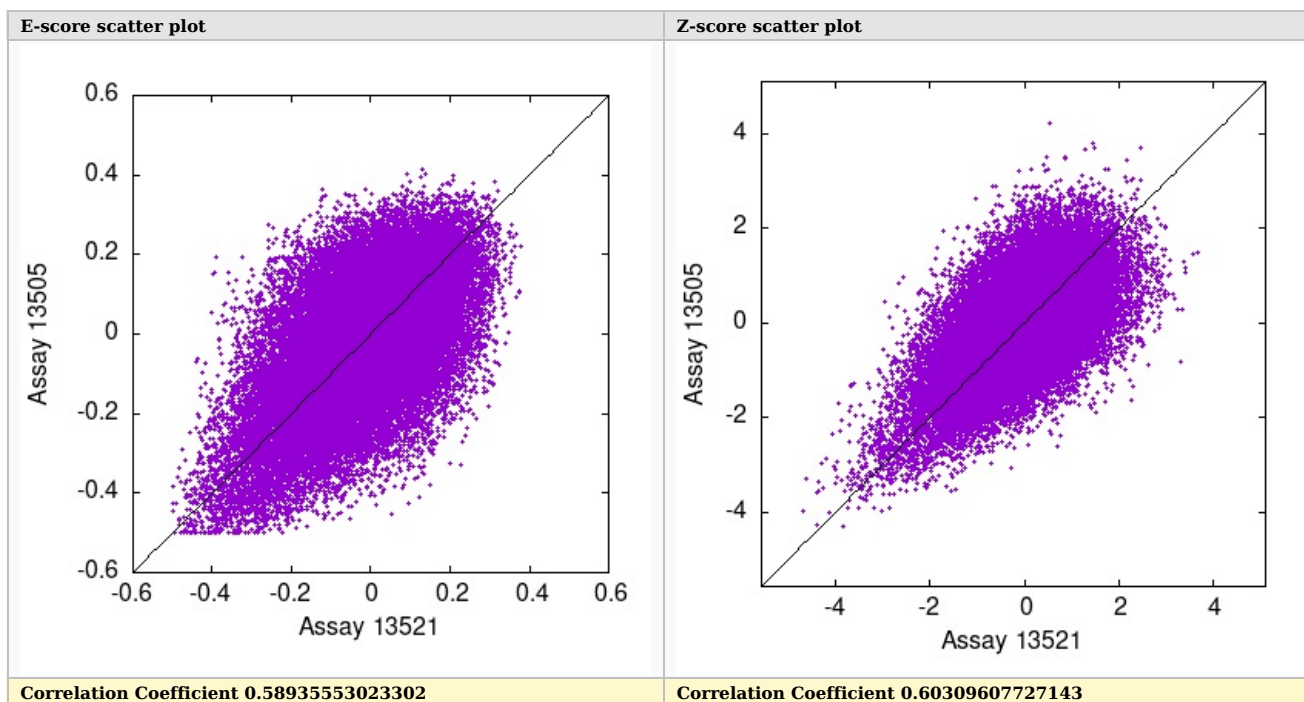




QC report



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-----
TACGTAA	0.36107	-TACGTAA-----
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---