





## QC report

Top scoring motifs for Assay 13863

Protein ID: pTH14270.1 Gene: MTERF4.FL Domain: mTERF Flag: Reject Array: 1M-ME

8 mer E-scores for probeset 'all'			8 mer Z-scores for probeset 'all'		
Forward:		Reverse:	Forward:		Reverse:
					
Top 10	Scores	Alignment	Top 10	Scores	Alignment
AAGCGCTT	0.33493	--AAGCGCTT----	ACCCGGGT	4.24762	---ACCCGGGT---
AGGGCGTA	0.32537	---AGGGCGTA----	CTGCCTCC	4.08919	-----CTGCCTCC-
GCTACCAC	0.32028	-----GTGGTAGC--	GATCGTTA	3.95171	--GATCGTTA----
GATCGTTA	0.30555	----GATCGTTA---	CAAAAGCA	3.83717	----TGCTTTTG--
CCTTCTCA	0.29866	----TGAGAAGG--	CGATGAAC	3.83207	----GTTTCATCG-
CGGATCTA	0.29379	---CGGATCTA----	CCCTTCTC	3.80020	----CCCTTCTC--
ACTCTAAG	0.29354	-----CTTAGAGT	ACTCTAAG	3.79711	-----ACTCTAAG
GGAGCAGA	0.29231	--GGAGCAGA----	AGGGCGTA	3.76979	TACGCCCT-----
CCCCTACC	0.29212	GGTAGGGG-----	AGTGATAG	3.75668	-----CTATCACT-
ACAGTGGC	0.29194	---ACAGTGGC----	CCTTCTCA	3.73192	-----CCTTCTCA-