





## QC report

Top scoring motifs for Assay 13768

Protein ID: pTH14317.1 Gene: PREB.FL Domain: Unknown 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
CCGTAACA	0.39362	--CCGTAACA-	CAATTCCC	5.63084	--CAATTCCC--
AAACACAC	0.34153	---AAACACAC	GCGCAACA	5.02635	GCGCAACA----
CGAGAGAA	0.33817	---CGAGAGAA	CAAATATC	4.63945	--CAAATATC--
GCGCAACA	0.33327	GCGCAACA---	ATTAATTA	4.48942	ATTAATTA----
CCAACACA	0.32398	--CCAACACA-	TATAAATA	4.45289	TATAAATA----
TATAAATA	0.32220	--TATAAATA-	TGGTGTAA	4.41668	---TTACACCA-
ACCAATAC	0.32182	-ACCAATAC--	AACACAGA	4.27230	AACACAGA----
TGGTGTAA	0.32022	---TTACACCA	AACAACGC	4.26383	----AACAACGC
CTCTAATC	0.31745	-CTCTAATC--	GCTAGAGA	4.16193	GCTAGAGA----
GGGTAA	0.31430	--TTTACCCC-	AGATTAAA	4.12995	-TTTAATCT---