND1E	BP, ME	1, 2, 3, 4, 5	TR	
19	6.30e-007	67	1.70e-006	65	8	0.89	0.78	0.66	MYOD_Q	BP, ME	1, 2, 3, 4, 5	PAL,TR	
20	1.30e-006	69	2.00e-006	67	6	0.67	0.71	0.64	ISRE	AA	4		
21	6.60e-006	74	6.60e-005	76	11	1.20	0.83	0.63	VDR	BP	4, 5	PAL	