

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	2.80e-013	2	2.60e-012	2	6	0.55	0.52	0.75	SOX9	AA, BP, MD, ME	1, 2, 3, 4, 5		
2	1.20e-012	5	1.20e-011	5	8	0.73	0.79	0.62	HNF3AL	AA, BP, MD, ME	1, 4, 5		
3	5.70e-012	7	2.70e-011	7	5	0.45	0.52	0.63	SP1_Q2	AA, BP, MD	1, 3	PAL	
4	5.50e-011	13	1.50e-010	9	4	0.36	0.50	0.62	MYOD	AA, ME	1, 2, 3, 4, 5		
5	6.90e-011	14	1.40e-009	19	9	0.82	0.75	0.65	SP1_Q2	AA, ME	1, 2, 3, 5	PAL,TR	
6	9.90e-011	17	9.00e-010	18	7	0.64	0.67	0.67	CP2	AA, BP, ME	1, 2, 3, 4, 5	PAL,TR	
7	9.60e-010	24	4.70e-008	29	12	1.10	1.10	0.76	AP2_Q6	AA, BP, MD, ME	1, 2, 4	PAL	
8	1.90e-009	26	3.70e-008	28	8	0.73	0.65	0.65	SMAD4	AA, ME	1, 2, 3, 4, 5		
9	3.60e-009	27	3.00e-008	26	8	0.73	0.90	0.65	EGR	AA, ME	1, 2, 3, 4, 5	TR	
10	4.90e-009	28	2.20e-008	25	7	0.64	0.81	0.62	CACBIN	AA, BP	3, 4	PAL,TR	
11	2.20e-007	38	1.70e-006	38	10	0.91	1.20	0.64	SP1_Q2	AA	2	TR	
12	3.00e-007	39	1.30e-006	37	8	0.73	1.00	0.67	CACBIN	BP, MD, ME	1, 2	TR	
13	4.60e-007	40	1.00e-004	50	19	1.70	1.30	0.76	SP1_Q2	BP, MD, ME	1, 2, 3, 5	PAL,TR	
14	1.10e-006	43	3.10e-005	45	13	1.20	1.30	0.73	SP1_Q2	BP	3, 4	TR	
15	3.30e-006	46	4.30e-005	46	9	0.82	0.75	0.64	EGR	AA	1, 4		
16	3.30e-006	46	2.30e-005	44	10	0.91	1.00	0.78	MAZ	BP	2, 5		
17	7.20e-006	48	1.70e-004	54	15	1.40	1.30	0.77	AP2_Q6	AA, ME	1, 2, 3, 4, 5		
18	1.30e-005	50	1.40e-004	52	16	1.50	1.60	0.61	CACBIN	AA	1	TR	
19	1.40e-005	51	9.30e-005	49	9	0.82	1.20	0.67	EGR	AA, ME	1, 2, 3, 4, 5		
20	3.40e-005	55	1.30e-004	51	13	1.20	2.10	0.62	MAZR	AA	1, 2, 3, 4		
21	7.30e-005	59	6.70e-004	60	7	0.64	0.50	0.73	E2F	AA, ME	1, 2, 3, 4, 5		
22	1.00e-004	60	1.80e-004	55	5	0.45	0.82	0.71	HFH4	AA	3, 5		
23	1.20e-004	61	2.90e-004	59	7	0.64	1.00	0.61	AHRARN	BP, ME	1, 2, 3, 4, 5		
24	2.20e-004	62	1.70e-003	62	11	1.00	1.00	0.74	ZIC2	BP, ME	3, 4, 5		
25	2.80e-004	63	5.40e-003	67	16	1.50	1.00	0.70	AP2ALP	BP, ME	1, 2, 3, 4, 5	PAL	