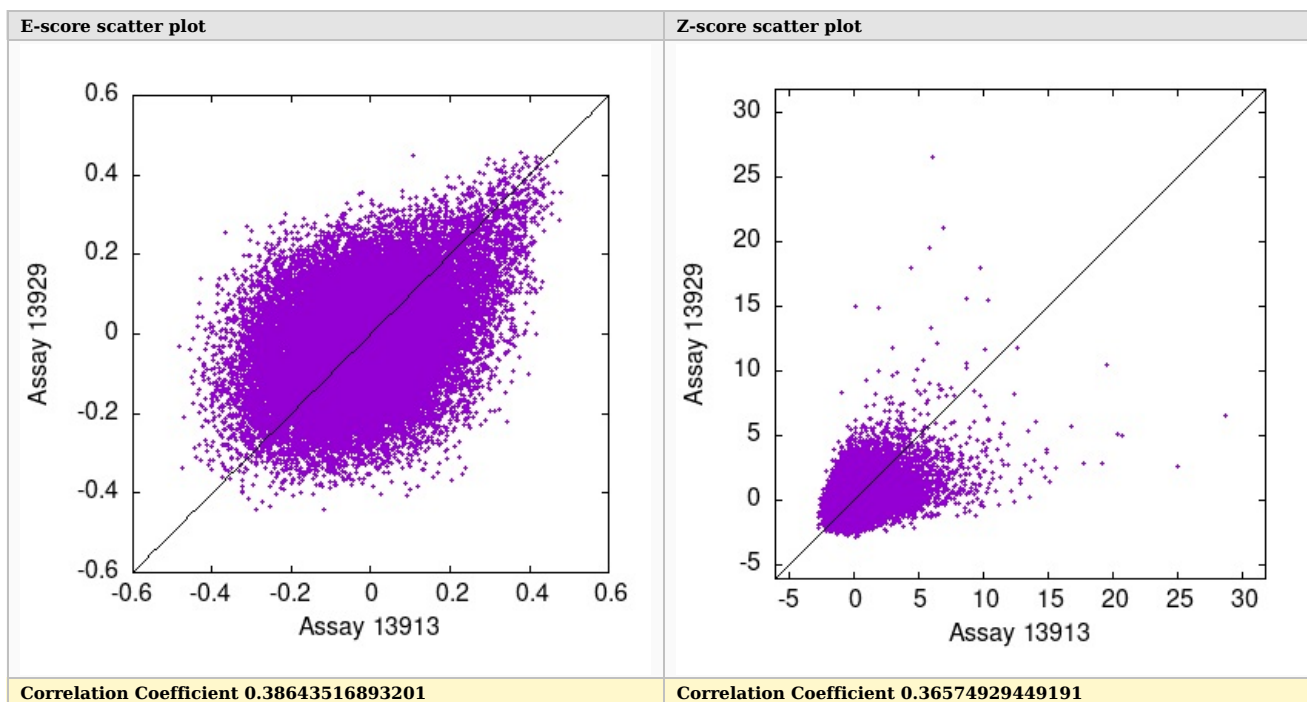




QC report





Top scoring motifs for Assay 13913

Protein ID: pTH14204.1 Gene: MBNL2 Domain: zf-CXXC Flag: Pass matched_pair Array: 1M-ME

8 mer E-scores for probeset 'all'

Forward:	Reverse:	
		
Top 10	Scores	Alignment
AGAAGCAA	0.47781	-AGAAGCAA--
AAGCAGAA	0.47411	---AAGCAGAA
AAAGCAGA	0.46771	---AAAGCAGA-
AAAGCAAT	0.45996	---AAAGCAAT-
AAGCAGGC	0.45539	---AAGCAGGC
AGAAGCGA	0.45438	-AGAAGCGA--
GAGAAGCA	0.44814	GAGAAGCA---
AAGCGAAG	0.44613	---AAGCGAAG
CGCTTATA	0.44485	TATAAGCG---
GAAGCGAA	0.43838	--GAAGCGAA-



8 mer Z-scores for probeset 'all'

Forward:		Reverse:	
			
Top 10	Scores	Alignment	
AGAAGCAA	28.64636	--	AGAAGCAA--
AAGCAGAA	25.01046	----	AAGCAGAA
AAAGCAGA	20.75437	----	AAAGCAGA-
CGCTTATA	20.39562	-	TATAAGCG---
AAGCGAAG	19.52657	----	AAGCGAAG
AAAGCAAT	19.16692	----	AAAGCAAT-
AGAAGCGA	17.75635	--	AGAAGCGA--
GAGAAGCA	16.82912	-	GAGAAGCA---
AGAGAAGC	15.65176		AGAGAAGC----
ATAAGGCA	15.13678	--	ATAAGGCA--



Top scoring motifs for Assay 13929

Protein ID: pTH14204.2 Gene: MBNL2 Domain: zf-CXXC Flag: Pass matched_pair Array: 1M-HK

8 mer E-scores for probeset 'all'

Forward:	Reverse:	
		
Top 10	Scores	Alignment
CGCTTATC	0.45514	-GATAAGCG---
GTAAGCGA	0.44717	--GTAAGCGA--
AAAAGCAG	0.44560	---AAAAGCAG--
CTGCTTTA	0.44467	---TAAAGCAG--
AAGCAAAG	0.44406	----AAGCAAAG
AAAGCAAA	0.44172	---AAAGCAAA-
ATAGAAGC	0.44098	---ATAGAAGC-
CGCTTACC	0.43985	-GGTAAGCG---
AAAAAAGC	0.43892	AAAAAAGC----
AAAGCAGA	0.43494	---AAAGCAGA-

8 mer Z-scores for probeset 'all'

Forward:		Reverse:
		
Top 10	Scores	Alignment
CGCTTATC	26.57682	-GATAAGCG---
ACGCCTCA	21.14358	--TGAGGCGT--
AATGAAGC	19.53108	AATGAAGC----
CGCTTACC	18.05208	-GGTAAGCG---
AAAAGCAG	18.02739	---AAAAGCAG--
AAAGCAAA	15.64107	---AAAGCAAA-
AAGCAAAG	15.48650	----AAGCAAAG
GTAAGCGA	15.00118	--GTAAGCGA--
AAGCGGGT	14.94552	---AAGCGGGT
CAAGCAAA	13.40709	---CAAGCAAA-