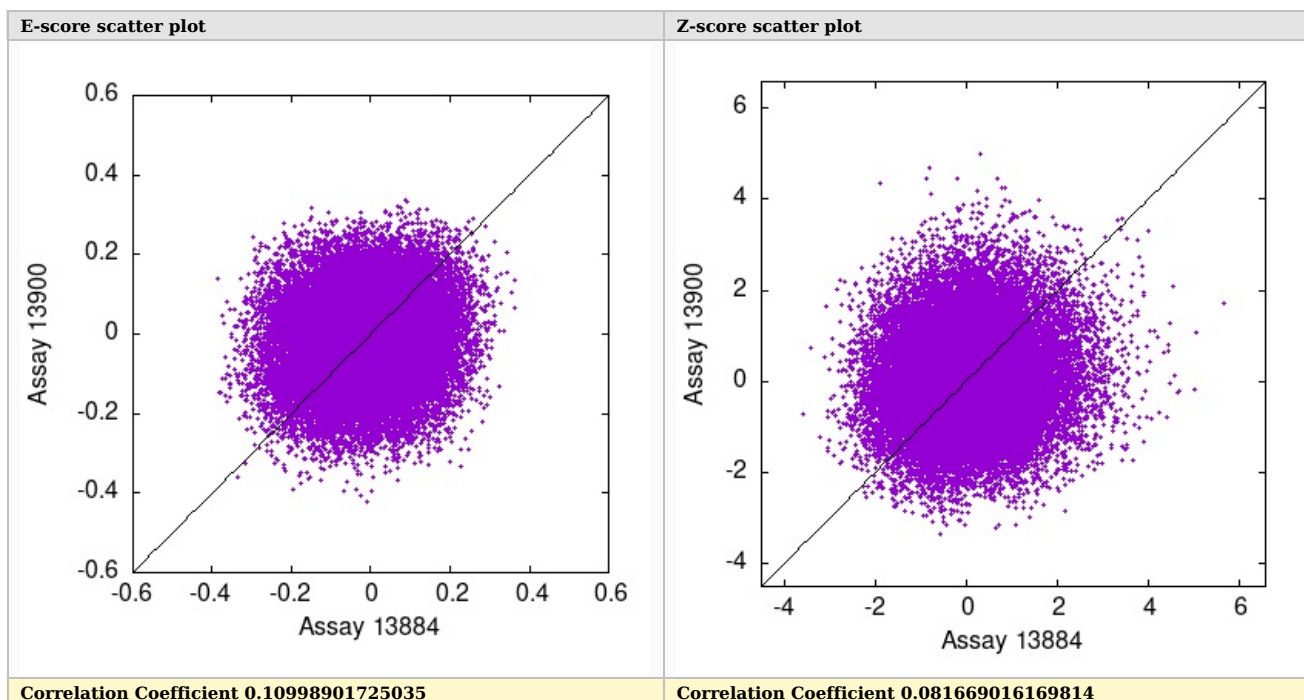


## QC report



### Top scoring motifs for Assay 13884

Protein ID: pTH14281.1 Gene: NFX1.DBD Domain: zf-NF-X1 Flag: Reject Array: 1M-ME

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

Forward:

Reverse:



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

Forward:

Reverse:



#### Top 10

#### Scores

#### Alignment

GTATAATA	0.36495	GTATAATA-----
AAGCAGAA	0.36195	-----AAGCAGAA
ACCAATAC	0.35716	--ACCAATAC----
AAACACAC	0.34352	--AAACACAC----
AAAGCAGA	0.34213	----AAAGCAGA--
ACAAAACC	0.33549	--ACAAAACC----
ATAATAAA	0.33005	--ATAATAAA----
GTACTTTA	0.32680	---TAAAGTAC---
CAAGAAAC	0.32383	--CAAGAAAC----
ATAAAAGC	0.32329	--ATAAAAGC----

#### Top 10

#### Scores

#### Alignment

AAGCAGAA	5.65334	-AAGCAGAA---
ACCAATAC	5.05933	-ACCAATAC---
ATAGTCTG	5.03017	--CAGACTAT--
ACAAATAA	4.69217	-ACAAATAA---
TATAACCA	4.66031	----TATAACCA
GTTTCTGA	4.57957	-TCAGAAAC---
AATAAAAT	4.54761	---AATAAAAT
GGAGAATA	4.49053	GGAGAATA----
GATATAAA	4.48701	--GATATAAA--
CCAACAAC	4.45079	--CCAACAAC--

### Top scoring motifs for Assay 13900

Protein ID: pTH14281.2 Gene: NFX1.DBD Domain: zf-NF-X1 Flag: Reject Array: 1M-HK

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

Forward:

Reverse:



Forward:

Reverse:



#### Top 10

#### Scores

#### Alignment

ACCCACCC	0.33653	-GGGTGGGT-
ATACAAAT	0.33250	-ATACAAAT-
ACAAACAG	0.32056	--ACAAACAG
GGGCAATA	0.31861	-GGGCAATA-
ACCACCAC	0.31559	--ACCACCAC
AGTGCGAT	0.31472	AGTGCGAT--
GGACGATA	0.31223	-GGACGATA-
GGGGACAC	0.30812	-GGGGACAC-
GAAATAAA	0.30757	--GAAATAAA
ATATTGAA	0.30721	--TTCAATAT

#### Top 10

#### Scores

#### Alignment

ACCCACCC	4.99902	--ACCCACCC---
ACATGTGG	4.67832	--CCACATGT---
GTGTACAC	4.46555	---GTGTACAC---
CCCTGTTA	4.46527	-----TAACAGGG
ATGAAGTA	4.44288	TACTTCAT-----
CACCACTA	4.36347	--CACCACATA---
CGTCATCA	4.25917	--CGTCATCA---
GGTATACA	4.25903	--GGTATACA---
CCAGGACC	4.18129	--GGTCCTGG---
CATCGACG	4.10327	--CATCGACG---