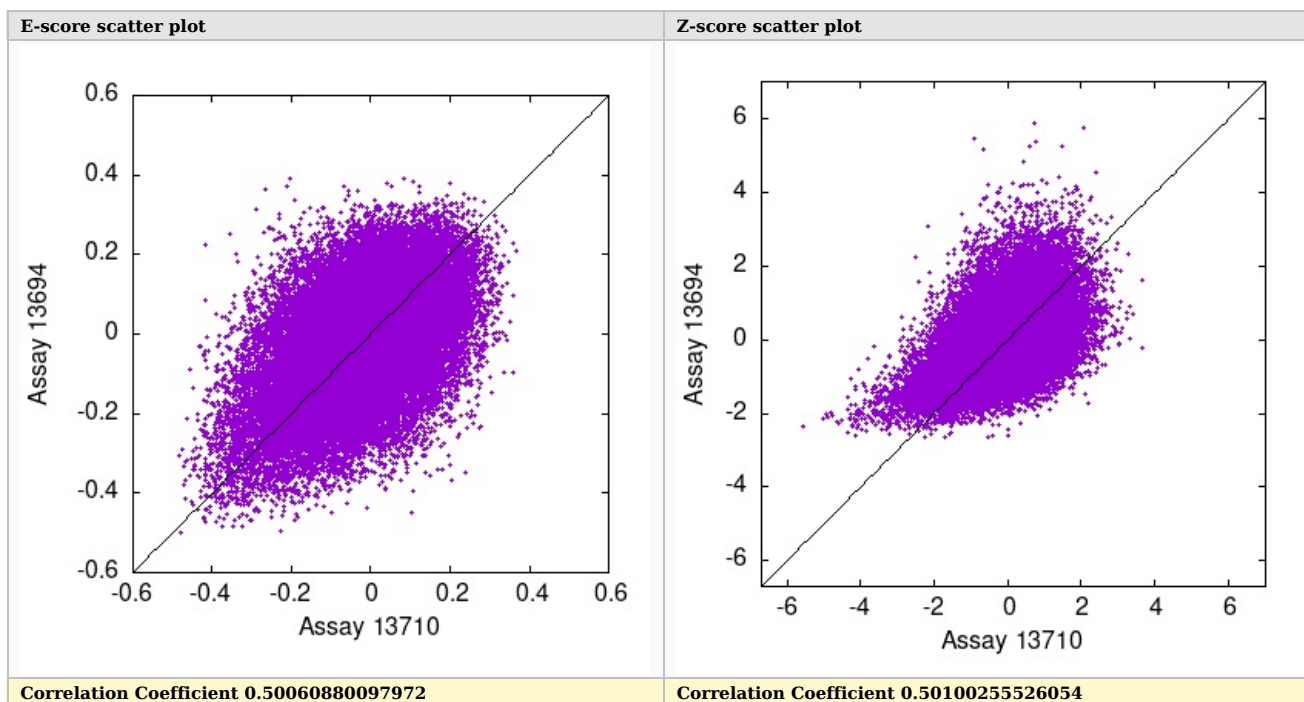


QC report





Top scoring motifs for Assay 13710

Protein ID: pTH13930.2 Gene: ZBED3.FL Domain: zf-BED Flag: Reject Array: 1M-HK

8 mer E-scores for probeset 'all'

Forward:		Reverse:	
Top 10	Scores	Alignment	
TAACGTAA	0.36728	TAACGTAA-----	
ATCAAGTG	0.35988	-CACTTGAT-----	
AGTTTTTC	0.35829	--AGTTTTTC-----	
ATATGTGC	0.35823	GCACATAT-----	
AAAATGTT	0.35272	-AACATTTT-----	
TTAACCAA	0.34751	-----TTGGTTAA	
ACTCCAAT	0.34532	-----ATTGGAGT-	
AAGCCTTA	0.34249	-TAAGCTTT-----	
ACGTATTG	0.34024	--ACGTATTG-----	
ATCAGTAC	0.33810	----GTA CTGAT---	



8 mer Z-scores for probeset 'all'

Forward:		Reverse:
		
Top 10	Scores	Alignment
AGTTTTTC	3.67993	---AGTTTTTC
ATATGTGC	3.67496	--GCACATAT--
ATACATAT	3.40541	--ATACATAT--
ACTCCAAT	3.33113	--ACTCCAAT--
ACTTGTA	3.30427	--TTACAAGT--
TAACGTAA	3.26487	--TAACGTAA--
TCAAACAA	3.22586	TCAAACAA---
TAATAATA	3.22175	---TAATAATA--
CTAATCTA	3.11797	---TAGATTAG--
GCATACAA	3.07649	GCATACAA---



Top scoring motifs for Assay 13694

Protein ID: pTH13930.1 Gene: ZBED3.FL Domain: zf-BED Flag: Reject Array: 1M-ME

8 mer E-scores for probeset 'all'

Forward:		Reverse:	
			
Top 10	Scores	Alignment	
ATTCGAAT	0.39135	---ATTCGAAT---	
ATGTACAT	0.38986	---ATGTACAT---	
GTATAATA	0.38194	-----GTATAATA-	
CTACAACA	0.37964	-----CTACAACA-	
ATAATGCA	0.37830	ATAATGCA-----	
AATTCGAA	0.37270	---AATTCGAA---	
TTGCGCAA	0.37224	---TTGCGCAA---	
CTGTATAC	0.37215	---CTGTATAC---	
GTGCGCAC	0.37086	---GTGCGCAC---	
AACATGTT	0.36545	-----AACATGTT	

8 mer Z-scores for probeset 'all'

Forward:		Reverse:
		
Top 10	Scores	Alignment
ATAATGCA	5.87392	ATAATGCA-----
ATGTACAT	5.76701	---ATGTACAT---
AACATGTT	5.47521	-----AACATGTT
CTGTATAC	5.37367	---CTGTATAC---
GTTTAAAC	5.25408	---GTTTAAAC---
TTGCGCAA	5.24393	---TTGCGCAA---
ATTCGAAT	5.16440	---ATTCGAAT---
GTATAATA	4.83038	-----GTATAATA-
AATTTGTA	4.55577	---AATTTGTA---
TATGTGAA	4.42042	--TATGTGAA----