

	motif_id	motif_alt_id	consensus	E-value	bin_location	sites_in_bin	total_sites	p_success
1	CMTCTCYCTTCTCMV	MEME-3	CMTCTCYCTTCTCMV	1.1e-11	0	67	136	0.17695
2	GHKGCCAVWSAGKGG	MEME-1	GHKGCCAVWSAGKGG	9.8e-10	0	35	50	0.18930
3	TKSCATGCTTSCYTG	MEME-2	TKSCATGCTTSCYTG	1.3e-05	0	15	16	0.17695
4	MA0739.1	Hic1	RTGCCAACY	2.3e-04	0	14	133	0.01220
5	MA1618.1	Ptf1a	NNACAGATGTTNN	4.6e-04	0	14	139	0.01230
6	MA0698.1	ZBTB18	NATCCAGATGTKB	1.1e-03	0	54	93	0.29098
7	MA0832.1	Tcf21	RYAACAGCTGTTRN	1.2e-03	0	7	26	0.01027
8	MA0521.1	Tcf12	RACAGCTGSNV	1.6e-03	0	11	85	0.01224
9	MA0101.1	REL	BGGRNWTTC	2.0e-03	0	83	116	0.45418
10	MA0783.1	PKNOX2	TGACAGGTGTCA	2.4e-03	0	6	17	0.01022
11	MA1522.1	MAZ	BBCCCCTCCCY	3.9e-03	0	34	91	0.13878
12	MA0738.1	HIC2	RTGCCCRSB	1.0e-02	0	13	151	0.01220
13	MA0500.2	MYOG	NDRCAGCTGYHN	1.2e-02	0	11	122	0.01022
14	MA1641.1	MYF5	NVACAGCTGTBN	1.5e-02	0	11	125	0.01022
15	MA0669.1	NEUROG2	RACATATGTC	1.5e-02	0	33	74	0.17719
16	MA0048.2	NHLH1	CGCAGCTGCK	1.5e-02	0	10	99	0.01018
17	MA1619.1	Ptf1a(var.2)	NNACAGCTGTNN	1.8e-02	0	11	127	0.01022
18	4-RCAAGGVAGCW	STREME-4	RCAAGGVAGCW	2.5e-02	0	94	126	0.51837
19	MA1635.1	BHLHE22(var.2)	NVCAGCTGBN	3.4e-02	0	8	60	0.01018

Showing 1 to 19 of 87 entries, 8 total columns