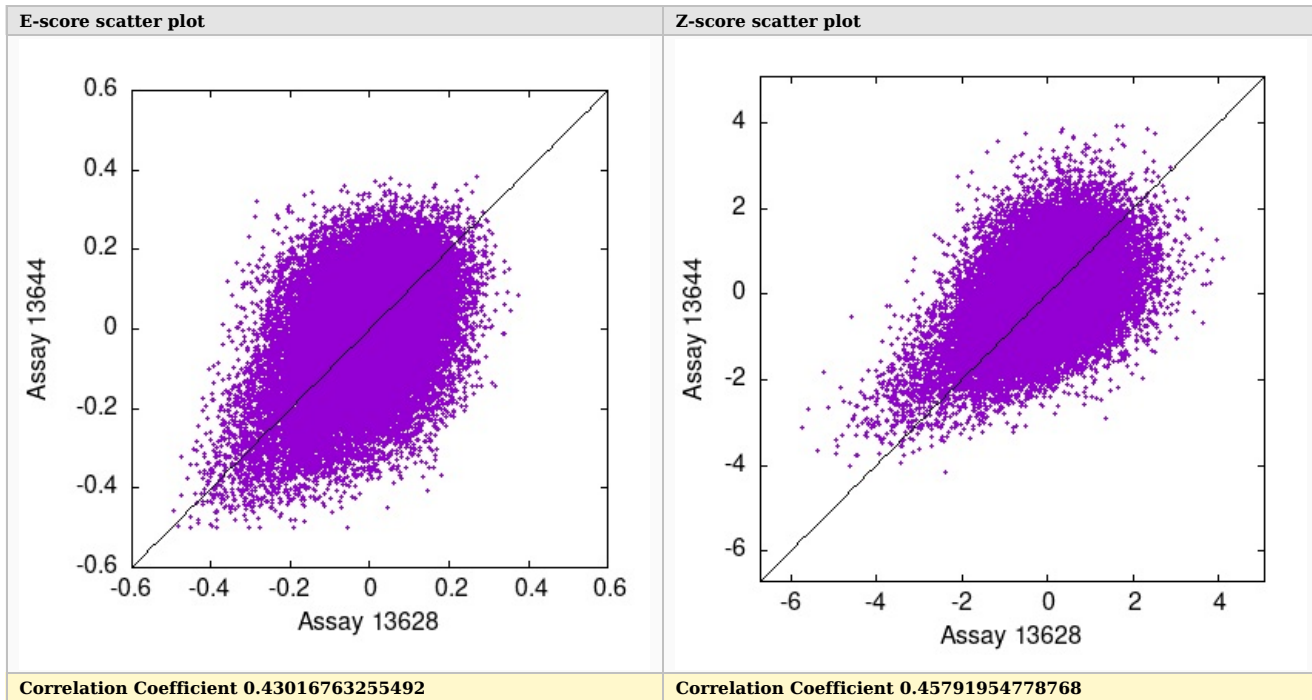


QC report



Top scoring motifs for Assay 13628

Protein ID: pTH14229.1 Gene: GATAD2B.FL Domain: GATA Flag: Reject 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
ACAACACG	0.37458	CGTGTGTG---	GTGTAGTA	4.09035	TACTACAC---
ACAACAAT	0.35726	ATTGTTGT---	ATGTTGTC	3.97185	GACAACAT---
ATGTTGTC	0.35208	-ATGTTGTC--	AAACCAAC	3.86439	-AAACCAAC--
ATGTTTCA	0.34812	-ATGTTTCA--	ACAACACG	3.79441	-ACAACACG--
ACAACCTGA	0.34046	TCAGTTGT---	ACAACAAT	3.73220	-ACAACAAT--
CACAACAC	0.33989	-GTGTTGTG--	ATTACGAC	3.70325	--ATTACGAC
CCAACAAC	0.33870	---GTTGTTGG	CCAACAAC	3.63409	-CCAACAAC--
GTGTAGTA	0.33837	-GTGTAGTA--	AACATGTT	3.63201	--AACATGTT
GAATCCAA	0.33479	-TTGGATTC--	AGTATTGA	3.62044	-TCAATACT--
AGTCAGGA	0.32015	--TCCTGACT-	CGAACAGA	3.57866	-CGAACAGA--

Top scoring motifs for Assay 13644

Protein ID: pTH14229.2 Gene: GATAD2B.FL Domain: GATA Flag: Reject 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
GTCATGAC	0.38185	-GTCATGAC----	ACAAACAG	3.93093	----ACAAACAG--
TATGCATA	0.37821	-TATGCATA----	TCGCGCGA	3.92404	----TCGCGCGA--
ACAATTCA	0.36977	-----TGAAGTGT	TATGCATA	3.85353	---TATGCATA---
TACGCGTA	0.36536	-TACGCGTA----	CGCGCGCG	3.84057	--CGCGCGCG---
CGCGCGCG	0.36499	-CGCGCGCG----	TATAACGA	3.77131	--TATAACGA----
CGCGCGAA	0.36466	-CGCGCGAA----	ACGAAACG	3.75244	--ACGAAACG---
TCGCGCGA	0.36282	TCGCGCGA-----	GAAATTTT	3.73879	GAAATTTT-----
ATCACACA	0.36058	-TGTGTGAT----	ATTTCAAC	3.70199	---ATTTCAAC---
GGTCTAGA	0.35857	--GGTCTAGA---	ACACTTCA	3.67082	-----ACACTTCA
ACTGTAAA	0.35813	-ACTGTAAA----	ATCACACA	3.62801	--ATCACACA---