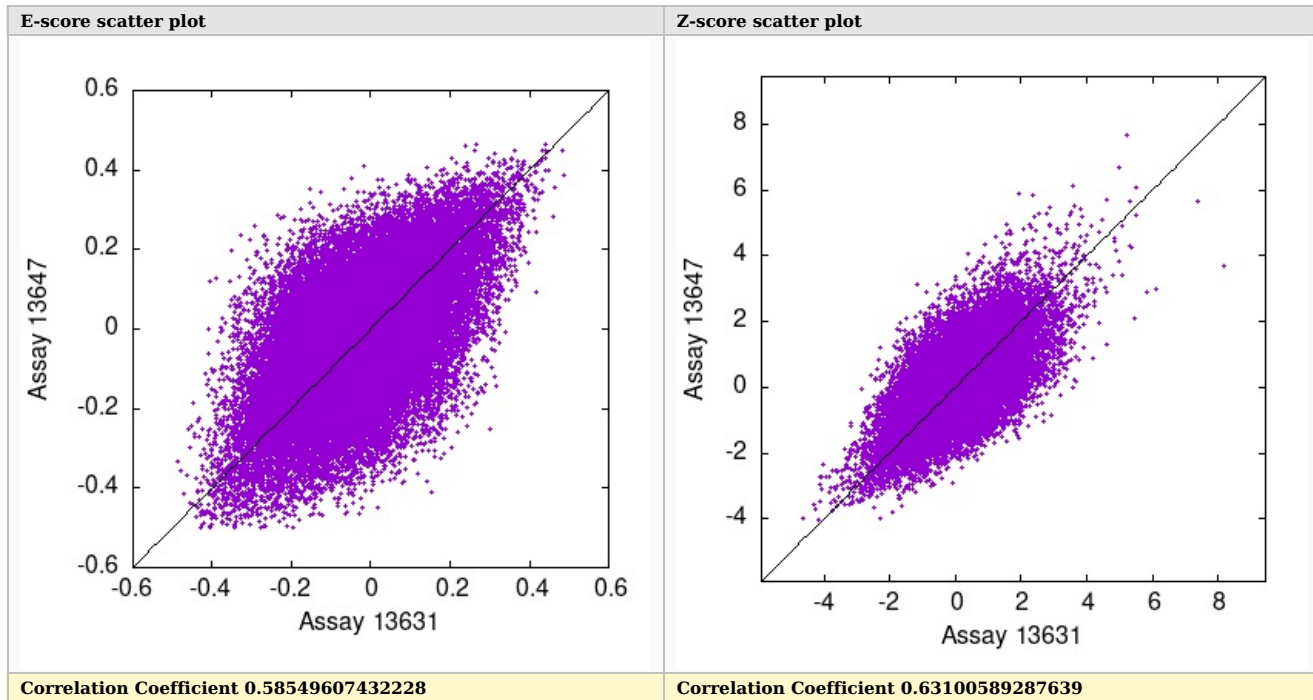


QC report



Top scoring motifs for Assay 13631

Protein ID: pTH14236.1 Gene: HMG20A.FL 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	0.48920	-ATATATAT-	ATATATAT	8.16685	-ATATATAT
TATATATA	0.48363	--TATATATA	TATATATA	7.40148	TATATATA-
ATAATATA	0.46321	ATAATATA--	ATAATATA	6.10451	ATAATATA-
ATTATATA	0.46108	ATTATATA--	ATTATATA	5.82810	ATTATATA-
TTATATAA	0.44932	-TTATATAA-	TATACATA	5.50008	TATACATA-
CTATATAG	0.44540	-CTATATAG-	ACATATAT	5.48242	-ACATATAT
ATATAATA	0.44197	-ATATAATA-	TGTATACA	5.44447	TGTATACA-
ACATATAT	0.44020	-ACATATAT-	ATATAATA	5.37069	-ATATAATA
CATATATG	0.43996	--CATATATG	CTATATAG	5.31455	-CTATATAG
TATACATA	0.43978	--TATACATA	TTATATAA	5.30903	-TTATATAA

Top scoring motifs for Assay 13647

Protein ID: pTH14236.2 Gene: HMG20A.FL 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
CATATATG	0.46528	CATATATG-	CATATATG	7.67344	-CATATATG-
ATGCGCAT	0.46429	-ATGCGCAT	TATGCATA	6.68412	-TATGCATA-
ATCTAGAT	0.46033	-ATCTAGAT	ATGCATAA	6.12691	--ATGCATAA
ACATATAT	0.45037	-ACATATAT	TATACATA	6.07157	-TATACATA-
TATGCATA	0.44906	TATGCATA-	ATGCGCAT	5.86720	--ATGCGCAT
TATCGATA	0.44891	TATCGATA-	TATCGATA	5.84034	-TATCGATA-
ATGCATAA	0.44752	-ATGCATAA	ATGTACAT	5.71924	ATGTACAT--
TATATATA	0.44738	TATATATA-	TATATATA	5.67751	-TATATATA-
CTATATAG	0.44509	-CTATATAG	CTATATAG	5.64479	--CTATATAG
ATATCGAT	0.44154	-ATATCGAT	TATTAATA	5.51782	-TATTAATA-