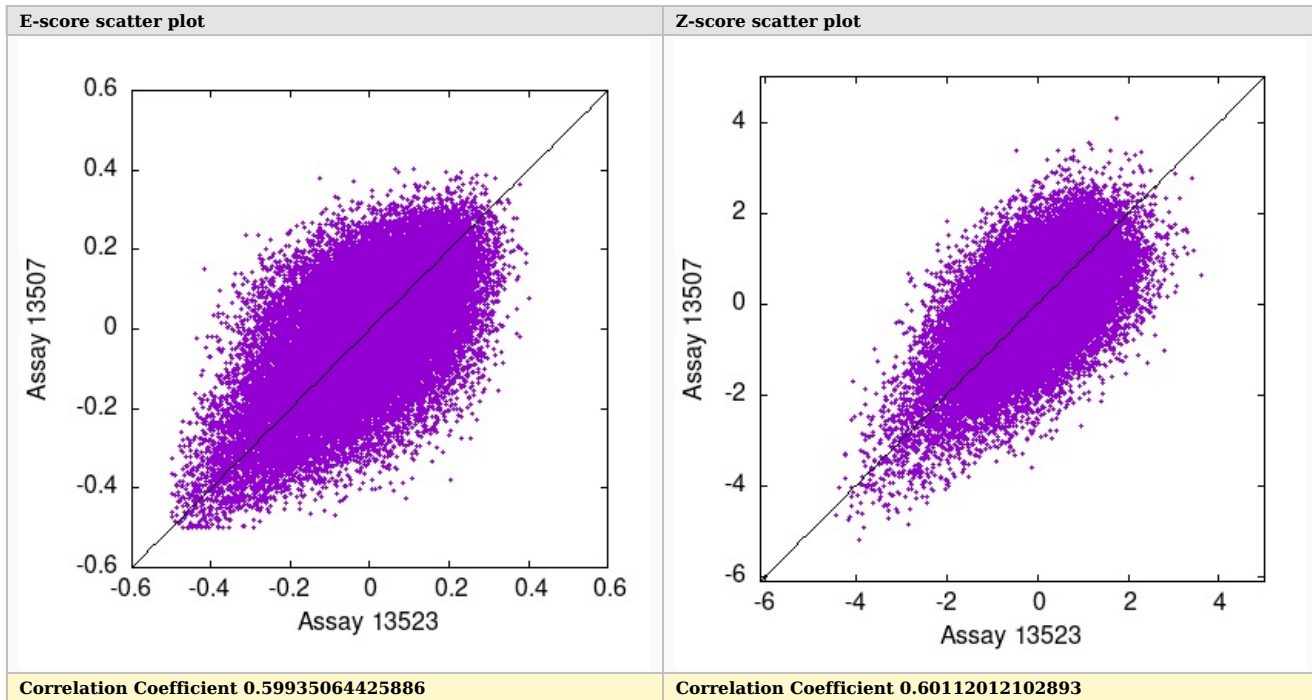


## QC report



## 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'

Forward:

Reverse:

## Top 10

## Scores

## Alignment

CGCGAAAA	0.40117	----TTTTCGCG--
AACCTATT	0.39439	--AACCTATT----
AACAATGT	0.39238	--ACATTGTT---
GTAACAAA	0.38143	---TTGTTAC----
AAACGTTT	0.38048	AAACGTTT-----
AACACTGT	0.37674	--AACACTGT----
ATATGTGC	0.37368	GCACATAT-----
TGTTAACA	0.37333	---TGTTAACA--
AACGTTTC	0.37310	--AACGTTTC----
TTTCGAAA	0.37201	-----TTTCGAAA

## 8 mer Z-scores for probeset 'all'

Forward:

Reverse:

## Top 10

## Scores

## Alignment

AACTAACG	3.61429	-AACTAACG-----
ATAACGTA	3.42914	---ATAACGTA----
AAACGTTT	3.41948	----AAACGTTT---
CAAAATAC	3.30742	-----CAAAATAC--
TGTTAACA	3.30185	-TGTTAACA-----
AACAATGT	3.26686	--AACAATGT-----
CCTATCAA	3.23829	-----CCTATCAA
GTATGAC	3.21805	GTATGAC-----
TTTCGAAA	3.14637	-----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'

Forward:

Reverse:

## Top 10

## Scores

## Alignment

CACATTCA	0.40307	TGAATGTG---
ATCGATAC	0.40288	---ATCGATAC
CAAATTTG	0.39610	CAAATTTG---
AACCAACA	0.39386	--TGTTGGTT-
AAATTTGT	0.39377	--AAATTTGT--
ACATCGAT	0.38650	-ACATCGAT--
AATTGTTA	0.38621	--AATTGTTA-
CATCGATA	0.38287	--CATCGATA-
AATTTGTA	0.38108	--AATTTGTA-
TAAATTTA	0.38081	TAAATTTA---

## 8 mer Z-scores for probeset 'all'

Forward:

Reverse:

## Top 10

## Scores

## Alignment

ATTTAAAT	4.07799	---ATTTAAAT
AACCAACA	3.53531	--TGTTGGTT-
ATCGATAC	3.50481	---ATCGATAC
CGAATTCG	3.40612	CGAATTCG---
CAAATTTG	3.38452	---CAAATTTG--
CATCGATA	3.38356	--CATCGATA-
ATAATTAT	3.38351	---ATAATTAT
AAATTTGT	3.36737	--AAATTTGT--
AATTGTTA	3.35179	--AATTGTTA-
AGTTGTTG	3.31475	--AGTTGTTG-