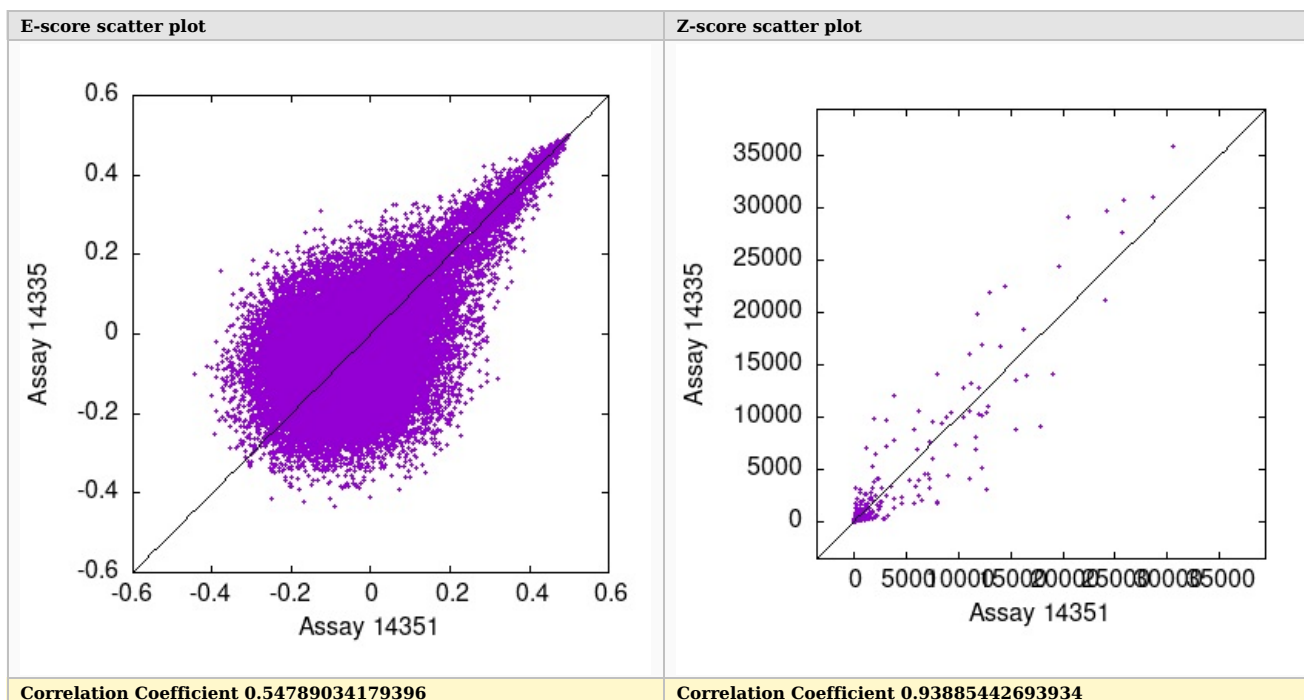




## QC report





### Top scoring motifs for Assay 14351

Protein ID: pTH15534.2 Gene: FLI1 Domain: Ets Flag: Pass\_matched\_pair Array: 1M-HK

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

Forward:	Reverse:	
		
Top 10	Scores	Alignment
ACTTCCGG	0.49786	-ACTTCCGG--
ACCGGAAA	0.49686	--TTTCCGGT-
ATTTCCGG	0.49678	-ATTTCCGG--
CCGGAAC	0.49606	-GTTTCCGG--
ACCGGAAG	0.49539	--CTTCCGGT-
TCCGAAA	0.49452	--TTTCCGGA-
TTCCGAA	0.49206	---TTCCGGAA
CTTCCGGA	0.49153	--CTTCCGGA-
CACTTCCG	0.49094	CAC TTCCG--
AAC TTCCG	0.49086	AAC TTCCG--



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

Forward:		Reverse:
		
Top 10	Scores	Alignment
ACTTCCGG	30,549.06547	-ACTTCCGG-
ACCGGAAA	28,647.37486	- -TTTCCGGT
ATTTCCGG	25,877.08096	-ATTTCCGG-
CCGGAAC	25,689.52793	-GTTTCCGG-
ACCGAAG	24,268.27159	- -CTTCCGGT
TCCGAAA	24,019.85036	- -TTTCCGGA
CTTCCGGA	20,449.49986	- -CTTCCGGA
CCGGAAGC	19,648.27174	-GCTTCCGG-
CTTCCGGC	18,968.99043	- -CTTCCGGC
AAC TTCCG	17,846.67565	AAC TTCCG - -



### Top scoring motifs for Assay 14335

Protein ID: pTH15534.1 Gene: FLI1 Domain: Ets Flag: Pass\_matched\_pair Array: 1M-ME

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

Forward:	Reverse:
	
Top 10	Scores
ACTTCCGG	0.49831
ACCGGAAA	0.49757
ACCGGAAG	0.49719
ATTTCCGG	0.49569
CGGAAGTC	0.49456
CTTCCGGA	0.49452
CCGGA AAC	0.49414
CCGGAAGC	0.49255
TTCCGGAA	0.49165
AACCGGAA	0.49079
Alignment	
-ACTTCCGG--	
--TTTCCGGT-	
-CTTCCGGT-	
-ATTTCCGG--	
GACTTCCG---	
--CTTCCGGA-	
-GTTTCCGG--	
-GCTTCCGG--	
---TTCCGGAA	
---TTCCGGTT	

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

Forward:		Reverse:
		
Top 10	Scores	Alignment
ACTTCCGG	35,854.96772	-ACTTCCGG--
ACCGGAAA	30,999.98294	--TTTCCGGT-
ATTTCCGG	30,696.80604	-ATTTCCGG--
ACCGGAAG	29,769.40466	--CTTCCGGT-
CTTCCGGA	29,074.97888	--CTTCCGGA-
CCGGAAC	27,701.19202	-GTTTCCGG--
CCGGAAGC	24,455.15412	-GCTTCCGG--
CGGAAGTC	22,492.17498	GACTTCCG---
TTCCGGAA	21,913.64367	---TTCCGGAA
TCCGAAA	21,190.82196	--TTTCCGGA-