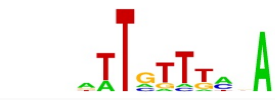


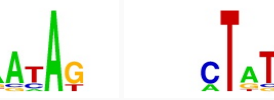


QC report

Top scoring motifs for Assay 13887

Protein ID: pTH14335.1 Gene: RFX8.DBD Domain: RFX_DNA_binding 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
ATTGTTTC	0.35186	-ATTGTTTC--	TTAAATAA	5.20392	TTAAATAA----
GTTGTAGA	0.34316	-GTTGTAGA--	AAAAACCA	5.13786	--AAAAACCA--
CACCAATA	0.33548	TATTGGTG----	AAATAATA	4.98430	--AAATAATA--
CGAGAGAA	0.33158	--TTCTCTCG--	TAACGAAA	4.85560	--TAACGAAA--
TGTTGTAA	0.32739	---TGTTGTAA-	CTGTTGTA	4.80158	---TACAACAG-
TATAAATA	0.32712	---TATTTATA-	ATTTTGAC	4.71663	-GTCAAAAT---
TAACGAAA	0.32669	-TTTCGTTA--	GTTGTAGA	4.65775	----GTTGTAGA
CTCTAATC	0.32515	----GATTAGAG	AGGTGATA	4.55854	--AGGTGATA--
AGTAATAT	0.32462	--ATATTACT--	AATAATAG	4.53899	---AATAATAG-
AAACATTC	0.32173	GAATGTTT----	ATATAAAT	4.52220	---ATTTATAT-