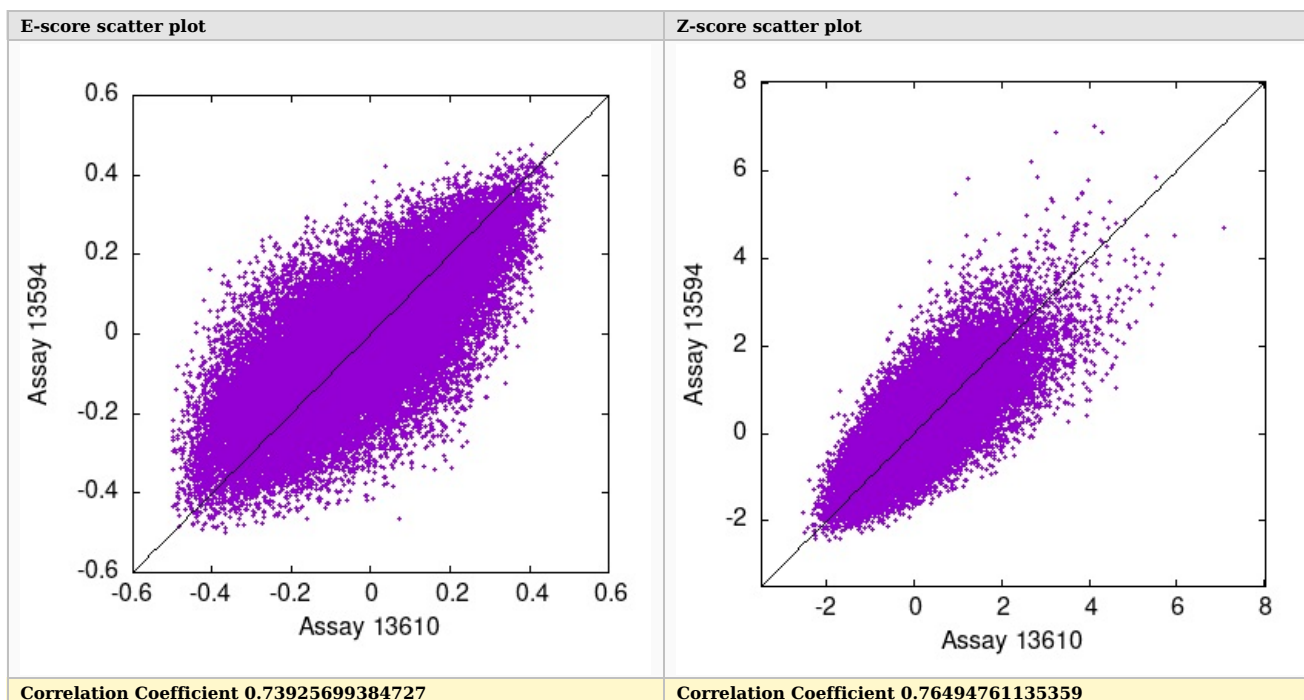


## QC report



### Top scoring motifs for Assay 13610

Protein ID: pTH14226.2 Gene: DMRTB1.DBD Domain: DM Flag: Pass\_matched\_pair Array: 1M-HK

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

Forward:

Reverse:



Top 10

Scores

Alignment

AATATATT	0.46852	--AATATATT
TATTAACA	0.45644	--TATTAACA
AATAATAA	0.45497	--AATAATAA
AAATAAAT	0.44900	-AAATAAAT-
AATAAATT	0.44573	--AATAAATT
ATTATTAA	0.44485	TTAATAAT--
AAAATATT	0.44380	--AAAATATT
TAAATATA	0.44267	-TAATATAA-
AAATATAT	0.44180	-AAATATAT-
GAATAATA	0.44050	-GAATAATA-

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

Forward:

Reverse:



Top 10

Scores

Alignment

AATATATT	7.07142	AATATATT----
AATAATAA	5.95528	---TTATTATT-
TATTAATA	5.66016	----TATTAATA
AATATAAT	5.59590	AATATAAT----
AAATATAT	5.52989	-ATATATTT--
TATTAACA	5.46498	----TATTAACA
ATTATTAA	5.42237	--ATTATTAA--
AATAAATT	5.41232	AATAAATT----
AACAATAA	5.34651	-AACAATAA---
ATATATTA	5.32693	-ATATATTA---

### Top scoring motifs for Assay 13594

Protein ID: pTH14226.1 Gene: DMRTB1.DBD Domain: DM Flag: Pass\_matched\_pair Array: 1M-ME

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

Forward:

Reverse:



Top 10

Scores

Alignment

ATAATATA	0.47699	---ATAATATA-
ATATAATA	0.47426	--TATTATAT--
ATTATATA	0.46641	---ATTATATA-
ATAATTAT	0.45588	ATAATTAT----
ATTATTTA	0.45358	---ATTATTTA-
AAATATAT	0.45283	--AAATATAT--
ATATTATA	0.44822	--TATAATAT--
AATAATAT	0.44243	--AATAATAT--
ATATATAT	0.44242	---ATATATAT
TATATATA	0.44191	---TATATATA-

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

Forward:

Reverse:



Top 10

Scores

Alignment

ATATAATA	7.01357	--TATTATAT--
ATAATTAT	6.88554	ATAATTAT----
ATAATATA	6.86015	---ATAATATA-
ATTATATA	6.22054	---ATTATATA-
ATTATTTA	5.85957	---ATTATTTA-
AAATATAT	5.83917	--AAATATAT--
ATGATCAT	5.80306	---ATGATCAT-
ATTGTTAA	5.77244	---ATTGTTAA-
ATATTATA	5.49699	--TATAATAT--
ACATATGT	5.48146	---ACATATGT