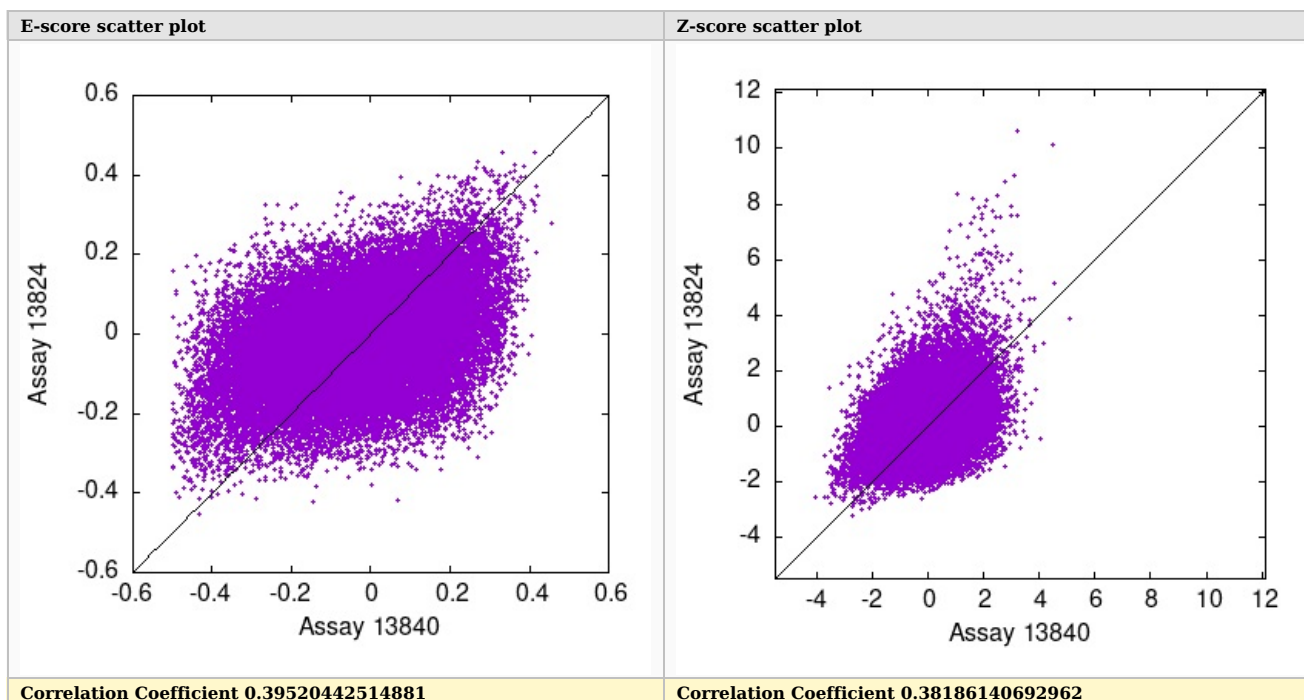


QC report



Top scoring motifs for Assay 13840

Protein ID: pTH13925.2 Gene: SETBP1.DBD.3 Domain: AT_hook Flag: Pass_matched_pair Array: 1M-HK

8 mer E-scores for probeset 'all'

Forward:	Reverse:
Top 10	Scores
AATAAATT	0.45633
AATATATT	0.42327
AAATTAAT	0.41668
ATAATGTA	0.41636
AAATAAAT	0.41447
AATTAATT	0.41297
AATTTAAA	0.41216
ACAAAATA	0.41084
ACATAGAC	0.40734
TATTAATA	0.40433

8 mer Z-scores for probeset 'all'

Forward:	Reverse:
Top 10	Scores
AATAAATT	5.10369
AATATATT	4.56762
AATTAATT	4.51073
AATTTAAA	4.17464
ACATAGAC	4.05674
ATAATGTA	3.90276
GTAATTAC	3.82645
ATATTAAT	3.81979
TATTAATA	3.80017
AATAATGT	3.71484

Top scoring motifs for Assay 13824

Protein ID: pTH13925.1 Gene: SETBP1.DBD.3 Domain: AT_hook Flag: Pass_matched_pair Array: 1M-ME

8 mer E-scores for probeset 'all'

Forward:	Reverse:
Top 10	Scores
AATTAATT	0.45763
ATTTAAAT	0.45514
TAAATTTA	0.43270
AAAAAATA	0.42640
AAAAATAA	0.42038
ATATTTTA	0.41913
AAATAAAA	0.41753
AAATATTA	0.41116
AAATAATT	0.40911
TAAATTAA	0.40909

8 mer Z-scores for probeset 'all'

Forward:	Reverse:
Top 10	Scores
AAAAAATA	10.65616
AATTAATT	10.13525
AAAAATAA	9.01262
ATTTAAAT	8.82092
ATTAATTA	8.35624
AATAAAAT	8.29293
AATTAAT	8.20997
TAAATTAA	8.13580
AAATAATT	7.92912
AAATAAAA	7.91230