





QC report

Top scoring motifs for Assay 13782

Protein ID: pTH14314.2 Gene: PHF19.FL Domain: Unknown Flag: Reject Array: 1M-HK

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
TCTCGAGA	0.42324	----TCTCGAGA--	GTGTACAC	7.18815	-GTGTACAC--
GTGTACAC	0.41292	----GTGTACAC--	TCTCGAGA	5.17014	TCTCGAGA---
TTTGCAAA	0.39163	----TTTGCAAA--	CGTTATAA	4.95466	-CGTTATAA--
AAGCGCTT	0.38382	AAGCGCTT-----	GCTTATAA	4.90507	---TTATAAGC
GCTTATAA	0.37951	----GCTTATAA--	TTTGCAAA	4.89658	--TTTGCAAA--
ATGCGCAT	0.36974	--ATGCGCAT----	CAATATTG	4.75730	-CAATATTG--
CAATATTG	0.36765	----CAATATTG--	AACAACAA	4.69877	TTGTTGTT---
AACAACAA	0.36711	-----TTGTTGTT	ATGTTACA	4.49534	ATGTTACA---
GTTATAAC	0.36549	----GTTATAAC--	GAAATTTC	4.49458	--GAAATTTC-
GTATACAA	0.36395	----GTATACAA--	ATACTAAC	4.46600	--GTTAGTAT-