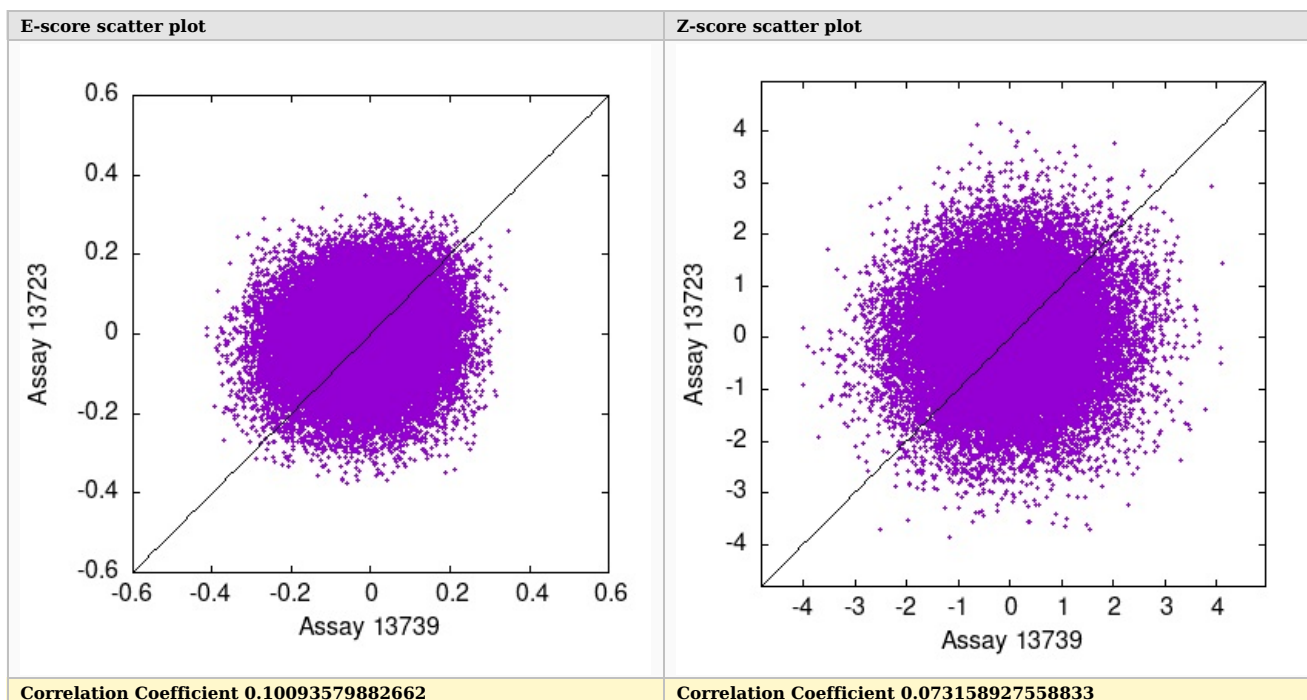


QC report



Top scoring motifs for Assay 13739

Protein ID: pTH14305.2 Gene: GPBP1L1.FL Domain: Unknown Flag: Reject Array: 1M-HK

8 mer E-scores for probeset 'all'

Forward:	Reverse:
Top 10	Scores
CGGATCCG	0.34938
AAAGTTGT	0.33947
GCAAAAAA	0.32680
CAGTACAA	0.32327
GATGCTTA	0.31941
ATGCAAAA	0.31539
GCAAATGA	0.31422
ACTGATAT	0.30938
AAAATATC	0.30629
GTAATTAC	0.30506

8 mer Z-scores for probeset 'all'

Forward:	Reverse:
Top 10	Scores
AGTTAACT	4.12263
AACAACCTA	4.09666
CAACTGCA	4.07481
CGGATCCG	3.89115
AAATAACG	3.77905
AACATACT	3.67281
GCAAAAAA	3.65919
AAAACCTTA	3.60710
CTGTTTTT	3.60045
CATCACTA	3.58557

Top scoring motifs for Assay 13723

Protein ID: pTH14305.1 Gene: GPBP1L1.FL Domain: Unknown Flag: Reject Array: 1M-ME

8 mer E-scores for probeset 'all'

Forward:	Reverse:
Top 10	Scores
ATGTCACG	0.34998
GTAAGTAC	0.34144
GCATCGAA	0.32143
ATAGAGAT	0.31899
AATCAATG	0.31312
AACACGAG	0.30679
TAACAGCA	0.30011
ATGTATAA	0.30000
TCCAGCCA	0.29782
ATTACAAC	0.29705

8 mer Z-scores for probeset 'all'

Forward:	Reverse:
Top 10	Scores
ATGTCACG	4.13534
ATAGAGAT	4.11204
GTAAGTAC	3.98725
AGATATAT	3.95398
ATAACAGC	3.79799
ATTACAAC	3.76845
ACGGTCAA	3.72603
GCATCGAA	3.69649
CAATGTTC	3.69192
TAACAGCA	3.62037