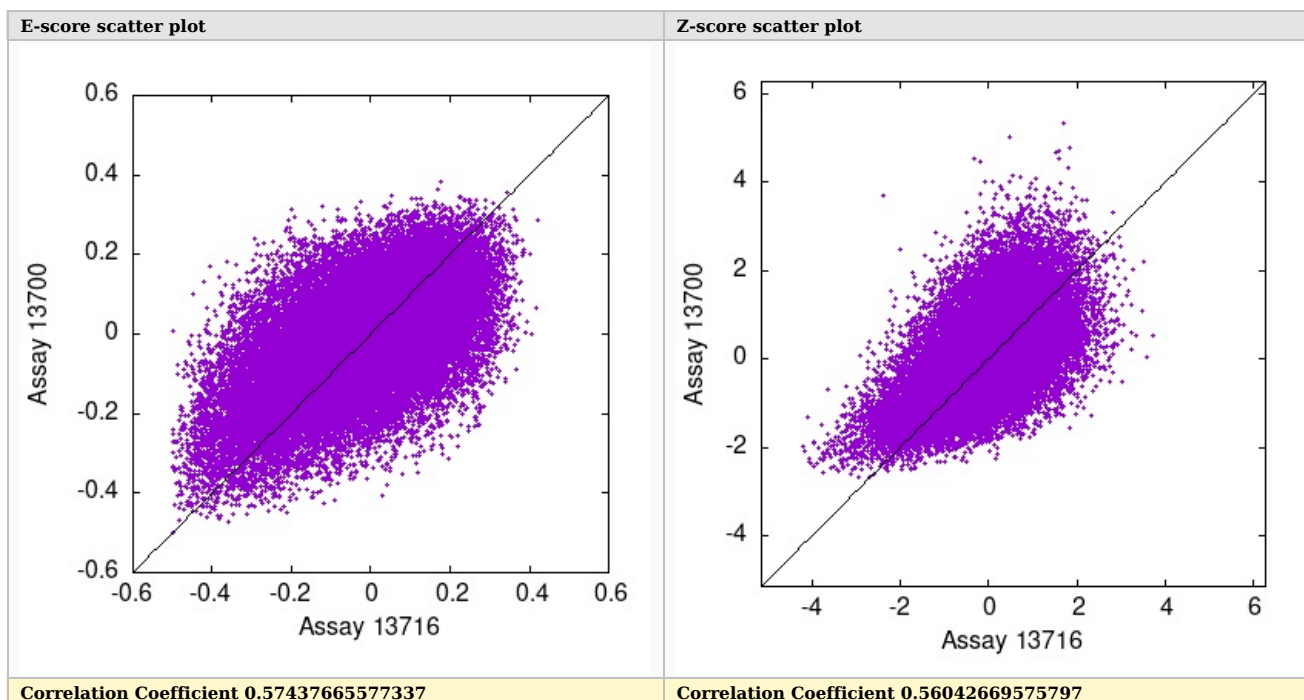


QC report



Top scoring motifs for Assay 13716

Protein ID: pTH14214.2 Gene: TIGD4.DBD Domain: CENP-B_N Flag: Reject Array: 1M-HK

8 mer E-scores for probeset 'all'

Forward:

Reverse:



Top 10

Scores

Alignment

CTATTATC
ATATCACA
GATATGTA
TATTAATA
CACTGATA
CATATCAC
TATTATCA
ATAGTAAT
CATTACTA
ACATTACT

0.42016
0.41708
0.40733
0.40234
0.39423
0.39047
0.38820
0.38755
0.38351
0.38309

-CTATTATC---
---ATATCACA-
-TACATATC---
--TATTAATA--
-----TATCAGTG
---CATATCAC-
---TATTATCA--
---ATTACTAT--
--CATTACTA---
AGTAATGT----

8 mer Z-scores for probeset 'all'

Forward:

Reverse:



Top 10

Scores

Alignment

ATATCACA
GATATGTA
ACATTACT
TATTAATA
TATTTGCA
CTGTATTA
AGAGATAT
AATAAATT
ATACTTTC
ACGTACTA

3.73177
3.60288
3.52855
3.47902
3.34073
3.28242
3.24400
3.23439
3.22002
3.17343

TGTGATAT-----
--GATATGTA---
--AGTAATGT---
--TATTAATA---
---TATTTGCA---
-----CTGTATTA
--AGAGATAT---
--AATAAATT---
----ATACTTTC--
-----ACGTACTA

Top scoring motifs for Assay 13700

Protein ID: pTH14214.1 Gene: TIGD4.DBD Domain: CENP-B_N Flag: Reject Array: 1M-ME

8 mer E-scores for probeset 'all'

Forward:

Reverse:



Top 10

Scores

Alignment

GTATAATA
CAATTTCA
AATATCAC
ATAAATAA
AAACAATA
CTGTATAC
CTCTAATC
ACACACAC
ACTGTATA
TAACGAAA

0.38300
0.36232
0.35430
0.34748
0.34293
0.34278
0.34249
0.34195
0.33937
0.33825

--GTATAATA--
-----CAATTTCA
---AATATCAC--
---ATAAATAA--
--AAACAATA---
GTATACAG-----
---GATTAGAG--
--ACACACAC---
-TATACAGT-----
--TAACGAAA---

8 mer Z-scores for probeset 'all'

Forward:

Reverse:



Top 10

Scores

Alignment

CATAATAC
GTATAATA
AATATCAC
ACAAAGAC
CAATTTCA
CAAGAAAC
AGTAATAT
ACGACAAC
GTACTATA
CGTTAATA

5.32287
5.01666
4.77952
4.70126
4.68611
4.53761
4.51692
4.46782
4.31487
4.16604

-CATAATAC-
GTATAATA--
-AATATCAC-
-ACAAAGAC-
-CAATTTCA-
-CAAGAAAC-
--ATATTACT
-ACGACAAC-
-TATAGTAC-
CGTTAATA--