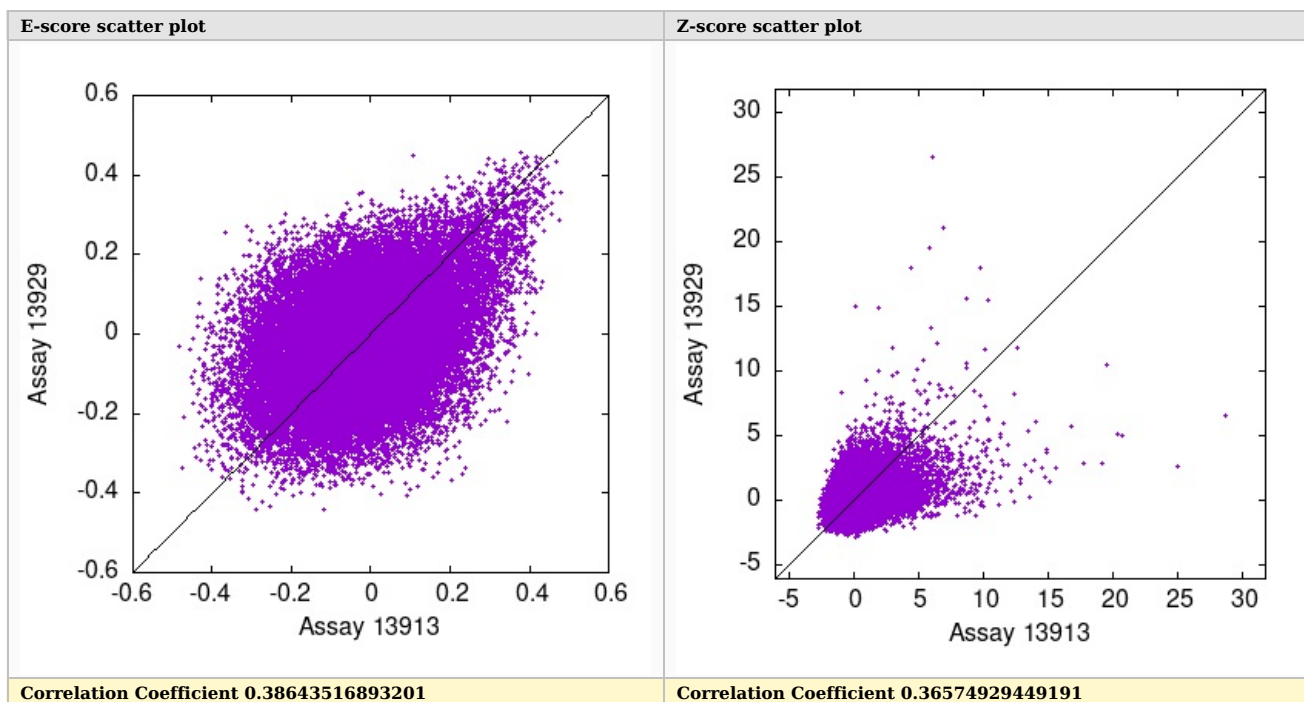


QC report



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-	