N ID)	Motif	W	Z	1	W	ZΙ	
1 12	217	PU.1:IRF8	0.38	6.33	0.24			GGAA_TGAAA
2 10	91	ISRE	0.26	6.32	0.08			ASTITCASTITC
3 99	92	RUNX1	1.00	6.21	1.00			AACCACA
4 84	14	NFKB1	0.63	5.95	0.60			GGG TTTCC
5 12	259	BATF3	0.62	5.75	0.43			TGACTCA
6 99	93	IRF4	0.94	5.39	0.16			-GAAAcc
7 95	55	bZIP:IRF	0.43	4.25	0.10			aTITCA T TAAT
8 10	76	Oct4	0.27	3.40	0.15			ATTIGCAT
9 24	ļ	V_IRF2_01	0.04	3.38	0.01			GAAAAG=GAAAS=
10 82	25	EBF1	0.17	3.24	0.04			-CcCxeGgGA
11 86	5	V_ELK1_01	0.26	2.70	0.08			ACEGGAAGT C.
12 94	12	ETS:RUNX	0.15	2.67	0.06			AGGAxGTGG.
13 10)33	EWS:FLI1-fusi	0.27	2.05	0.07			- CaGGAAai
14 10	74	T1ISRE	0.06	2.04	0.01			ACTITCETT
15 97	76	PAX3:FKHR-fus	0.14	1.73	0.02			SCSTGACTAAT.
16 19	905	USF1	0.20	1.73	0.07			G-CATGTGAC
17 18	374	TFAP2A(var.3)	0.04	1.65	0.00			cCc cGGc
18 18	359	Tcf21	0.07	1.56	0.05			AACAGCTGTT
19 16	597	PAX9	0.12	1.48	0.06			GTCACGCATGA TGS
20 9		V_MEF2_04	0.06	1.36	0.03			TGTT CTAAAAATAGAA
		Znf281	0.05					TGGGGGA=g
		CREM	0.11					_s TGAcgTCA _c
			-0.15					ECATAAT TAT
24 19			-0.26					TCCAGA
25 3			-0.01					TCCCAGcTACTTTGGGA
26 17		·	-0.14					ATGGGATGE
			-0.20					TAATCAAATTA
28 59	99	V_NFAT_Q6	-0.64	-1.65	0.18			GGAAA
29 69	93	V_IK2_01	-1.48	-1.72	0.44			-gGGA
30 11	L50	TEAD	-0.64	-3.26	0.35			= TGGAAT ==