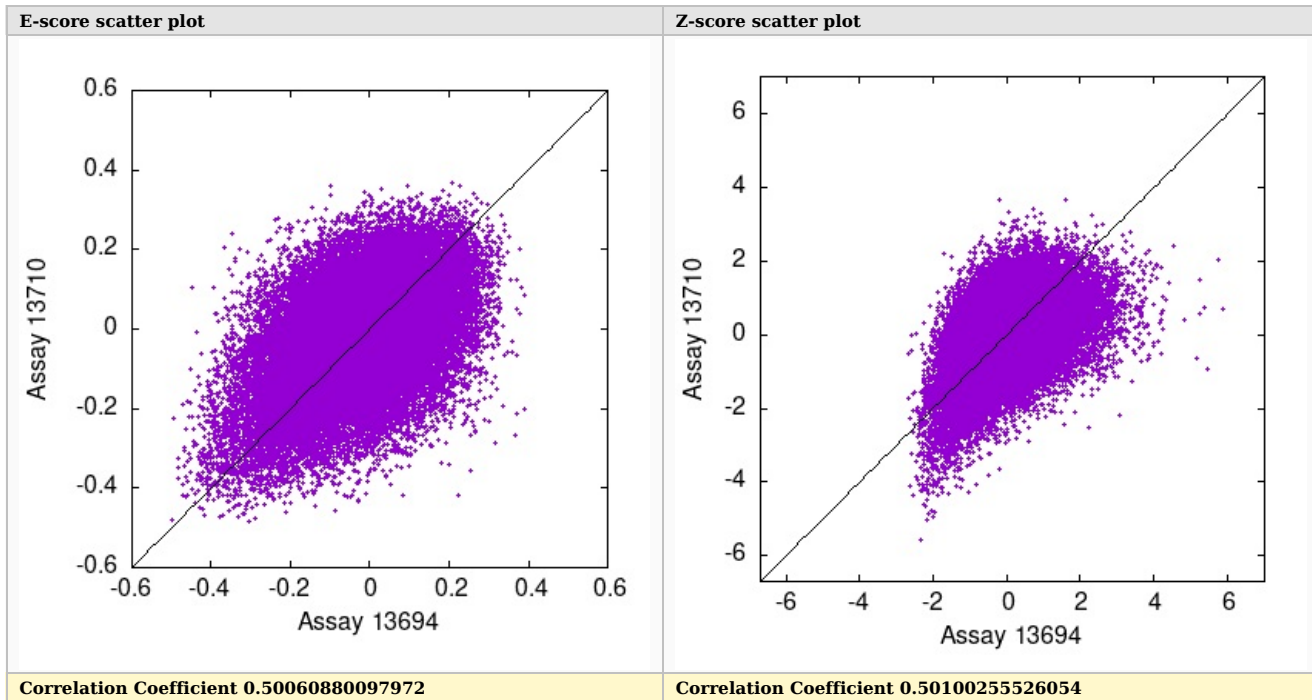


QC report



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'			8 mer Z-scores for probeset 'all'		
Forward:		Reverse:	Forward:		Reverse:
Top 10	Scores	Alignment	Top 10	Scores	Alignment
ATTCGAAT	0.39135	---ATTCGAAT---	ATAATGCA	5.87392	ATAATGCA-----
ATGTACAT	0.38986	---ATGTACAT---	ATGTACAT	5.76701	---ATGTACAT---
GTATAATA	0.38194	-----GTATAATA-	AACATGTT	5.47521	-----AACATGTT
CTACAACA	0.37964	-----CTACAACA-	CTGTATAC	5.37367	---CTGTATAC---
ATAATGCA	0.37830	ATAATGCA-----	GTTTAAAC	5.25408	---GTTTAAAC---
AATTCGAA	0.37270	---AATTCGAA---	TTGCGCAA	5.24393	---TTGCGCAA---
TTGCGCAA	0.37224	---TTGCGCAA---	ATTCGAAT	5.16440	---ATTCGAAT---
CTGTATAC	0.37215	---CTGTATAC---	GTATAATA	4.83038	-----GTATAATA-
GTGCGCAC	0.37086	---GTGCGCAC---	AATTTGTA	4.55577	--AATTTGTA----
AACATGTT	0.36545	-----AACATGTT	TATGTGAA	4.42042	--TATGTGAA----

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'			8 mer Z-scores for probeset 'all'		
Forward:		Reverse:	Forward:		Reverse:
Top 10	Scores	Alignment	Top 10	Scores	Alignment
TAACGTAA	0.36728	TAACGTAA-----	AGTTTTTC	3.67993	----AGTTTTTC
ATCAAGTG	0.35988	-CACTTGAT-----	ATATGTGC	3.67496	--GCACATAT--
AGTTTTTC	0.35829	--AGTTTTTC-----	ATACATAT	3.40541	--ATACATAT--
ATATGTGC	0.35823	GCACATAT-----	ACTCCAAT	3.33113	--ACTCCAAT--
AAAATGTT	0.35272	---AACATTTT-----	ACTTGTA	3.30427	--TTACAAGT--
TTAACCAA	0.34751	-----TTGGTTAA	TAACGTAA	3.26487	--TAACGTAA--
ACTCCAAT	0.34532	-----ATTGGAGT-	TCAAACAA	3.22586	TCAAACAA----
AAAGCTTA	0.34249	-TAAGCTTT-----	TAATAATA	3.22175	---TAATAATA-
ACGTATTG	0.34024	--ACGTATTG-----	CTAATCTA	3.11797	---TAGATTAG-
ATCAGTAC	0.33810	----GTAAGTAT---	GCATACAA	3.07649	GCATACAA----