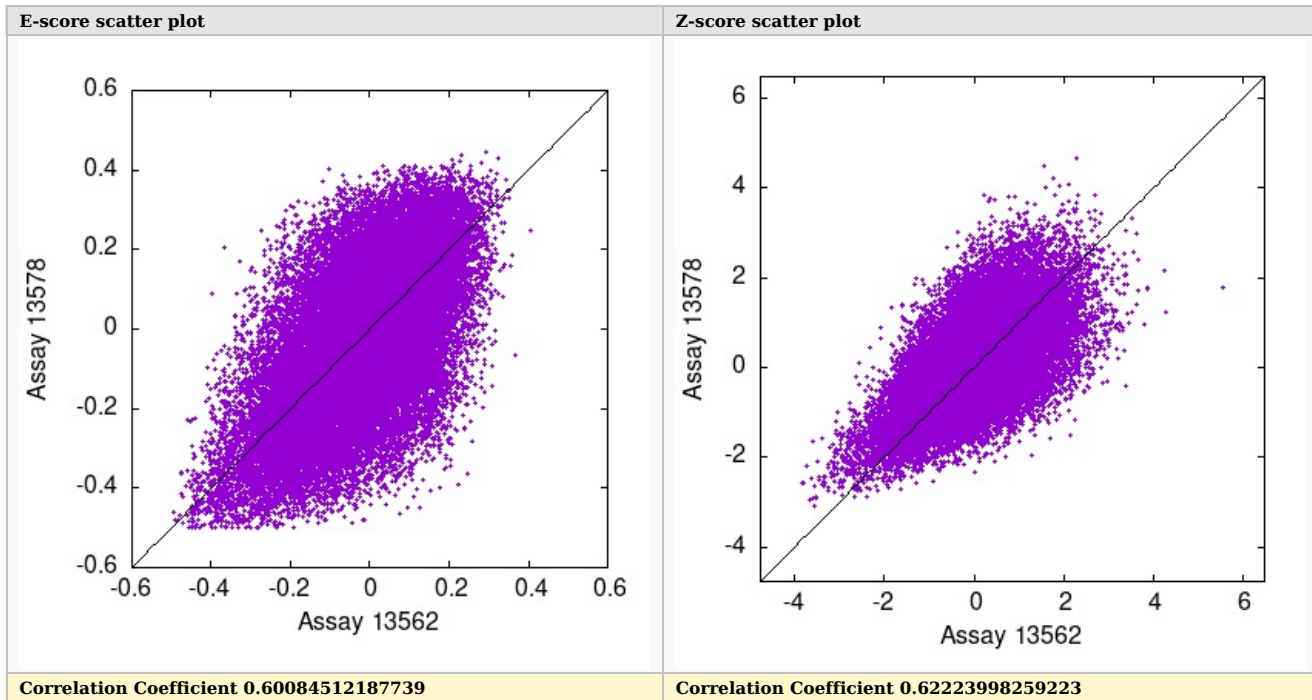


QC report



Top scoring motifs for Assay 13562

Protein ID: pTH13917.1 Gene: DNTTIP1.DBD Domain: AT_hook 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
ATGGTTTA	0.40562	----TAAACCAT	ATGGTTTA	5.54310	-TAAACCAT----
ATGAACTG	0.36517	---ATGAACTG-	ACATATAC	4.28014	ACATATAC----
ACATATAC	0.36044	---ACATATAC-	CGTAATAA	4.24388	---CGTAATAA--
GATGAGTA	0.35390	--GATGAGTA--	ATACAATA	3.87864	--ATACAATA--
GATATTAA	0.35129	GATATTAA----	ATAATGCA	3.87218	---TGCATTAT--
AATAATAG	0.34954	---AATAATAG-	ATACTCTA	3.80799	--ATACTCTA--
CGTAATAA	0.34836	----TTATTACG	ACTGATTG	3.67477	-----CAATCAGT
AAATATAT	0.34221	---AAATATAT-	ATTAATCG	3.65778	----CGATTAAT-
TATAAATA	0.34149	--TATAAATA--	AATCAGTA	3.63829	--AATCAGTA--
ACGAAATG	0.34056	--ACGAAATG--	AATATATG	3.61372	-CATATATT----

Top scoring motifs for Assay 13578

Protein ID: pTH13917.2 Gene: DNTTIP1.DBD Domain: AT_hook 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
ATATTAAT	0.44349	---ATATTAAT-	ATATTAAT	4.66670	-----ATATTAAT-
ATAGTAAT	0.43812	---ATAGTAAT-	CATATAAC	4.48736	----CATATAAC--
AATATTAA	0.43018	--AATATTAA--	GCATATAA	4.23368	---GCATATAA---
TATATGAA	0.42628	--TATATGAA--	TATTAATA	4.08793	-----TATTAATA
CTGTATTA	0.42310	-CTGTATTA---	TATTTGCA	4.06263	TGCAAATA-----
TATTAACA	0.41815	---TATTAACA	CATATATG	4.00870	----CATATATG--
TATTAATA	0.41593	----TATTAATA	AATTTAAA	3.84191	----AATTTAAA--
CATATAAC	0.41383	--CATATAAC--	AATATTAA	3.83866	----AATATTAA--
AATAATAA	0.41112	--AATAATAA--	AATAATAA	3.83644	----AATAATAA--
AATATAGA	0.41064	TCTATATT----	ATATAAAC	3.82957	-----ATATAAAC-