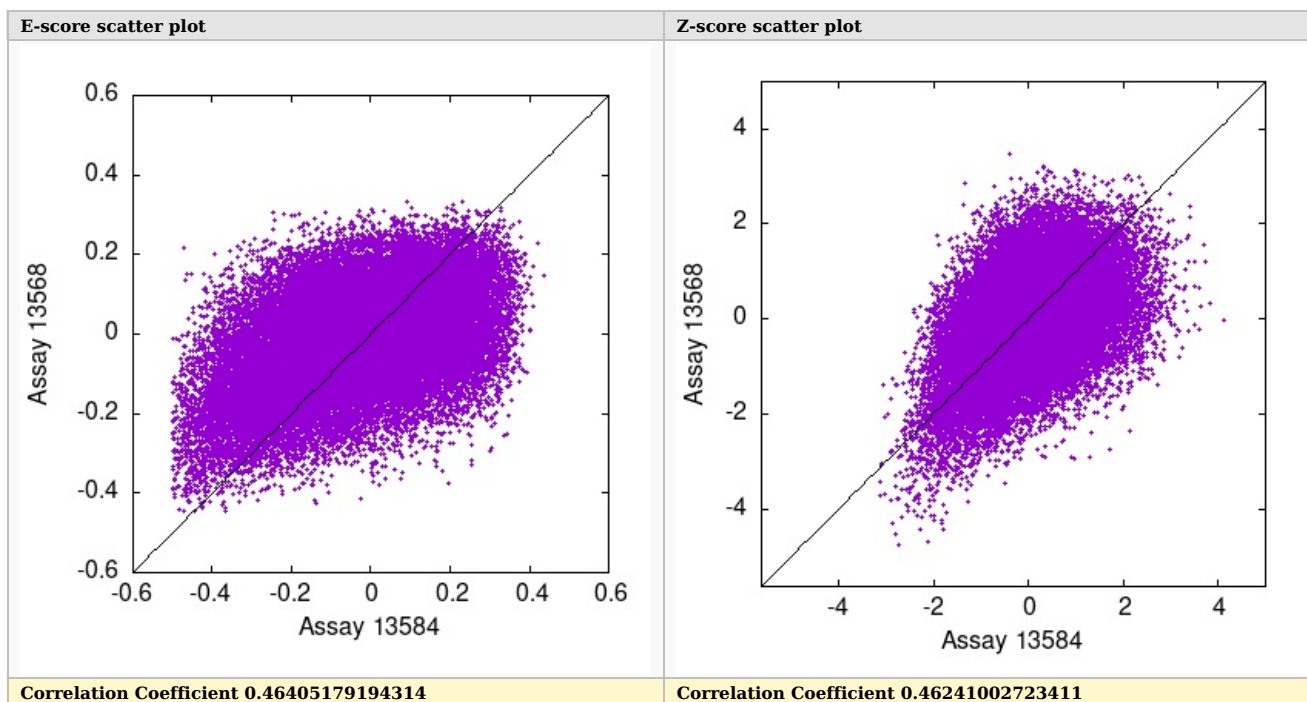


QC report



Top scoring motifs for Assay 13584

Protein ID: pTH13960.2 Gene: DZIP1.DBD Domain: zf-C2H2 Flag: Reject Array: 1M-HK

8 mer E-scores for probeset 'all'

Forward:

Reverse:

ATATT AT

AT AATAT

Top 10

Scores

Alignment

CTGTATTA 0.43754
ATAGTAAT 0.42068
ATATTAAT 0.41142
TATTTGCA 0.40929
TATATGAA 0.40583
AAAACCTTA 0.40281
TACACACA 0.40278
GATATGTA 0.40184
GAAATTTT 0.39711
ACAAATA 0.39597

CTGTATTA--
--ATAGTAAT
--ATATTAAT
-TATTTGCA-
-TATATGAA-
-AAAACCTTA-
-TGTGTGTA-
-GATATGTA-
-GAAATTTT-
-TATTTTGT-

8 mer Z-scores for probeset 'all'

Forward:

Reverse:

ATTAATA

TAATAATA

Top 10

Scores

Alignment

CTGTATTA 4.10800
GCATATAA 3.81530
TAATATAA 3.72981
ATAGTAAT 3.69828
TAATAATA 3.63271
ATATCTTA 3.61135
GATATGTA 3.59688
ATATTAAT 3.49932
TACACACA 3.48835
CATATAAC 3.48615

--TAATACAG-
--GCATATAA-
--TAATATAA-
ATTACTAT--
--TAATAATA-
--TAAGATAT-
--GATATGTA
ATTAATAT--
-TGTGTGTA-
--CATATAAC

Top scoring motifs for Assay 13568

Protein ID: pTH13960.1 Gene: DZIP1.DBD Domain: zf-C2H2 Flag: Reject Array: 1M-ME

8 mer E-scores for probeset 'all'

Forward:

Reverse:

TACATTA

TATGTA

Top 10

Scores

Alignment

ACAATGTA 0.33318
CTGGTAGA 0.33170
TACCAATA 0.32532
ACAAATAT 0.31669
ACTTTAAT 0.31649
ACAAAACA 0.31631
AATCATTA 0.31544
ACCGGTTA 0.31326
AATGTAGT 0.31139
AGTAGTAT 0.31119

---ACAATGTA--
TCTACCAG-----
--TACCAATA--
---ACAAATAT---
---ACTTTAAT---
---ACAAAACA---
--AATCATTA---
---ACCGGTTA---
-----AATGTAGT
-ATACTACT-----

8 mer Z-scores for probeset 'all'

Forward:

Reverse:

TACCTTA

TAGCTTA

Top 10

Scores

Alignment

AAGCGCGA 3.48286
CAACATTG 3.22502
AATAGTGT 3.19100
GTTACTAA 3.12008
ATGACAAC 3.11093
ATGTAGAC 3.09874
ACCCAAAA 3.09682
ACTGCGAA 3.08951
TAACACTA 3.08930
AATTTTAC 3.03811

---AAGCGCGA---
---CAACATTG---
-----ACACTATT-
TGTAGTAA-----
-----ATGACAAC--
-----GTCTACAT
-----ACCCAAAA-
---TTCGCAGT---
---TAACACTA---
---GTAAATT---