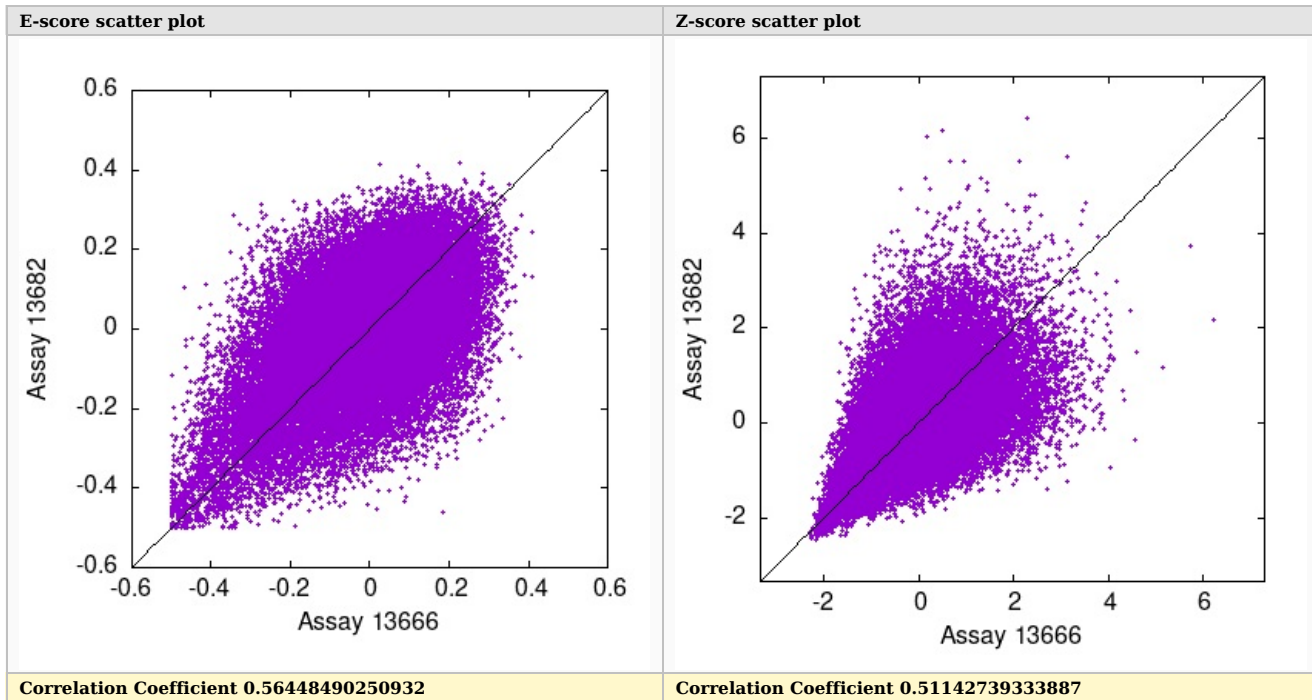




QC report





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'

Forward:		Reverse:	
			
Top 10	Scores	Alignment	
TGTATACA	0.41062	TGTATACA----	
TAACGAAA	0.40818	--TAACGAAA--	
ATGTACAT	0.38847	-ATGTACAT---	
GTACTTTA	0.38380	--TAAAGTAC--	
AGACGAAA	0.37736	---AGACGAAA--	
CAATTTCA	0.37161	--CAATTTCA---	
AATAATAA	0.36880	--AATAATAA--	
TATACCAA	0.36817	--TATACCAA--	
GTACTATA	0.36716	--TATAGTAC--	
TAATAAGA	0.36535	----TAATAAGA	



8 mer Z-scores for probeset 'all'

Forward:		Reverse:	
			
Top 10	Scores	Alignment	
TGTATACA	6.22062	--TGTATACA-	
GTACTTTA	5.75202	--TAAAGTAC-	
CAATTTCA	5.14586	--CAATTTCA--	
ATACGTAT	4.59862	---ATACGTAT	
AAGCAGAA	4.56592	--AAGCAGAA--	
ACATATAT	4.47957	--ACATATAT--	
AATAATAA	4.34606	--AATAATAA--	
GAATAATA	4.29485	--GAATAATA--	
AGTATACA	4.16118	TGTATACT---	
TACAAATA	4.12400	-TACAAATA--	



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'

Forward:		Reverse:	
			
Top 10	Scores	Alignment	
TGTTAACA	0.41795	---TGTTAACA	
GTGTACAC	0.41440	---GTGTACAC-	
TACACACA	0.40878	--TGTGTGTA--	
ATGTTAAC	0.39118	---ATGTTAAC-	
GTACAATA	0.38913	--TATTGTAC--	
TGTCACAA	0.38224	---TGTCACAA	
AAGTACTT	0.38088	---AAGTACTT-	
AACAATGA	0.37470	TCATTGTT----	
AGATATCT	0.37379	---AGATATCT	
GTAAAATA	0.37080	--TATTTTAC--	

8 mer Z-scores for probeset 'all'

Forward:		Reverse:
		
Top 10	Scores	Alignment
TGTTAACA	6.41318	--- -TGTTAACA
TACACACA	6.14902	--TGTGTGTA--
GTGTACAC	6.04620	---GTGTACAC-
ATGTTAAC	5.61527	---ATGTTAAC-
AACAATGA	5.52916	TCATTGTT---
ATATTAAT	5.51458	---ATATTAAT-
ATGTATTA	5.50955	---ATGTATTA-
CAATATTG	5.16943	--CAATATTG--
AATTTAAA	5.15846	---AATTTAAA-
ATACATTA	5.04870	-ATACATTA---