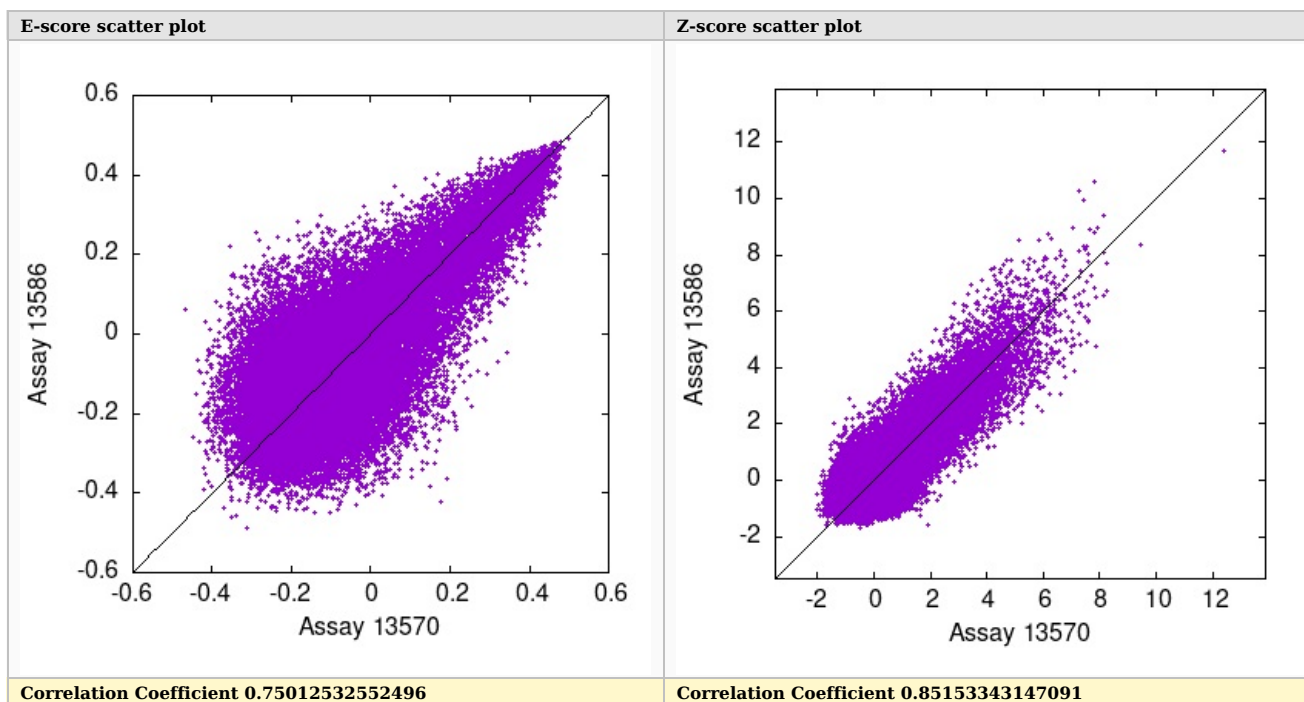


QC report



Top scoring motifs for Assay 13570

Protein ID: pTH14221.1 Gene: zf-CXXC4.DBD Domain: zf-CXXC Flag: Pass_matched_pair Array: 1M-ME

8 mer E-scores for probeset 'all'

Forward:

Reverse:

?

8 mer Z-scores for probeset 'all'

Forward:

Reverse:

Top 10	Scores	Alignment	Top 10	Scores	Alignment
ACGTACGT	0.49725	----ACGTACGT	ACGTACGT	12.43537	---ACGTACGT-
ACGTCACG	0.48667	CGTGACGT----	ACGTCACG	9.46349	---ACGTCACG-
ACGACGTC	0.47953	---ACGACGTC---	CACGTGCG	8.27013	CGCACGTG----
CACGTACG	0.47843	---CACGTACG---	ACGTAACG	8.25909	---ACGTAACG-
ACGTGACG	0.47790	CGTCACGT----	ACGTAGCG	8.14898	---ACGTAGCG-
CACGTGCG	0.47761	---CACGTGCG-	TGACGTCA	8.14374	-TGACGTCA---
ACGCACGC	0.47633	---ACGCACGC	ACGTGACG	8.12515	---ACGTGACG-
CGACGTCG	0.47628	--CGACGTCG--	GCGTACGC	7.90629	---GCGTACGC-
AACGCGTT	0.47626	-AACGCGTT---	ACGCACGA	7.88820	---ACGCACGA-
AACGTACG	0.47614	---AACGTACG-	CGTGCGCA	7.86670	---CGTGCGCA

Top scoring motifs for Assay 13586

Protein ID: pTH14221.2 Gene: zf-CXXC4.DBD Domain: zf-CXXC Flag: Pass_matched_pair Array: 1M-HK

8 mer E-scores for probeset 'all'

Forward:

Reverse:

Forward:

Reverse:

?

Top 10	Scores	Alignment	Top 10	Scores	Alignment
ACGTACGT	0.48971	-ACGTACGT--	ACGTACGT	11.69088	----ACGTACGT
CACGTACG	0.48196	CACGTACG---	CACGTACG	10.59068	---CACGTACG-
GCGTACGC	0.48126	---GCGTACGC--	ACGCACGT	10.29530	---ACGCACGT
ACGTACGC	0.48016	-ACGTACGC--	GCACGTGC	9.97501	--GCACGTGC--
ACGCACGT	0.47903	-ACGCACGT--	ACGTGACG	9.40951	CGTCACGT----
CGTACGTA	0.47701	--CGTACGTA-	GCGTACGC	8.97791	---GCGTACGC
GCACGTGC	0.47671	---GCACGTGC	ACGCAACG	8.96734	---ACGCAACG
ACGTAACG	0.47568	-ACGTAACG--	ACGTACGC	8.90466	---ACGTACGC
ACGCAACG	0.47478	-ACGCAACG--	ACGACGTC	8.78493	-ACGACGTC---
GTACGTAC	0.47388	---GTACGTAC	CTACGTAG	8.74858	--CTACGTAG--