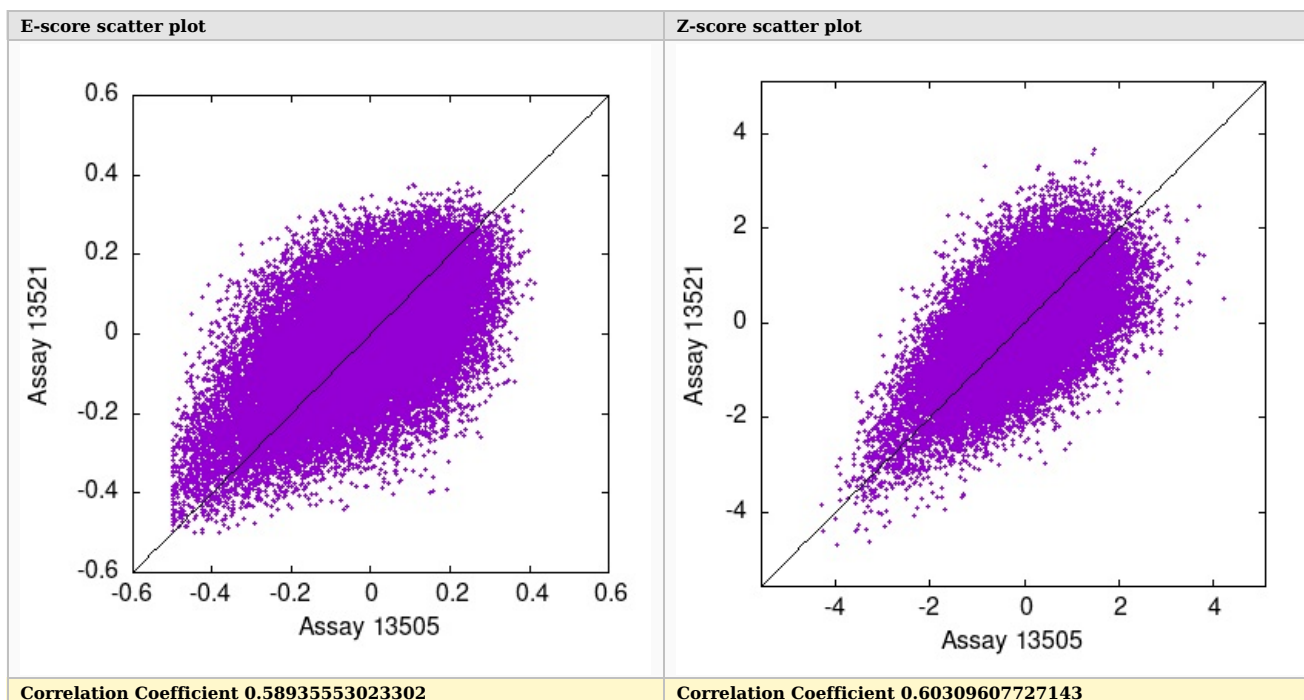


QC report



Top scoring motifs for Assay 13505

Protein ID: pTH13950.1 Gene: AKAP8.DBD Domain: zf-C2H2 Flag: Reject 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
ATACAAAA	0.41519	---ATACAAAA	ATACAAAA	4.21983	----ATACAAAA
TAACGAAA	0.40401	-TAACGAAA--	ATGTATAA	3.80391	--TTATACAT--
CATAATAC	0.40147	--CATAATAC-	ACAAACAT	3.71843	--ACAAACAT--
ATGTATAA	0.39679	-ATGTATAA--	AACATACA	3.69719	-AACATACA---
ATTTATAA	0.39401	-ATTTATAA--	ATAATATA	3.66692	-ATAATATA---
ACAAATAA	0.38562	-ACAAATAA--	ATTTATAA	3.51304	--ATTTATAA--
ACAAACAT	0.38289	-ACAAACAT--	TAACGAAA	3.51252	TAACGAAA----
ATAATGCA	0.38133	ATAATGCA---	ATAATGCA	3.48686	-ATAATGCA---
ACATACAA	0.37976	-ACATACAA--	ACATACAA	3.45821	--ACATACAA--
TACATACA	0.37832	TACATACA---	AAATTATT	3.36110	-AAATTATT---

Top scoring motifs for Assay 13521

Protein ID: pTH13950.2 Gene: AKAP8.DBD Domain: zf-C2H2 Flag: Reject Array: 1M-HK

8 mer E-scores for probeset 'all'

Forward:

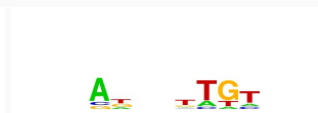
Reverse:



8 mer Z-scores for probeset 'all'

Forward:

Reverse:



Top 10	Scores	Alignment	Top 10	Scores	Alignment
ATACGTTA	0.38022	-ATACGTTA----	ACAACGAT	3.66777	--ACAACGAT-
ATGCGCAT	0.37600	-ATGCGCAT----	ACATTACT	3.58169	--ACATTACT-
TACTAGTA	0.37331	--TACTAGTA--	TACTAGTA	3.42066	-TACTAGTA--
CAACAATG	0.36870	-CAACAATG----	AACGTTGA	3.41134	--TCAACGTT-
ACATTACT	0.36664	---ACATTACT--	AGTATGTG	3.38288	---CACATACT
ATACATAA	0.36392	-ATACATAA----	ATGCGCAT	3.33451	--ATGCGCAT-
ACAACGAT	0.36173	ACAACGAT-----	CAGCGCTG	3.30292	---CAGCGCTG
TAACGTAA	0.36107	-TTACGTTA----	GCGCAAAA	3.24371	GCGCAAAA----
GTTGCAAC	0.35599	-----GTTGCAAC	AAACGCGC	3.16900	---AAACGCGC
ACTTGACA	0.35419	---ACTTGACA--	ATACGTTA	3.15481	---ATACGTTA