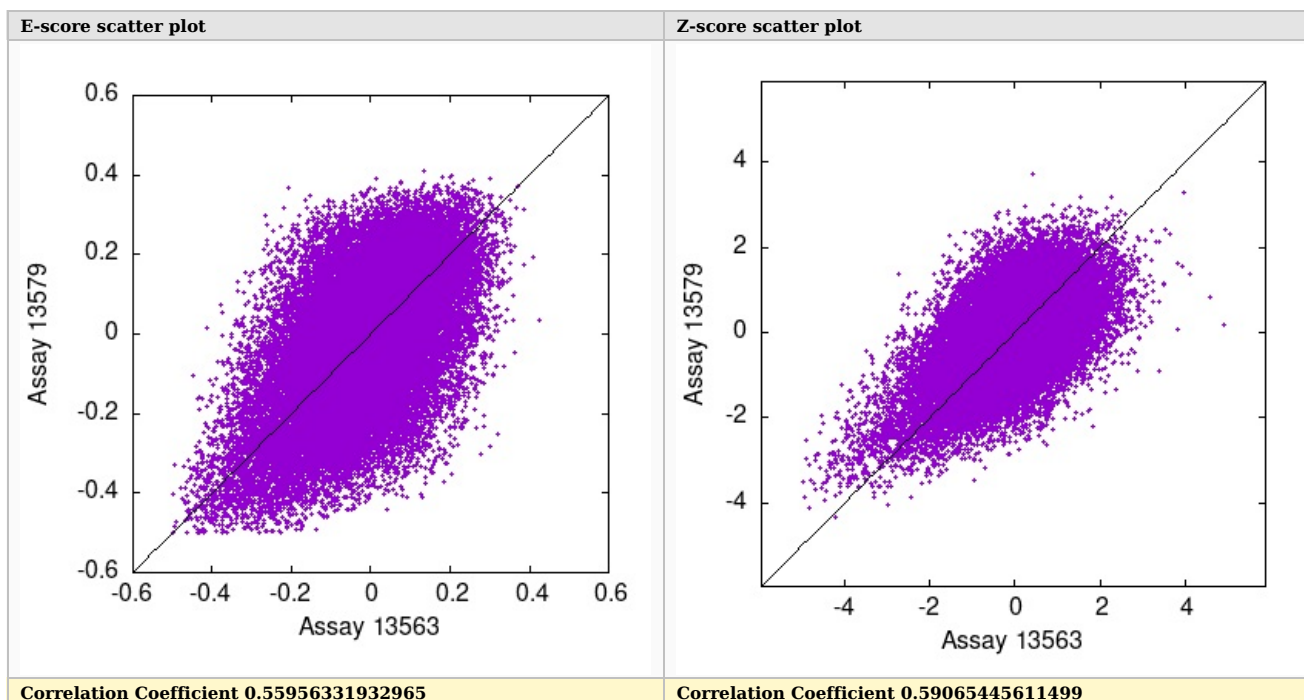


QC report



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'			8 mer Z-scores for probeset 'all'		
Forward:		Reverse:	Forward:		Reverse:
Top 10	Scores	Alignment	Top 10	Scores	Alignment
TGTGCACA	0.42399	TGTGCACA----	TGTGCACA	4.89892	TGTGCACA-----
TTTCGAAA	0.40783	TTTCGAAA----	TTTCGAAA	4.57592	---TTTCGAAA---
AGCAATTG	0.38768	--AGCAATTG--	AGCAATTG	4.12439	-----CAATTGCT
AATGCATT	0.38611	AATGCATT----	GTCATGAC	3.95142	----GTCATGAC--
GTCATGAC	0.37255	-GTCATGAC---	CACAACAC	3.91770	----CACAACAC--
ATACAACA	0.36821	ATACAACA----	ATGTACAT	3.81340	----ATGTACAT--
GTGCGCAC	0.36490	-GTGCGCAC---	GACGCGTC	3.80980	----GACGCGTC--
AACAGACA	0.36287	AACAGACA----	AAGTACTT	3.63414	----AAGTACTT--
AACATCCA	0.35946	----AACATCCA	TCGCGCGA	3.54531	---TCGCGCGA---
GACGCGTC	0.35768	GACGCGTC----	ACAACAAT	3.49360	-----ACAACAAT-

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'			8 mer Z-scores for probeset 'all'		
Forward:		Reverse:	Forward:		Reverse:
Top 10	Scores	Alignment	Top 10	Scores	Alignment
ACCAATTG	0.41072	---ACCAATTG	ATATCACA	3.71966	---TGTGATAT
TAACAAAA	0.39864	--TAACAAAA--	GTCATGAC	3.27108	-GTCATGAC--
AATATTAA	0.39676	AATATTAA---	AATATTAA	3.19082	-AATATTAA--
GAATATTC	0.39346	--GAATATTC--	TAACAAAA	3.16284	--TAACAAAA--
ACTAATTA	0.39203	---ACTAATTA	ATTATAAT	2.99754	-ATTATAAT--
ATATCACA	0.39043	-ATATCACA--	CTTATAAG	2.96180	-CTTATAAG--
TACGCGTA	0.38609	--TACGCGTA--	GATTAATA	2.95164	GATTAATA---
GATTAATA	0.38315	--GATTAATA--	ACTATAAT	2.93707	-ACTATAAT--
TATTAATA	0.38120	--TATTAATA--	TACGCGTA	2.92415	--TACGCGTA--
AGTTAACT	0.38024	--AGTTAACT--	AGTTAACT	2.90106	-AGTTAACT--