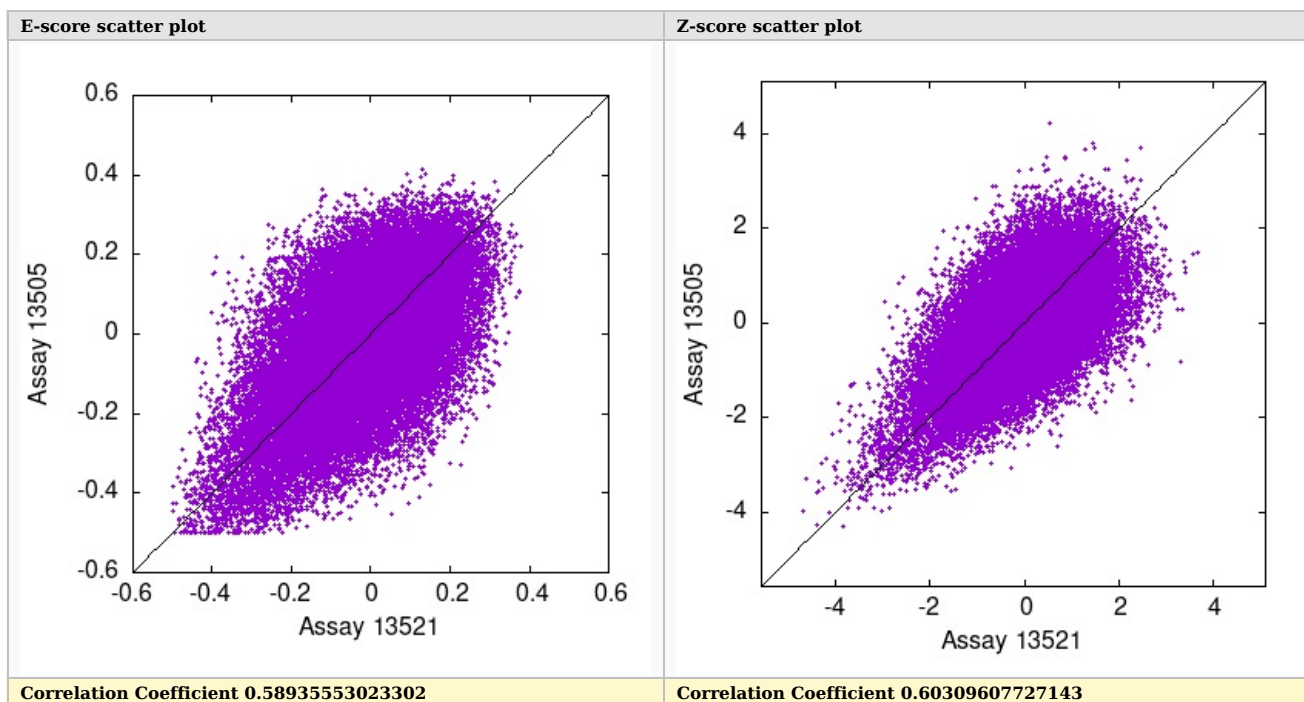


QC report



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:



Top 10

Scores

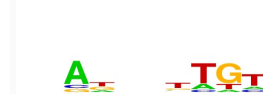
Alignment

ATACGTTA	0.38022	-ATACGTTA----
ATGCGCAT	0.37600	-ATGCGCAT----
TACTAGTA	0.37331	--TACTAGTA---
CAACAATG	0.36870	-CAACAATG----
ACATTACT	0.36664	---ACATTACT--
ATACATAA	0.36392	-ATACATAA----
ACAACGAT	0.36173	ACAACGAT-----
TACGTTA	0.36107	-TACGTTA-----
GTTGCAAC	0.35599	-----GTTGCAAC
ACTTGACA	0.35419	---ACTTGACA--

8 mer Z-scores for probeset 'all'

Forward:

Reverse:



Top 10

Scores

Alignment

ACAACGAT	3.66777	--ACAACGAT--
ACATTACT	3.58169	--ACATTACT--
TACTAGTA	3.42066	-TACTAGTA--
AACGTTGA	3.41134	--TCAACGTT--
AGTATGTG	3.38288	---CACATACT
ATGCGCAT	3.33451	--ATGCGCAT--
CAGCGCTG	3.30292	---CAGCGCTG
GCGCAAAA	3.24371	GCGCAAAA---
AAACGCGC	3.16900	---AAACGCGC
ATACGTTA	3.15481	---ATACGTTA

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:



Top 10

Scores

Alignment

ATACAAAA	0.41519	---ATACAAAA
TAACGAAA	0.40401	--TAACGAAA--
CATAATAC	0.40147	--CATAATAC--
ATGTATAA	0.39679	-ATGTATAA--
ATTTATAA	0.39401	-ATTTATAA--
ACAAATAA	0.38562	-ACAAATAA--
ACAAACAT	0.38289	-ACAAACAT--
ATAATGCA	0.38133	ATAATGCA---
ACATACAA	0.37976	-ACATACAA--
TACATACA	0.37832	TACATACA---

8 mer Z-scores for probeset 'all'

Forward:

Reverse:



Top 10

Scores

Alignment

ATACAAAA	4.21983	---ATACAAAA
ATGTATAA	3.80391	--TTATACAT--
ACAAACAT	3.71843	--ACAAACAT--
AACATACA	3.69719	-AACATACA--
ATAATATA	3.66692	-ATAATATA--
ATTTATAA	3.51304	--ATTTATAA--
TAACGAAA	3.51252	TAACGAAA----
ATAATGCA	3.48686	-ATAATGCA---
ACATACAA	3.45821	--ACATACAA--
AAATTATT	3.36110	-AAATTATT---