





## QC report

Top scoring motifs for Assay 13518

Protein ID: pTH13946.2 Gene: AC008770.FL Domain: zf-C2H2 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
AGTTGGTA	0.40575	----AGTTGGTA--	ATTGCAAT	4.41502	ATTGCAAT---
ATTGCAAT	0.40061	-----ATTGCAAT	CTTGCAAG	4.37539	CTTGCAAG---
CTTGCAAG	0.39127	--CTTGCAAG----	CGTAACAA	3.80740	--TTGTTACG-
ATGTATGC	0.36991	---ATGTATGC---	ATGCATAC	3.76075	-ATGCATAC--
CGATAACA	0.36370	----TGTATCG--	AAGTACTT	3.74581	-AAGTACTT--
GTACATTA	0.35500	-TAATGTAC-----	AGTTGGTA	3.70703	--TACCAACT-
AACACGTA	0.35404	AACACGTA-----	ATGTATGC	3.70664	---GCATACAT
TATAAGCA	0.35403	----TATAAGCA--	ATACATAT	3.70418	-ATACATAT--
CGTAACAA	0.35256	----CGTAACAA--	ACGTACTA	3.64852	---ACGTACTA
CTGTATTA	0.35216	---CTGTATTA---	CTCTAGAG	3.64092	-CTCTAGAG--