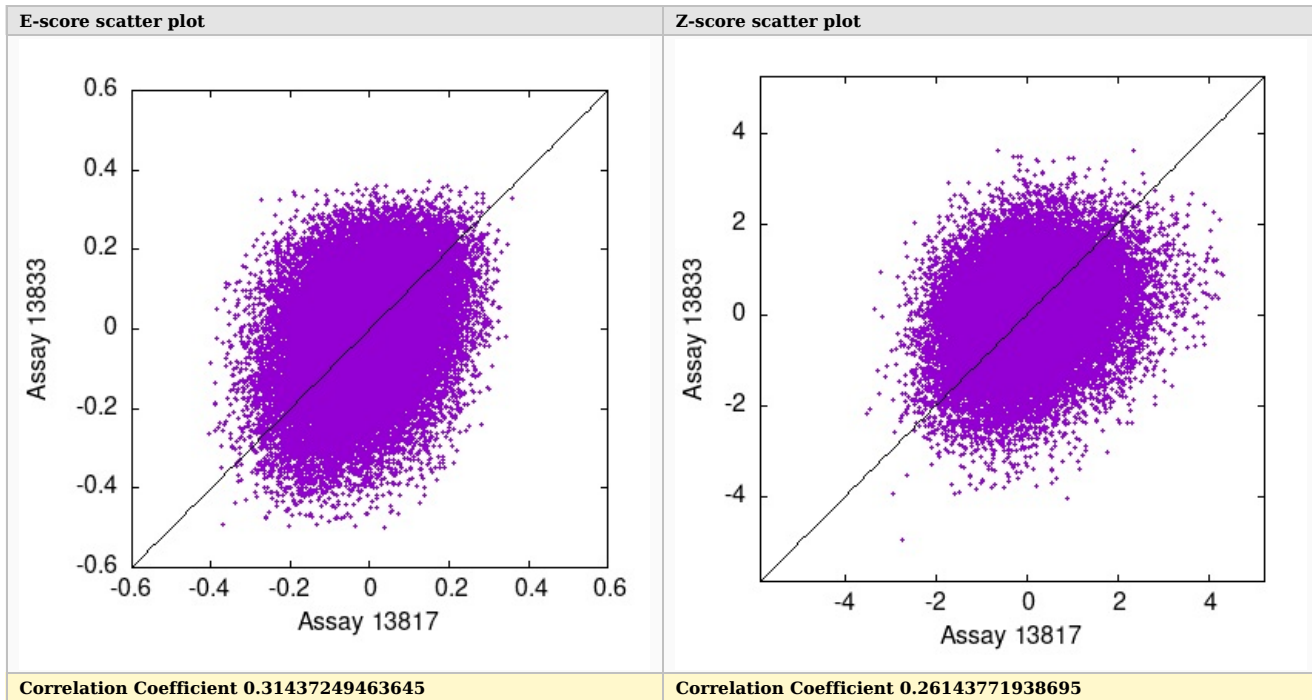


QC report



Top scoring motifs for Assay 13817

Protein ID: pTH13911.1 Gene: KDM5B.DBD Domain: ARID Flag: Reject Array: 1M-ME

8 mer E-scores for probeset 'all'			8 mer Z-scores for probeset 'all'		
Forward:		Reverse:	Forward:		Reverse:
Top 10	Scores	Alignment	Top 10	Scores	Alignment
ATACAAAA	0.35921	--ATACAAAA--	GAATATTC	4.31402	---GAATATTC
CGCGAATA	0.34854	TATTCGCG--	GTCGAATA	4.27925	GTCGAATA---
GACTAATA	0.34176	--GACTAATA	ACTACTAA	4.24660	--ACTACTAA--
ATATATAA	0.33164	--ATATATAA	CATATTAA	4.23189	--TTAATATG-
AACGTAAA	0.32639	--AACGTAAA	CGCGAATA	4.18441	CGCGAATA---
GATGTATA	0.32574	TATACATC--	AAATAGTC	4.17357	---AAATAGTC
GTA CTCTA	0.32486	--TAGAGTAC	GTA CTCTA	4.14316	--TAGAGTAC--
GTGCGCAC	0.32460	-GTGCGCAC-	TACAAACA	4.12471	TACAAACA---
TACAAACA	0.32429	--TACAAACA	GAAGTGTA	4.10976	GAAGTGTA---
TAAAAAAA	0.32288	--TAAAAAAA	CGTAAAAA	4.10851	--CGTAAAAA-

Top scoring motifs for Assay 13833

Protein ID: pTH13911.2 Gene: KDM5B.DBD Domain: ARID Flag: Reject Array: 1M-HK

8 mer E-scores for probeset 'all'			8 mer Z-scores for probeset 'all'		
Forward:		Reverse:	Forward:		Reverse:
Top 10	Scores	Alignment	Top 10	Scores	Alignment
AGATATCT	0.37174	--AGATATCT--	ATAAGATA	3.61234	---ATAAGATA-
CACATATA	0.36489	--CACATATA----	ACTATAAT	3.60608	---ATTATAGT-
ATAGTTTA	0.35868	----ATAGTTTA-	AAATAAAT	3.48988	---AAATAAAT--
AAATCGAT	0.35868	--AAATCGAT--	AATAAATT	3.48988	---AATAAATT-
ACAAAATA	0.35819	-----TATTTTGT	ACATAGAC	3.47886	--ACATAGAC--
TAAGATAA	0.35716	----TTATCTTA-	ATAGTTTA	3.45387	----ATAGTTTA
ACAAATTA	0.35492	--ACAAATTA---	TAAGATAA	3.44840	TAAGATAA----
ATACATAT	0.35417	ATACATAT----	ATAACGAT	3.40689	--ATAACGAT--
ATAAGATA	0.35283	-----TATCTTAT	ATCATGAT	3.37403	--ATCATGAT--
ACATATCT	0.35278	--ACATATCT---	AAATCGAT	3.36970	--AAATCGAT--