

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	2.60e-013	2	2.60e-014	2	6	0.38	0.62	0.67	ARP1	AA, BP	1, 2, 3, 4, 5	PAL,TR	
2	3.60e-011	6	2.90e-012	6	8	0.50	0.63	0.68	HNF1	AA, BP, MD, ME	1, 3, 4		
3	3.90e-011	7	3.10e-012	7	8	0.50	0.63	0.61	LHX3	BP, MD, ME	1, 2, 3, 4, 5	PAL,TR	
4	1.80e-009	20	2.70e-010	21	5	0.31	0.60	0.63	AFP1	AA, BP, ME	2, 3, 5	PAL	
5	2.90e-009	21	9.30e-010	26	21	1.30	1.40	0.65	SRY	AA, BP, MD, ME	1, 2, 3, 4, 5		
6	2.90e-009	21	2.90e-010	22	14	0.88	1.00	0.66	GF11	AA, BP	1, 2, 3, 4, 5		
7	5.80e-008	36	6.20e-009	37	6	0.38	0.62	0.66	HNF3	BP	1, 3, 5		
8	8.90e-008	37	7.20e-009	39	7	0.44	0.51	0.67	VBP	BP, MD	2, 5		
9	1.10e-007	38	1.00e-008	41	23	1.40	1.80	0.63	NKX62	AA	1, 2, 4	PAL,TR	
10	4.50e-007	45	3.40e-008	45	13	0.81	0.91	0.72	LHX3	AA, BP, ME	1, 2, 4, 5		
11	8.50e-007	46	7.80e-008	50	25	1.60	2.70	0.66	HNF1	AA, BP	1, 2	PAL	
12	9.30e-007	48	7.50e-008	49	10	0.62	0.72	0.65	HNF4_0	AA	1, 2, 4, 5	TR	
13	9.40e-007	49	7.90e-008	51	14	0.88	0.89	0.62	HNF1	BP, ME	1, 2, 3, 4, 5		
14	1.50e-006	53	1.70e-007	57	5	0.31	0.48	0.62	APOLYA	BP	1, 4		
15	1.60e-006	54	1.80e-007	58	7	0.44	0.81	0.62	FXR	BP	1, 3, 4, 5	PAL	
16	1.80e-006	55	1.40e-007	55	9	0.56	0.51	0.73	ZID	AA, BP, ME	1, 2		
17	2.30e-006	56	1.50e-007	56	15	0.94	1.50	0.61	AMEF2	AA, BP	2	PAL,TR	
18	5.80e-006	65	1.10e-001	159	67	4.20	5.80	0.64	TAL1BE	ME	1, 2, 3, 4, 5		
19	6.30e-006	66	5.10e-007	66	11	0.69	1.10	0.73	FREAC2	BP	3, 4, 5	PAL	
20	7.40e-006	67	9.30e-007	71	20	1.20	1.10	0.66	LHX3	AA, BP	1, 3, 4, 5		
21	8.00e-006	69	6.80e-007	67	12	0.75	0.68	0.91	HNF4_0	AA, ME	1, 2, 3, 4, 5		
22	8.20e-006	70	7.50e-007	69	9	0.56	0.81	0.68	POU1F1	BP	1, 3, 5	PAL	
23	2.30e-005	75	5.20e-006	86	28	1.80	1.90	0.83	XFD1	BP	1, 2, 3, 5	PAL	
24	2.40e-005	78	1.30e-001	161	34	2.10	2.40	0.65	CACBIN	BP, ME	1, 2, 3, 4, 5		
25	3.60e-005	83	4.90e-006	85	26	1.60	1.90	0.70	HNF1	BP	2, 4, 5		
26	3.80e-005	84	2.90e-006	83	9	0.56	0.51	0.68	TBP	BP, MD, ME	1, 2, 3, 4, 5		
27	3.90e-005	86	1.20e-001	160	38	2.40	3.00	0.83	MYOGEN	ME	1, 2, 3, 4, 5	PAL	
28	4.20e-005	87	3.10e-006	84	11	0.69	0.70	0.62	HNF4	BP	1, 2		
29	1.00e-004	94	1.10e-005	90	21	1.30	1.30	0.78	CDPCR3	MD	1, 2, 3, 4, 5		
30	1.30e-004	96	1.10e-005	90	13	0.81	1.10	0.63	HNF3AL	BP	1, 2, 4		
31	1.30e-004	96	1.10e-005	90	15	0.94	0.85	0.79	COUP	BP, MD, ME	1, 2, 4, 5		
32	1.60e-004	100	1.20e-005	95	26	1.60	2.40	0.71	HNF1	BP	1, 3, 4, 5	PAL,TR	