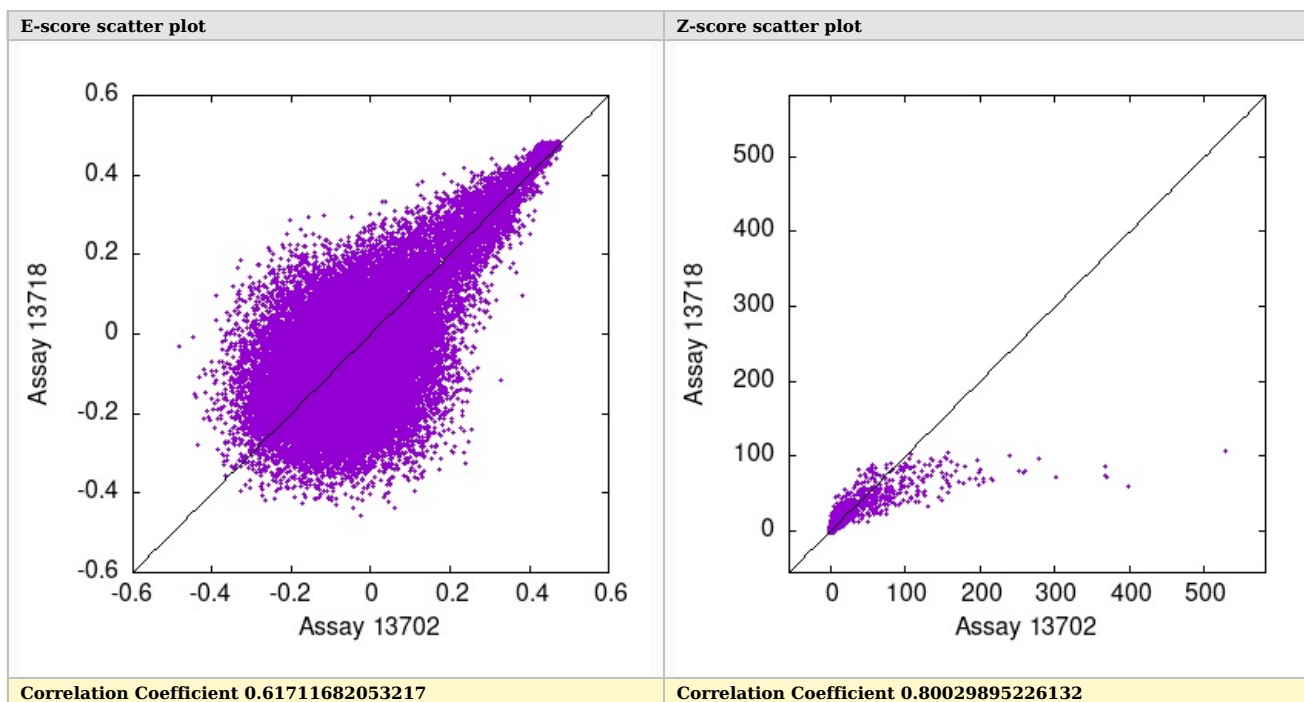


## QC report



## 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'

Forward:		Reverse:	
Top 10	Scores	Alignment	
ACGATCGT	0.47792	--ACGATCGT--	
ATCGATCG	0.47661	-ATCGATCG--	
CGATCGGA	0.47639	---CGATCGGA-	
GATCGATC	0.47568	GATCGATC----	
ATGATCAT	0.47527	--ATGATCAT--	
GATGATCA	0.47189	---TGATCATC-	
GCGATCGA	0.46965	---TCGATCGC--	
GATCATCA	0.46917	----GATCATCA	
CGATCGCA	0.46907	---CGATCGCA--	
ATCGATCA	0.46883	-ATCGATCA---	

## 8 mer Z-scores for probeset 'all'

Forward:		Reverse:	
Top 10	Scores	Alignment	
GATCGATC	528.82813	---GATCGATC	
ATCGATCG	397.56767	--CGATCGAT-	
ACGATCGT	369.66832	-ACGATCGT--	
ATCGATCA	368.03588	--TGATCGAT-	
ATGATCAT	367.16833	-ATGATCAT--	
GATGATCA	367.16833	GATGATCA---	
GCGATCGA	300.66634	-GCGATCGA--	
GCGATCGC	278.29637	-GCGATCGC--	
CGATCGCG	259.18618	CGCGATCG--	
CGATCGCA	258.52797	TGCGATCG--	

## 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'

Forward:		Reverse:	
Top 10	Scores	Alignment	
GATCGATC	0.48419	---GATCGATC	
GATGATCA	0.48404	GATGATCA---	
CGATCACG	0.48360	CGTGATCG---	
CGATCGCC	0.48360	GGCGATCG---	
GCGATCGC	0.48221	-GCGATCGC--	
TTGATCAA	0.48195	-TTGATCAA--	
CGTGATCA	0.48155	CGTGATCA---	
GTGATCAA	0.48102	-GTGATCAA--	
TGATCAAA	0.48014	--TGATCAAA--	
AGCGATCG	0.48001	AGCGATCG---	

## 8 mer Z-scores for probeset 'all'

Forward:		Reverse:	
Top 10	Scores	Alignment	
GATCGATC	106.37461	GATCGATC---	
TTGATCAA	104.45268	--TTGATCAA-	
CGTGATCA	102.28186	-CGTGATCA--	
AGCGATCG	100.08916	-AGCGATCG--	
CGATCGCC	98.86468	-GCGATCGC--	
GTGATCAC	97.39004	--GTGATCAC-	
CACGATCG	96.82296	---GATCGTGT	
GCGATCGC	96.70618	--GCGATCGC-	
AGCGATCA	96.50692	-AGCGATCA--	
CGATCATC	95.43301	---CGATCATC	