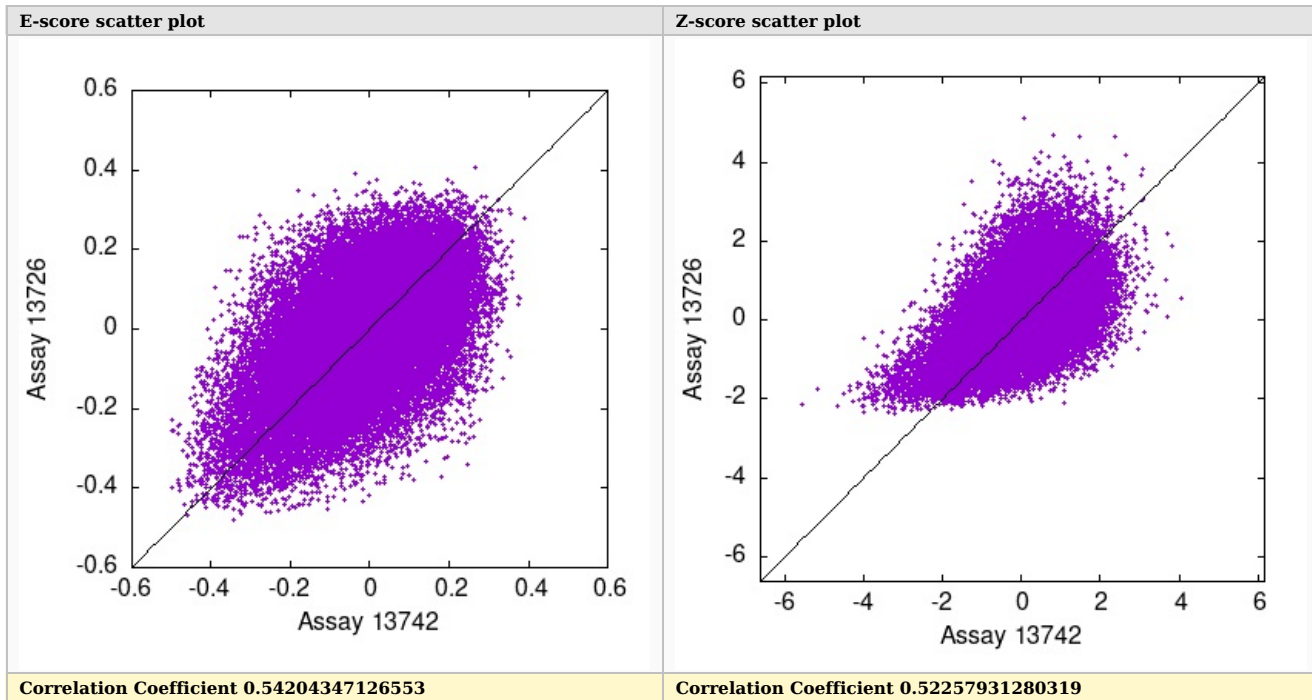


QC report



Top scoring motifs for Assay 13742

Protein ID: pTH14309.2 Gene: NKRF.DBD Domain: Unknown 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
CACATATA	0.39109	--CACATATA----	TAACGTAA	4.05640	--TAACGTAA--
TAACGTAA	0.38005	--TAACGTAA----	CACATATA	3.84130	-TATATGTG--
ACCATACT	0.37489	--ACCATACT----	GATTTGTA	3.68761	-GATTTGTA--
CACTGATA	0.37326	--CACTGATA----	AGAATTCT	3.68577	AGAATTCT---
ACTCTACT	0.36115	---ACTCTACT--	ACCATACT	3.65246	AGTATGGT---
ATGTATGC	0.36091	---GCATACAT--	ATTTACAG	3.41585	--ATTTACAG--
ACATTATA	0.35935	-----TATAATGT	ATTTGTAG	3.38508	--ATTTGTAG--
AGTTTTTC	0.35701	---AGTTTTTC---	ACAAACAG	3.32198	CTGTTTGT---
TACGTCAA	0.35700	TTGACGTA-----	CACTGATA	3.30977	-CACTGATA--
TATAAGCA	0.35357	--TGCTTATA----	ACTCTACT	3.23233	---ACTCTACT

Top scoring motifs for Assay 13726

Protein ID: pTH14309.1 Gene: NKRF.DBD Domain: Unknown 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
TAACGAAA	0.40582	-TAACGAAA--	ATGTACAT	5.10341	-ATGTACAT--
ATGTACAT	0.39108	ATGTACAT--	TATACAAA	4.67532	--TATACAAA--
ATAAACAT	0.37685	--ATAAACAT	GTACTATA	4.66453	--TATAGTAC--
GTAATACT	0.37637	-TATAGTAC--	TAACGAAA	4.66062	--TAACGAAA--
ATATACAG	0.36815	ATATACAG--	AATGTTCA	4.25678	AATGTTCA---
CTACAACA	0.36772	--CTACAACA	AATAATAA	4.18596	--AATAATAA--
ATTGGAAT	0.36697	--ATTGGAAT	CTACAACA	4.14201	---CTACAACA
ATAAACAC	0.36344	--ATAAACAC	TATATGCA	4.12802	--TATATGCA--
TATACAAA	0.35629	-TATACAAA--	ATAAACAT	4.11442	---ATAAACAT
AACGAAAT	0.34983	--AACGAAAT	ACATACAA	4.03627	-ACATACAA--