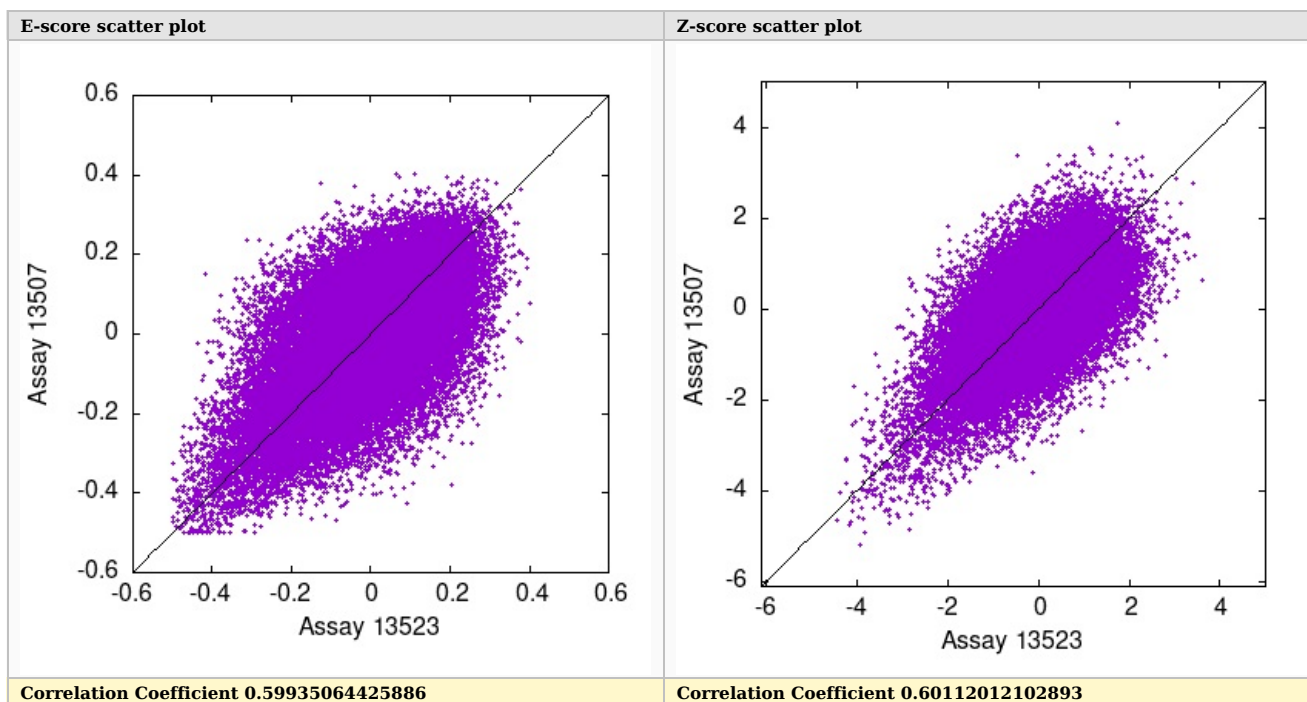


QC report



Top scoring motifs for Assay 13523

Protein ID: pTH13952.2 Gene: ANKZF1.DBD Domain: zf-C2H2 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
CGCGAAAA	0.40117	----TTTTCGCG--	AACTAACG	3.61429	-AACTAACG-----
AACCTATT	0.39439	-AACCTATT----	ATAACGTA	3.42914	---ATAACGTA----
AACAATGT	0.39238	--ACATTGTT---	AAACGTTT	3.41948	----AAACGTTT---
GTAACAAA	0.38143	-TTTGTTCAC---	CAAAATAC	3.30742	----CAAAATAC---
AAACGTTT	0.38048	AAACGTTT-----	TGTTAACA	3.30185	-TGTTAACA-----
AACACTGT	0.37674	-AACACTGT----	AACAATGT	3.26686	--AACAATGT-----
ATATGTGC	0.37368	GCACATAT-----	CCTATCAA	3.23829	-----CCTATCAA
TGTTAACA	0.37333	---TGTTAACA--	GTATGAC	3.21805	GTATGAC-----
AACGTTTC	0.37310	-AACGTTTC----	TTTCGAAA	3.14637	----TTTCGAAA---
TTTCGAAA	0.37201	-----TTTCGAAA	GAATATTC	3.07929	----GAATATTC---

Top scoring motifs for Assay 13507

Protein ID: pTH13952.1 Gene: ANKZF1.DBD Domain: zf-C2H2 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
CACATTCA	0.40307	TGAATGTG---	ATTTAAAT	4.07799	---ATTTAAAT---
ATCGATAC	0.40288	---ATCGATAC	AACCAACA	3.53531	--TGTTGGTT---
CAAATTTG	0.39610	CAAATTTG---	ATCGATAC	3.50481	---ATCGATAC---
AACCAACA	0.39386	--TGTTGGTT---	CGAATTCG	3.40612	CGAATTCG---
AAATTTGT	0.39377	-AAATTTGT---	CAAATTTG	3.38452	-CAAATTTG---
ACATCGAT	0.38650	-ACATCGAT---	CATCGATA	3.38356	--CATCGATA---
AATTGTTA	0.38621	--AATTGTTA---	ATAATTAT	3.38351	---ATAATTAT---
CATCGATA	0.38287	--CATCGATA---	AAATTTGT	3.36737	-AAATTTGT---
AATTTGTA	0.38108	--AATTTGTA---	AATTGTTA	3.35179	--AATTGTTA---
TAAATTTA	0.38081	TAAATTTA---	AGTTGTTG	3.31475	--AGTTGTTG---