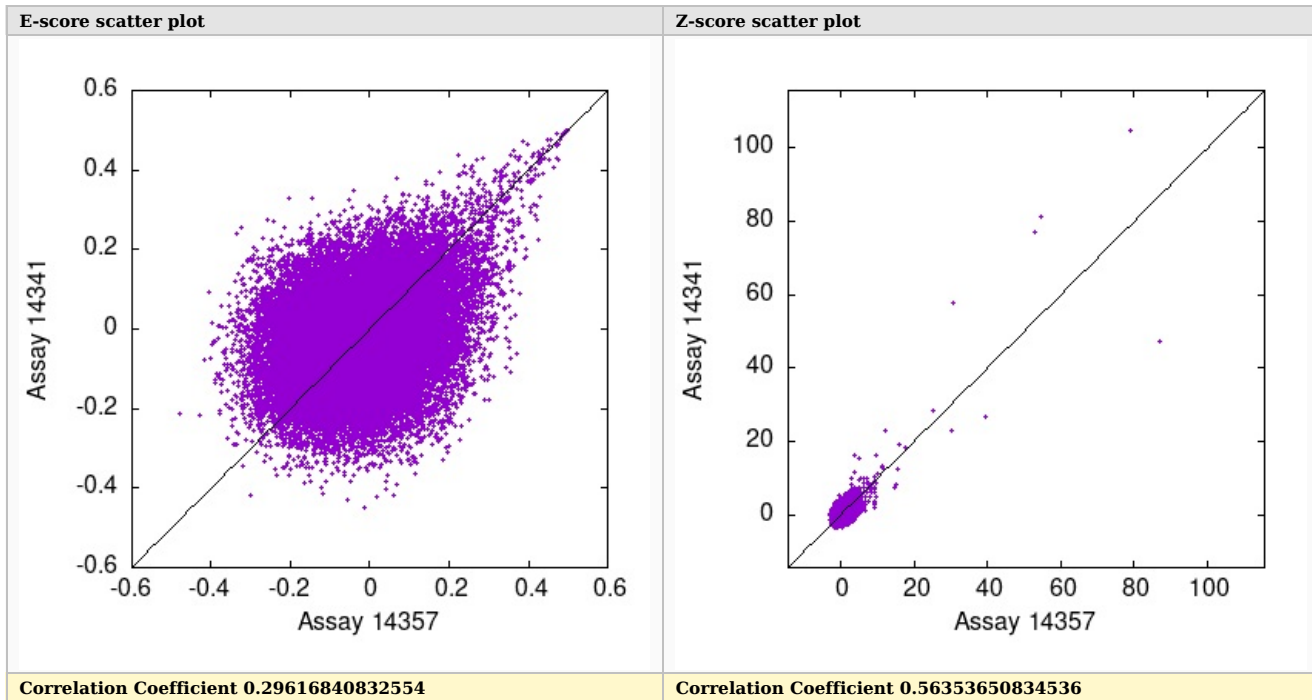


QC report



Top scoring motifs for Assay 14357

Protein ID: pTH15540.2 Gene: cJUN Domain: bZIP_1 Flag: Pass_matched_pair 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
ATGACTCA	0.49771	ATGACTCA-	ATGACGCA	86.76620	ATGACGCA-
ATGACGCA	0.49700	ATGACGCA-	ATGACTCA	79.00299	ATGACTCA-
ATGAGTCA	0.49658	-TGACTCAT	ATGAGTCA	54.36302	-TGACTCAT
TGACGTCA	0.49494	-TGACGTCA	TGACGTCA	52.94354	-TGACGTCA
ATGCGTCA	0.49280	-TGACGCAT	ATGCGTCA	39.37808	-TGACGCAT
GTGAGTCA	0.49081	-TGACTCAC	GTGAGTCA	30.65331	-TGACTCAC
GTGACTCA	0.48844	GTGACTCA-	GTGACTCA	30.29523	GTGACTCA-
ATGACTAA	0.48816	ATGACTAA-	ATGACTAA	25.10138	ATGACTAA-
TGACGTAA	0.48163	-TGACGTAA	GTGCGTCA	17.44397	-TGACGCAC
GTGCGTCA	0.48094	-TGACGCAC	TGACGTAA	16.06702	-TGACGTAA

Top scoring motifs for Assay 14341

Protein ID: pTH15540.1 Gene: cJUN Domain: bZIP_1 Flag: Pass_matched_pair 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
ATGAGTCA	0.49855	-TGACTCAT	ATGACTCA	104.73234	ATGACTCA-
ATGACTCA	0.49810	ATGACTCA-	ATGAGTCA	80.97356	-TGACTCAT
TGACGTCA	0.49695	-TGACGTCA	TGACGTCA	76.95390	-TGACGTCA
ATGACGCA	0.49538	ATGACGCA-	GTGAGTCA	57.76842	-TGACTCAC
GTGAGTCA	0.49505	-TGACTCAC	ATGACGCA	47.18297	ATGACGCA-
GTGACTCA	0.49197	GTGACTCA-	ATGACTAA	28.36665	ATGACTAA-
ATGCGTCA	0.49196	-TGACGCAT	ATGCGTCA	26.52485	-TGACGCAT
CTGACTCA	0.49124	CTGACTCA-	GTGACTCA	22.77214	GTGACTCA-
ATGACTAA	0.48711	ATGACTAA-	CTGACTCA	22.69827	CTGACTCA-
TGACGTAA	0.48338	-TGACGTAA	TGACGTAA	18.96248	-TGACGTAA