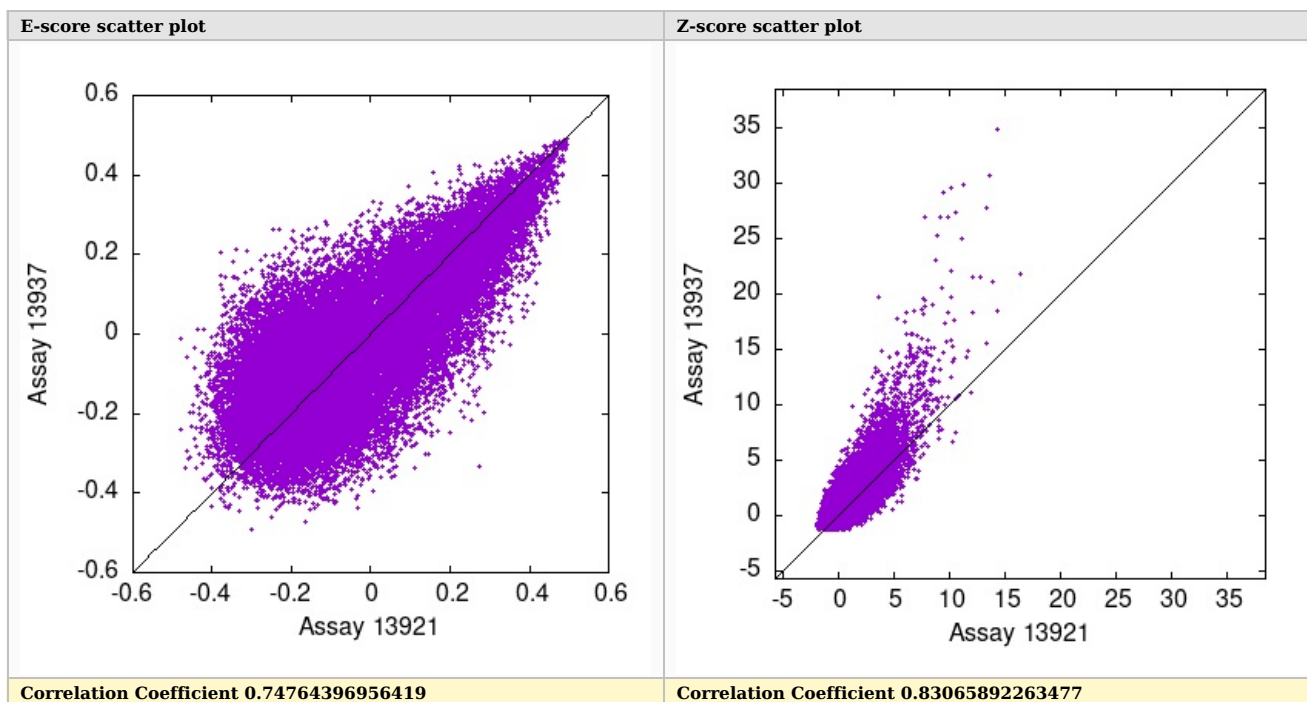


QC report



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'			8 mer Z-scores for probeset 'all'		
Forward:		Reverse:	Forward:		Reverse:
Top 10	Scores	Alignment	Top 10	Scores	Alignment
ACGTACGT	0.49608	ACGTACGT---	ACGTACGT	16.41565	---ACGTACGT---
TTACGTAA	0.49435	--TTACGTAA-	TACGCGTA	14.30965	-----TACGCGTA
TACGCGTA	0.49220	-TACGCGTA--	TTACGTAA	14.29846	-TTACGTAA-----
CGTACGTA	0.49145	-CGTACGTA--	GCGTACGC	13.93642	---GCGTACGC---
GTTACGTA	0.49071	---TACGTAAC	GTTACGTA	13.60979	GTTACGTA-----
GCGTACGC	0.48936	GCGTACGC---	GCGGCGCG	13.39153	---GCGGCGCG---
CGCGCGCG	0.48725	-CGCGCGCG--	CGTACGTA	13.36270	--TACGTACG----
ATTACGTA	0.48598	---TACGTAAT	ACGTACGC	12.78945	---ACGTACGC---
GCGATCGC	0.48528	GCGATCGC---	ACGATCGT	12.12672	---ACGATCGT---
CGTATACG	0.48504	-CGTATACG--	GCGGTACG	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'			8 mer Z-scores for probeset 'all'		
Forward:		Reverse:	Forward:		Reverse:
Top 10	Scores	Alignment	Top 10	Scores	Alignment
TTACGTAA	0.49053	--TTACGTAA-	TTACGTAA	34.79042	-TTACGTAA-
ATTACGTA	0.48879	---TACGTAAT	GTTACGTA	30.76147	--TACGTAAC
GTTACGTA	0.48481	---TACGTAAC	ATTACGTA	29.82187	--TACGTAAT
GTACGTAA	0.48363	--GTACGTAA-	CTACGTAC	29.51934	-GTACGTAG-
CACGTACG	0.48237	-CGTACGTG--	CACGTACG	29.17913	CGTACGTG--
TACGTAAA	0.48182	---TACGTAAA	CGTACGTA	27.72453	CGTACGTA--
GCGATCGC	0.48166	GCGATCGC---	GCTACGTA	27.31137	GCTACGTA--
CGTACGTA	0.47772	-CGTACGTA--	GTACGTAA	26.90148	-GTACGTAA-
CTACGTAC	0.47738	--GTACGTAG-	TACGTAAA	26.90148	--TACGTAAA
CTACGTAG	0.47560	--CTACGTAG-	CCTACGTA	26.89397	CCTACGTA--