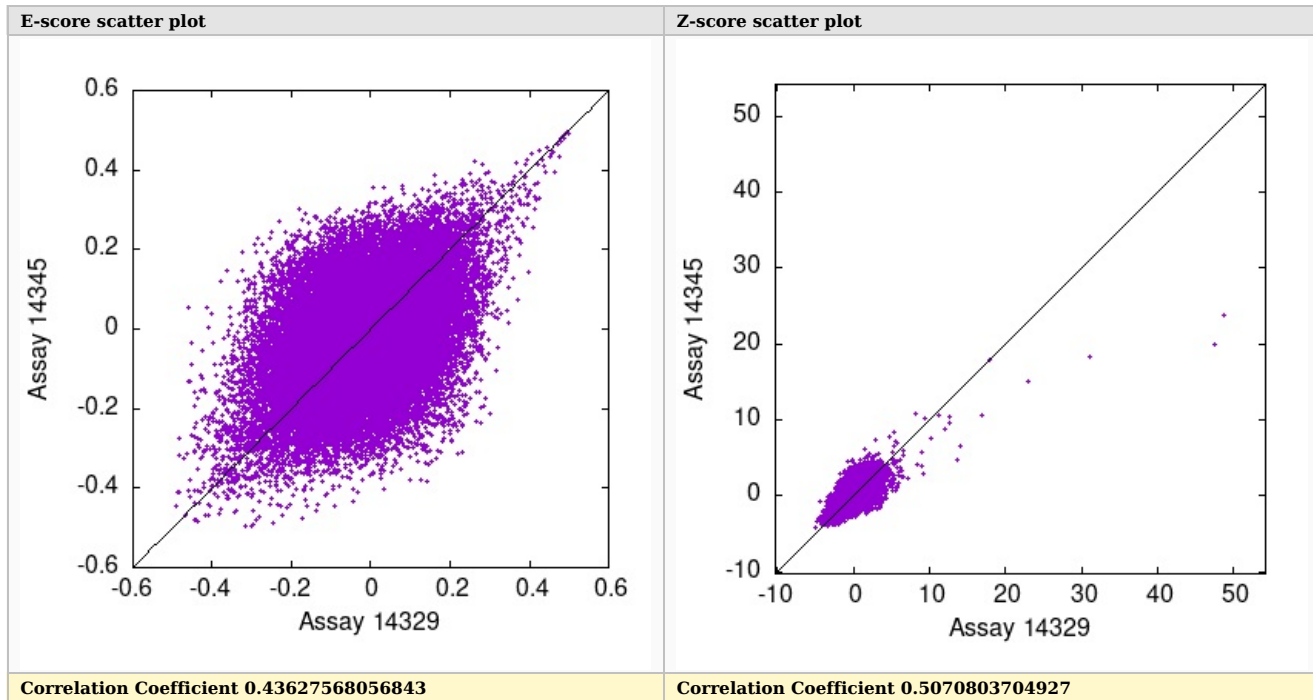


QC report



Top scoring motifs for Assay 14329

Protein ID: pTH15523.1 Gene: JUNB 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	0.49748	-TGACTCAT	ATGACTCA	48.79270	ATGACTCA-
ATGACTCA	0.49678	ATGACTCA-	ATGAGTCA	47.38880	-TGACTCAT
GTGAGTCA	0.48759	-TGACTCAC	GTGAGTCA	31.11947	-TGACTCAC
ATGACTAA	0.48658	ATGACTAA-	ATGACTAA	22.90196	ATGACTAA-
GTGACTCA	0.48136	GTGACTCA-	GTGACTCA	17.87174	GTGACTCA-
ATTAGTCA	0.47817	-TGACTAAT	ATTAGTCA	16.76451	-TGACTAAT
CTGACTCA	0.47533	CTGACTCA-	CTGACTCA	14.11243	CTGACTCA-
TGACGTCA	0.47360	-TGACGTCA	CTGAGTCA	13.54571	-TGACTCAG
ATGACGCA	0.47119	ATGACGCA-	GTGACTAA	12.70878	GTGACTAA-
CTGAGTCA	0.46459	-TGACTCAG	ATGACGCA	12.62905	ATGACGCA-

Top scoring motifs for Assay 14345

Protein ID: pTH15523.2 Gene: JUNB 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	0.49500	ATGACTCA-	ATGACTCA	23.73626	ATGACTCA-
ATGAGTCA	0.49049	-TGACTCAT	ATGAGTCA	19.84882	-TGACTCAT
GTGAGTCA	0.48682	-TGACTCAC	GTGAGTCA	18.22112	-TGACTCAC
GTGACTCA	0.48170	GTGACTCA-	GTGACTCA	17.88284	GTGACTCA-
ATGACTAA	0.47923	ATGACTAA-	ATGACTAA	15.05573	ATGACTAA-
ATTAGTCA	0.47385	-TGACTAAT	ATTACTCA	10.83359	ATTACTCA-
ATGACGCA	0.46438	ATGACGCA-	ATTAGTCA	10.60495	-TGACTAAT
TGACGTCA	0.46361	-TGACGTCA	TGACGTCA	10.56336	-TGACGTCA
ATTACTCA	0.45495	ATTACTCA-	ATGACGCA	10.31939	ATGACGCA-
ATGCGTCA	0.45049	-TGACGCAT	ATGCGTCA	10.14866	-TGACGCAT