



QC report

Top scoring motifs for Assay 14451

Protein ID: pTH15569.2 Gene: CTCF Domain: zf-C2H2 Flag: Unsure 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
GCGCCCCC	0.46749	-GCGCCCCC---	AATATATT	6.36113	---AATATATT---
AAAATAAA	0.46161	AAAATAAA----	GCGCCCCC	6.36026	-----GCGCCCCC
GTGGCGCA	0.45191	---GTGGCGCA-	AATAAATT	6.23340	---AATAAATT---
ATGCGCCA	0.44977	---ATGCGCCA-	GTGGCGCA	6.10206	-----TGCGCCAC-
AATATTAA	0.44784	AATATTAA----	AATATTAA	5.92103	--TTAATATT----
AATAAATT	0.44762	--AATAAATT--	AATTTAAA	5.72159	TTTAAATT-----
AATATATT	0.44692	--AATATATT--	ATGCGCCA	5.60508	-----ATGCGCCA--
GGGCGCA	0.44217	---TGCGCCCC	CCACTAGG	5.56340	-----CCACTAGG
ATATTAAT	0.43926	-ATATTAAT---	GGGCGCA	5.47540	-----TGCGCCCC-
AAATAAAT	0.43877	-AAATAAAT---	ATAAAAAA	5.35767	-----ATAAAAAA--