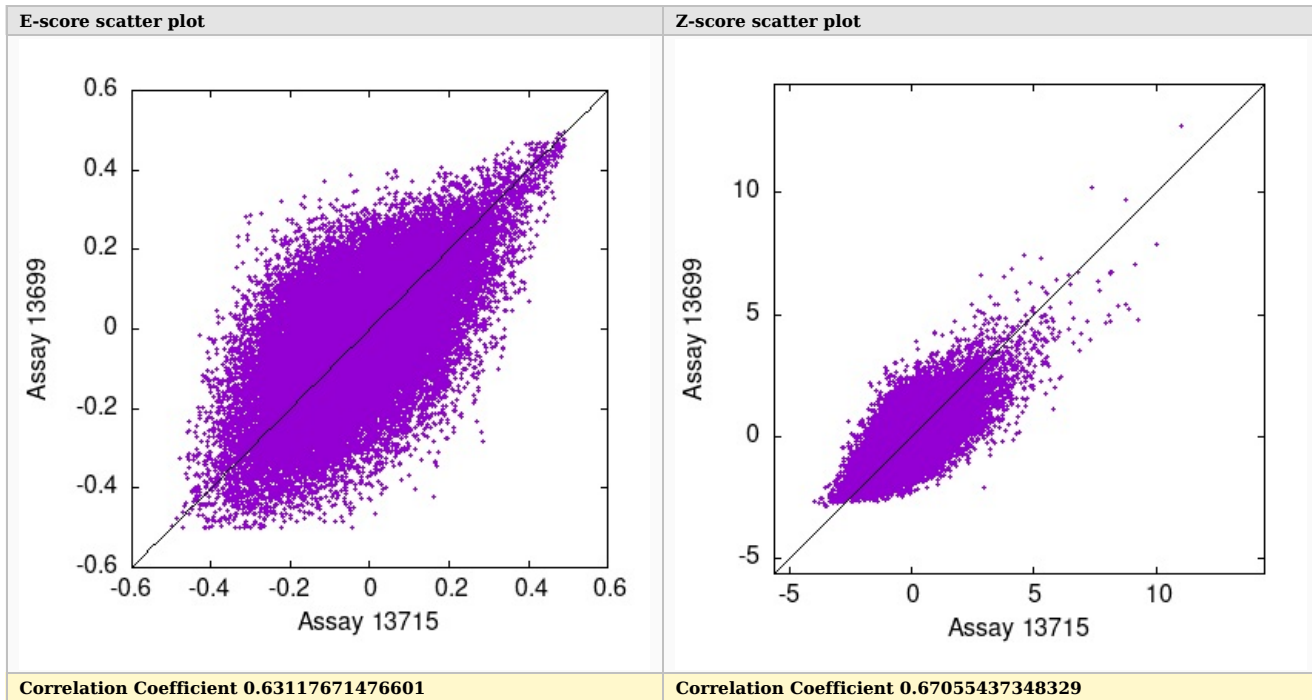


QC report



Top scoring motifs for Assay 13715

Protein ID: pTH14212.2 Gene: TIGD3.DBD Domain: CENP-B_N Flag: Pass_matched_pair 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
TATGCATA	0.49246	TATGCATA--	TATGCATA	11.02486	TATGCATA--
ATTATGCA	0.49228	--TGCATAAT	ATTATGCA	10.03849	--TGCATAAT
GTTATTCA	0.49115	--TGAATAAC	GTTATTCA	9.24222	--TGAATAAC
GTTATGCA	0.48814	--TGCATAAC	GTTATGCA	9.10610	--TGCATAAC
GTGAATAA	0.48596	-GTGAATAA-	CATGAATA	8.87183	CATGAATA--
CATGAATA	0.48471	CATGAATA--	TTATGCAA	8.75568	-TTGCATAA-
CTGAATAA	0.48398	-CTGAATAA-	ATGCATAA	8.74709	-ATGCATAA-
ATGCATAA	0.48186	-ATGCATAA-	GTGAATAA	8.43279	-GTGAATAA-
ACTGAATA	0.48099	ACTGAATA--	ATTATTCA	8.16283	--TGAATAAT
TTATGCAA	0.47977	-TTGCATAA-	TATTCACA	8.14408	TATTCACA--

Top scoring motifs for Assay 13699

Protein ID: pTH14212.1 Gene: TIGD3.DBD Domain: CENP-B_N Flag: Pass_matched_pair 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
TATGCATA	0.49540	TATGCATA---	TATGCATA	12.74771	---TATGCATA
GAATATTC	0.49119	---GAATATTC	GAATATTC	10.17627	GAATATTC---
ATGCATAA	0.48664	-ATGCATAA--	ATGCATAA	9.69973	--TTATGCAT-
AGTGAATA	0.48134	AGTGAATA---	ATTATGCA	7.84913	-ATTATGCA--
ATTATGCA	0.47668	--TGCATAAT-	TATATTCA	7.46007	-TATATTCA--
GTTATGCA	0.47546	--TGCATAAC-	TATATGCA	7.32461	-TATATGCA--
GCATATGC	0.46845	---GCATATGC	GTTATGCA	7.02562	-GTTATGCA--
TATATTCA	0.46806	--TGAATATA-	ATATGCAC	6.85615	--ATATGCAC-
TATTCACA	0.46801	-TATTCACA--	GCATATGC	6.76697	GCATATGC---
ATGCGCAT	0.46731	-ATGCGCAT--	ATTATTCA	6.72857	-ATTATTCA--