





## QC report

Top scoring motifs for Assay 13842

Protein ID: pTH13931.2 Gene: ZBED4 Domain: zf-BED 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
CGGTACCG	0.43244	--CGGTACCG---	AATCGATT	3.63411	---AATCGATT--
ACGGTACC	0.40299	-ACGGTACC----	ACTTATTG	3.61788	-----ACTTATTG
ACTTATTG	0.39097	--CAATAAGT---	CGGTACCA	3.50811	TGGTACCG-----
TACGCGTA	0.37617	-----TACGCGTA	TACCGGTA	3.37047	---TACCGGTA--
CAATATTG	0.37309	--CAATATTG---	ACGGTACC	3.34301	-GGTACCGT----
CATAATAG	0.37248	CATAATAG-----	TACGCGTA	3.31034	--TACGCGTA---
ACAATAAG	0.37043	-ACAATAAG----	ATAACGTA	3.21844	--ATAACGTA---
ACAAAATA	0.36601	-ACAAAATA----	ACATTATG	3.18100	-CATAATGT----
CGGTACCA	0.36393	--CGGTACCA---	AAACAGTT	3.13977	--AAACAGTT---
GGTTCCGA	0.36183	-TCGGAACC----	ATATGTGC	3.12733	---GCACATAT--