





QC report

Top scoring motifs for Assay 13743

Protein ID: pTH14311.2 Gene: PA2G4.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
GATATGTA	0.36111	-GATATGTA--	ACTTGCAT	4.59686	---ACTTGCAT--
TACGCGTA	0.35750	-TACGCGTA--	CGGTGGAA	4.35914	-----TTCCACCG
ATAGTAAT	0.35284	--ATAGTAAT--	TACGCGTA	4.26927	--TACGCGTA---
ACACAATC	0.34540	GATTGTGT---	ATAGTAAT	4.18666	---ATTACTAT--
GCTGTGAA	0.33503	-GCTGTGAA--	TATTTGCA	4.16153	--TATTTGCA---
AACGGCTC	0.33433	-GAGCCGTT--	GTTCGAAC	4.06180	---GTTCGAAC--
TATTTGCA	0.33137	-TATTTGCA--	ACACAATC	4.03266	--GATTGTGT---
ACTTGCAT	0.32984	---ATGCAAGT	ACAAATCC	3.94185	-GGATTTGT----
ACATTACA	0.32978	TGTAATGT----	CATATATG	3.88100	CATATATG-----
ACAATTAG	0.32443	--ACAATTAG--	AACGGCTC	3.82945	---AACGGCTC--