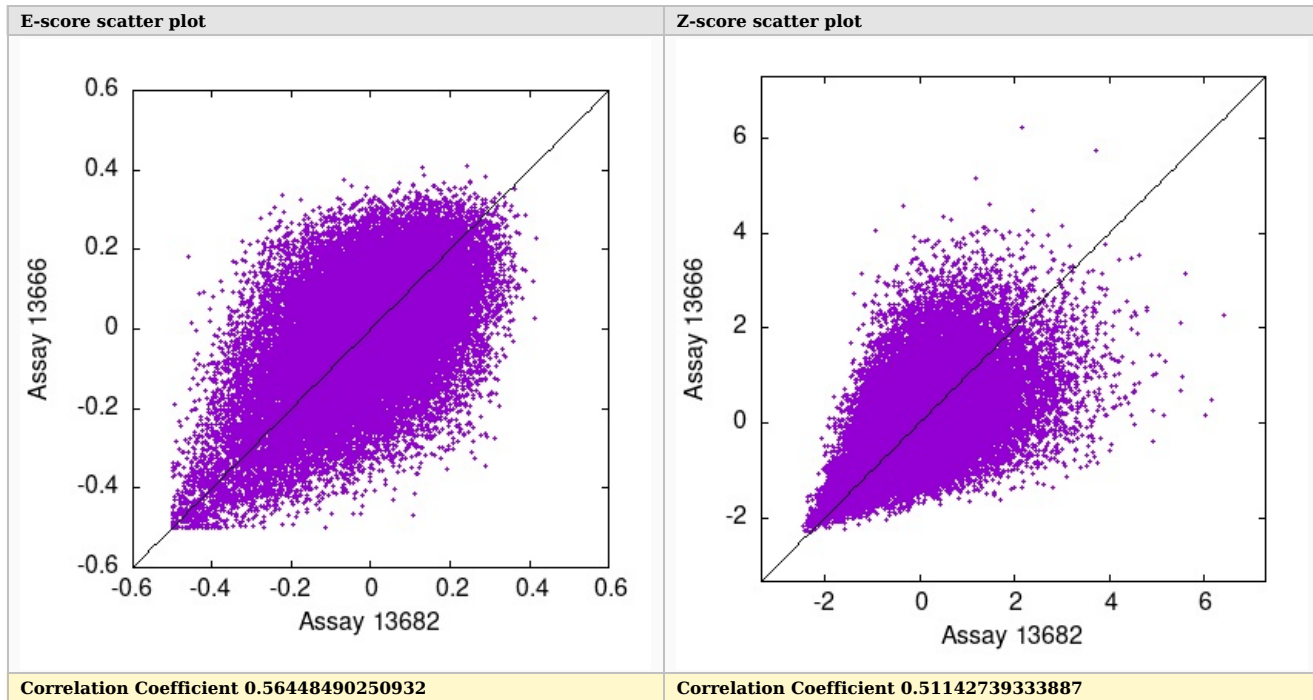


QC report



Top scoring motifs for Assay 13682

Protein ID: pTH14276.2 Gene: TERB1.DBD Domain: Myb_DNA-binding Flag: Reject 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
TGTTAACA	0.41795	----TGTTAACA	TGTTAACA	6.41318	----TGTTAACA
GTGTACAC	0.41440	---GTGTACAC-	TACACACA	6.14902	--TGTGTGTA--
TACACACA	0.40878	--TGTGTGTA--	GTGTACAC	6.04620	---GTGTACAC-
ATGTTAAC	0.39118	---ATGTTAAC-	ATGTTAAC	5.61527	---ATGTTAAC-
GTACAATA	0.38913	--TATTGTAC--	AACAATGA	5.52916	TCATTGTT----
TGTCACAA	0.38224	---TGTCACAA	ATATTAAT	5.51458	---ATATTAAT-
AAGTACTT	0.38088	---AAGTACTT-	ATGTATTA	5.50955	---ATGTATTA-
AACAATGA	0.37470	TCATTGTT----	CAATATTG	5.16943	---CAATATTG--
AGATATCT	0.37379	---AGATATCT	AATTTAAA	5.15846	---AATTTAAA-
GTAAATAA	0.37080	--TATTTTAC--	ATACATTA	5.04870	---ATACATTA--

Top scoring motifs for Assay 13666

Protein ID: pTH14276.1 Gene: TERB1.DBD Domain: Myb_DNA-binding Flag: Reject 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
TGTATACA	0.41062	TGTATACA----	TGTATACA	6.22062	--TGTATACA-
TAACGAAA	0.40818	--TAACGAAA--	GTACTTTA	5.75202	--TAAAGTAC-
ATGTACAT	0.38847	-ATGTACAT---	CAATTTC	5.14586	---CAATTTC--
GTACTTTA	0.38380	--TAAAGTAC--	ATACGTAT	4.59862	---ATACGTAT
AGACGAAA	0.37736	---AGACGAAA-	AAGCAGAA	4.56592	---AAGCAGAA-
CAATTTC	0.37161	--CAATTTC---	ACATATAT	4.47957	---ACATATAT-
AATAATAA	0.36880	---AATAATAA-	AATAATAA	4.34606	---AATAATAA-
TATACCAA	0.36817	--TATACCAA--	GAATAATA	4.29485	---GAATAATA-
GTACTATA	0.36716	--TATAGTAC--	AGTATACA	4.16118	TGTATACT---
TAATAAGA	0.36535	----TAATAAGA	TACAAATA	4.12400	---TACAAATA--