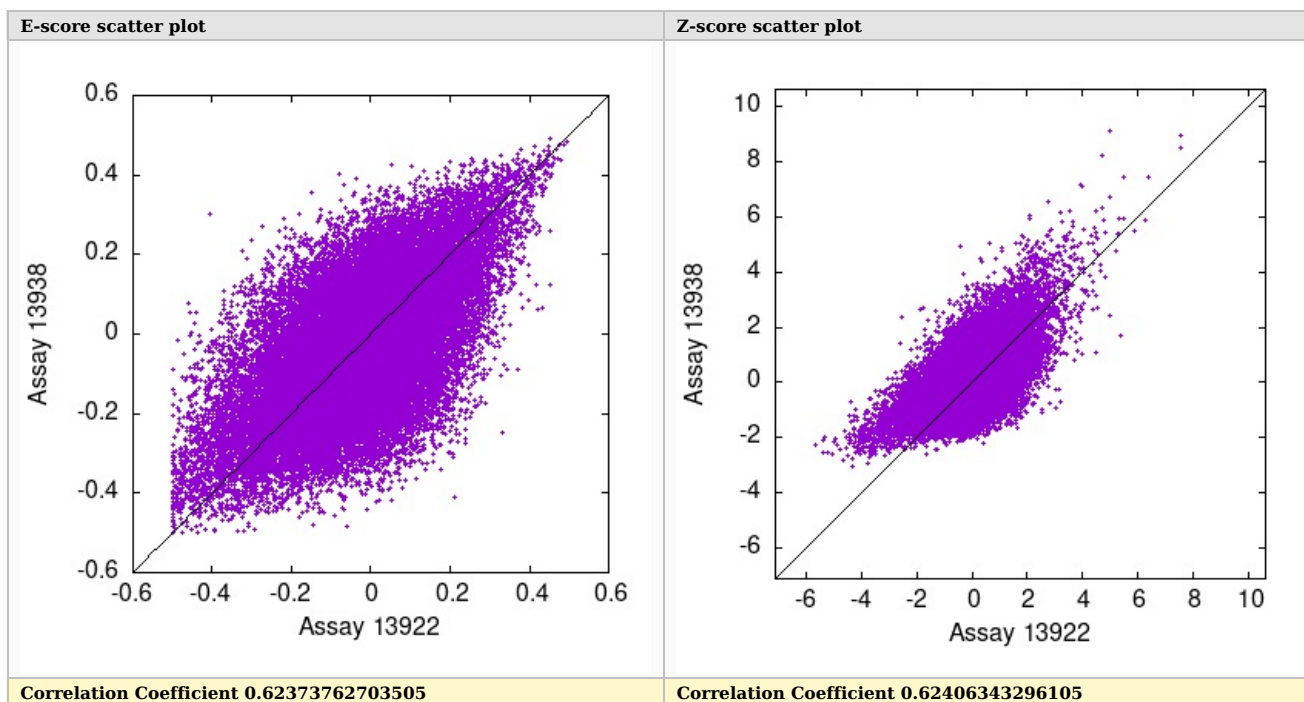


QC report



Top scoring motifs for Assay 13922

Protein ID: pTH14223.1 Gene: KDM2A.DBD Domain: zf-CXXC Flag: Pass_matched_pair Array: 1M-ME

8 mer E-scores for probeset 'all'

Forward:	Reverse:
Top 10	Scores
CGCGCGCG	0.49329
TACGCGTA	0.48626
ACGTACGT	0.48059
GC CGCGCG	0.47718
CGCGTAAC	0.47211
GCGATCGC	0.46795
ATCGCGCG	0.46528
ACGCGTAA	0.46023
ATACGTAT	0.46004
CGATCGCG	0.45733

8 mer Z-scores for probeset 'all'

Forward:	Reverse:
Top 10	Scores
TACGCGTA	7.56474
CGCGCGCG	7.56077
GCGCGCGC	6.36709
CGCGTAAC	6.27990
ACGTACGT	5.88930
ACGCGTAA	5.52059
GCGATCGC	5.48057
CGAATTCG	5.38351
AATACGTA	5.32068
ATACGTAT	5.32068

Top scoring motifs for Assay 13938

Protein ID: pTH14223.2 Gene: KDM2A.DBD Domain: zf-CXXC Flag: Pass_matched_pair Array: 1M-HK

8 mer E-scores for probeset 'all'

Forward:	Reverse:
Top 10	Scores
TCGCGCGA	0.49053
CGCGCGCG	0.48578
GCGCGCGC	0.47717
TACGCGTA	0.47512
TTACGTAA	0.47389
CGCGCGAA	0.47256
GCGATCGC	0.46556
TACGTAAA	0.46267
GCGTACGC	0.46143
CGCGCGTA	0.46034

8 mer Z-scores for probeset 'all'

Forward:	Reverse:
Top 10	Scores
TCGCGCGA	9.15609
CGCGCGCG	8.97293
TACGCGTA	8.51206
TTACGTAA	8.21350
GCGCGCGC	7.47813
GCGATCGC	7.43904
TACGTAAA	7.17926
CGCGCGAA	7.10477
CGTACGTA	6.72384
ACGTACGA	6.54474