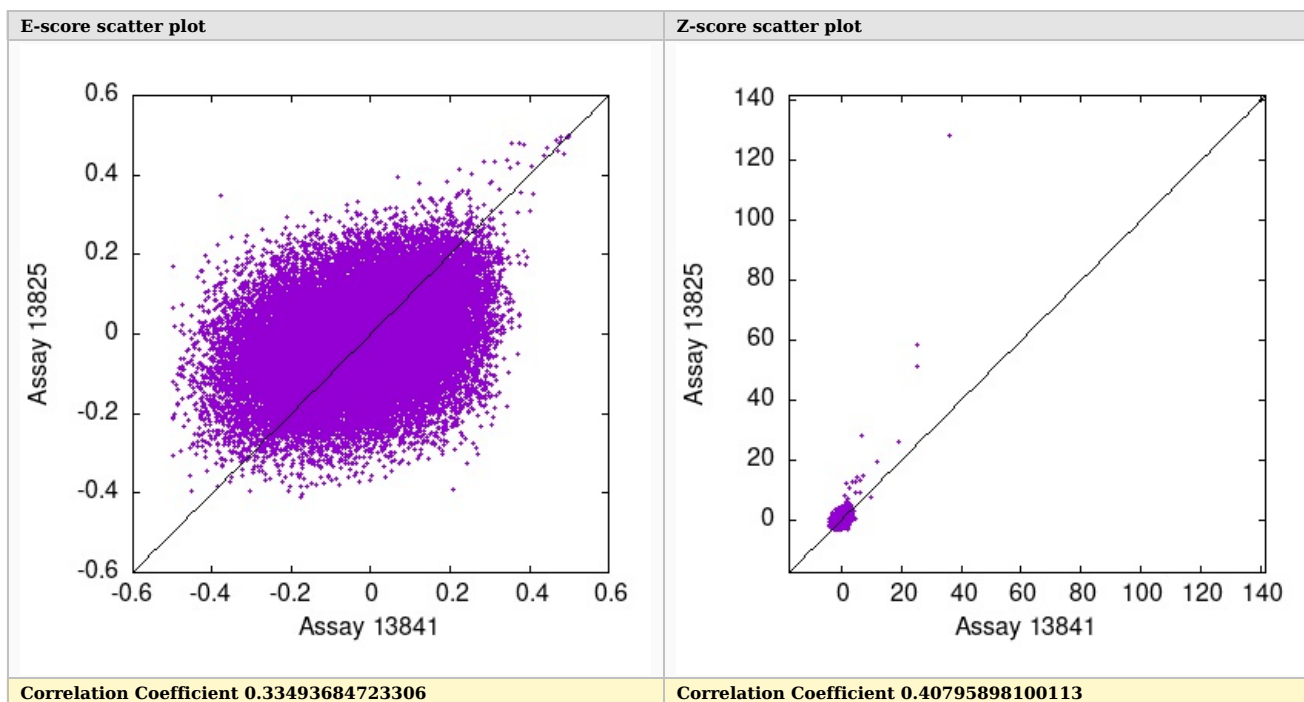


QC report



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'			8 mer Z-scores for probeset 'all'		
Forward:	Reverse:		Forward:	Reverse:	
Top 10	Scores	Alignment	Top 10	Scores	Alignment
GGGGCCCC	0.49936	-GGGGCCCC-	GGGGCCCC	36.34755	-GGGGCCCC-
GGGCCACA	0.49823	--GGGCCACA	GGGCCACA	25.22974	--GGGCCACA
GGGGCCAC	0.49802	-GGGGCCAC-	GGGGCCAC	25.22974	-GGGGCCAC-
AGGGCCAC	0.49785	-AGGGCCAC-	AGGGCCAC	18.91028	-AGGGCCAC-
AGGGGCCC	0.49355	AGGGGCCC--	AGGGGCCC	11.88342	AGGGGCCC--
CGTGGCCC	0.48567	CGTGGCCC--	CGTGGCCC	9.72461	CGTGGCCC--
CGGGGCCC	0.48150	CGGGGCCC--	CGGGGCCC	7.54496	CGGGGCCC--
GGGGCCCA	0.47790	--GGGGCCCA	GGGGCCCA	6.92087	--GGGGCCCA
AGGGCCCC	0.47696	-AGGGCCCC-	AGGGCCCC	6.41836	-AGGGCCCC-
AGTGGCCC	0.47368	AGTGGCCC--	AGTGGCCC	6.22574	AGTGGCCC--

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'			8 mer Z-scores for probeset 'all'		
Forward:	Reverse:		Forward:	Reverse:	
Top 10	Scores	Alignment	Top 10	Scores	Alignment
GGGGCCCC	0.49913	-GGGGCCCC-	GGGGCCCC	128.37896	-GGGGCCCC-
GGGCCACA	0.49841	--GGGCCACA	GGGGCCAC	58.33786	-GGGGCCAC-
GGGGCCAC	0.49821	-GGGGCCAC-	GGGCCACA	51.19141	--GGGCCACA
GGGGCCCA	0.49628	--GGGGCCCA	GGGGCCCA	28.07789	--GGGGCCCA
AGGGCCAC	0.49533	-AGGGCCAC-	AGGGCCAC	26.04493	-AGGGCCAC-
AGGGGCCC	0.49362	AGGGGCCC--	AGGGGCCC	19.37504	AGGGGCCC--
GGGGCCCC	0.48843	-GGGGCCCC-	CGGGGCCC	14.87884	CGGGGCCC--
CGGGGCCC	0.48355	CGGGGCCC--	GGGGCCCC	14.60970	-GGGGCCCC-
GGGGCCAA	0.48173	-GGGGCCAA-	AGGGCCCC	13.20249	-AGGGCCCC-
AGGGGCCA	0.48114	AGGGGCCA--	GGGGCCCA	12.64959	-GGGGCCCA-