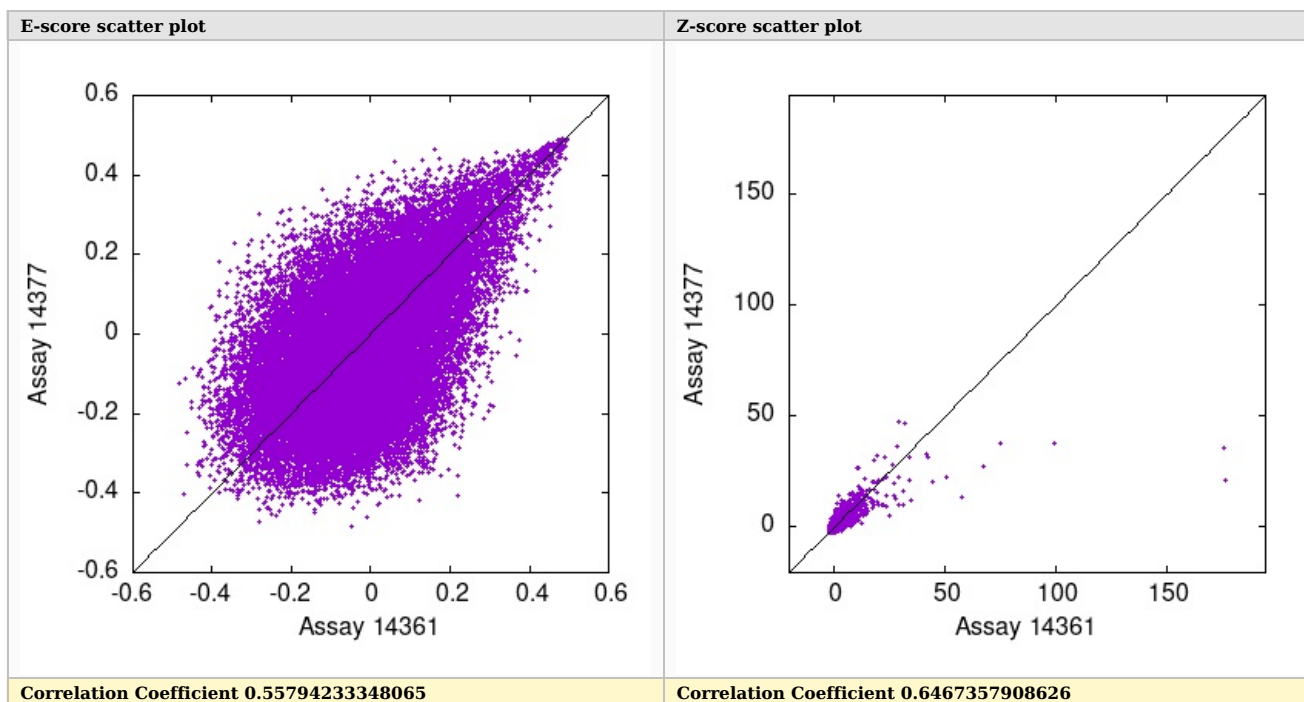


## QC report



### Top scoring motifs for Assay 14361

Protein ID: pTH15521.1 Gene: MYF6 Domain: HLH 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
AACAGTTG	0.49656	-CAACTGTT	AACAGTTG	176.31624	-CAACTGTT
CAGCTGTC	0.49536	-CAGCTGTC	CAGCTGTC	175.37721	-CAGCTGTC
AACAGCTG	0.49532	-CAGCTGTT	ACAGCTGT	98.74419	ACAGCTGT-
ACAGCTGT	0.49504	ACAGCTGT-	AACAGCTG	75.02473	-CAGCTGTT
ACAACCTGT	0.49300	ACAACCTGT-	ACAACCTGT	66.79021	ACAACCTGT-
ACACGTGT	0.49104	ACACGTGT-	ACACGTGT	57.27163	ACACGTGT-
AACAACCTG	0.49053	-CAGTTGTT	AACAACCTG	50.65873	-CAGTTGTT
CAGTTGTC	0.48940	-CAGTTGTC	CAGTTGTC	43.86993	-CAGTTGTC
CAGCTGAC	0.48753	-CAGCTGAC	CAGCTGAC	42.05138	-CAGCTGAC
ACAGCTGC	0.48750	GCAGCTGT-	ACAGCTGC	41.66044	GCAGCTGT-

### Top scoring motifs for Assay 14377

Protein ID: pTH15521.2 Gene: MYF6 Domain: HLH 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
ACAGCTGT	0.49211	-ACAGCTGT	AACAGGTG	47.38325	-CACCTGTT
CAACTGTC	0.49161	GACAGTTG-	CAACTGTC	46.38727	-CAACTGTC
AACAGCTG	0.49132	AACAGCTG-	ACAGCTGT	37.61352	ACAGCTGT-
AACAGGTG	0.49121	AACAGGTG-	AACAGCTG	37.52505	-CAGCTGTT
ACACCTGT	0.48899	-ACACCTGT	ACACCTGT	36.01456	ACACCTGT-
ACAGTTGC	0.48884	-ACAGTTGC	CAGCTGTC	35.38388	-CAGCTGTC
CAGCTGTC	0.48803	GACAGCTG-	ACAGCTGC	32.54027	GCAGCTGT-
ACAGCTGC	0.48778	-ACAGCTGC	ACAGTTGC	32.33513	GCAACTGT-
AACAGTTG	0.48691	AACAGTTG-	CACCTGTC	31.51514	-CACCTGTC
AACACCTG	0.48672	AACACCTG-	CAGCTGAC	31.46222	-CAGCTGAC