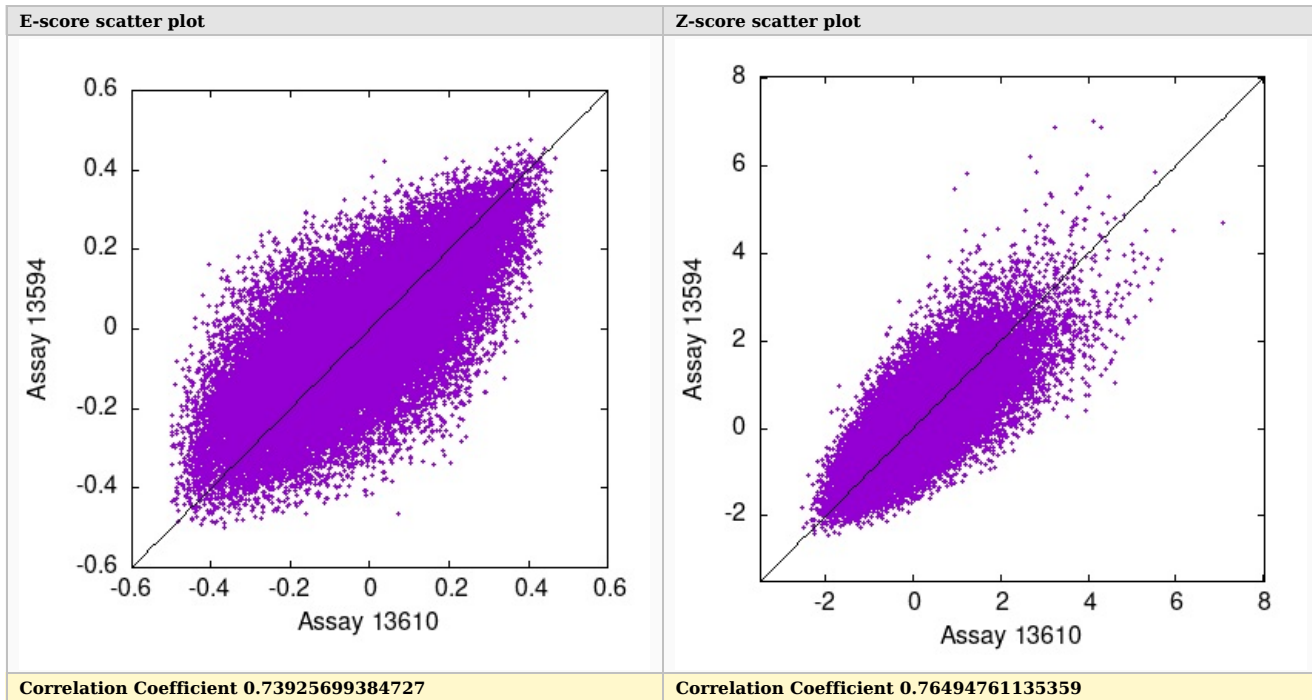


QC report



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'			8 mer Z-scores for probeset 'all'		
Forward:		Reverse:	Forward:		Reverse:
Top 10	Scores	Alignment	Top 10	Scores	Alignment
AATATATT	0.46852	--AATATATT	AATATATT	7.07142	AATATATT----
TATTAACA	0.45644	--TATTAACA	AATAATAA	5.95528	---TTATTATT--
AATAATAA	0.45497	--AATAATAA	TATTAATA	5.66016	----TATTAATA
AAATAAAT	0.44900	-AAATAAAT-	AAATAAAT	5.59590	AATATAAT----
AATAAATT	0.44573	--AATAAATT	AAATATAT	5.52989	-ATATATTT---
ATTATTAA	0.44485	TTAATAAT--	TATTAACA	5.46498	----TATTAACA
AAAATATT	0.44380	--AAAATATT	ATTATTAA	5.42237	---ATTATTAA--
TAAATATA	0.44267	-TAATATAA-	AATAAATT	5.41232	AATAAATT----
AAATATAT	0.44180	-AAATATAT-	AACAATAA	5.34651	-AACAATAA---
GAATAATA	0.44050	-GAATAATA-	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'			8 mer Z-scores for probeset 'all'		
Forward:		Reverse:	Forward:		Reverse:
Top 10	Scores	Alignment	Top 10	Scores	Alignment
ATAATATA	0.47699	---ATAATATA-	ATATAATA	7.01357	--TATTATAT--
ATATAATA	0.47426	--TATTATAT--	ATAATTAT	6.88554	ATAATTAT----
ATTATATA	0.46641	---ATTATATA-	ATAATATA	6.86015	---ATAATATA-
ATAATTAT	0.45588	ATAATTAT----	ATTATATA	6.22054	---ATTATATA-
ATTATTTA	0.45358	---ATTATTTA-	ATTATTTA	5.85957	---ATTATTTA-
AAATATAT	0.45283	--AAATATAT--	AAATATAT	5.83917	---AAATATAT--
ATATTATA	0.44822	--TATAATAT--	ATGATCAT	5.80306	---ATGATCAT-
AATAATAT	0.44243	--AATAATAT--	ATTGTTAA	5.77244	---ATTGTTAA-
ATATATAT	0.44242	---ATATATAT	ATATTATA	5.49699	--TATAATAT--
TATATATA	0.44191	---TATATATA-	ACATATGT	5.48146	---ACATATGT