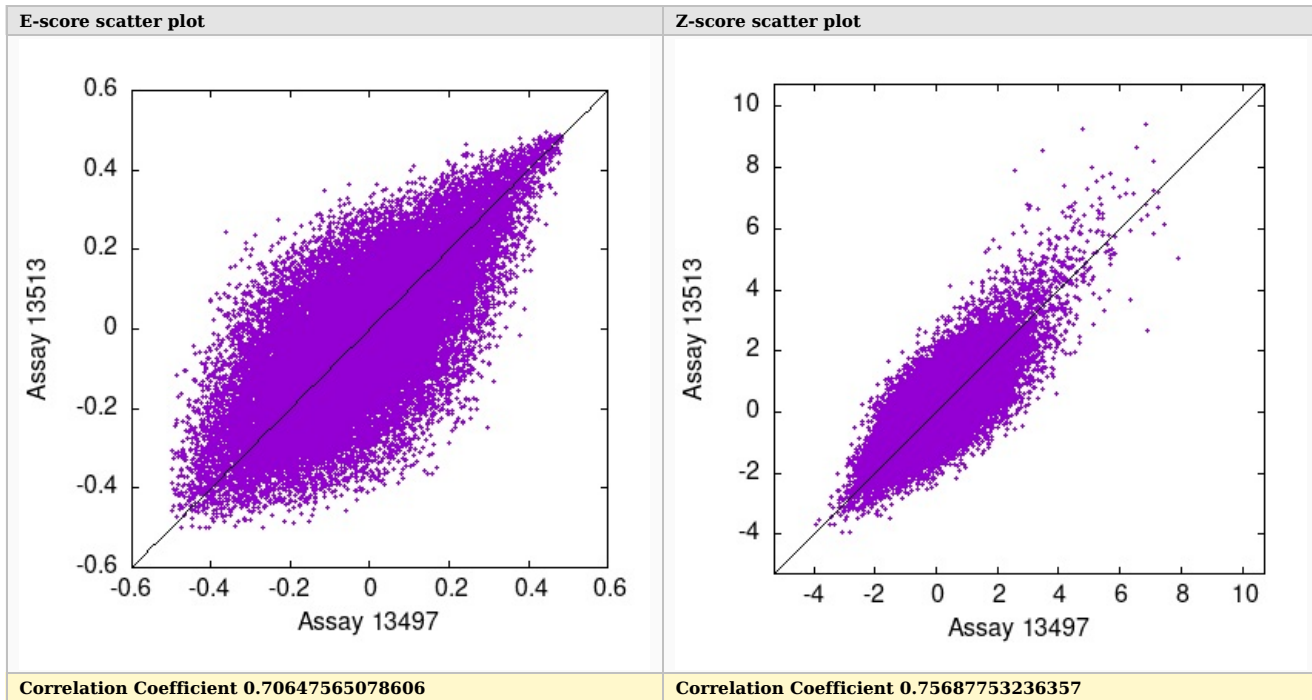


QC report



Top scoring motifs for Assay 13497

Protein ID: pTH13913.1 Gene: AHCTF1.DBD Domain: AT_hook 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 |
| AAATATTT | 0.48374 | --AAATATTT-- | AAAAAAT | 7.89610 | -AAAAAAT- |
| AAAAAATA | 0.48304 | --AAAAAATA-- | AAATATTT | 7.43353 | -AAATATTT- |
| ATTTAAAT | 0.48291 | --ATTTAAAT-- | AAAAATAA | 7.25444 | -AAAAATAA- |
| AAAAATAA | 0.48082 | AAAAATAA---- | ATTTAAAT | 7.22518 | -ATTTAAAT- |
| AAAAAAAT | 0.48026 | --AAAAAAAT-- | AAAAAATA | 7.11045 | --AAAAAATA |
| AATTAATT | 0.47835 | --AATTAATT-- | AATTTAAT | 7.10444 | -AATTTAAT- |
| ATAATATA | 0.47644 | ----ATAATATA | ATTATTTA | 7.07541 | -TAAATAAT- |
| ATTATTTA | 0.47640 | -TAAATAAT--- | CAAAAAAA | 6.91058 | CAAAAAAA-- |
| AAATAATT | 0.47460 | --AAATAATT-- | AAATAATT | 6.90748 | -AAATAATT- |
| AAAATATT | 0.47356 | ---AATATTTT- | AATTAAT | 6.86585 | -AATTAAT- |

Top scoring motifs for Assay 13513

Protein ID: pTH13913.2 Gene: AHCTF1.DBD Domain: AT_hook 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 |
| AAATAAAT | 0.49357 | --AAATAAAT- | AATTAATT | 9.39101 | -AATTAATT- |
| AATTAATT | 0.48760 | --AATTAATT- | AAATAAAT | 9.28751 | AAATAAAT-- |
| AATAAATT | 0.48584 | ---AATAAATT | AAAAATAT | 8.66712 | --ATATTTTT |
| AATTTAAA | 0.48509 | ---TTTAAATT | AATTTAAA | 8.54261 | -TTTAAATT- |
| AAAAAATA | 0.48473 | ---AAAAAATA | AAAAAATA | 8.21743 | -AAAAAATA- |
| AAAAATAT | 0.48382 | --AAAAATAT- | AATATATT | 8.01288 | -AATATATT- |
| AAAAATAA | 0.48167 | AAAAATAA---- | ATTTTTC | 7.91754 | GAAAAAAT-- |
| AATAATAA | 0.48042 | ---AATAATAA | AATAATAA | 7.81197 | -AATAATAA- |
| ATTTAAAT | 0.47921 | --ATTTAAAT- | AAATAATT | 7.70647 | -AAATAATT- |
| AAAAAAAT | 0.47898 | -AAAAAAAT- | AAAATATT | 7.60208 | -AAAATATT- |