

## General Motif Characteristics:

### Top Similar Looking Motifs:

|                                       |       |  |
|---------------------------------------|-------|--|
| ZNF143.p2                             | 0.000 |  |
| HCMC.ZN143_si.wm                      | 0.075 |  |
| JASPAR.znf143.wm                      | 0.083 |  |
| ENCODE.ZNF143                         | 0.134 |  |
| HTSELEX.ZNF143.C2H2.DBD.monomeric.wm1 | 0.198 |  |
| TP53.p2                               | 0.241 |  |
| EBF1.p2                               | 0.267 |  |
| ENCODE.ZNF143-ext                     | 0.288 |  |
| STAT1,3.p3                            | 0.313 |  |
| UNIPROBE.Zic2_secondary.wm            | 0.320 |  |

### Statistics:

- Motif name: ZNF143.p2
- Enrichment: 2.29550220034
- log-Likelihood Ratio: 413.813918338
- Area under precision recall curve: 0.1787

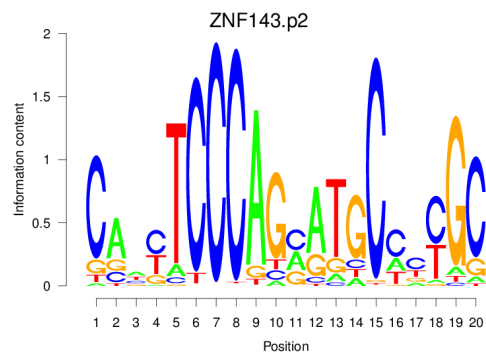


Figure 1: Logo of Weight Matrix

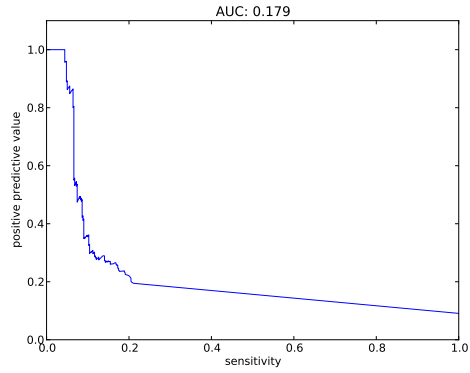


Figure 2: Precision and Recall curve

## Statistics for TFBS - Peak Centering and Peak Posterior - Peak Height Correlation:

### 0.1

Overall statistics:

- Number of true peaks out of total number of peaks: 870/19318
- 4.50 percent are true.
- Cut-off: minimum summed posterior of 0.20
- Peak plots contain TFBS of posterior  $\geq 0.20$

Statistic for centering of TFBSs at peak centers:

- Enrichment at binding sites: 1.8131

Correlation between peak Z-score and number of binding sites at peak: 0.0528

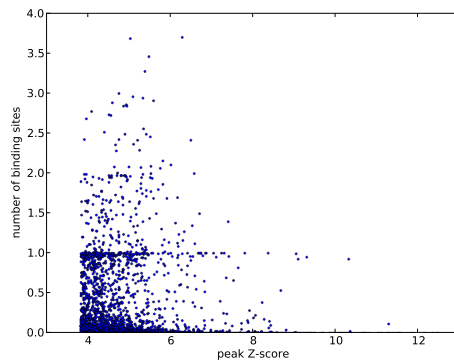


Figure 3: Scatter plot of peak Z-score versus number of binding sites at peak

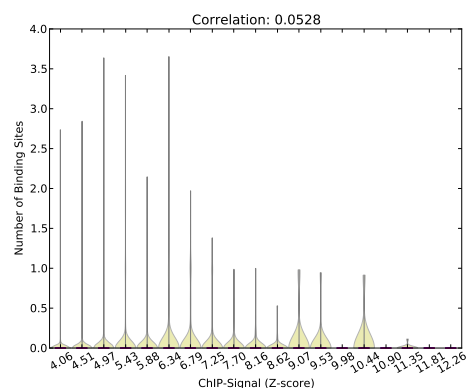


Figure 4: Violin plot of peak Z-score versus number of binding sites at peak

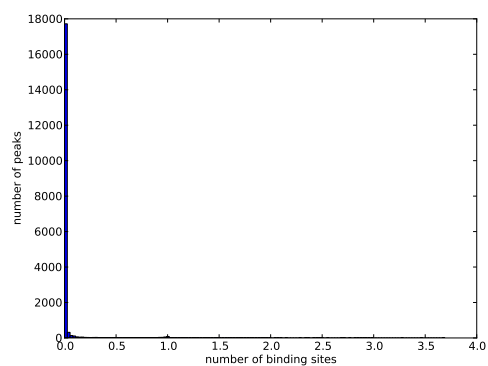


Figure 5: Histogram of number of binding sites at peaks

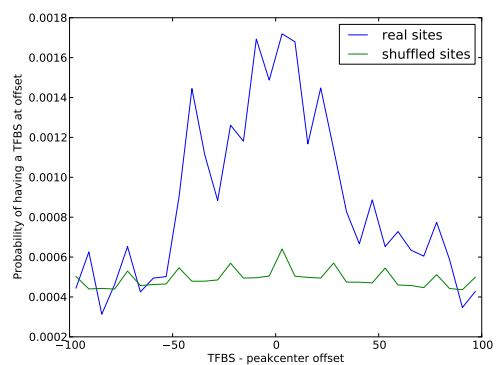


Figure 6: Histogram of distances of TFBS to its nearest peak center

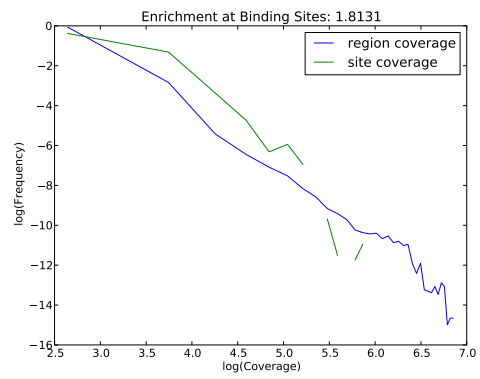


Figure 7: Histograms of coverage at sites and in total regions