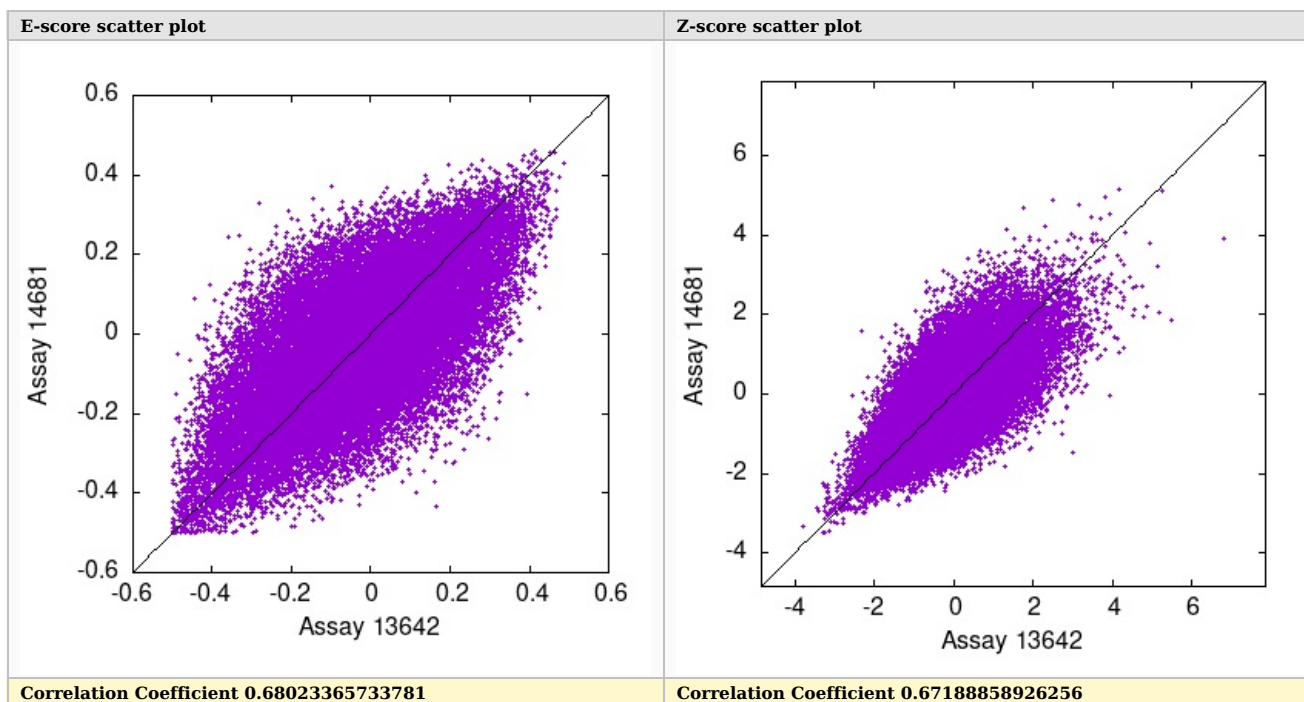




QC report





Top scoring motifs for Assay 13642

Protein ID: pTH13919.2 Gene: GLYR1.DBD Domain: AT_hook Flag: Backup_matched_pair Array: 1M-HK

8 mer E-scores for probeset 'all'

Forward:		Reverse:	
			
Top 10	Scores	Alignment	
AATATATT	0.48709	--AATATATT	
AATAAATT	0.47669	--AATAAATT	
AATTTAAA	0.46807	--AATTTAAA	
AAATATTT	0.46748	-AAATATTT-	
ACAAAATA	0.46683	ACAAAATA--	
AAATATAT	0.46457	-AAATATAT-	
ATATATTG	0.45898	-CAATATAT-	
TATTAATA	0.45591	--TATTAATA	
AAATTGAT	0.45416	-AAATTGAT-	
ATTAAAT	0.45305	-ATTAAAT-	

8 mer Z-scores for probeset 'all'

Forward:		Reverse:	
			
Top 10	Scores	Alignment	
AATATATT	6.80307	- -AATATATT - -	
AATTTAAA	5.49601	- -TTTAAATT - -	
AAATATAT	5.25725	- -AAATATAT - -	
GAAATTTT	5.16522	- - - -GAAATTTT	
AATAAATT	5.14720	- -AATAAATT - -	
TATTAATA	4.96037	- -TATTAATA - -	
AAAATAAA	4.88125	AAAATAAA - - - -	
ATAATTTT	4.79419	- - - -ATAATTTT - -	
GAATATTC	4.77199	- - - -GAATATTC - -	
AAATTAAT	4.58082	- -AAATTAAT - - -	

Top scoring motifs for Assay 14681

Protein ID: pTH13919.3 Gene: GLYR1.DBD Domain: AT_hook Flag: Backup_matched_pair Array: 1M-ME

8 mer E-scores for probeset 'all'

Forward:		Reverse:	
Top 10	Scores	Alignment	
ATATATAT	0.46168	-ATATATAT-	
AAATATAT	0.45536	-AAATATAT-	
ATTTAAAT	0.45509	-ATTTAAAT-	
ATAATATA	0.45099	ATAATATA--	
TAAATTTA	0.44980	TAAATTTA--	
AATAATAA	0.44467	-AATAATAA-	
ATATATTA	0.44118	-TAATATAT-	
TTATATAA	0.43882	-TTATATAA-	
ATAATTAT	0.43746	-ATAATTAT-	
ATTATATA	0.43596	--ATTATATA	

8 mer Z-scores for probeset 'all'

Forward:		Reverse:
Top 10	Scores	Alignment
ATTTAAAT	5.13180	-ATTTAAAT--
AAATATAT	5.10548	-AAATATAT--
ATAATATA	4.94166	--ATAATATA-
TTATATAA	4.87884	---TTATATAA
ATATATTA	4.76412	---TAATATAT
ATTATATA	4.68903	--ATTATATA-
ATATATAT	4.53674	---ATATATAT
AATAATAA	4.46489	TTATTATT---
AAAATTTT	4.42557	--AAAATTTT--
TTTATAAA	4.23567	--TTTATAAA-