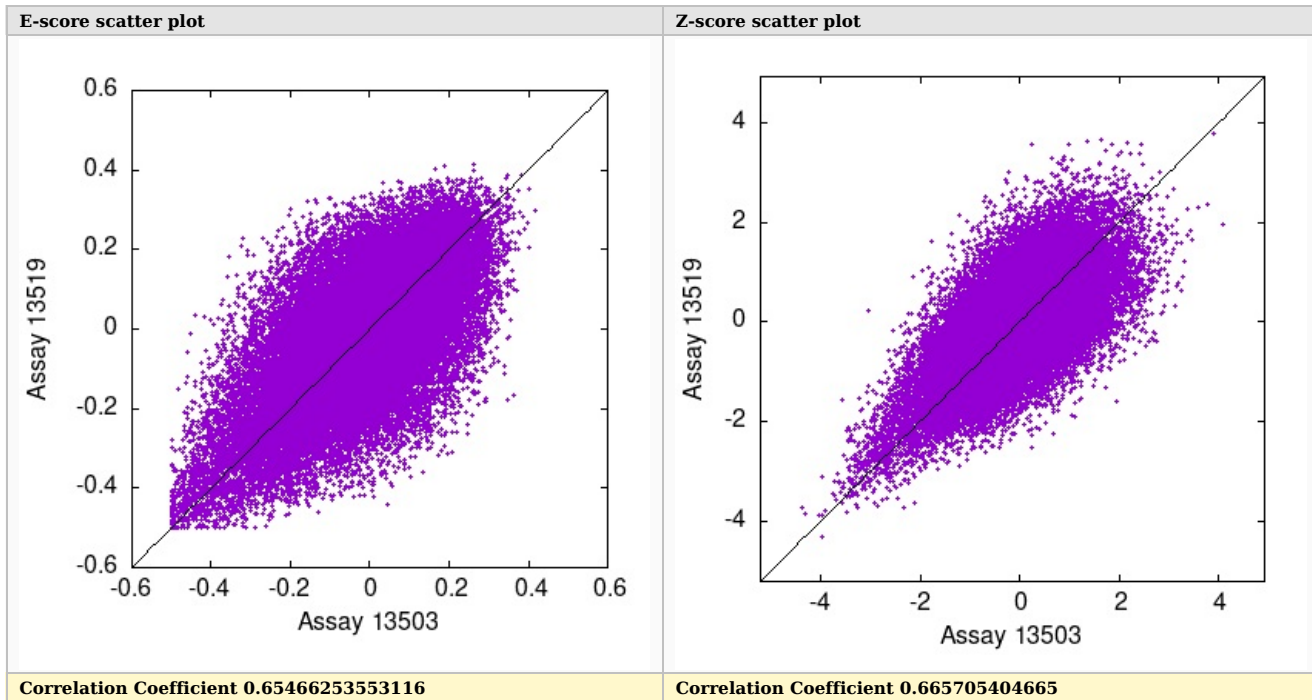


## QC report



## Top scoring motifs for Assay 13503

Protein ID: pTH13947.1 Gene: AC008770.DBD Domain: zf-C2H2 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
ATACAAAA	0.41604	---ATACAAAA	ATACAAAA	4.08514	-----ATACAAAA
CATAATAC	0.40353	--CATAATAC-	ACAAACAT	3.89211	---ACAAACAT--
TAACGAAA	0.40239	-TAACGAAA--	AACATACA	3.78678	--AACATACA---
ATGTATAA	0.39900	-ATGTATAA--	ATAATATA	3.58098	--ATAATATA---
ATTTATAA	0.39027	-ATTTATAA--	ATGTATAA	3.47000	---TTATACAT--
ACAAATAA	0.38519	-ACAAATAA--	AAATTATT	3.45276	--AAATTATT---
ACAAACAT	0.38375	-ACAAACAT--	ACACATAC	3.38279	---ACACATAC---
TACATACA	0.37772	TACATACA---	ATAACGAA	3.38162	ATAACGAA----
ACATACAA	0.37736	-ACATACAA--	ATAATGCA	3.30602	--ATAATGCA---
AGTACAAA	0.37610	--AGTACAAA-	TGTATACA	3.28442	--TGTATACA---

## Top scoring motifs for Assay 13519

Protein ID: pTH13947.2 Gene: AC008770.DBD Domain: zf-C2H2 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
ATAGTAAT	0.41344	---ATAGTAAT---	ACAAACAT	3.77497	ACAAACAT-----
GATATGTA	0.40971	--GATATGTA----	GATATGTA	3.65999	-GATATGTA-----
AATATATT	0.40313	AATATATT-----	TGTTAACA	3.62520	-----TGTTAACA
ATACATTA	0.39887	-ATACATTA----	ATATATTG	3.60485	--ATATATTG----
ACAAACAT	0.38650	---ATGTTTGT---	AATTTAAA	3.58148	-AATTTAAA-----
ATAATGTA	0.38561	--TACATTAT----	AACGTATT	3.57699	-AACGTATT-----
ACAATTCA	0.38088	-----TGAATTGT	TATTTGCA	3.55747	---TATTTGCA----
ATGTTAAC	0.38002	---ATGTTAAC---	ATAGTAAT	3.55438	-ATAGTAAT-----
TAATAATA	0.37834	---TAATAATA---	AATATATT	3.43093	-AATATATT-----
ATATTAAT	0.37817	---ATATTAAT---	ATGTTAAC	3.40403	----ATGTTAAC--