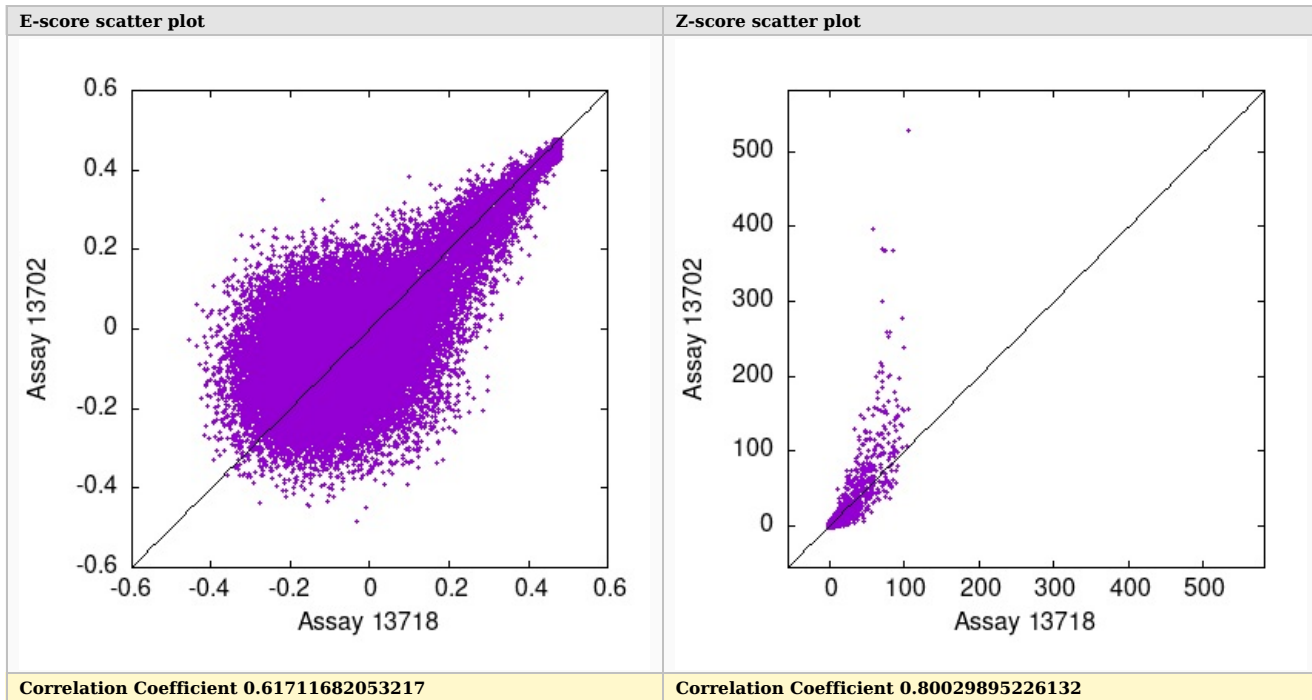


QC report



Top scoring motifs for Assay 13718

Protein ID: pTH14232.2 Gene: ZGLP1.DBD Domain: GATA 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
GATCGATC	0.48419	---GATCGATC	GATCGATC	106.37461	GATCGATC---
GATGATCA	0.48404	GATGATCA---	TTGATCAA	104.45268	--TTGATCAA-
CGATCAGC	0.48360	CGTGATCG---	CGTGATCA	102.28186	-CGTGATCA--
CGATCGCC	0.48360	GCGGATCG---	AGCGATCG	100.08916	-AGCGATCG--
GCGATCGC	0.48221	-GCGATCGC--	CGATCGCC	98.86468	-GCGATCGC--
TTGATCAA	0.48195	--TTGATCAA--	GTGATCAC	97.39004	--GTGATCAC-
CGTGATCA	0.48155	CGTGATCA---	CACGATCG	96.82296	---CGATCGTG
GTGATCAA	0.48102	-GTGATCAA--	GCGATCGC	96.70618	--GCGATCGC-
TGATCAAA	0.48014	--TGATCAAA--	AGCGATCA	96.50692	-AGCGATCA--
AGCGATCG	0.48001	AGCGATCG---	CGATCATC	95.43301	---CGATCATC

Top scoring motifs for Assay 13702

Protein ID: pTH14232.1 Gene: ZGLP1.DBD Domain: GATA 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
ACGATCGT	0.47792	--ACGATCGT--	GATCGATC	528.82813	---GATCGATC
ATCGATCG	0.47661	-ATCGATCG---	ATCGATCG	397.56767	--CGATCGAT-
CGATCGGA	0.47639	---CGATCGGA-	ACGATCGT	369.66832	-ACGATCGT--
GATCGATC	0.47568	GATCGATC----	ATCGATCA	368.03588	--TGATCGAT-
ATGATCAT	0.47527	--ATGATCAT--	ATGATCAT	367.16833	-ATGATCAT--
GATGATCA	0.47189	---TGATCATC-	GATGATCA	367.16833	GATGATCA---
GCGATCGA	0.46965	--TCGATCGC--	GCGATCGA	300.66634	-GCGATCGA--
GATCATCA	0.46917	----GATCATCA	GCGATCGC	278.29637	-GCGATCGC--
CGATCGCA	0.46907	---CGATCGCA-	CGATCGCG	259.18618	CGCGATCG---
ATCGATCA	0.46883	-ATCGATCA---	CGATCGCA	258.52797	TGCGATCG---