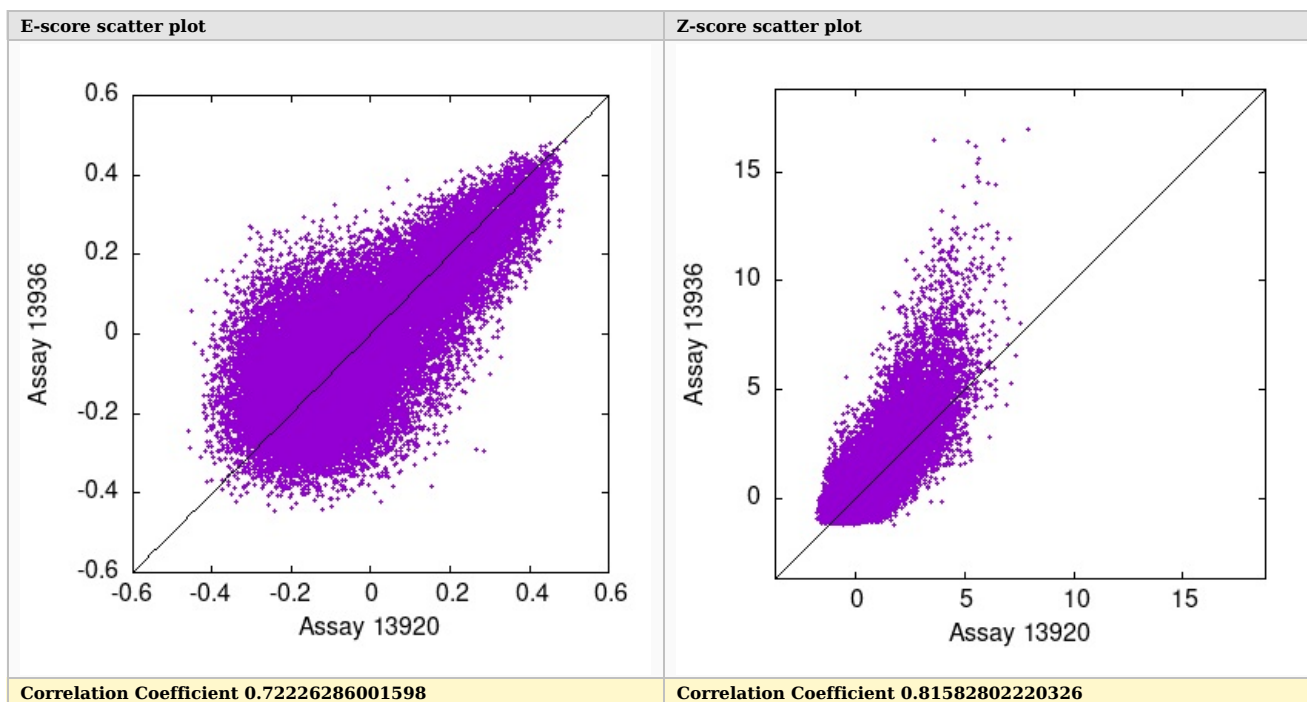




QC report





Top scoring motifs for Assay 13920

Protein ID: pTH14220.1 Gene: CXXC4 Domain: zf-CXXC Flag: Pass matched_pair Array: 1M-ME

8 mer E-scores for probeset 'all'

Forward:		Reverse:
		
Top 10	Scores	Alignment
ACGTACGT	0.49282	----ACGTACGT--
CGCGCGCG	0.48515	--CGCGCGCG---
CACGTGCG	0.47790	---CACGTGCG---
CGACGTCG	0.47636	--CGACGTCG----
ACGTCGCG	0.47485	CGCGACGT-----
CGTCGCGC	0.47440	-----GCGCGACG
ACGTGCGC	0.47427	-----ACGTGCGC--
ACGCGTTA	0.47383	---ACGCGTTA--
CGTAACGC	0.47382	-----CGTAACGC--
ACGCACGC	0.47308	-----GCGTGCCT--



8 mer Z-scores for probeset 'all'

Forward:		Reverse:
		
Top 10	Scores	Alignment
ACGTACGT	7.94023	---ACGTACGT--
CACGTGCG	7.55875	---CACGTGCG---
ACGCACGC	7.32779	---GCGTGCGT---
CGCGCGCG	7.11353	---CGCGCGCG---
ACGTGACG	7.04952	CGTCACGT-----
GCACGTGC	7.01928	--GCACGTGC----
CGACGTCG	7.01769	--CGACGTCG----
CGTCGCGC	6.92845	-----GCGCGACG
CGTACGTC	6.86325	---GACGTACG---
CGTAACGC	6.85282	-----CGTAACGC--



Top scoring motifs for Assay 13936

Protein ID: pTH14220.2 Gene: CXXC4 Domain: zf-CXXC Flag: Pass matched_pair Array: 1M-HK

8 mer E-scores for probeset 'all'

Forward:		Reverse:
		
Top 10	Scores	Alignment
ACGTACGT	0.48399	--ACGTACGT---
CACGTACG	0.48061	-CACGTACG----
ACGTTGCG	0.47242	--ACGTTGCG---
ACGCAACG	0.47230	---ACGCAACG---
ACGACGTC	0.46311	-----GACGTCGT
CTACGTAC	0.46180	CTACGTAC-----
GCGTACGC	0.46094	---GCGTACGC---
CGCGTACG	0.45588	-CGCGTACG----
ACGTAAAT	0.45489	---ACGTAAAT---
GTACGCAA	0.45481	-----GTACGCAA--

8 mer Z-scores for probeset 'all'

Forward:		Reverse:
		
Top 10	Scores	Alignment
ACGTACGT	16.91192	----ACGTACGT
ACGACGTC	16.48418	---ACGACGTC---
CGACGTCC	16.46381	---CGACGTCC---
CTACGTAC	16.37540	---CTACGTAC---
ACGCAACG	16.20114	ACGCAACG----
CACGTACG	15.63391	----CACGTACG--
ACGTTGCG	15.37336	---CGCAACGT----
ACGTCGTA	14.74815	---TACGACGT---
ACGCGACG	14.52302	ACGCGACG----
CGTGACGC	14.46683	---CGTGACGC---