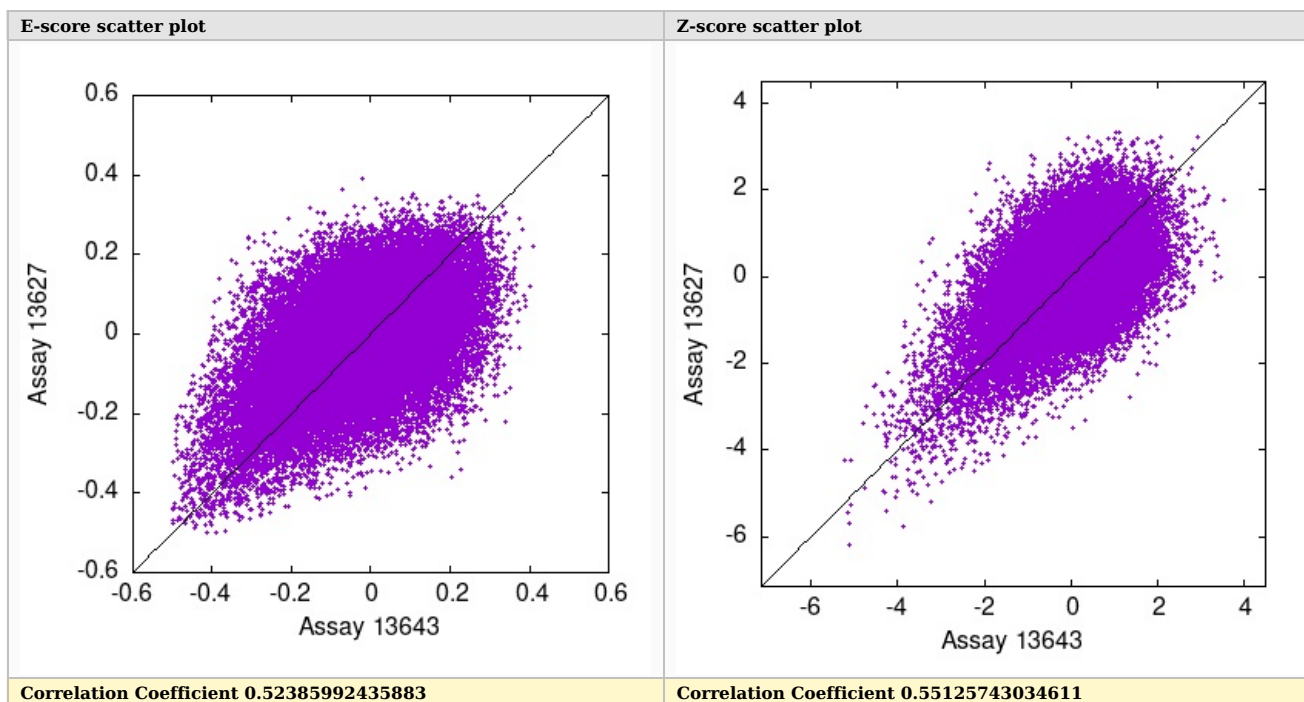


QC report



Top scoring motifs for Assay 13643

Protein ID: pTH14228.2 Gene: GATAD2A.DBD 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
TACGCGTA	0.41137	----TACGCGTA	AACGTTTC	3.51649	----GAAACGTT----
GATGCATC	0.40236	--GATGCATC--	AATTTAAA	3.44841	AATTTAAA-----
GCGCGCGC	0.39127	-GCGCGCGC---	AAACGCGC	3.39672	-----AAACGCGC--
AAACGCGC	0.39122	-AAACGCGC---	GTACAACA	3.33225	--GTACAACA-----
CAATTGTC	0.38803	----GACAATTG	TTCATGAA	3.30641	-----TTCATGAA--
AACGGTTG	0.38113	--CAACGGTT--	GCGCGCGC	3.29938	-----GCGCGCGC--
TTCATGAA	0.37461	TTCATGAA----	AGTATACT	3.28963	--AGTATACT-----
ACATACAA	0.37452	-ACATACAA---	CGCAACAA	3.26796	-----CGCAACAA
ATTTAAAA	0.37448	-ATTTAAAA---	ATCAGTAC	3.21310	-----ATCAGTAC--
AAACAAC	0.37425	---AAACAAC--	CGCGCGCG	3.18081	-----CGCGCGCG--

Top scoring motifs for Assay 13627

Protein ID: pTH14228.1 Gene: GATAD2A.DBD 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
CCACAACA	0.39037	-CCACAACA---	AATCGAAC	3.30045	AATCGAAC----
TACAAC	0.36552	--TACAAC--	TATACAAA	3.29949	TATACAAA-----
CACAACAC	0.35298	--CACAACAC--	CAACAATC	3.22444	----CAACAATC
TATACAAA	0.34910	TATACAAA----	ACACATAC	3.22136	---ACACATAC-
ATTAGTTG	0.34514	----CAACTAAT	AACGCGTT	3.21166	--AACGCGTT--
GATCCGAA	0.34465	GATCCGAA----	AACAGACA	3.19092	--AACAGACA--
CGTTTTGC	0.34162	-GCAAAACG---	GACAAC	3.18982	--GACAAC--
GTTTTGTA	0.33994	--TACAAAAC--	TACAAC	3.18541	--TACAAC--
AAATGCAT	0.33980	-ATGCATTT---	ACTGATTG	3.16697	----CAATCAGT
GACAAC	0.33273	--GACAAC--	AAACAACG	3.14770	-AAACAACG---