## Motifs in ER

forward <b>n</b>	1 <b>1</b> RC	forward <b>I</b>	n2 RC	forward <b>n</b>	n3 RC
A.CA. A.CA. + A.A. + A.A.A.A.A.A.A.A.A.A.A.A.A.A.	TOTOTOTOTOTOTO	_AGG-CATGACCI	ACGTCATGCCCT.	CAGES C COUTG	CACCCC S SECTION
AAALTGAAACC	<b>IRF1</b> 1.2054e-02	AGGTCA TGACCT	<b>ESR1</b> 0e+00	_cC_CATG_Cc_	<b>ESR1</b> 1.3965e–04
GGAAGGAAGGAAGG	<b>EWSR1-FLI1</b> 2.1894e-02	ACCIVA IC.CCI	<b>ESR2</b> 0e+00	AGGTCA TG_CCT	<b>ESR2</b> 2.3777e-04
c_TGT	<b>SOX10</b> 8.0076e-02	TAGTCA TCAQUA T	<b>PPARG</b> 1.1102e-15	IAGTCA LEACCEA I	<b>PPARG</b> 2.1509e-03
<b>A</b> SGGAA	<b>SPIB</b> 8.8257e–02	AGGTÇA	NR4A2 8.5007e-06	GG_cAAAGGTCA	<b>PPARG::RXRA</b> 2.6645e-03
AGGGTA.CAGC	<b>Spz1</b> 1.3698e–01	TGGCA GCCAA	TLX1::NFIC 1.0486e-03	AGGTCAS	<b>NR4A2</b> 3.0525e-03
forward <b>m4</b> RC		forward <b>m5</b> RC			
SCTGSC SSCOOAG	CTCCC CAGE	CCTGGGG	CCCCAGG		
TGCCAGCCAA	<b>TLX1::NFIC</b> 5.4367e-07	CCCAGGGA	<b>EBF1</b> 1.5332e-05		
TGLSAGGGGSS	<b>INSM1</b> 3.0891e-04	GCC	<b>TFAP2A</b> 7.5218e-04		
AGGTCA TGACCT	ESR1 8.1143e-03	SERAWAA WATE S	<b>Zfp423</b> 1.6471e–03		
TICC GGAAS	<b>Stat3</b> 1.063e-02	<u>TGTSAGGGGSS</u>	<b>INSM1</b> 4.5059e-03		
RICTGG	Hand1::Tcfe2a 1.8439e-02	GGGGCC_AAGGGGG	<b>PLAG1</b> 1.0278e-02		