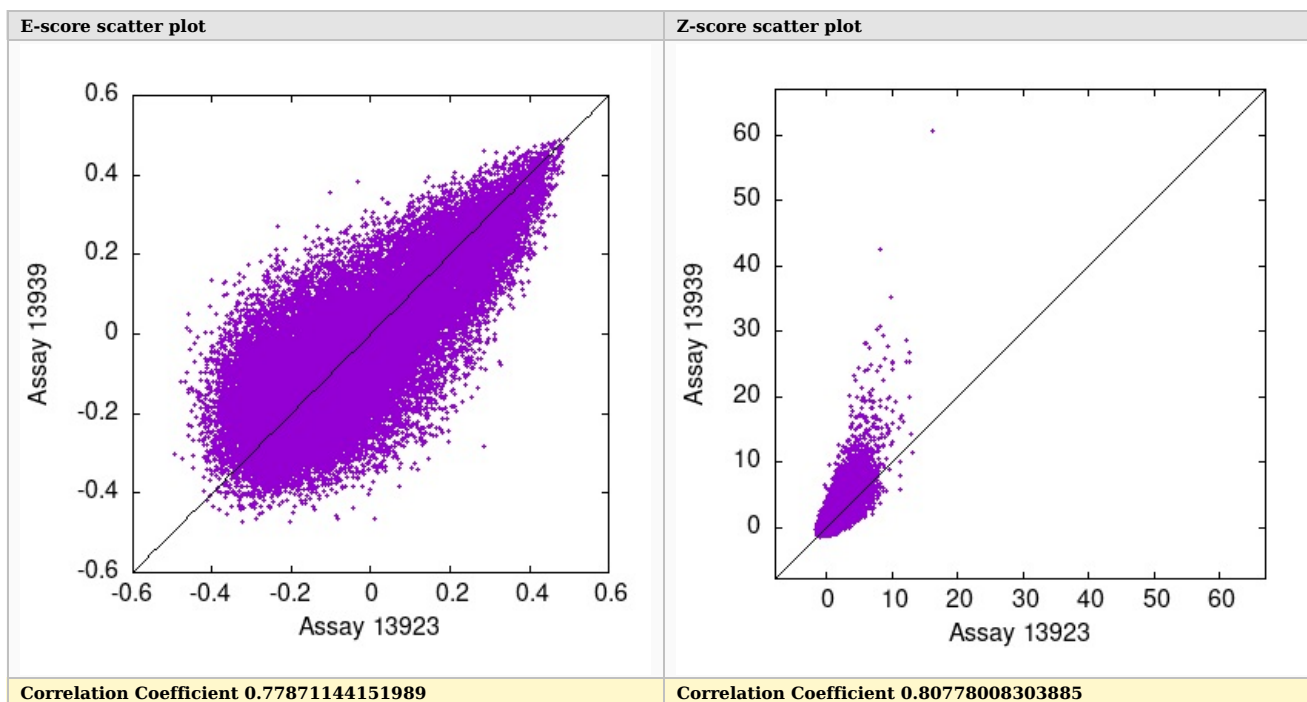


## QC report



### Top scoring motifs for Assay 13923

Protein ID: pTH14224.1 Gene: TET3 Domain: zf-CXXC Flag: Pass\_matched\_pair Array: 1M-ME

#### 8 mer E-scores for probeset 'all'

Forward:	Reverse:
Top 10	Scores
ACGTACGT	0.49355
GGGGGGGA	0.48511
CCCCCCCC	0.48468
ACGTACGC	0.48323
CGTACGTC	0.48256
CGTACGTA	0.48172
GTTACGTA	0.47916
CGCGTACG	0.47870
CGACGTAC	0.47785
ACGTAGCG	0.47701

#### 8 mer Z-scores for probeset 'all'

Forward:	Reverse:
Top 10	Scores
ACGTACGT	16.27967
GGGGGGGA	13.12314
CGCGTACG	12.82485
CGTACGTC	12.73670
ACGTACGC	12.64737
CGTACGTA	12.62628
GTACGTAC	12.20438
GTTACGTA	12.16962
ACGTAGCG	11.70842
GCGTACGC	11.60819

### Top scoring motifs for Assay 13939

Protein ID: pTH14224.2 Gene: TET3 Domain: zf-CXXC Flag: Pass\_matched\_pair Array: 1M-HK

#### 8 mer E-scores for probeset 'all'

Forward:	Reverse:
Top 10	Scores
ACGTACGT	0.49348
CACGTACG	0.48775
CTACGTAC	0.48552
ACGTAACG	0.47964
CCTACGTA	0.47897
TACGTAAA	0.47806
GTACGTAA	0.47642
CTACGTAG	0.47586
ATACGTAA	0.47507
CGTACGTC	0.47244

#### 8 mer Z-scores for probeset 'all'

Forward:	Reverse:
Top 10	Scores
ACGTACGT	60.77964
CACGTACG	42.50194
CTACGTAC	35.25481
TTACGTAA	30.80576
GTACGTAA	30.39577
TACGTAAA	29.38805
GTTACGTA	28.61336
AGGGGGGA	28.27380
ACGTTGCG	28.12513
ACGTAACG	27.83867