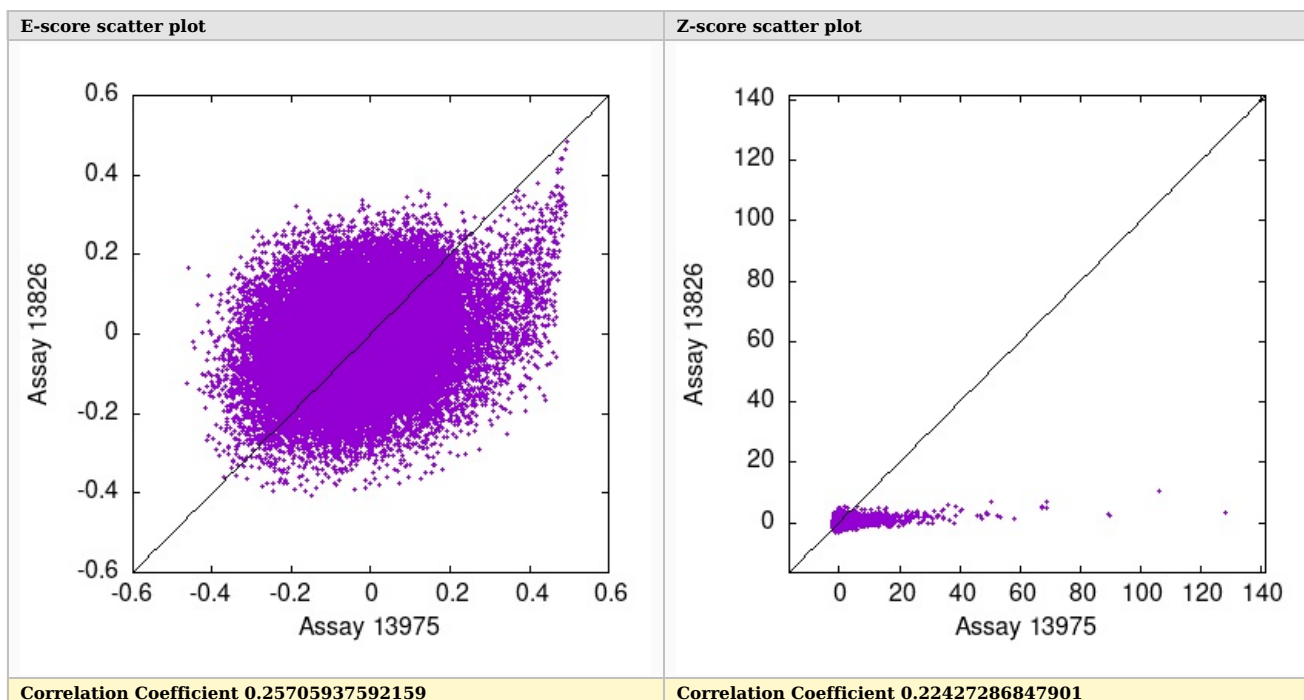


QC report



Top scoring motifs for Assay 13975

Protein ID: pTH13931.3 Gene: ZBED4 Domain: zf-BED Flag: Pass matched_pair Array: 1M-HK

8 mer E-scores for probeset 'all'

Forward:	Reverse:
Top 10	Scores
CGGTACCG	0.49340
TGGTACCA	0.49241
CGGTCAA	0.49079
CGGTACCA	0.49026
ATTGGACC	0.48954
CTGGTACC	0.48734
AGGTTCCG	0.48486
GCCGAACC	0.48390
CCGGTACC	0.48389
CCGAACCG	0.48200

8 mer Z-scores for probeset 'all'

Forward:	Reverse:
Top 10	Scores
CGGTCCAA	128.23762
CGGTACCG	105.93106
TGGTACCA	89.93481
ATTGGACC	89.49647
CGGTACCA	69.00148
CCGAACCA	68.77158
CCGGTACC	67.43938
GCCGAACC	67.14504
ATTGGGTC	58.01877
CTGGTACC	53.46846

Top scoring motifs for Assay 13826

Protein ID: pTH13931.1 Gene: ZBED4 Domain: zf-BED Flag: Pass matched_pair Array: 1M-ME

8 mer E-scores for probeset 'all'

Forward:	Reverse:
Top 10	Scores
CGGTACCG	0.48308
CGGTACCA	0.46619
GGTTCCGA	0.44051
CCGGTACC	0.44047
CCGAACCA	0.41598
CGGAACCG	0.41530
GGTACCGA	0.40596
ACCGGTAC	0.38076
GGTTCCGA	0.36996
GGTACCAA	0.36985

8 mer Z-scores for probeset 'all'

Forward:	Reverse:
Top 10	Scores
CGGTACCG	10.57487
GGTTCCGA	6.86183
CGGTACCA	6.65803
GGTACCGA	5.74843
CCGGTACC	5.55751
CCGAACCA	5.51458
CATAATAC	5.44774
ACCCAAC	5.14440
GCCGAACC	5.07613
ATTATTAC	4.92334