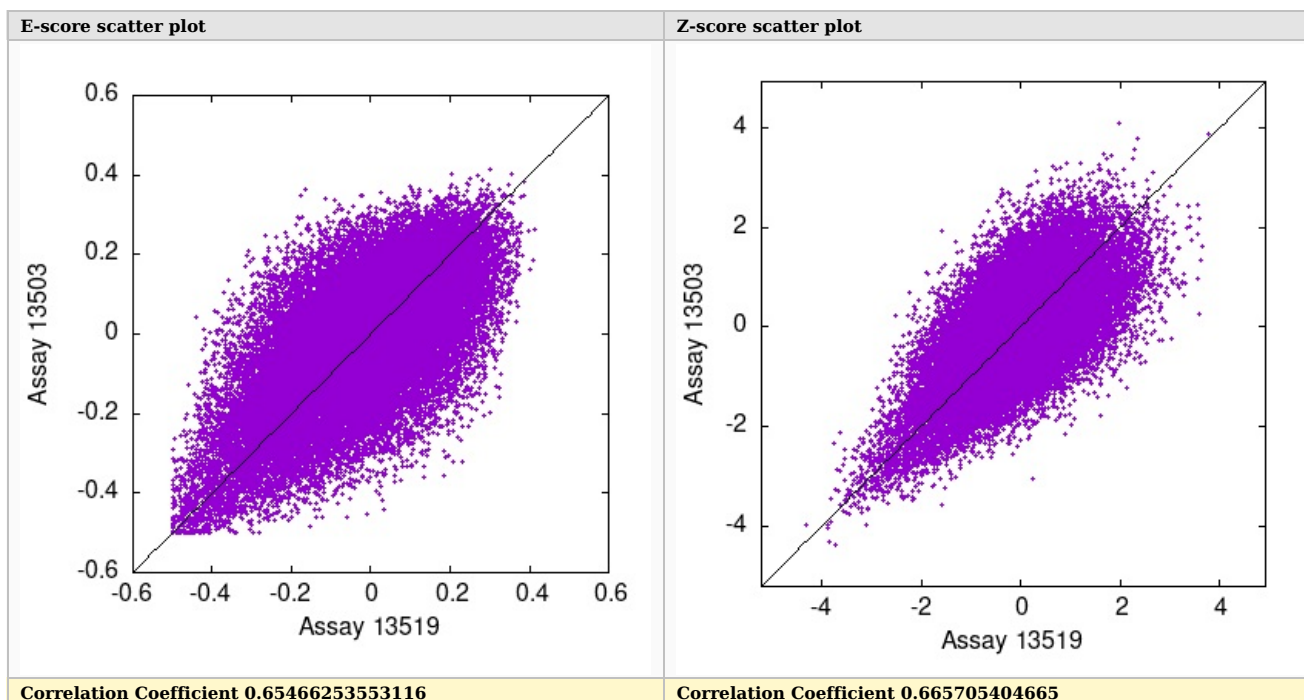


## QC report



### 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-

### 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---