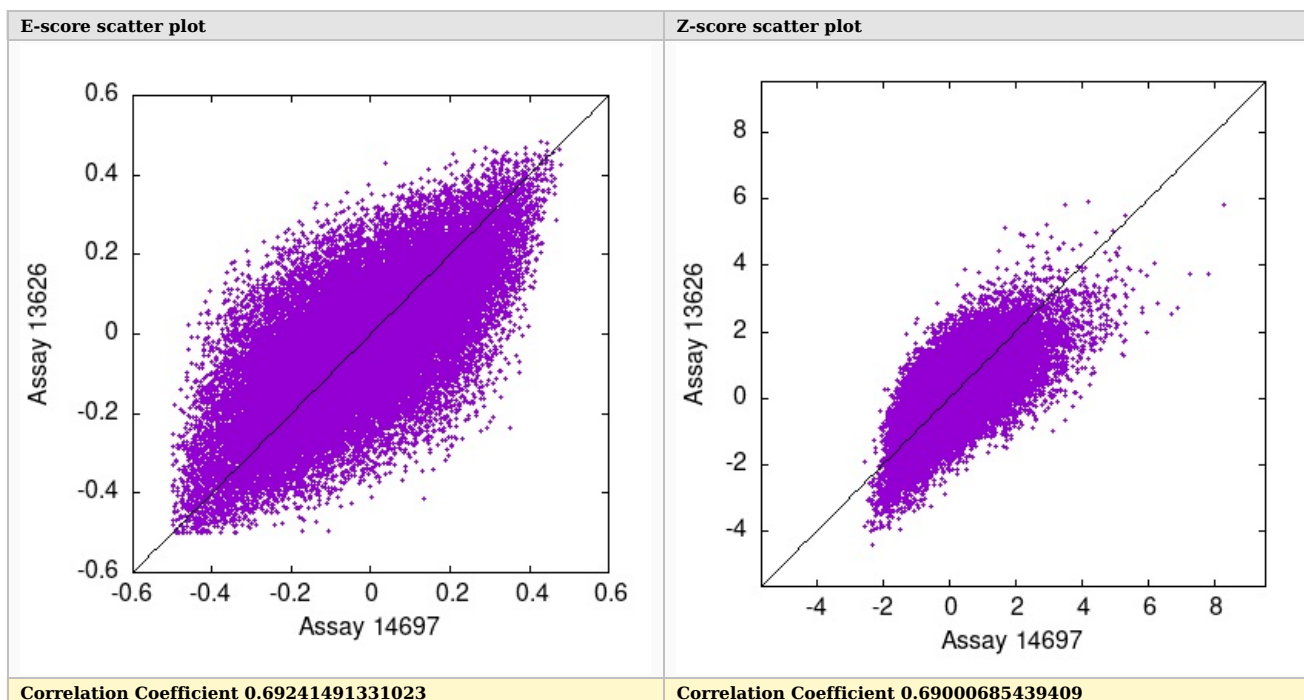


## QC report



### Top scoring motifs for Assay 14697

Protein ID: pTH13919.4 Gene: GLYR1.DBD Domain: AT\_hook Flag: Pass\_matched\_pair Array: 1M-HK

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

| Forward: | Reverse: |
|----------|----------|
|          |          |
| Top 10   | Scores   |
| AATAAATT | 0.47966  |
| AATATATT | 0.47744  |
| ACAAAATA | 0.46651  |
| AATATTAA | 0.46531  |
| ATAAATTA | 0.46460  |
| AAAATTAA | 0.46227  |
| AAATAAAT | 0.46153  |
| ATATTAAT | 0.45767  |
| AAAATAAA | 0.45670  |
| AAATATTT | 0.45578  |

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

| Forward: | Reverse: |
|----------|----------|
|          |          |
| Top 10   | Scores   |
| AATATATT | 8.26500  |
| AATAAATT | 7.80140  |
| AAAATTAA | 7.27601  |
| AATTTAAA | 6.86334  |
| AATATTAA | 6.67980  |
| ATATTAAT | 6.28896  |
| AAATAAAT | 6.17234  |
| AAATTAAT | 5.95404  |
| ACAAAATA | 5.93722  |
| ATAAATTA | 5.80870  |

### Top scoring motifs for Assay 13626

Protein ID: pTH13919.1 Gene: GLYR1.DBD Domain: AT\_hook Flag: Pass\_matched\_pair Array: 1M-ME

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

| Forward: | Reverse: |
|----------|----------|
|          |          |
| Top 10   | Scores   |
| ATTTAAAT | 0.48199  |
| AATTAATT | 0.48119  |
| ATATATTA | 0.46896  |
| ATATATAT | 0.46791  |
| AATATATT | 0.46550  |
| ATAAATAT | 0.46146  |
| AAAATAAT | 0.46119  |
| AACGAATT | 0.46104  |
| AATATTTA | 0.45701  |
| AAATAAAT | 0.45661  |

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

| Forward: | Reverse: |
|----------|----------|
|          |          |
| Top 10   | Scores   |
| ATTTAAAT | 5.90596  |
| AATATATT | 5.82386  |
| ATATATTA | 5.82386  |
| AATTAATT | 5.51849  |
| AATATTTA | 5.23363  |
| AACGAATT | 5.14920  |
| AAAATAAT | 5.02124  |
| AATAAATA | 4.98625  |
| ATATATAT | 4.96139  |
| TAAATTTA | 4.94597  |