


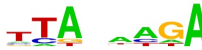


## QC report

Top scoring motifs for Assay 13784

Protein ID: pTH14317.2 Gene: PREB.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
CTCTAGAG	0.43441	----CTCTAGAG-	CTCTAGAG	7.38349	--CTCTAGAG-
GTGTACAC	0.38438	----GTGTACAC-	GTGTACAC	5.97532	---GTGTACAC
GTACAACA	0.37429	--TGTTGTAC--	TATTAATA	5.86924	---TATTAATA
ATACGTTA	0.36976	TAACGTAT----	ATTATAAT	5.76819	--ATTATAAT-
GATGCATC	0.36905	-----GATGCATC	CATATATG	5.67998	--CATATATG-
GTTTAAAC	0.36767	-----GTTTAAAC-	CTTGTAAC	5.49173	--CTTGTAAC-
TATTAATA	0.36133	-----TATTAATA	GTCGACTA	5.01016	GTCGACTA---
CGTATACG	0.35565	--CGTATACG--	AACAACAA	4.81344	---TTGTTGTT
GTATACAC	0.35264	----GTATACAC-	TTTGCAAA	4.80991	--TTTGCAAA-
GTTTATAA	0.35260	----TTATAAAC-	TCAGCTGA	4.77278	-TCAGCTGA--