





QC report

Top scoring motifs for Assay 13771

Protein ID: pTH14328.2 Gene: SNAPC2.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
GATGCATC	0.44212	GATGCATC-	GTGTACAC	6.90771	-GTGTACAC-
TGTTAACA	0.41453	TGTTAACA-	GATGCATC	6.74121	GATGCATC--
TGTGCACA	0.41011	TGTGCACA-	TGTTAACA	6.26495	--TGTTAACA
GTTGCAAC	0.40709	GTTGCAAC-	GTACAACA	5.69451	--TGTTGTAC
TACACACA	0.39602	TACACACA-	ACGTTAAC	5.58789	-ACGTTAAC-
GTTATAAC	0.39593	-GTTATAAC	ATTTAAAT	5.43782	-ATTTAAAT-
GTGTACAC	0.39011	-GTGTACAC	TCTCGAGA	5.42818	--TCTCGAGA
TATGCATA	0.38792	TATGCATA-	AGATATCT	5.29208	--AGATATCT
GTACAACA	0.38667	-GTACAACA	ATGTTAAC	5.25880	-ATGTTAAC-
AACAACAA	0.38143	-AACAACAA	TACACACA	5.23098	TGTGTGTA--