





## QC report

Top scoring motifs for Assay 13736

Protein ID: pTH14325.1 Gene: SAFB2.DBD 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
TGTGCACA	0.38775	TGTGCACA-	TGTGCACA	6.96751	--TGTGCACA-
AAATATTT	0.37547	-AAATATTT	GCACGTGC	5.57851	--GCACGTGC--
GCACGTGC	0.36833	GCACGTGC-	ATAATTAT	5.50643	--ATAATTAT--
TAAATTTA	0.36710	TAAATTTA-	TGTATACA	5.28067	--TGTATACA-
AATTTCAA	0.36692	-AATTTCAA	ATCAAATA	5.22730	--TATTTGAT-
AAACGTTT	0.36430	AAACGTTT-	TAACGAAA	5.13500	-TTTCGTTA---
AAAAAATA	0.35892	AAAAAATA-	TTGTACAA	4.97341	----TTGTACAA
CAAAGCAA	0.35556	CAAAGCAA-	AGAATTCT	4.75329	--AGAATTCT--
TTACGTAA	0.35287	TTACGTAA-	ATATTTAC	4.74791	--ATATTTAC--
CAAATTTG	0.35005	CAAATTTG-	AAATATTT	4.73545	AAATATTT----