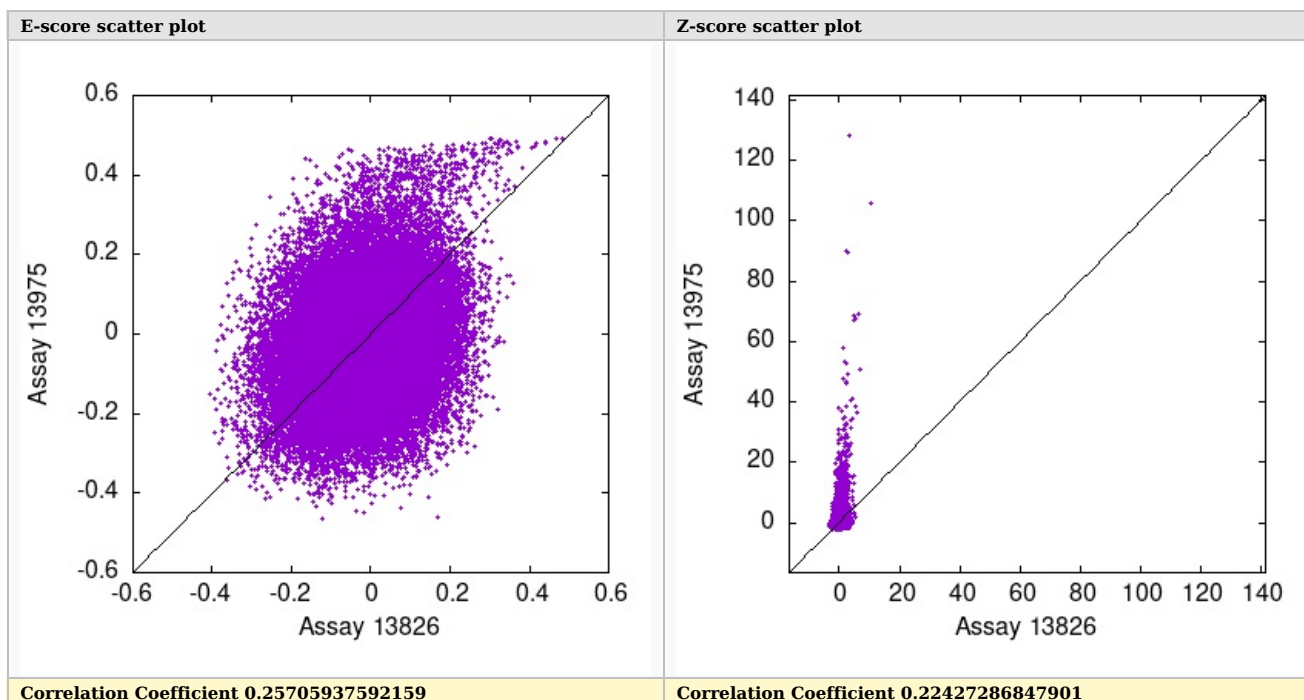




QC report





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	Alignment
CGGTACCG	0.48308	CGGTACCG--
CGGTACCA	0.46619	TGGTACCG--
GGTTCGGA	0.44051	-GGTTCGGA-
CCGGTACC	0.44047	-GGTACCGG-
CGGAACCA	0.41598	TGGTTCGG--
CGGAACCG	0.41530	CGGAACCG--
GGTACCGA	0.40596	-GGTACCGA-
ACCGGTAC	0.38076	--GTACCGGT
GGTTCGGA	0.36996	-GGTTCGGA-
GGTACCAA	0.36985	-GGTACCAA-



8 mer Z-scores for probeset 'all'

Forward:		Reverse:
		
Top 10	Scores	Alignment
CGGTACCG	10.57487	---CGGTACCG--
GGTTCCGA	6.86183	---TCGGAACC---
CGGTACCA	6.65803	---CGGTACCA--
GGTACCGA	5.74843	---TCGGTACC---
CCGGTACC	5.55751	---CCGGTACC---
CGGAACCA	5.51458	---CGGAACCA--
CATAATAC	5.44774	-----GTATTATG
ACCCAAC	5.14440	GTTTGGGT-----
GCCGAACC	5.07613	---GCCGAACC---
ATTATTAC	4.92334	-----GTAATAAT



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	Alignment
CGGTACCG	0.49340	-CGGTACCG-
TGGTACCA	0.49241	-TGGTACCA-
CGGTCCAA	0.49079	-CGGTCCAA-
CGGTACCA	0.49026	-CGGTACCA-
ATTGGACC	0.48954	--GGTCCAAT
CTGGTACC	0.48734	--GGTACCAG
AGGTTCCG	0.48486	-CCGAACCT-
GCCGAACC	0.48390	GCCGAACC--
CCGGTACC	0.48389	CCGGTACC--
CCGAACCG	0.48200	-CCGAACCG-

8 mer Z-scores for probeset 'all'

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