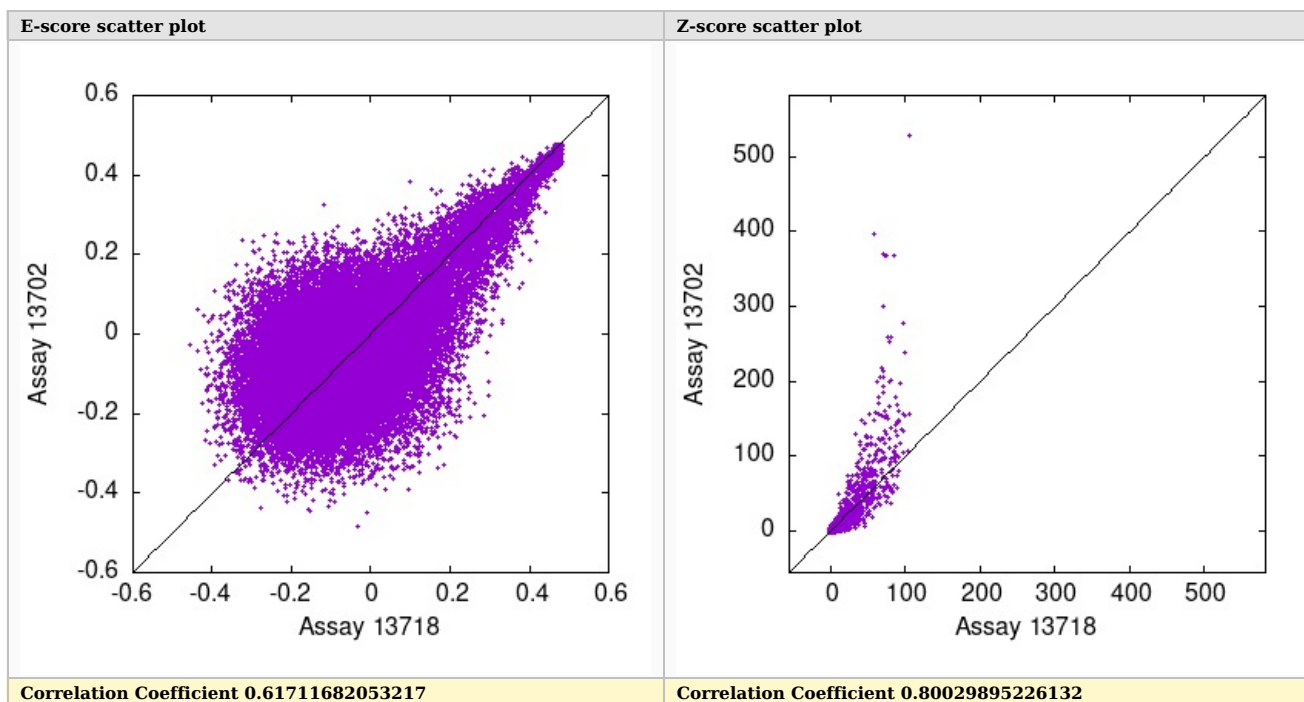


## QC report



## Top scoring motifs for Assay 13718

Protein ID: pTH14232.2 Gene: ZGLP1.DBD Domain: GATA Flag: Pass\_matched\_pair Array: 1M-HK

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

Forward:

Reverse:



## Top 10

## Scores

## Alignment

GATCGATC	0.48419	---GATCGATC
GATGATCA	0.48404	GATGATCA---
CGATCAG	0.48360	CGTGATCG---
CGATCGCC	0.48360	GCGATCG---
GCGATCGC	0.48221	-GCGATCGC--
TTGATCAA	0.48195	--TTGATCAA--
CGTGATCA	0.48155	CGTGATCA---
GTGATCAA	0.48102	-GTGATCAA--
TGATCAAA	0.48014	--TGATCAAA--
AGCGATCG	0.48001	AGCGATCG---

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

Forward:

Reverse:



## Top 10

## Scores

## Alignment

GATCGATC	106.37461	GATCGATC---
TTGATCAA	104.45268	--TTGATCAA--
CGTGATCA	102.28186	-CGTGATCA--
AGCGATCG	100.08916	-AGCGATCG--
CGATCGCC	98.86468	-GCGATCGC--
GTGATCAC	97.39004	--GTGATCAC--
CACGATCG	96.82296	---CGATCGTG
GCGATCGC	96.70618	--GCGATCGC--
AGCGATCA	96.50692	-AGCGATCA--
CGATCATC	95.43301	---CGATCATC

## Top scoring motifs for Assay 13702

Protein ID: pTH14232.1 Gene: ZGLP1.DBD Domain: GATA Flag: Pass\_matched\_pair Array: 1M-ME

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

Forward:

Reverse:



## Top 10

## Scores

## Alignment

ACGATCGT	0.47792	--ACGATCGT--
ATCGATCG	0.47661	-ATCGATCG---
CGATCGGA	0.47639	---CGATCGGA---
GATCGATC	0.47568	GATCGATC----
ATGATCAT	0.47527	--ATGATCAT--
GATGATCA	0.47189	---TGATCATC--
GCGATCGA	0.46965	--TCGATCGC--
GATCATCA	0.46917	----GATCATCA
CGATCGCA	0.46907	---CGATCGCA---
ATCGATCA	0.46883	-ATCGATCA---

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

Forward:

Reverse:



## Top 10

## Scores

## Alignment

GATCGATC	528.82813	---GATCGATC
ATCGATCG	397.56767	--CGATCGAT--
ACGATCGT	369.66832	-ACGATCGT--
ATCGATCA	368.03588	--TGATCGAT--
ATGATCAT	367.16833	-ATGATCAT--
GATGATCA	367.16833	GATGATCA---
GCGATCGA	300.66634	-GCGATCGA--
GCGATCGC	278.29637	-GCGATCGC--
CGATCGCG	259.18618	CGCGATCG---
CGATCGCA	258.52797	TGCGATCG---