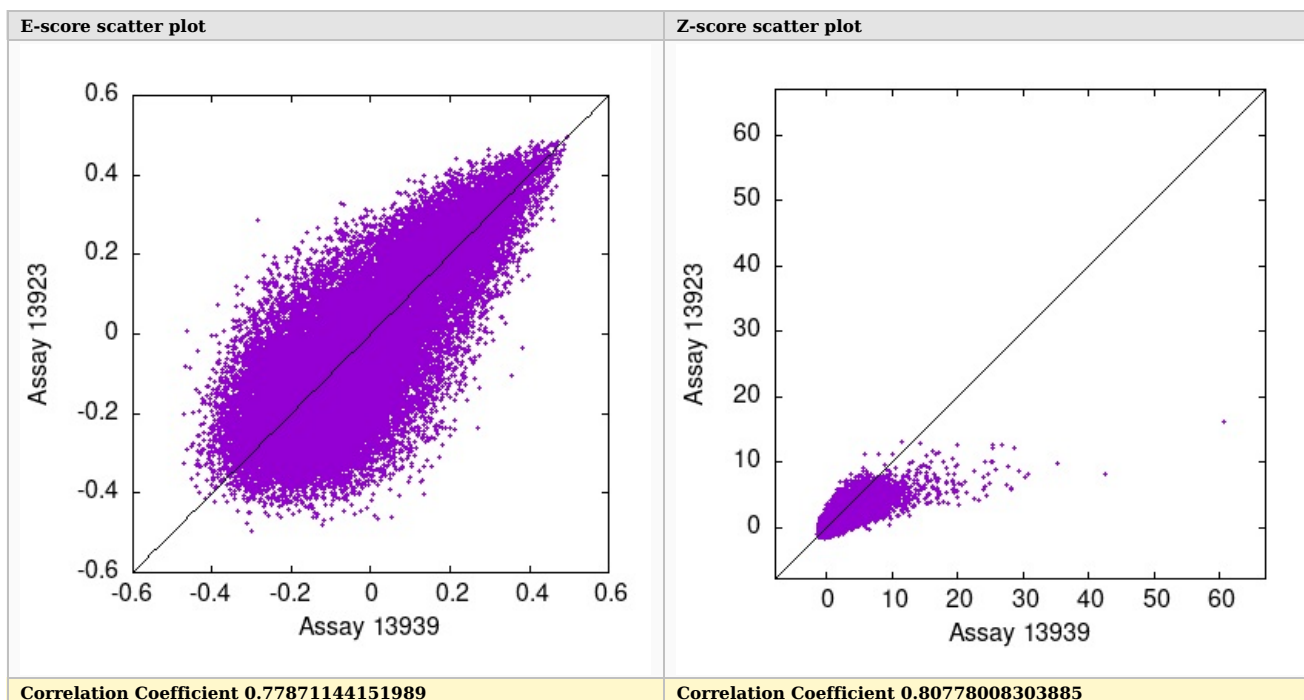


QC report



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	Alignment
ACGTACGT	0.49348	ACGTACGT----
CACGTACG	0.48775	-CGTACGTG---
CTACGTAC	0.48552	--GTACGTAG--
ACGTAACG	0.47964	----ACGTAACG
CCTACGTA	0.47897	-CCTACGTA---
TACGTAAA	0.47806	---TACGTAAA--
GTACGTAA	0.47642	--GTACGTAA--
CTACGTAG	0.47586	--CTACGTAG--
ATACGTAA	0.47507	--ATACGTAA--
CGTACGTC	0.47244	-CGTACGTC---

8 mer Z-scores for probeset 'all'

Forward:	Reverse:

Top 10	Scores	Alignment
ACGTACGT	60.77964	ACGTACGT-----
CACGTACG	42.50194	-CGTACGTG----
CTACGTAC	35.25481	--GTACGTAG----
TTACGTAA	30.80576	---TTACGTAA----
GTACGTAA	30.39577	--GTACGTAA----
TACGTAAA	29.38805	---TACGTAAA----
GTTACGTA	28.61336	---TACGTAA-----
AGGGGGGA	28.27380	-AGGGGGGA-----
ACGTTGCG	28.12513	----CGCAACGT---
ACGTAACG	27.83867	----ACGTAACG---

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	Alignment
ACGTACGT	0.49355	---ACGTACGT-
GGGGGGGA	0.48511	---TCCCCCCC-
CCCCCCCC	0.48468	---CCCCCCCC-
ACGTACGC	0.48323	---ACGTACGC-
CGTACGTC	0.48256	--GACGTACG--
CGTACGTA	0.48172	--TACGTACG--
GTTACGTA	0.47916	GTTACGTA----
CGCGTACG	0.47870	----CGTACGCG
CGACGTAC	0.47785	--CGACGTAC---
ACGTAGCG	0.47701	---ACGTAGCG-

8 mer Z-scores for probeset 'all'

Forward:	Reverse:

Top 10	Scores	Alignment
ACGTACGT	16.27967	---ACGTACGT-
GGGGGGGA	13.12314	---TCCCCCCC-
CGCGTACG	12.82485	---CGTACGCG-
CGTACGTC	12.73670	--GACGTACG--
ACGTACGC	12.64737	---ACGTACGC-
CGTACGTA	12.62628	--TACGTACG--
GTACGTAC	12.20438	-GTACGTAC---
GTTACGTA	12.16962	GTTACGTA----
ACGTAGCG	11.70842	---ACGTAGCG-
GCGTACGC	11.60819	---GCGTACGC-