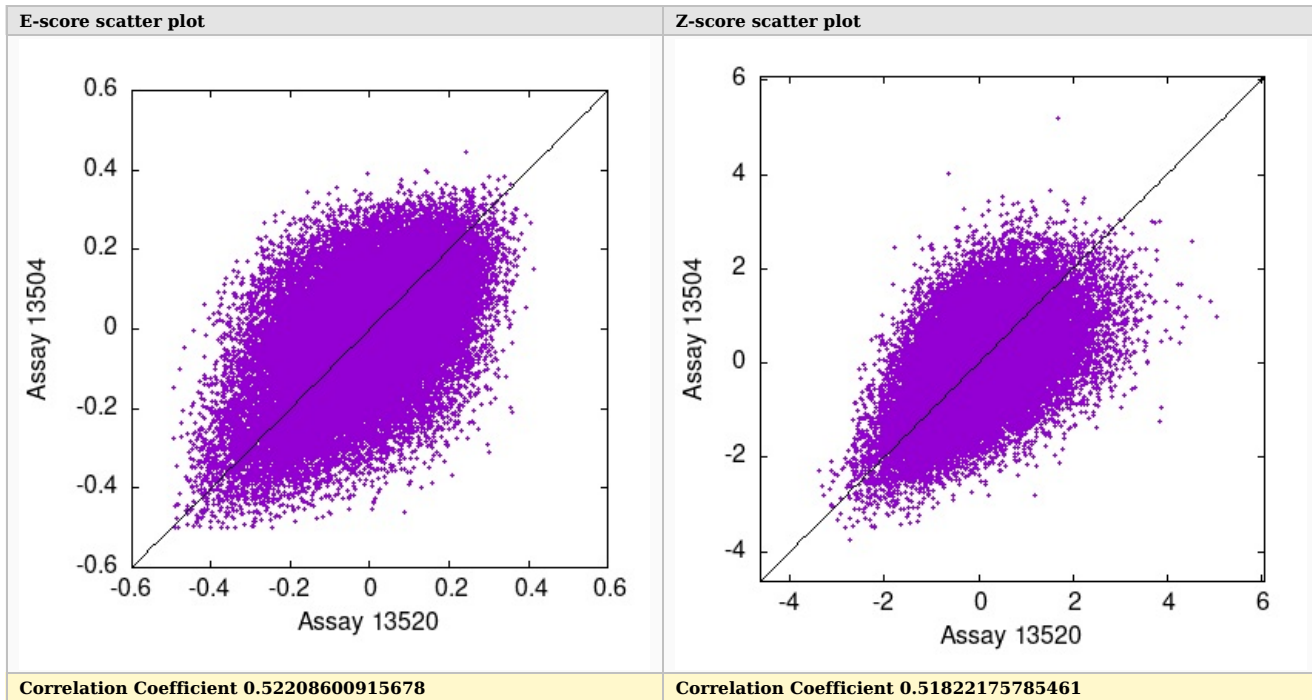


QC report



Top scoring motifs for Assay 13520

Protein ID: pTH13949.2 Gene: AC092835.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
TATTTGCA	0.41178	--TATTTGCA--	TATTTGCA	5.07006	---TATTTGCA---
AAGGTAGT	0.40733	--AAGGTAGT---	ATAGTAAT	4.92865	-----ATTACTAT
AAGTACTT	0.39527	--AAGTACTT--	ATACATTA	4.70828	--ATACATTA----
ATAGTAAT	0.39521	----ATAGTAAT	ATGTATAC	4.54790	GTATACAT-----
CTGTATTA	0.39184	--CTGTATTA--	GTACAACA	4.41926	--GTACAACA----
CATTTACA	0.38409	-CATTTACA---	AAGTACTT	4.32677	AAGTACTT-----
ATACATTA	0.37601	TAATGTAT----	AAGGTAGT	4.31198	-ACTACCTT-----
AAGTAAAA	0.37467	--AAGTAAAA--	CTACCAAC	4.28250	--CTACCAAC----
ATACTAGC	0.37037	---GCTAGTAT-	CTGTATTA	4.25188	TAATACAG-----
ATGTAGTA	0.36986	--ATGTAGTA--	ATGTATGC	4.23578	GCATACAT-----

Top scoring motifs for Assay 13504

Protein ID: pTH13949.1 Gene: AC092835.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
AGACTACC	0.44642	--GGTAGTCT---	AGACTACC	5.18131	---AGACTACC
AAACATAA	0.39862	-----AAACATAA	AGACGAAA	4.00419	-AGACGAAA--
TAACGAAA	0.39646	--TTTCGTTA---	ACATACAA	3.65041	-ACATACAA--
AGACGAAA	0.38962	--TTTCGTTCT---	ATAATGCA	3.53294	ATAATGCA---
GGTAGTCA	0.38493	--GGTAGTCA---	GTAGTCAA	3.49244	-GTAGTCAA--
ACTACTAT	0.38369	ATAGTAGT-----	AACATGCA	3.44270	AACATGCA---
ATAATGCA	0.37660	---ATAATGCA---	AACATACA	3.42844	AACATACA---
GTAGTCAA	0.37425	---GTAGTCAA---	ATAGTCTG	3.42629	--CAGACTAT-
ATGAGTAG	0.36711	---CTACTCAT--	TAACGAAA	3.40187	-TAACGAAA--
AAAACATA	0.36594	-----AAAACATA-	ATATACAG	3.39470	-ATATACAG--