





QC report

Top scoring motifs for Assay 13745

Protein ID: pTH14313.2 Gene: PCGF6.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
AAGTAAGT	0.35783	-----AAGTAAGT	AAGTAAGT	5.79221	--ACTTACTT--
ATTTGTAA	0.34898	---ATTTGTAA--	GTAATGTA	4.76361	TACATTAC----
CTCAACAC	0.34461	---GTGTTGAG--	ATTTGTAA	4.72843	----TTACAAAT
AATAGCAA	0.34393	---AATAGCAA--	ATATCACA	4.70874	---ATATCACA--
TTCGAAAA	0.34262	----TTCGAAAA--	CATCACTC	4.70605	--CATCACTC--
GCTACTAA	0.33872	----TTAGTAGC--	ATTACTCA	4.70195	---ATTACTCA--
AAAATAAC	0.33753	GTTATTTT-----	GATAACTA	4.67057	--GATAACTA--
AGTTATCA	0.33721	---AGTTATCA--	AGTTATCA	4.63120	-TGATAACT---
AAGCGCTT	0.33710	---AAGCGCTT--	ACATTACA	4.61731	-ACATTACA---
CGCTACAA	0.32925	-----TTGTAGCG	TTCGAAAA	4.60142	---TTTTCGAA--