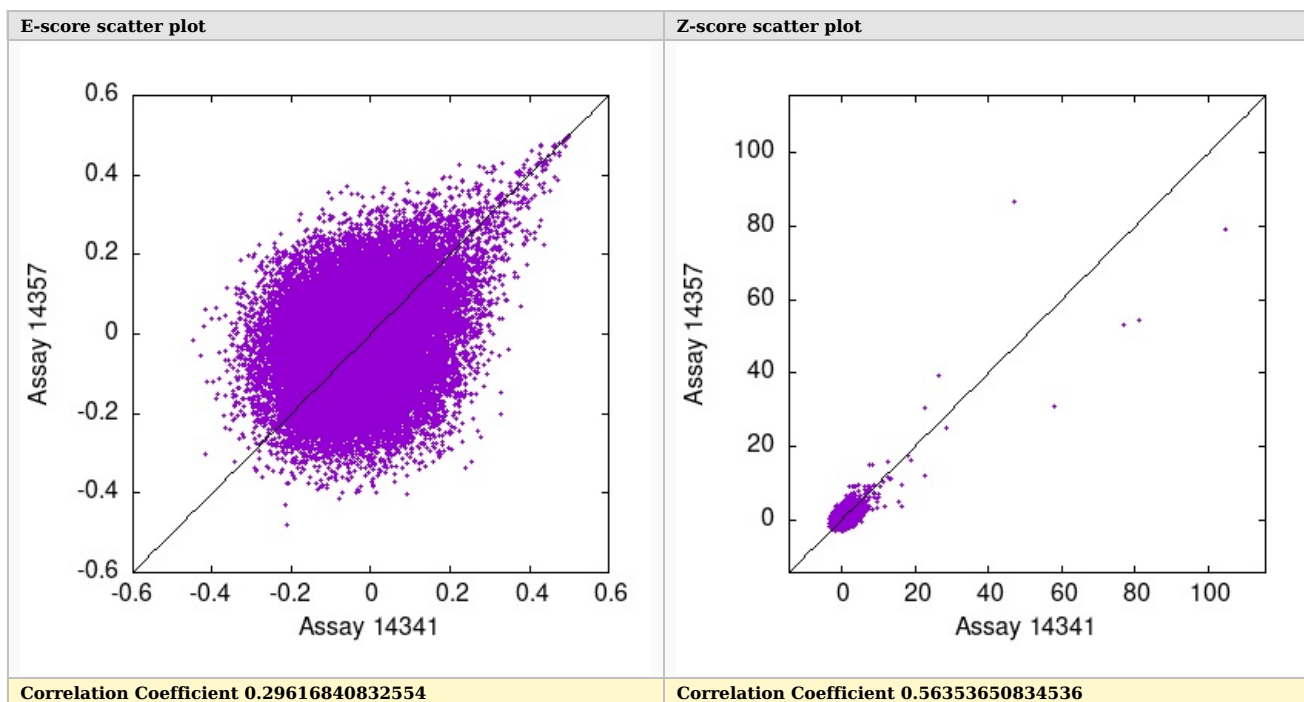




QC report





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'

Forward:		Reverse:	
			
Top 10	Scores	Alignment	
ATGAGTCA	0.49855	-TGACTCAT	
ATGACTCA	0.49810	ATGACTCA-	
TGACGTCA	0.49695	-TGACGTCA	
ATGACGCA	0.49538	ATGACGCA-	
GTGAGTCA	0.49505	-TGACTCAC	
GTGACTCA	0.49197	GTGACTCA-	
ATGCGTCA	0.49196	-TGACGCAT	
CTGACTCA	0.49124	CTGACTCA-	
ATGACTAA	0.48711	ATGACTAA-	
TGACGTAA	0.48338	-TGACGTAA	



8 mer Z-scores for probeset 'all'

Forward:		Reverse:	
			
Top 10	Scores	Alignment	
ATGACTCA	104.73234	ATGACTCA-	
ATGAGTCA	80.97356	-TGACTCAT	
TGACGTCA	76.95390	-TGACGTCA	
GTGAGTCA	57.76842	-TGACTCAC	
ATGACGCA	47.18297	ATGACGCA-	
ATGACTAA	28.36665	ATGACTAA-	
ATGCGTCA	26.52485	-TGACGCAT	
GTGACTCA	22.77214	GTGACTCA-	
CTGACTCA	22.69827	CTGACTCA-	
TGACGTAA	18.96248	-TGACGTAA	



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'

Forward:		Reverse:
		
Top 10	Scores	Alignment
ATGACTCA	0.49771	ATGACTCA-
ATGACGCA	0.49700	ATGACGCA-
ATGAGTCA	0.49658	-TGACTCAT
TGACGTCA	0.49494	-TGACGTCA
ATGCGTCA	0.49280	-TGACGCAT
GTGAGTCA	0.49081	-TGACTCAC
GTGACTCA	0.48844	GTGACTCA-
ATGACTAA	0.48816	ATGACTAA-
TGACGTAA	0.48163	-TGACGTAA
GTGCGTCA	0.48094	-TGACGCAC

8 mer Z-scores for probeset 'all'

Forward:		Reverse:	
			
Top 10	Scores	Alignment	
ATGACGCA	86.76620	ATGACGCA-	
ATGACTCA	79.00299	ATGACTCA-	
ATGAGTCA	54.36302	-TGACTCAT	
TGACGTCA	52.94354	-TGACGTCA	
ATGCGTCA	39.37808	-TGACGCAT	
GTGAGTCA	30.65331	-TGACTCAC	
GTGACTCA	30.29523	GTGACTCA-	
ATGACTAA	25.10138	ATGACTAA-	
GTGCGTCA	17.44397	-TGACGCAC	
TGACGTAA	16.06702	-TGACGTAA	