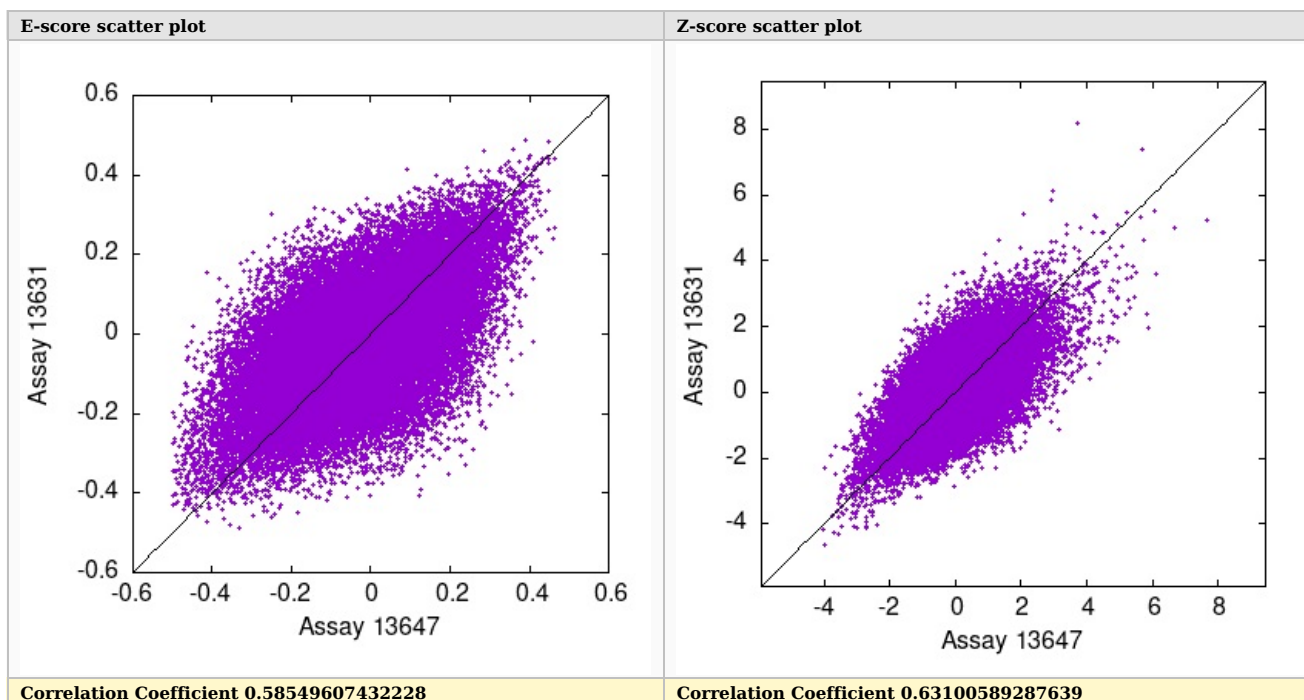


## QC report



### 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'

Forward:		Reverse:	
Top 10	Scores	Alignment	
CATATATG	0.46528	CATATATG-	
ATGCGCAT	0.46429	-ATGCGCAT	
ATCTAGAT	0.46033	-ATCTAGAT	
ACATATAT	0.45037	-ACATATAT	
TATGCATA	0.44906	TATGCATA-	
TATCGATA	0.44891	TATCGATA-	
ATGCATAA	0.44752	-ATGCATAA	
TATATATA	0.44738	TATATATA-	
CTATATAG	0.44509	-CTATATAG	
ATATCGAT	0.44154	-ATATCGAT	



#### 8 mer Z-scores for probeset 'all'

Forward:		Reverse:	
Top 10	Scores	Alignment	
CATATATG	7.67344	-CATATATG-	
TATGCATA	6.68412	-TATGCATA-	
ATGCATAA	6.12691	--ATGCATAA	
TATACATA	6.07157	-TATACATA-	
ATGCGCAT	5.86720	--ATGCGCAT	
TATCGATA	5.84034	-TATCGATA-	
ATGTACAT	5.71924	ATGTACAT--	
TATATATA	5.67751	-TATATATA-	
CTATATAG	5.64479	--CTATATAG	
TATTAATA	5.51782	-TATTAATA-	

### 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'

Forward:	Reverse:	
		
Top 10	Scores	Alignment
ATATATAT	0.48920	-ATATATAT-
TATATATA	0.48363	--TATATATA
ATAATATA	0.46321	ATAATATA--
ATTATATA	0.46108	ATTATATA--
TTATATAA	0.44932	-TTATATAA-
CTATATAG	0.44540	-CTATATAG-
ATATAATA	0.44197	-ATATAATA-
ACATATAT	0.44020	-ACATATAT-
CATATATG	0.43996	--CATATATG
TATACATA	0.43978	--TATACATA

#### 8 mer Z-scores for probeset 'all'

Forward:		Reverse:	
Top 10	Scores	Alignment	
ATATATAT	8.16685	-ATATATAT	
TATATATA	7.40148	TATATATA-	
ATAATATA	6.10451	ATAATATA-	
ATTATATA	5.82810	ATTATATA-	
TATACATA	5.50008	TATACATA-	
ACATATAT	5.48242	-ACATATAT	
TGTATACA	5.44447	TGTATACA-	
ATATAATA	5.37069	-ATATAATA	
CTATATAG	5.31455	-CTATATAG	
TTATATAA	5.30903	-TTATATAA	