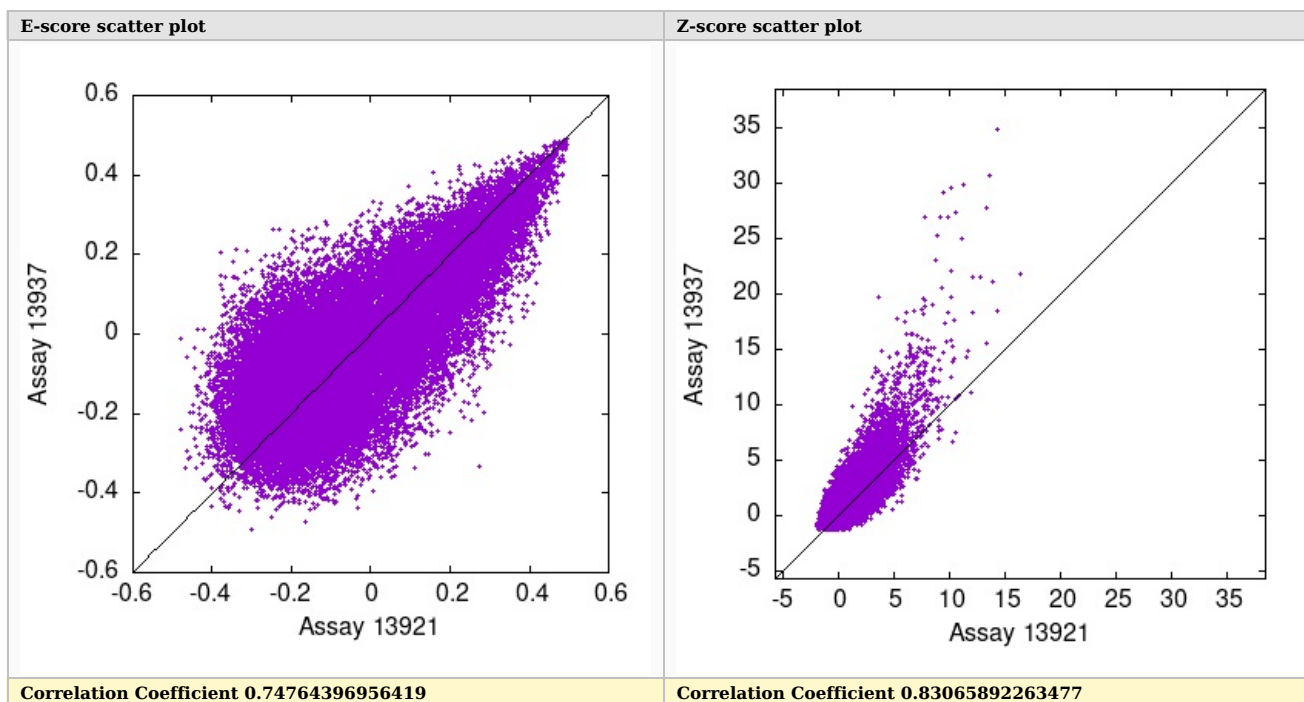


QC report



Top scoring motifs for Assay 13921

Protein ID: pTH14222.1 Gene: FBXL19 Domain: zf-CXXC Flag: Pass matched_pair Array: 1M-ME

8 mer E-scores for probeset 'all'

Forward:	Reverse:

Top 10	Scores	Alignment
ACGTACGT	0.49608	ACGTACGT---
TTACGTAA	0.49435	--TTACGTAA-
TACGCGTA	0.49220	-TACGCGTA--
CGTACGTA	0.49145	-CGTACGTA--
GTTACGTA	0.49071	---TACGTAAC
GCGTACGC	0.48936	GCGTACGC---
CGCGCGCG	0.48725	-CGCGCGCG--
ATTACGTA	0.48598	---TACGTAAT
GCGATCGC	0.48528	GCGATCGC---
CGTATACG	0.48504	-CGTATACG--

8 mer Z-scores for probeset 'all'

Forward:	Reverse:

Top 10	Scores	Alignment
ACGTACGT	16.41565	---ACGTACGT---
TACGCGTA	14.30965	-----TACGCGTA
TTACGTAA	14.29846	-TTACGTAA-----
GCGTACGC	13.93642	---GCGTACGC---
GTTACGTA	13.60979	GTTACGTA-----
CGCGCGCG	13.39153	---CGCGCGCG---
CGTACGTA	13.36270	--TACGTACG----
ACGTACGC	12.78945	---ACGTACGC---
ACGATCGT	12.12672	---ACGATCGT---
CGCGTACG	12.10318	---CGTACGCG--

Top scoring motifs for Assay 13937

Protein ID: pTH14222.2 Gene: FBXL19 Domain: zf-CXXC Flag: Pass matched_pair Array: 1M-HK

8 mer E-scores for probeset 'all'

Forward:	Reverse:

Top 10	Scores	Alignment
TTACGTAA	0.49053	--TTACGTAA-
ATTACGTA	0.48879	---TACGTAAT
GTTACGTA	0.48481	---TACGTAAC
GTACGTAA	0.48363	--GTACGTAA-
CACGTACG	0.48237	-CGTACGTG--
TACGTAAA	0.48182	---TACGTAAA
GCGATCGC	0.48166	GCGATCGC---
CGTACGTA	0.47772	-CGTACGTA--
CTACGTAC	0.47738	--GTACGTAG-
CTACGTAG	0.47560	--CTACGTAG-

8 mer Z-scores for probeset 'all'

Forward:	Reverse:

Top 10	Scores	Alignment
TTACGTAA	34.79042	-TTACGTAA-
GTTACGTA	30.76147	--TACGTAAC
ATTACGTA	29.82187	--TACGTAAT
CTACGTAC	29.51934	-GTACGTAG-
CACGTACG	29.17913	CGTACGTG--
CGTACGTA	27.72453	CGTACGTA--
GCTACGTA	27.31137	GCTACGTA--
GTACGTAA	26.90148	-GTACGTAA-
TACGTAAA	26.90148	--TACGTAAA
CCTACGTA	26.89397	CCTACGTA--