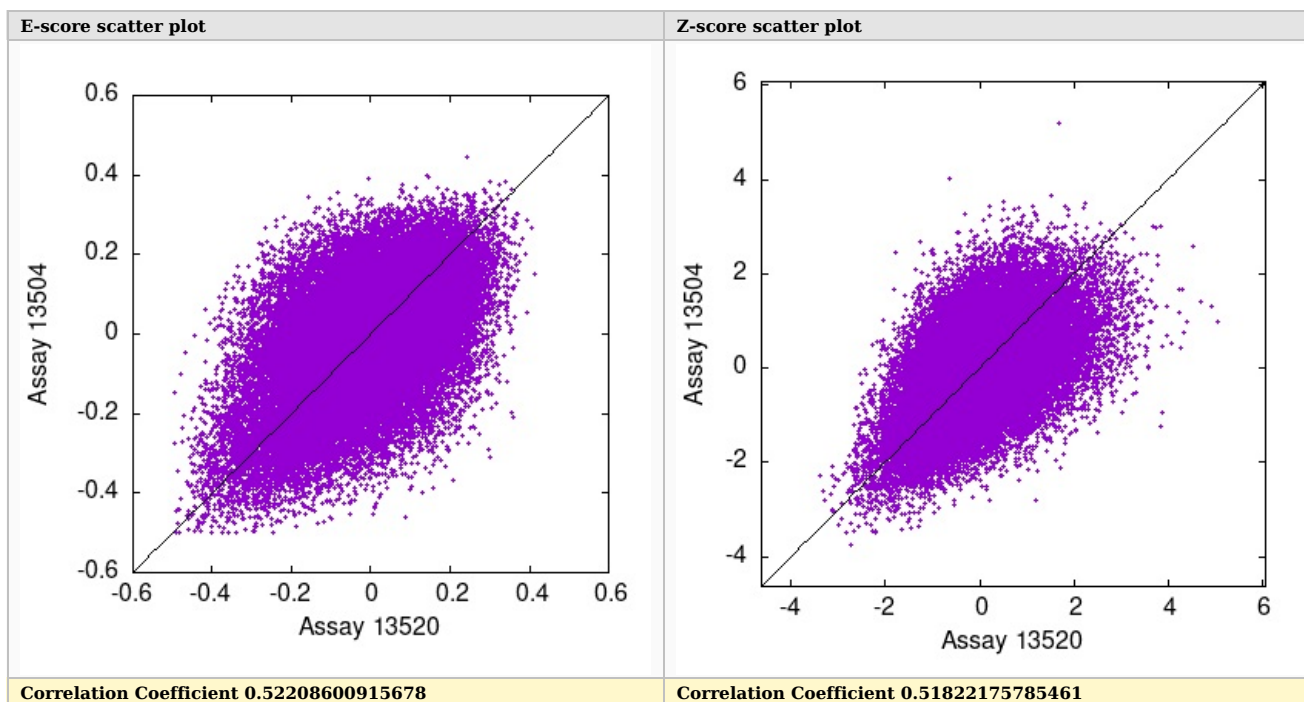


QC report



Top scoring motifs for Assay 13520

Protein ID: pTH13949.2 Gene: AC092835.DBD Domain: zf-C2H2 Flag: Reject Array: 1M-HK

8 mer E-scores for probeset 'all'

Forward:

Reverse:



Top 10

Scores

Alignment

TATTTGCA	0.41178	--TATTTGCA--
AAGGTAGT	0.40733	--AAGGTAGT---
AAGTACTT	0.39527	--AAGTACTT--
ATAGTAAT	0.39521	----ATAGTAAT
CTGTATTA	0.39184	--CTGTATTA--
CATTTACA	0.38409	--CATTTACA---
ATACATTA	0.37601	TAATGTAT----
AAGTAAAA	0.37467	--AAGTAAAA--
ATACTAGC	0.37037	---GCTAGTAT-
ATGTAGTA	0.36986	--ATGTAGTA--

8 mer Z-scores for probeset 'all'

Forward:

Reverse:



Top 10

Scores

Alignment

TATTTGCA	5.07006	---TATTTGCA---
ATAGTAAT	4.92865	-----ATTACTAT
ATACATTA	4.70828	--ATACATTA----
ATGTATAC	4.54790	GTATACAT-----
GTACAACA	4.41926	--GTACAACA----
AAGTACTT	4.32677	AAGTACTT-----
AAGGTAGT	4.31198	-ACTACCTT-----
CTACCAAC	4.28250	--CTACCAAC----
CTGTATTA	4.25188	TAATACAG-----
ATGTATGC	4.23578	GCATACAT-----

Top scoring motifs for Assay 13504

Protein ID: pTH13949.1 Gene: AC092835.DBD Domain: zf-C2H2 Flag: Reject Array: 1M-ME

8 mer E-scores for probeset 'all'

Forward:

Reverse:



Top 10

Scores

Alignment

AGACTACC	0.44642	--GGTAGTCT---
AAACATAA	0.39862	-----AAACATAA
TAAAGAAA	0.39646	--TTTCGTGA---
AGACGAAA	0.38962	--TTTCGTCT---
GGTAGTCA	0.38493	--GGTAGTCA---
ACTACTAT	0.38369	ATAGTAGT-----
ATAATGCA	0.37660	---ATAATGCA---
GTAGTCAA	0.37425	---GTAGTCAA---
ATGAGTAG	0.36711	---CTACTCAT--
AAAACATA	0.36594	-----AAAACATA-

8 mer Z-scores for probeset 'all'

Forward:

Reverse:



Top 10

Scores

Alignment

AGACTACC	5.18131	---AGACTACC
AGACGAAA	4.00419	-AGACGAAA--
ACATACAA	3.65041	-ACATACAA--
ATAATGCA	3.53294	ATAATGCA---
GTAGTCAA	3.49244	-GTAGTCAA--
AACATGCA	3.44270	AACATGCA---
AACATACA	3.42844	AACATACA---
ATAGTCTG	3.42629	--CAGACTAT-
TAACGAAA	3.40187	-TAACGAAA--
ATATACAG	3.39470	-ATATACAG--