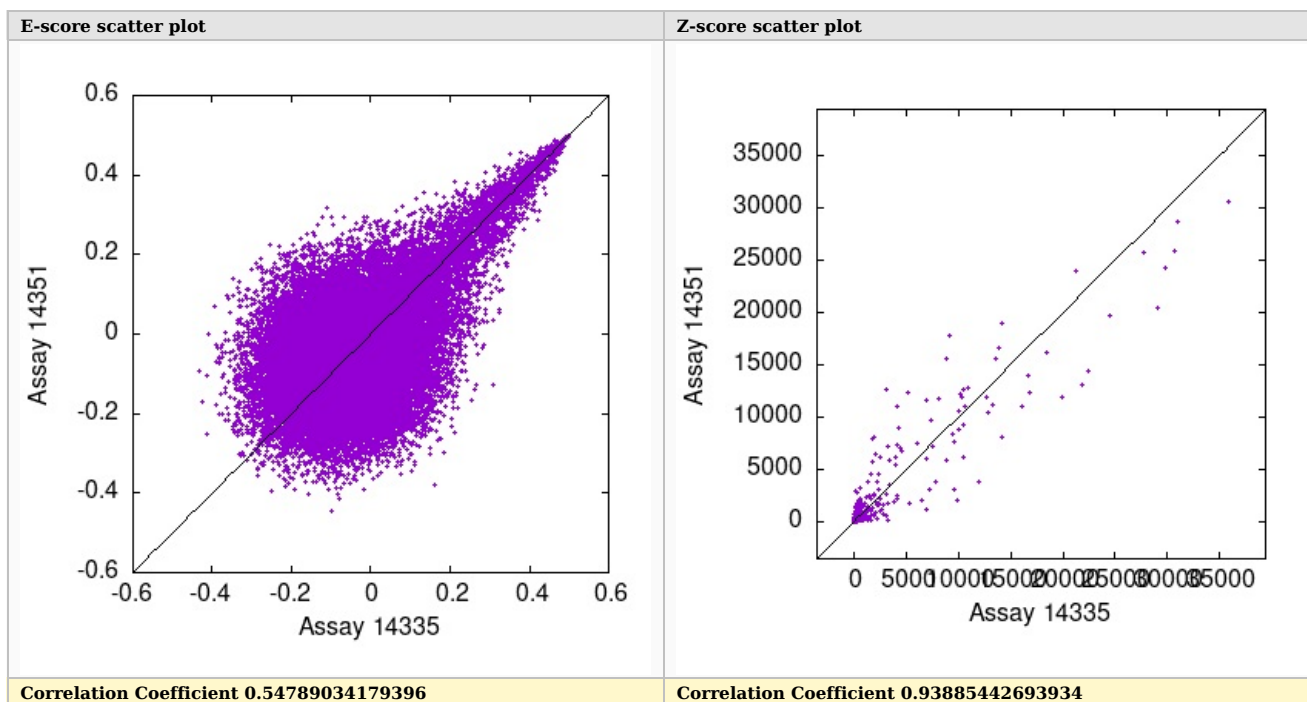


## QC report



### 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' |         |             | 8 mer Z-scores for probeset 'all' |              |             |
|-----------------------------------|---------|-------------|-----------------------------------|--------------|-------------|
| Forward:                          |         | Reverse:    | Forward:                          |              | Reverse:    |
|                                   |         |             |                                   |              |             |
| Top 10                            | Scores  | Alignment   | Top 10                            | Scores       | Alignment   |
| ACTTCCGG                          | 0.49831 | -ACTTCCGG-- | ACTTCCGG                          | 35,854.96772 | -ACTTCCGG-- |
| ACCGGAAA                          | 0.49757 | --TTTCCGGT- | ACCGGAAA                          | 30,999.98294 | --TTTCCGGT- |
| ACCGGAAG                          | 0.49719 | --CTTCCGGT- | ATTTCGGG                          | 30,696.80604 | -ATTTCGGG-- |
| ATTTCGGG                          | 0.49569 | -ATTTCGGG-- | ACCGGAAG                          | 29,769.40466 | --CTTCCGGT- |
| CGGAAGTC                          | 0.49456 | GACTTCCG--- | CTTCCGGA                          | 29,074.97888 | --CTTCCGGA- |
| CTTCCGGA                          | 0.49452 | --CTTCCGGA- | CCGGAAC                           | 27,701.19202 | -GTTTCCGG-- |
| CCGGAAC                           | 0.49414 | -GTTTCCGG-- | CCGGAAGC                          | 24,455.15412 | -GCTTCCGG-- |
| CCGGAAGC                          | 0.49255 | -GCTTCCGG-- | CGGAAGTC                          | 22,492.17498 | GACTTCCG--- |
| TTCCGGAA                          | 0.49165 | ---TTCCGGAA | TTCCGGAA                          | 21,913.64367 | ---TTCCGGAA |
| AACCGGAA                          | 0.49079 | ---TTCCGGTT | TCCGAAA                           | 21,190.82196 | --TTTCCGGA- |

### 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' |         |              | 8 mer Z-scores for probeset 'all' |              |              |
|-----------------------------------|---------|--------------|-----------------------------------|--------------|--------------|
| Forward:                          |         | Reverse:     | Forward:                          |              | Reverse:     |
|                                   |         |              |                                   |              |              |
| Top 10                            | Scores  | Alignment    | Top 10                            | Scores       | Alignment    |
| ACTTCCGG                          | 0.49786 | -ACTTCCGG--  | ACTTCCGG                          | 30,549.06547 | -ACTTCCGG--  |
| ACCGGAAA                          | 0.49686 | --TTTCCGGT-  | ACCGGAAA                          | 28,647.37486 | --TTTCCGGT-  |
| ATTTCGGG                          | 0.49678 | -ATTTCGGG--  | ATTTCGGG                          | 25,877.08096 | -ATTTCGGG--  |
| CCGGAAC                           | 0.49606 | -GTTTCCGG--  | CCGGAAC                           | 25,689.52793 | -GTTTCCGG--  |
| ACCGGAAG                          | 0.49539 | --CTTCCGGT-  | ACCGGAAG                          | 24,268.27159 | --CTTCCGGT-  |
| TCCGAAA                           | 0.49452 | ---TTTCCGGA- | TCCGAAA                           | 24,019.85036 | ---TTTCCGGA- |
| TTCCGGAA                          | 0.49206 | ---TTCCGGAA  | CTTCCGGA                          | 20,449.49986 | --CTTCCGGA-  |
| CTTCCGGA                          | 0.49153 | --CTTCCGGA-  | CCGGAAGC                          | 19,648.27174 | -GCTTCCGG--  |
| CAC TTCG                          | 0.49094 | CAC TTCG---  | CTTCCGGC                          | 18,968.99043 | --CTTCCGGC   |
| AAC TTCG                          | 0.49086 | AAC TTCG---  | AAC TTCG                          | 17,846.67565 | AAC TTCG---  |