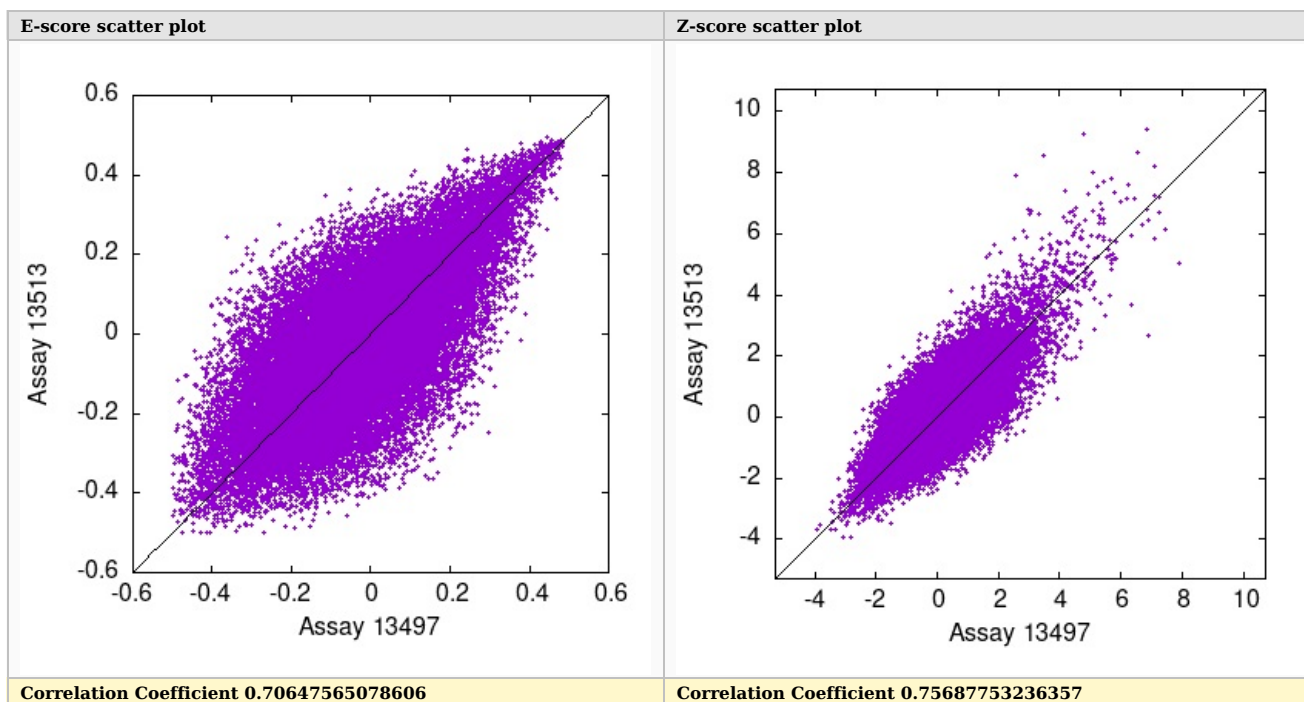


## QC report



## Top scoring motifs for Assay 13497

Protein ID: pTH13913.1 Gene: AHCTF1.DBD Domain: AT\_hook Flag: Pass\_matched\_pair Array: 1M-ME

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

Forward:

Reverse:

## Top 10

## Scores

## Alignment

AAATATTT	0.48374	--AAATATTT--
AAAAAATA	0.48304	--AAAAAATA--
ATTTAAAT	0.48291	--ATTTAAAT--
AAAAATAA	0.48082	AAAAATAA----
AAAAAAAT	0.48026	--AAAAAAAT--
AATTAATT	0.47835	--AATTAATT--
ATAATATA	0.47644	----ATAATATA
ATTATTTA	0.47640	-TAAATAAT---
AAATAATT	0.47460	--AAATAATT--
AAAATATT	0.47356	---AATATTTT-

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

Forward:

Reverse:

## Top 10

## Scores

## Alignment

AAAAAAAT	7.89610	-AAAAAAAT-
AAATATTT	7.43353	-AAATATTT-
AAAAATAA	7.25444	-AAAAATAA-
ATTTAAAT	7.22518	-ATTTAAAT-
AAAAAATA	7.11045	--AAAAAATA
AATTTAAT	7.10444	-AATTTAAT-
ATTATTTA	7.07541	-TAAATAAT-
CAAAAAAA	6.91058	CAAAAAAA--
AAATAATT	6.90748	-AAATAATT-
AATTAAT	6.86585	-AATTAAT-

## Top scoring motifs for Assay 13513

Protein ID: pTH13913.2 Gene: AHCTF1.DBD Domain: AT\_hook Flag: Pass\_matched\_pair Array: 1M-HK

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

Forward:

Reverse:

## Top 10

## Scores

## Alignment

AAATAAAT	0.49357	--AAATAAAT-
AATTAATT	0.48760	--AATTAATT-
AATAAATT	0.48584	---AATAAATT
AATTTAAA	0.48509	---TTTAAATT
AAAAAATA	0.48473	---AAAAAATA
AAAAATAT	0.48382	--AAAAATAT-
AAAAATAA	0.48167	AAAAATAA----
AATAATAA	0.48042	---AATAATAA
ATTTAAAT	0.47921	--ATTTAAAT-
AAAAAATA	0.47898	--AAAAAATA-

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

Forward:

Reverse:

## Top 10

## Scores

## Alignment

AATTAATT	9.39101	-AATTAATT-
AAATAAAT	9.28751	AAATAAAT--
AAAAATAT	8.66712	--ATATTTTT
AATTTAAA	8.54261	-TTTAAATT-
AAAAAATA	8.21743	-AAAAAATA-
AATATATT	8.01288	-AATATATT-
ATTTTTC	7.91754	GAAAAAAT--
AATAATAA	7.81197	-AATAATAA-
AAATAATT	7.70647	-AAATAATT-
AAAATATT	7.60208	-AAAATATT-