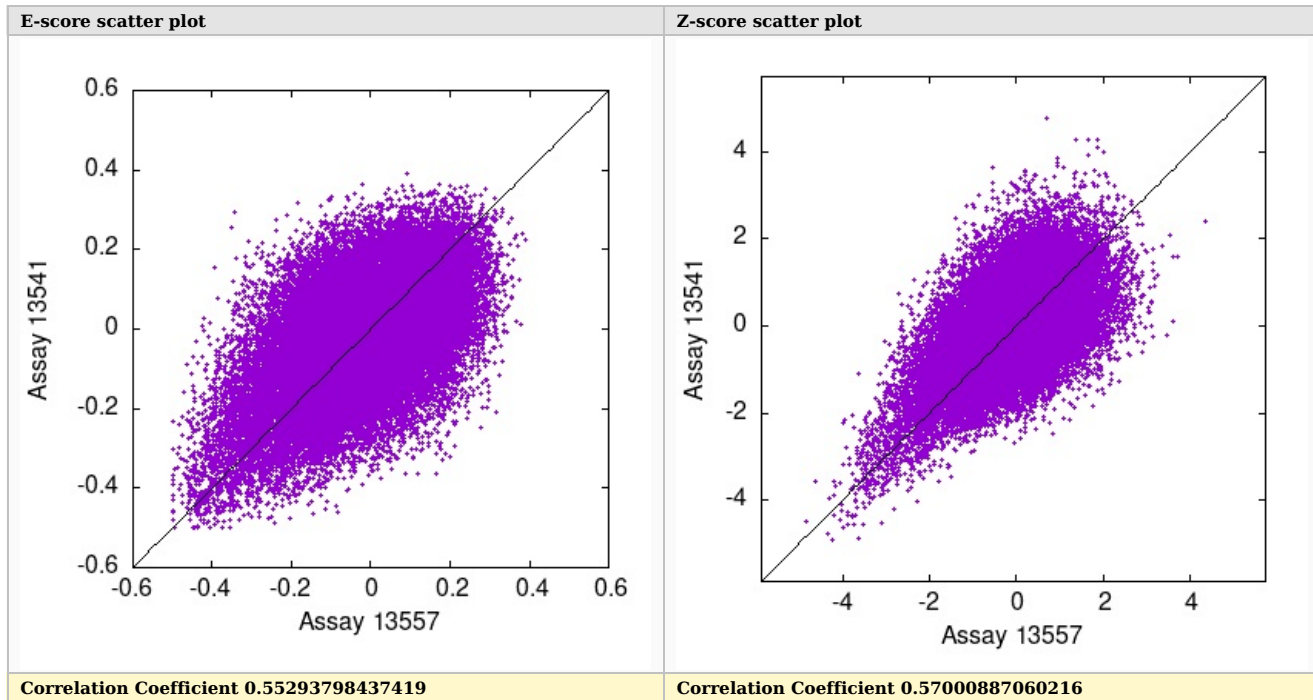


## QC report



### Top scoring motifs for Assay 13557

Protein ID: pTH14288.2 Gene: CENPT.FL Domain: Unknown Flag: Reject Array: 1M-HK

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

Forward:

Reverse:



##### Top 10

##### Scores

##### Alignment

|          |         |                |
|----------|---------|----------------|
| AAGTACTT | 0.38890 | AAGTACTT-----  |
| TACGCGTA | 0.38234 | ---TACGCGTA--- |
| GAAATTTT | 0.37691 | --GAAATTTT---  |
| TCGCGCGA | 0.37565 | ---TCGCGCGA--- |
| TTGCGCAA | 0.37157 | -----TTGCGCAA  |
| CGTCGGAA | 0.36867 | ----CGTCGGAA-- |
| TAGTTGCA | 0.36505 | ---TAGTTGCA--- |
| ATAATGTA | 0.36428 | ---ATAATGTA--- |
| GTACAGTA | 0.35961 | -----GTACAGTA  |
| GTATACTA | 0.35768 | ---TAGTATAC--- |

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

Forward:

Reverse:



##### Top 10

##### Scores

##### Alignment

|          |         |                |
|----------|---------|----------------|
| TACGCGTA | 4.35058 | -----TACGCGTA  |
| TTGCGCAA | 3.69983 | ---TTGCGCAA--- |
| ATAATGTA | 3.60263 | ATAATGTA-----  |
| CACAGTAG | 3.58399 | CTACTGTG-----  |
| ATATGTGC | 3.53777 | -----GCACATAT- |
| TCGCGCGA | 3.45373 | ---TCGCGCGA--- |
| ATTGTACA | 3.35293 | ---ATTGTACA--- |
| TATTTGCA | 3.25446 | ----TGCAAATA-- |
| GTATTGTA | 3.17908 | GTATTGTA-----  |
| GAATATTC | 3.15232 | -GAATATTC----- |

### Top scoring motifs for Assay 13541

Protein ID: pTH14288.1 Gene: CENPT.FL Domain: Unknown Flag: Reject Array: 1M-ME

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

Forward:

Reverse:



##### Top 10

##### Scores

##### Alignment

|          |         |                |
|----------|---------|----------------|
| TGTGCACA | 0.39149 | ---TGTGCACA--  |
| AAAGTACA | 0.36464 | -----TGTA CTTT |
| ACGTAACA | 0.36305 | ---ACGTAACA--- |
| ATATTGTA | 0.36116 | ----ATATTGTA-  |
| AACAAAGC | 0.36001 | -AACAAAGC----  |
| ATGTACAT | 0.35918 | ----ATGTACAT-  |
| ATATATGA | 0.35887 | --TCATATAT---  |
| ACAACAAT | 0.35445 | --ACAACAAT---  |
| ATTCGTTA | 0.35364 | TAACGAAT-----  |
| ACATATAC | 0.35255 | --ACATATAC---  |

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

Forward:

Reverse:



##### Top 10

##### Scores

##### Alignment

|          |         |             |
|----------|---------|-------------|
| ATATTGTA | 4.75439 | ATATTGTA--- |
| ATTCGTTA | 4.27385 | -ATTCGTTA-- |
| ACAACAAT | 4.27157 | -ATTGTTGT-- |
| TGTGCACA | 4.26695 | --TGTGCACA- |
| ATGTACAT | 4.09826 | ---ATGTACAT |
| ATTTATAA | 4.00548 | -ATTTATAA-- |
| ATTTGTAC | 3.91244 | -ATTTGTAC-- |
| AACAAAGC | 3.84111 | GCTTTGTT--- |
| AACATGTT | 3.72651 | AACATGTT--- |
| ATGTTAAT | 3.67272 | -ATGTTAAT-- |