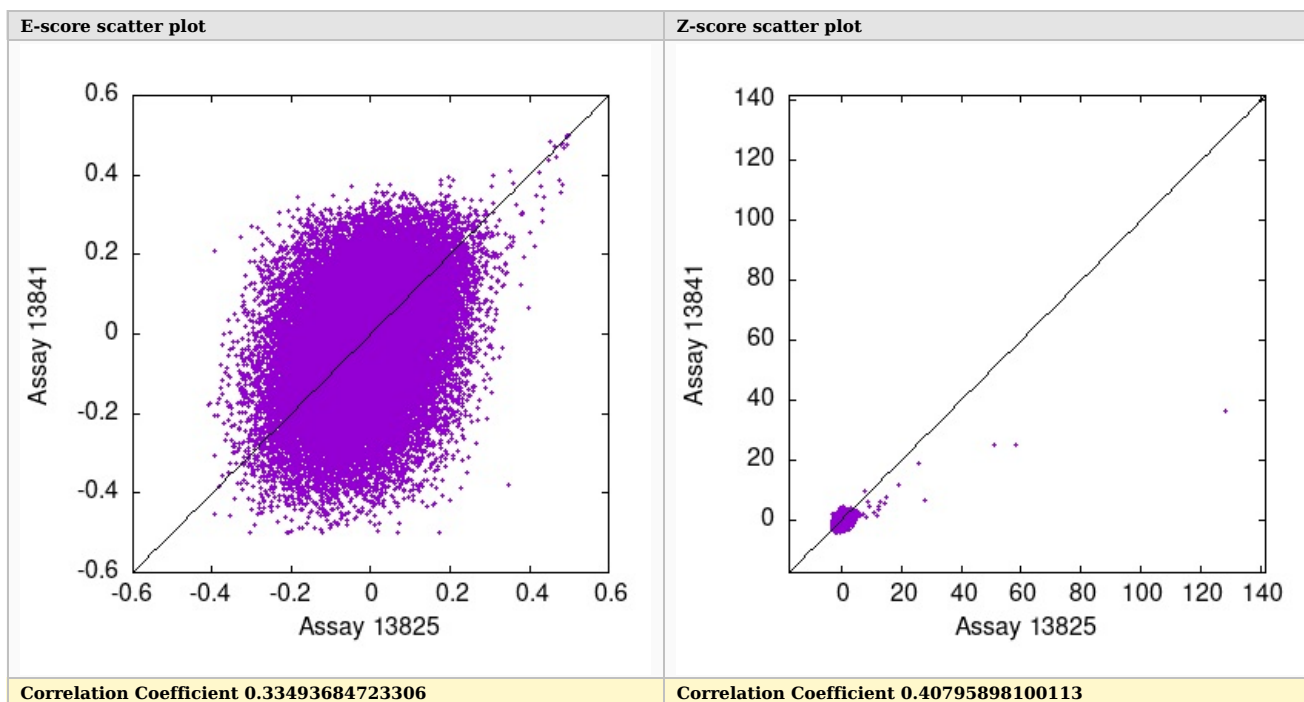




## QC report





### Top scoring motifs for Assay 13825

Protein ID: pTH13926.1 Gene: C11orf95 Domain: zf-BED Flag: Pass matched\_pair Array: 1M-ME

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

Forward:		Reverse:	
			
Top 10	Scores	Alignment	
GGGGCCCC	0.49913	-GGGGCCCC-	
GGGCCACA	0.49841	--GGGCCACA	
GGGGCCAC	0.49821	-GGGGCCAC-	
GGGCCCCA	0.49628	--GGGCCCCA	
AGGGCCAC	0.49533	-AGGGCCAC-	
AGGGGCCC	0.49362	AGGGGCCC--	
GGGCCCCC	0.48843	-GGGCCCCC-	
CGGGGCCC	0.48355	CGGGGCCC--	
GGGGCCAA	0.48173	-GGGGCCAA-	
AGGGGCCA	0.48114	AGGGGCCA--	



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

Forward:		Reverse:
		
Top 10	Scores	Alignment
GGGGCCCC	128.37896	-GGGGCCCC-
GGGGCCAC	58.33786	-GGGGCCAC-
GGGCCACA	51.19141	--GGGCCACA
GGGCCCCA	28.07789	--GGGCCCCA
AGGGCCAC	26.04493	-AGGGCCAC-
AGGGGCCC	19.37504	AGGGGCCC--
CGGGGCCC	14.87884	CGGGGCCC--
GGGCCCCC	14.60970	-GGGCCCCC-
AGGGCCCC	13.20249	-AGGGCCCC-
GGGGCCA	12.64959	-GGGGCCA-



### Top scoring motifs for Assay 13841

Protein ID: pTH13926.2 Gene: C11orf95 Domain: zf-BED Flag: Pass matched\_pair Array: 1M-HK

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

Forward:	Reverse:	
		
Top 10	Scores	Alignment
GGGGCCCC	0.49936	-GGGGCCCC-
GGGCCACA	0.49823	--GGGCCACA
GGGGCCAC	0.49802	-GGGGCCAC-
AGGGCCAC	0.49785	-AGGGCCAC-
AGGGGCCC	0.49355	AGGGGCCC--
CGTGGCCC	0.48567	CGTGGCCC--
CGGGGCCC	0.48150	CGGGGCCC--
GGGCCCCA	0.47790	--GGGCCCCA
AGGGCCCC	0.47696	-AGGGCCCC-
AGTGGCCC	0.47368	AGTGGCCC--

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

Forward:		Reverse:	
			
Top 10	Scores	Alignment	
GGGGCCCC	36.34755	-GGGGCCCC-	
GGGCCACA	25.22974	--GGGCCACA	
GGGGCCAC	25.22974	-GGGGCCAC-	
AGGGCCAC	18.91028	-AGGGCCAC-	
AGGGGCCC	11.88342	AGGGGCCC--	
CGTGGCCC	9.72461	CGTGGCCC--	
CGGGGCCC	7.54496	CGGGGCCC--	
GGGCCCCA	6.92087	--GGGCCCCA	
AGGGCCCC	6.41836	-AGGGCCCC-	
AGTGGCCC	6.22574	AGTGGCCC--	