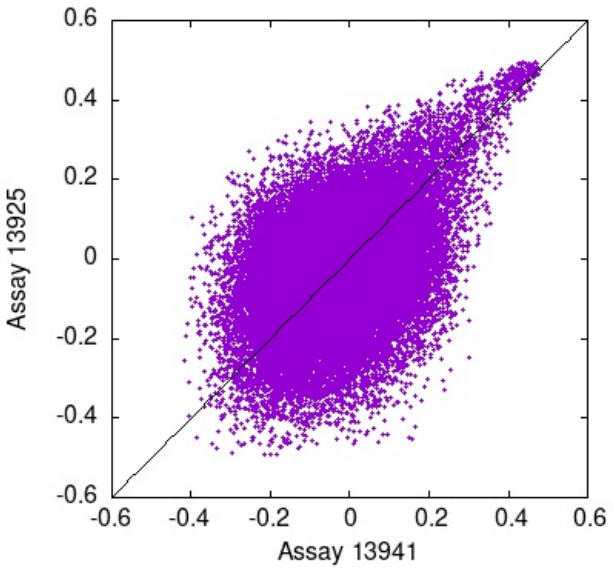
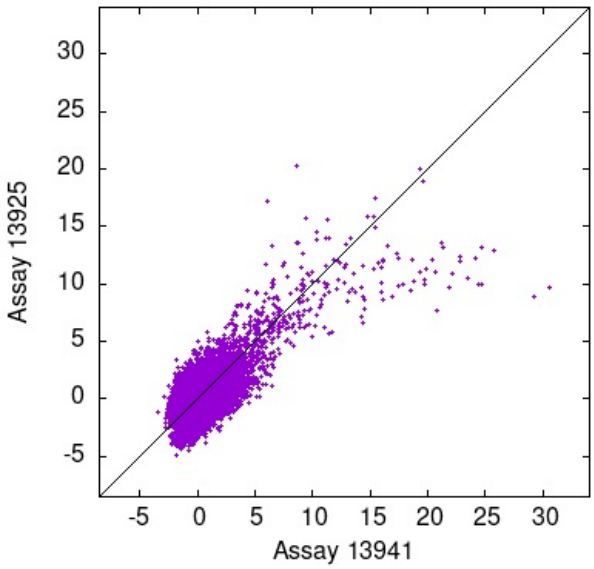
QC report

E-score scatter plot	Z-score scatter plot
	
Correlation Coefficient 0.45119520983161	**Correlation Coefficient 0.62722090749018**

Top scoring motifs for Assay 13941
Protein ID: pTH14231.2 **Gene:** ZGLP1.FL **Domain:** GATA **Flag:** Pass_matched_pair **Array:** 1M-HK

8 mer E-scores for probeset 'all'

Forward:

Reverse:

Top 10	Scores	Alignment
TTGATCAA	0.48027	-TTGATCAA-
GTGATCAA	0.47646	-GTGATCAA-
TGATCACA	0.47464	TGTGATCA--
TGATCAAA	0.47426	--TGATCAAA
CGTGATCA	0.47405	CGTGATCA--
CGATCACG	0.47365	CGTGATCG--
ATTGATCA	0.47348	--TGATCAAT
ACCGATCA	0.47326	ACCGATCA--
GTGATCAC	0.47224	-GTGATCAC-
CGATCAAA	0.47220	--CGATCAAA

8 mer Z-scores for probeset 'all'

Forward:

Reverse:

Top 10	Scores	Alignment
TGATCACA	30.50083	--TGATCACA
GTGATCAA	29.26744	-TTGATCAC-
TCGATCGA	25.79891	-TCGATCGA-
ACGATCAC	24.68139	-ACGATCAC-
ATTGATCA	24.67758	ATTGATCA--
AACGATCA	24.34985	AACGATCA--
CACGATCG	24.12365	CACGATCG--
CACGATCA	23.45950	CACGATCA--
CGTGATCA	22.83875	--TGATCACG
GTGATCAC	22.83875	-GTGATCAC-

Top scoring motifs for Assay 13925
Protein ID: pTH14231.1 **Gene:** ZGLP1.FL **Domain:** GATA **Flag:** Pass_matched_pair **Array:** 1M-ME

8 mer E-scores for probeset 'all'

Forward:

Reverse:

Top 10	Scores	Alignment
GATCGATC	0.49644	GATCGATC---
ATCGATCG	0.49532	-ATCGATCG--
GCGATCGC	0.49319	--GCGATCGC-
ATGATCAT	0.49264	--ATGATCAT-
ATCGATCA	0.49120	-ATCGATCA--
CGATCGCG	0.49115	---CGATCGCG
ACGATCGT	0.49109	--ACGATCGT-
CGATCGGA	0.49042	---CGATCGGA
GATGATCA	0.49016	---TGATCATC
GCGATCGA	0.48718	--TCGATCGC-

8 mer Z-scores for probeset 'all'

Forward:

Reverse:

Top 10	Scores	Alignment
ATCGATCG	20.30647	-ATCGATCG--
GATCGATC	19.99364	GATCGATC---
GCGATCGC	18.87720	--GCGATCGC-
ATGATCAT	17.44901	--ATGATCAT-
ACGATCGT	17.10858	--ACGATCGT-
GATGATCA	15.81041	---TGATCATC
ATCGATCA	15.77328	-ATCGATCA--
CGATCGGA	15.76326	---CGATCGGA
CGATCGCG	15.55654	---CGATCGCG
CGATCGCA	14.88697	---CGATCGCA