





QC report

Top scoring motifs for Assay 13502

Protein ID: pTH13946.1 Gene: AC008770.FL Domain: zf-C2H2 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
AACGTTGT	0.40743	--AACGTTGT--	TGTATACA	3.68924	-----TGTATACA
TTGTACAA	0.40132	---TTGTACAA-	AACATGTT	3.47191	-AACATGTT----
ATGATCAT	0.39346	--ATGATCAT--	ACACTAAT	3.31274	-ATTAGTGT----
ACAACAAT	0.38943	--ATTGTTGT--	AGACGTCT	3.25876	AGACGTCT-----
AATTGTTA	0.38872	-AATTGTTA---	ACAACAAT	3.24955	-ATTGTTGT----
TAACGAAA	0.38761	-TTTCGTTA---	AACGTTGT	3.20450	-AACGTTGT-----
AATTTGTA	0.38705	-AATTTGTA---	AATTTGTA	3.17455	--AATTTGTA----
AAATTGTT	0.38656	AAATTGTT----	AATTAATT	3.13390	-AATTAATT-----
ATGTACAT	0.38633	--ATGTACAT--	CAACGTTG	3.12170	CAACGTTG-----
CAGAAACA	0.38366	----TGTTTCTG	ACAAAATA	3.03125	-TATTTTGT-----