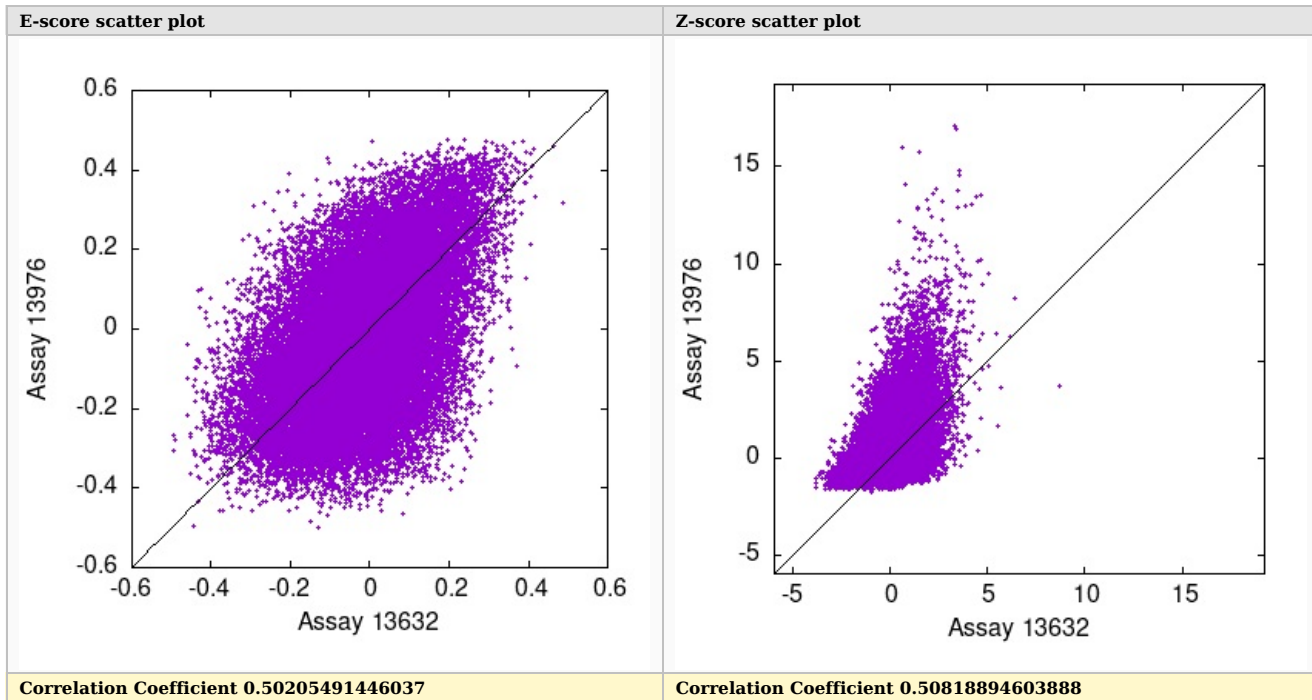


QC report



Top scoring motifs for Assay 13632

Protein ID: pTH14237.1 Gene: HMG20A.DBD Domain: HMG_box 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
ATATATAT	0.48561	-ATATATAT-	ATATATAT	8.68579	-ATATATAT-
TATATATA	0.46359	--TATATATA	TATATATA	6.38855	--TATATATA
ATACACAT	0.41870	-ATACACAT-	ACATATAT	6.20349	-ACATATAT-
ACATATAT	0.41817	-ACATATAT-	ATACACAT	5.73987	-ATACACAT-
GTACTATA	0.41213	GTACTATA--	ATATGGAT	5.54216	-ATATGGAT-
ATAATATA	0.41132	ATAATATA--	TACATATA	5.48360	--TACATATA
TATACATA	0.40523	--TATACATA	ATACGATA	5.04404	-ATACGATA-
TACATATA	0.40442	--TACATATA	ATATATTA	5.03339	-ATATATTA-
ATATGGAT	0.40403	-ATATGGAT-	AGATATCT	4.79274	-AGATATCT-
CATATATA	0.39479	CATATATA--	ATAATATA	4.77013	ATAATATA--

Top scoring motifs for Assay 13976

Protein ID: pTH14237.3 Gene: HMG20A.DBD Domain: HMG_box 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
TTATATAA	0.47701	-TTATATAA-	TTATATAA	17.13004	-TTATATAA-
ACTATAAT	0.47609	--ACTATAAT	CTTATATA	16.96272	CTTATATA--
CTCTTATA	0.47322	CTCTTATA--	CTCTTATA	15.94716	CTCTTATA--
CTTATATA	0.47067	CTTATATA--	ACTATAAT	15.77200	--ACTATAAT
ACTATATA	0.47039	ACTATATA--	TATATACA	14.83302	--TATATACA
TACTATAA	0.46912	-TACTATAA-	ACTATATA	14.53085	ACTATATA--
TATATACA	0.46698	--TATATACA	TATTATAA	14.10460	-TATTATAA-
GTTATATA	0.46399	GTTATATA--	GTTATATA	13.85595	GTTATATA--
TATATATA	0.46241	--TATATATA	ATATATAG	13.77679	-ATATATAG-
ACTATGTA	0.46054	ACTATGTA--	CTATATTA	13.62514	-CTATATTA-