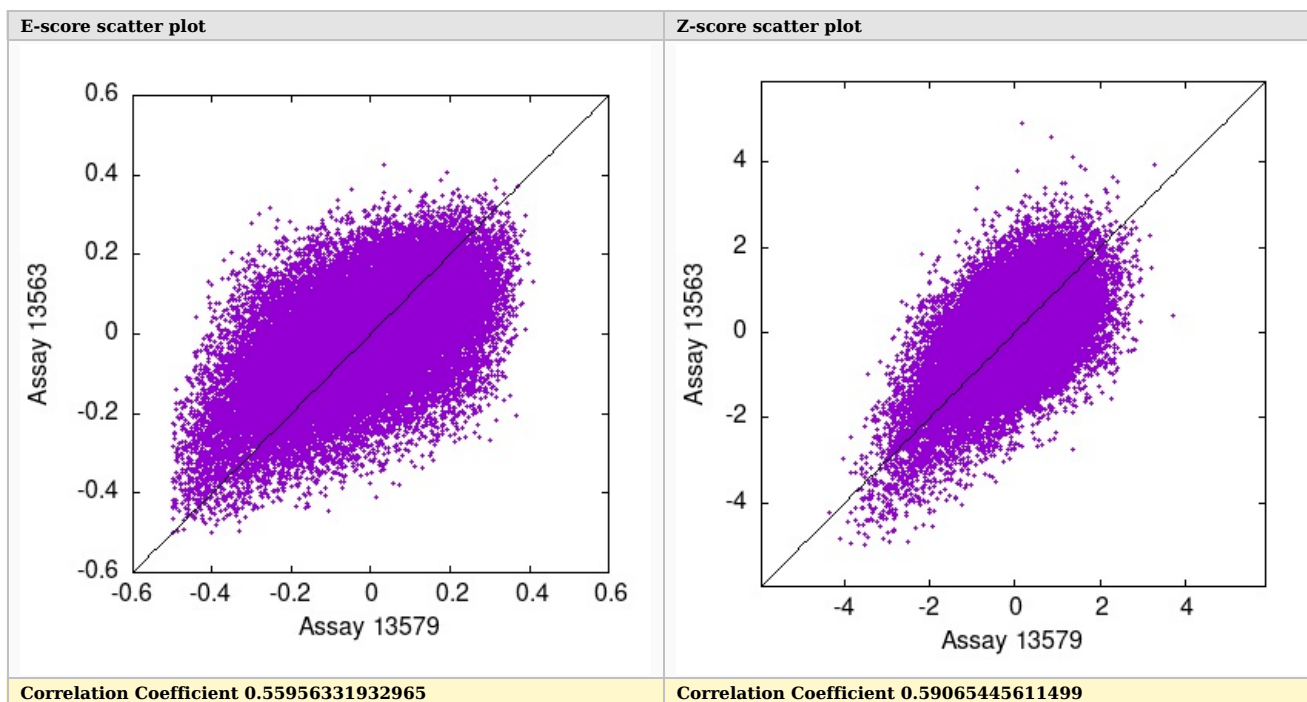


QC report



Top scoring motifs for Assay 13579

Protein ID: pTH13944.2 Gene: CREB3L3.FL Domain: bZIP_1 Flag: Reject Array: 1M-HK

8 mer E-scores for probeset 'all'

Forward:

Reverse:



Top 10

Scores

Alignment

ACCAATTG	0.41072	---ACCAATTG
TAACAAAA	0.39864	--TAACAAAA--
AATATTAA	0.39676	AATATTAA---
GAATATTC	0.39346	--GAATATTC--
ACTAATTA	0.39203	---ACTAATTA
ATATCACA	0.39043	-ATATCACA--
TACGCGTA	0.38609	--TACGCGTA--
GATTAATA	0.38315	--GATTAATA--
TATTAATA	0.38120	--TATTAATA--
AGTTAACT	0.38024	--AGTTAACT--

8 mer Z-scores for probeset 'all'

Forward:

Reverse:



Top 10

Scores

Alignment

ATATCACA	3.71966	---TGTGATAT
GTCATGAC	3.27108	-GTCATGAC--
AATATTAA	3.19082	-AATATTAA--
TAACAAAA	3.16284	--TAACAAAA--
ATTATAAT	2.99754	-ATTATAAT--
CTTATAAG	2.96180	-CTTATAAG--
GATTAATA	2.95164	GATTAATA---
ACTATAAT	2.93707	-ACTATAAT--
TACGCGTA	2.92415	--TACGCGTA--
AGTTAACT	2.90106	-AGTTAACT--

Top scoring motifs for Assay 13563

Protein ID: pTH13944.1 Gene: CREB3L3.FL Domain: bZIP_1 Flag: Reject Array: 1M-ME

8 mer E-scores for probeset 'all'

Forward:

Reverse:



Top 10

Scores

Alignment

TGTGCACA	0.42399	TGTGCACA----
TTTCGAAA	0.40783	TTTCGAAA----
AGCAATTG	0.38768	--AGCAATTG--
AATGCATT	0.38611	AATGCATT----
GTCATGAC	0.37255	-GTCATGAC--
ATACAACA	0.36821	ATACAACA----
GTGCGCAC	0.36490	-GTGCGCAC--
AACAGACA	0.36287	AACAGACA----
AACATCCA	0.35946	----AACATCCA
GACGCGTC	0.35768	GACGCGTC----

8 mer Z-scores for probeset 'all'

Forward:

Reverse:



Top 10

Scores

Alignment

TGTGCACA	4.89892	TGTGCACA-----
TTTCGAAA	4.57592	---TTTCGAAA---
AGCAATTG	4.12439	-----CAATTGCT
GTCATGAC	3.95142	----GTCATGAC--
CACAACAC	3.91770	----CACAACAC--
ATGTACAT	3.81340	----ATGTACAT--
GACGCGTC	3.80980	----GACGCGTC--
AAGTACTT	3.63414	----AAGTACTT--
TCGCGCGA	3.54531	---TCGCGCGA---
ACAACAAT	3.49360	-----ACAACAAT-