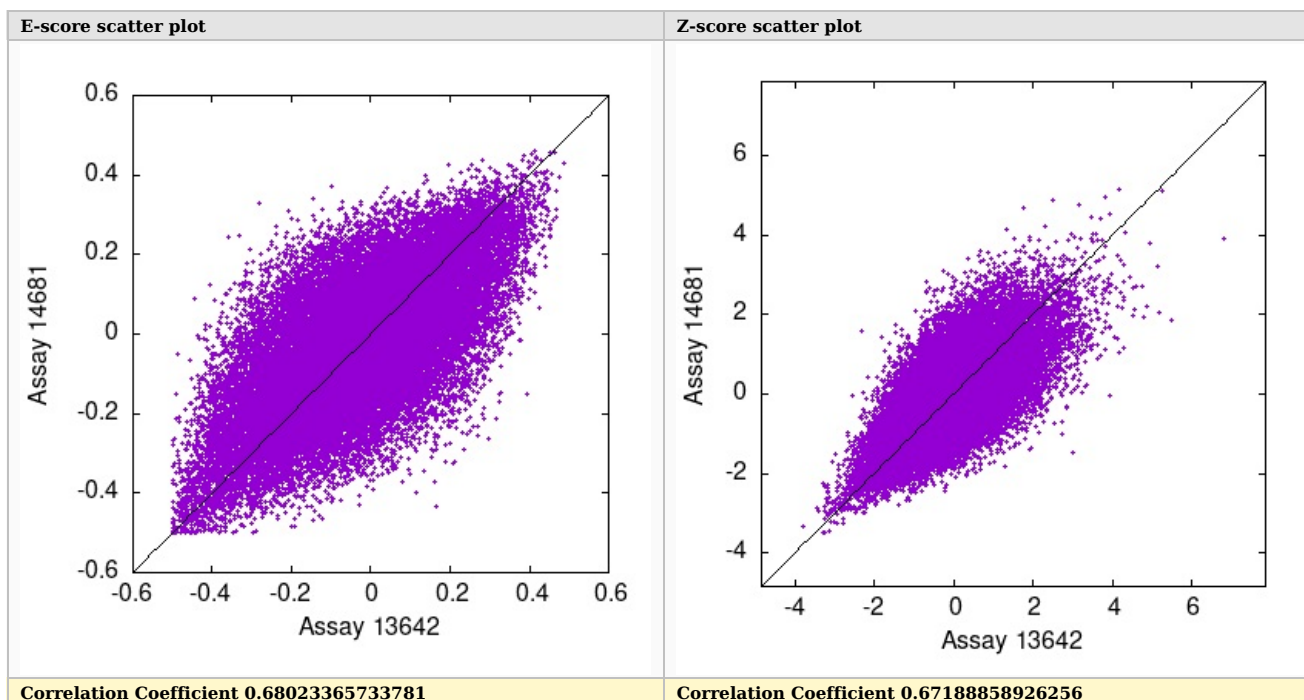


QC report



Correlation Coefficient 0.68023365733781

Correlation Coefficient 0.67188858926256

Top scoring motifs for Assay 13642

Protein ID: pTH13919.2 Gene: GLYR1.DBD Domain: AT_hook Flag: Backup_matched_pair Array: 1M-HK

8 mer E-scores for probeset 'all'

Forward:

Reverse:



Top 10

Scores

Alignment

AATATATT	0.48709	--AATATATT
AATAAATT	0.47669	--AATAAATT
AATTTAAA	0.46807	--AATTTAAA
AAATATTT	0.46748	--AAATATTT
ACAAATA	0.46683	ACAAATA--
AAATATAT	0.46457	--AAATATAT-
ATATATTG	0.45898	-CAATATAT-
TATTAATA	0.45591	--TATTAATA
AAATTGAT	0.45416	-AAATTGAT-
ATTTAAAT	0.45305	-ATTTAAAT-

8 mer Z-scores for probeset 'all'

Forward:

Reverse:



Top 10

Scores

Alignment

AATATATT	6.80307	--AATATATT--
AATTTAAA	5.49601	--TTTAAATT--
AAATATAT	5.25725	--AAATATAT---
GAAATTC	5.16522	----GAAATTC
AATAAATT	5.14720	--AATAAATT--
TATTAATA	4.96037	--TATTAATA--
AAAATAAA	4.88125	AAAATAAA----
ATAATTC	4.79419	--ATAATTC-
GAATATTC	4.77199	--GAATATTC-
AAATTAAT	4.58082	-AAATTAAT--

Top scoring motifs for Assay 14681

Protein ID: pTH13919.3 Gene: GLYR1.DBD Domain: AT_hook Flag: Backup_matched_pair Array: 1M-ME

8 mer E-scores for probeset 'all'

Forward:

Reverse:



Top 10

Scores

Alignment

ATATATAT	0.46168	-ATATATAT-
AAATATAT	0.45536	-AAATATAT-
ATTTAAAT	0.45509	-ATTTAAAT-
ATAATATA	0.45099	ATAATATA--
TAAATTTA	0.44980	TAAATTTA--
AATAATAA	0.44467	-AATAATAA-
ATATATTA	0.44118	-TAATATAT-
TTATATAA	0.43882	-TTATATAA-
ATAATTAT	0.43746	-ATAATTAT-
ATTATATA	0.43596	--ATTATATA

8 mer Z-scores for probeset 'all'

Forward:

Reverse:



Top 10

Scores

Alignment

ATTTAAAT	5.13180	-ATTTAAAT--
AAATATAT	5.10548	-AAATATAT--
AATAATA	4.94166	--ATAATATA-
TTATATAA	4.87884	--TTATATAA
ATATATTA	4.76412	--TAATATAT
ATTATATA	4.68903	--ATTATATA-
ATATATAT	4.53674	--ATATATAT
AATAATAA	4.46489	TTATTATT---
AAAATTTT	4.42557	---AAAATTTT-
TTTATAAA	4.23567	--TTTATAAA-