

Visualization of the relationship between ChIP-seq peak and attention weights.

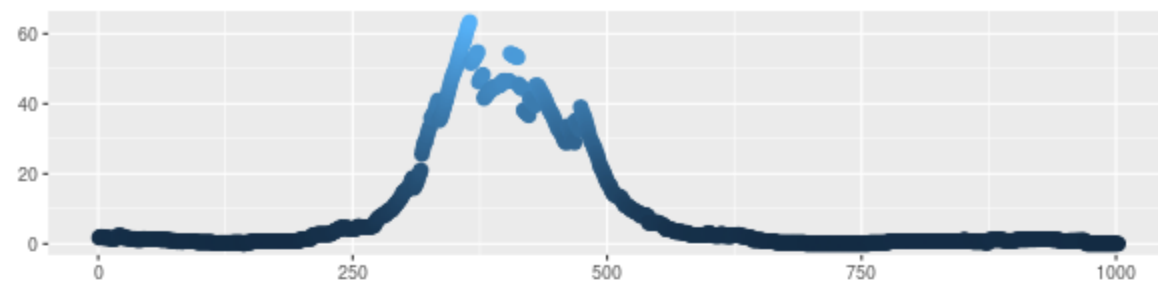
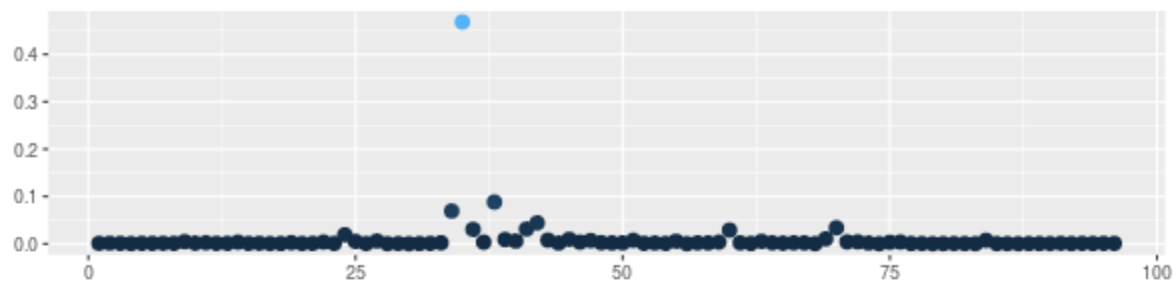
For each genomic region, the figure on the left represents the attention weights and the figure on the right represents the enrichment of fold changes in ChIP-seq BigWig file in the same region. Since the lengths of attention weights are reduced by the convolution and pooling layers, their lengths are less than the fold change values. Thus, the plots are aligned on the X-axis to represent the relative position of fold change and averaged attention weights.

The attention scores are extracted from both single and pairwise models. For single module, the dimension of attention scores is $L \times d_{model}$, and the channel that contains the highest attention weight is used. Similarly, for pairwise module, the dimension of attention scores is $L \times L$, and the row sum of these weights are always 1. As a result, we use the row that contains the highest attention weight.

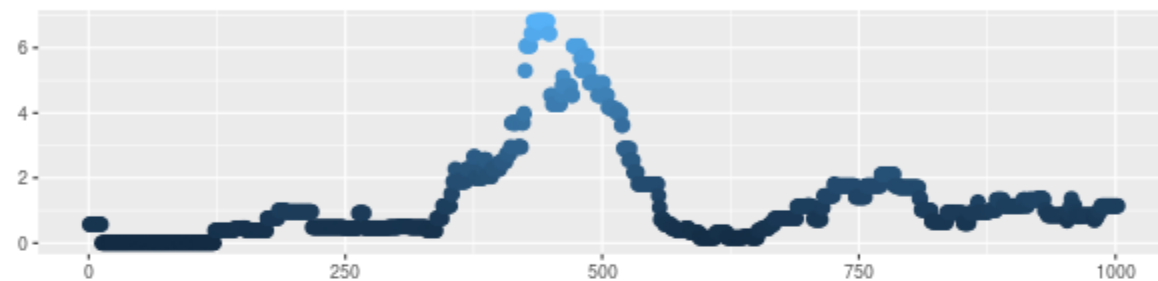
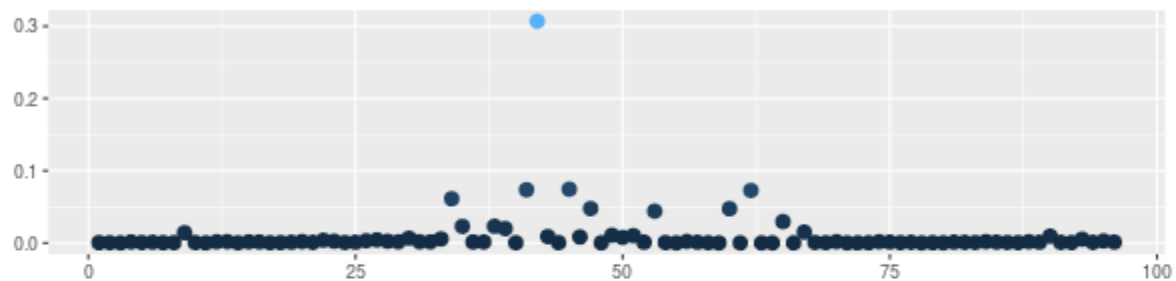
Single attention module

CTCF/induced pluripotent stem cell

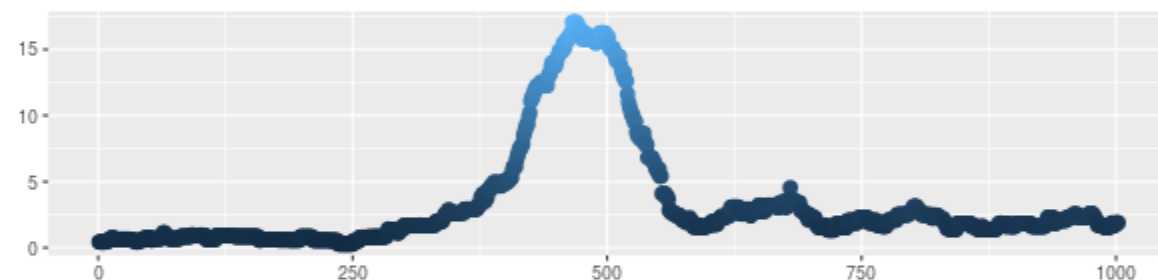
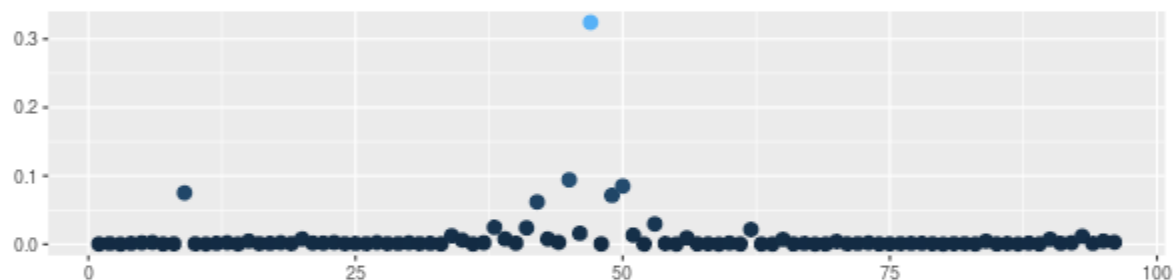
chr1 176528348 - 176529350



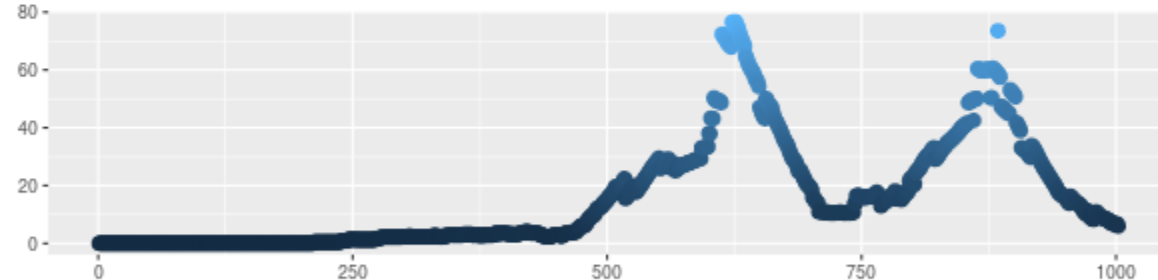
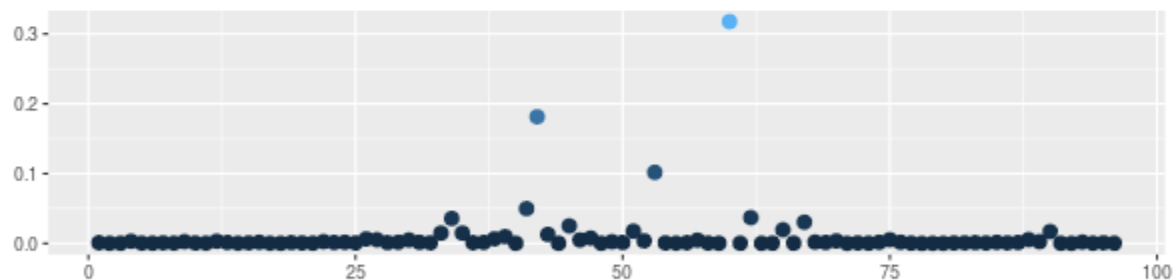
chr8 82541998 - 82543000



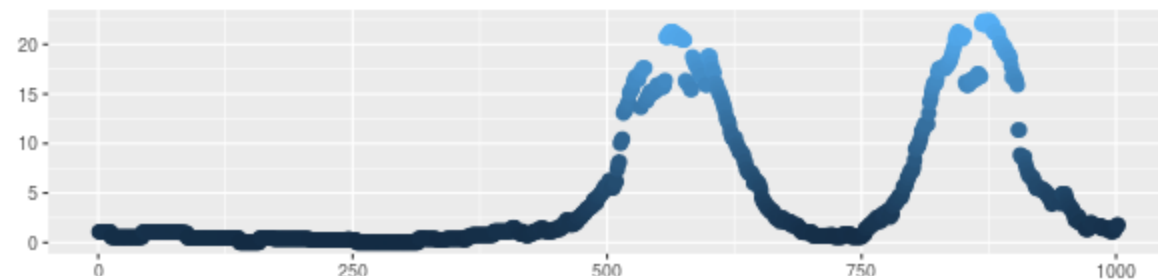
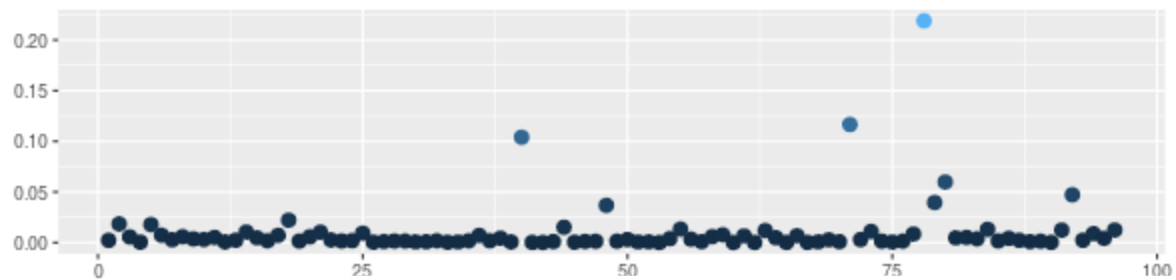
chr1 26360998 - 26362000



chr1 85409548 - 85410550



chr8 48422648 - 48423650

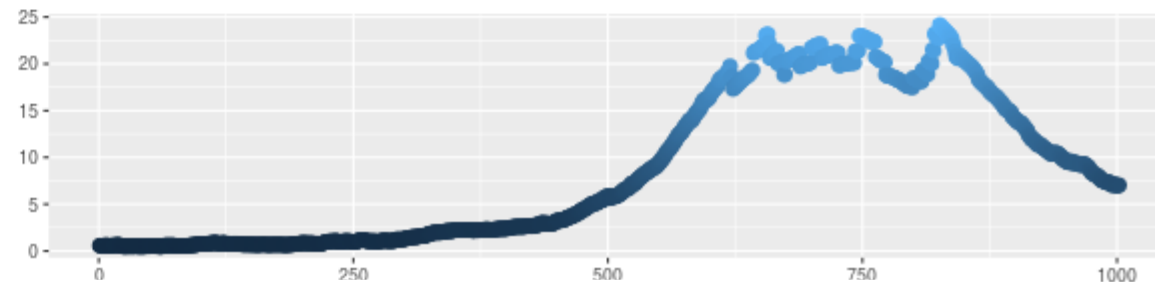
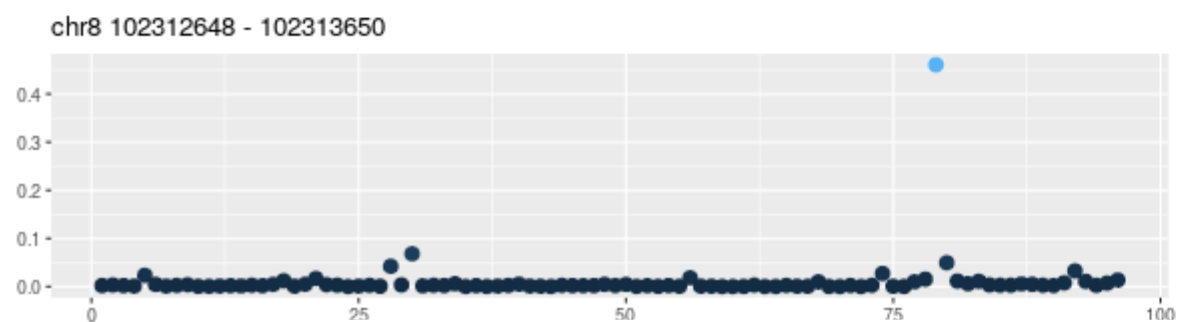
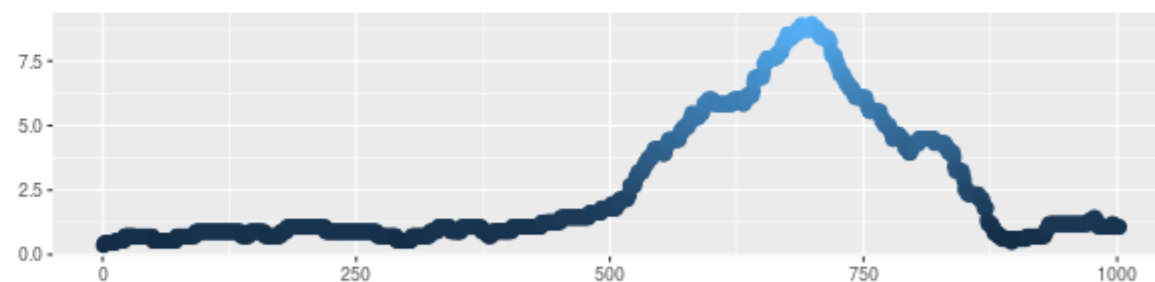
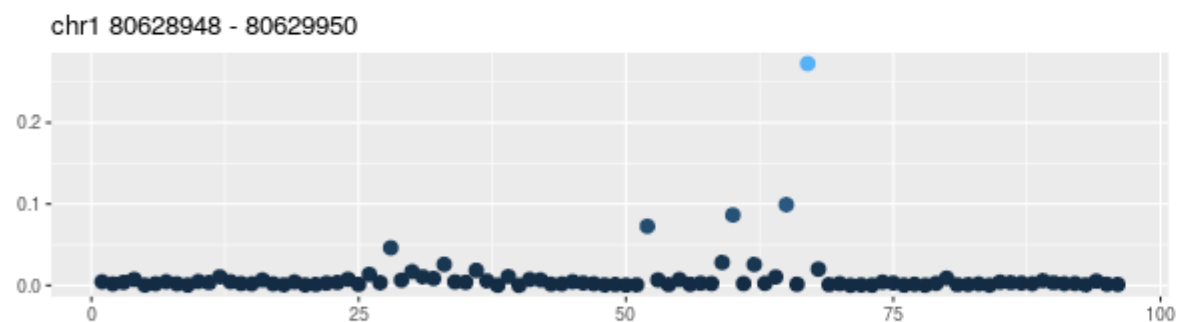
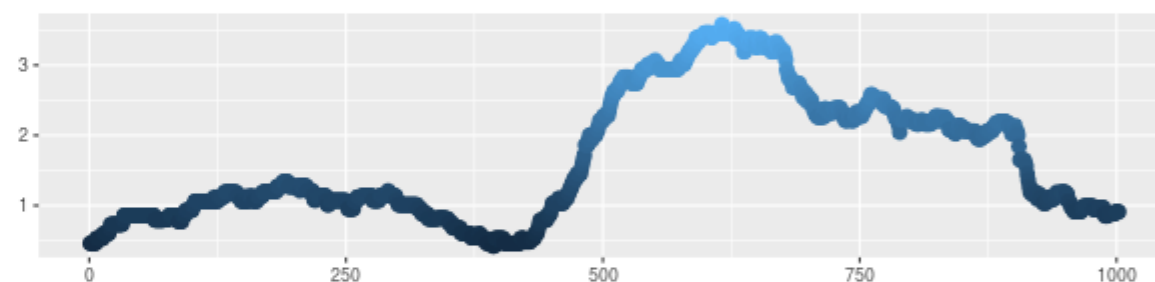
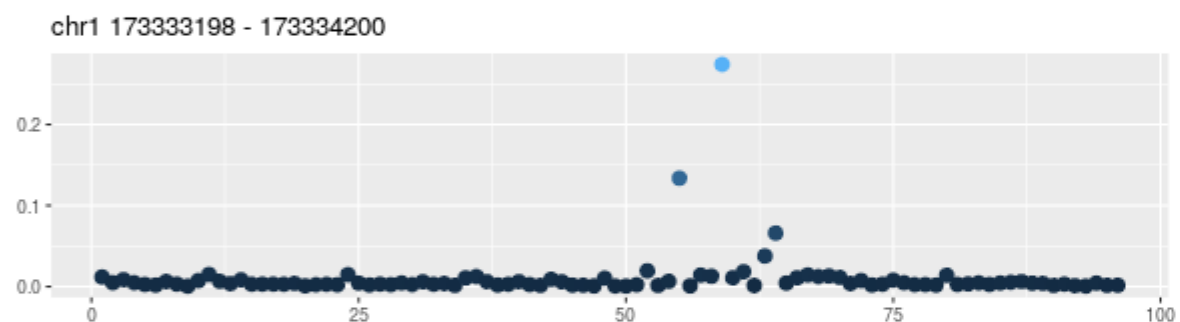
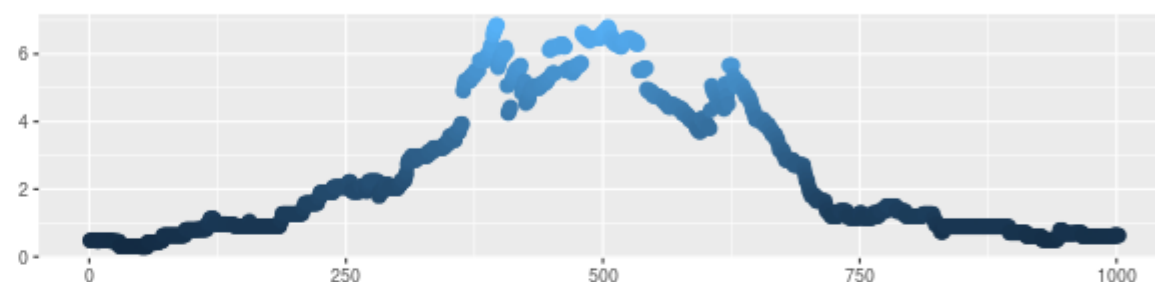
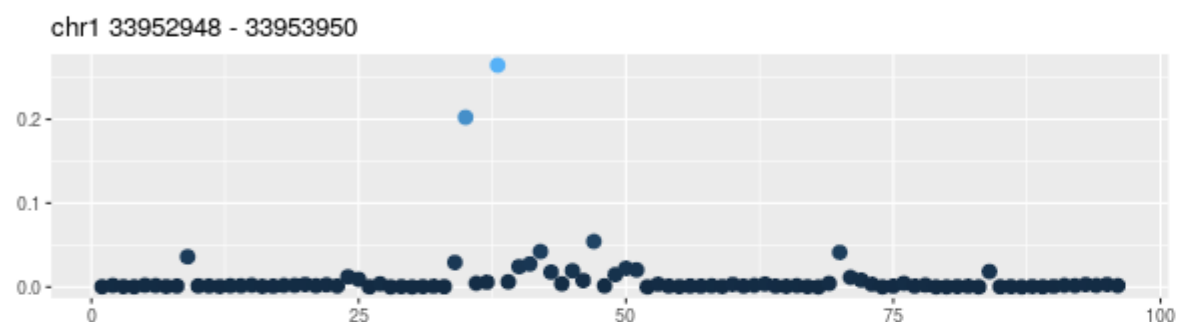
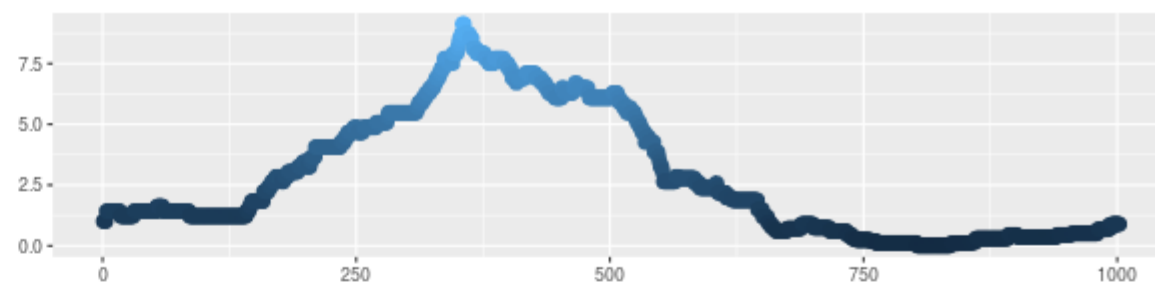
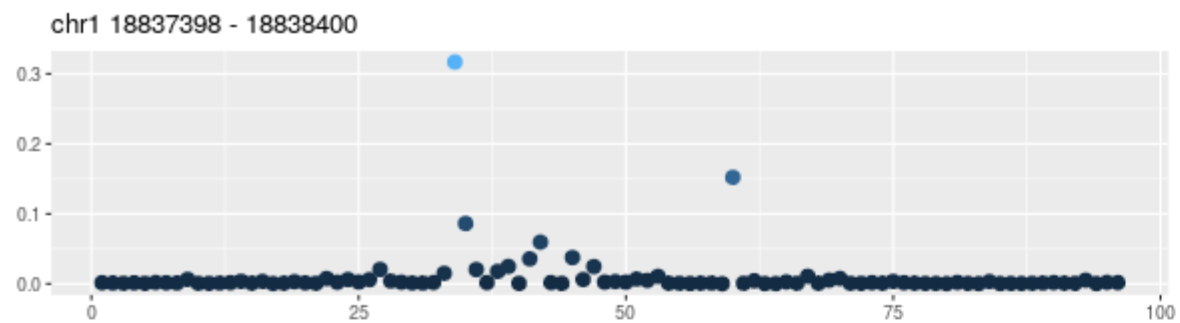


Attention weights

ChIP-seq score

Single attention module

CTCF/PC-3

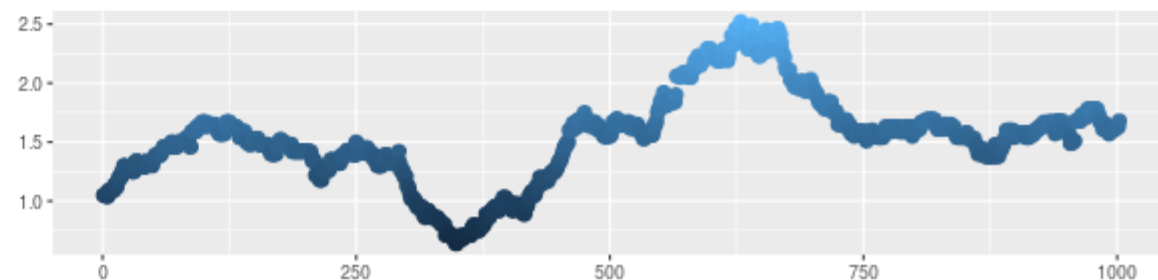
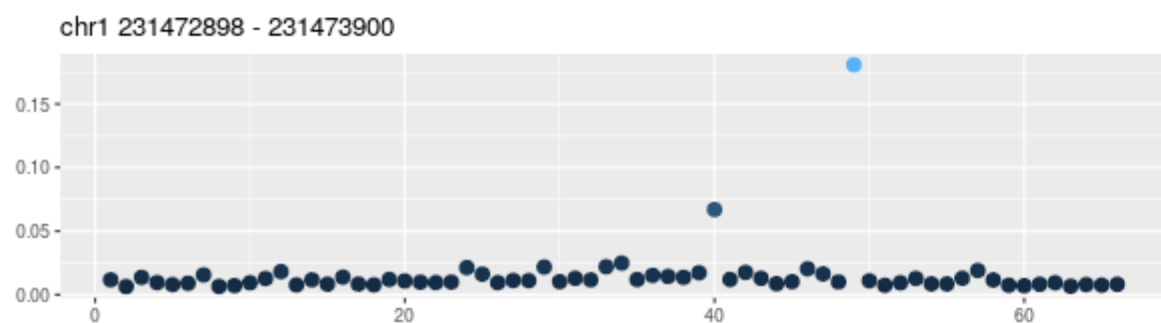
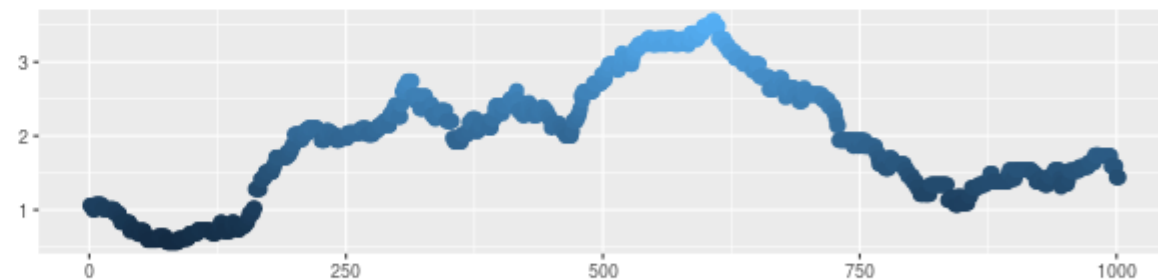
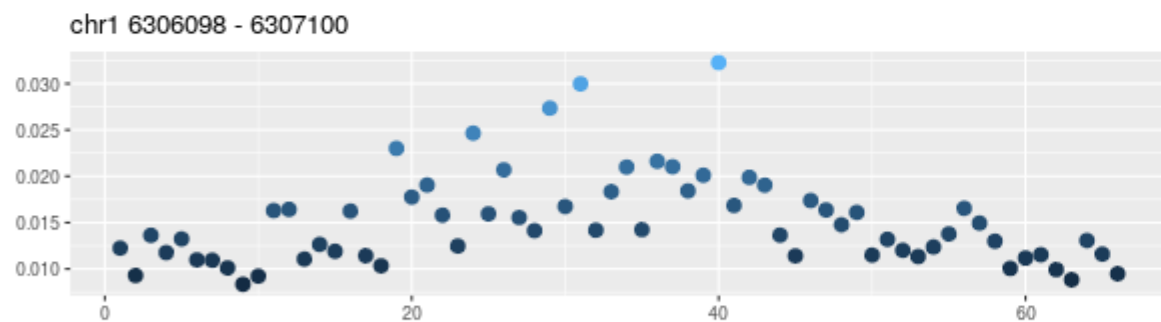
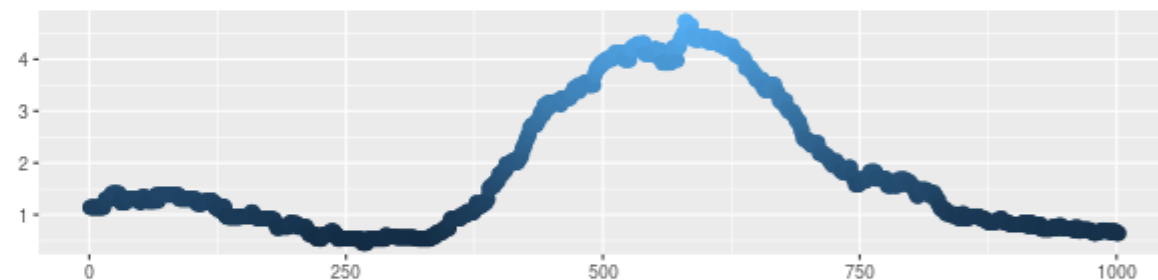
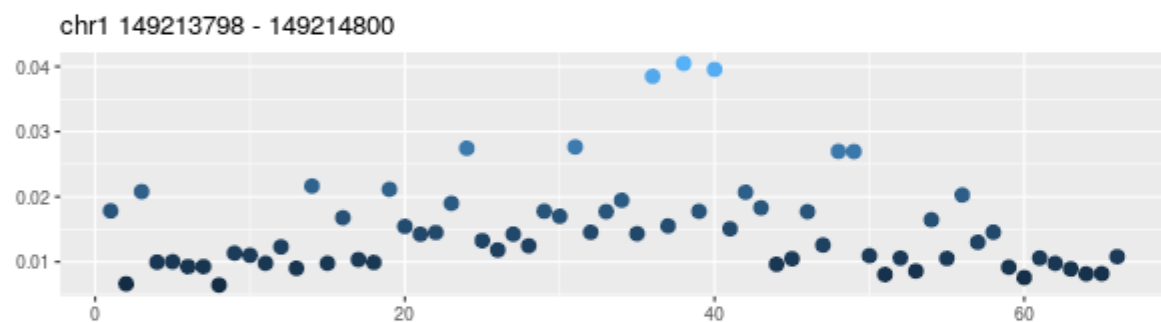
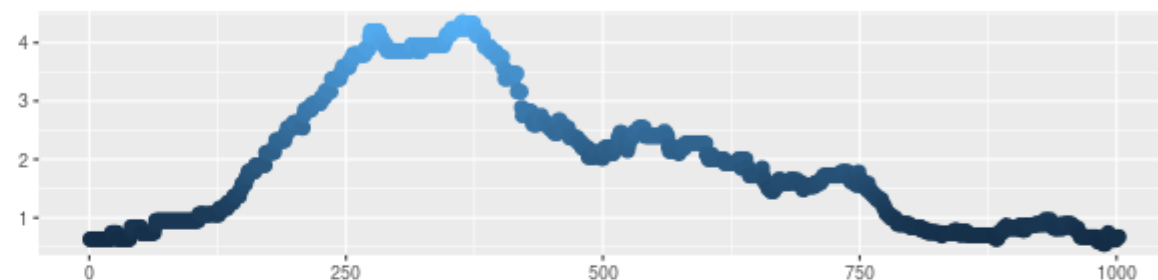
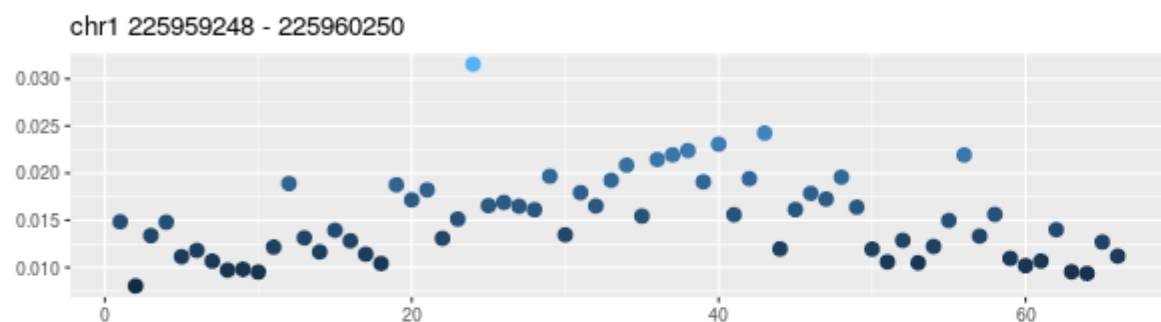
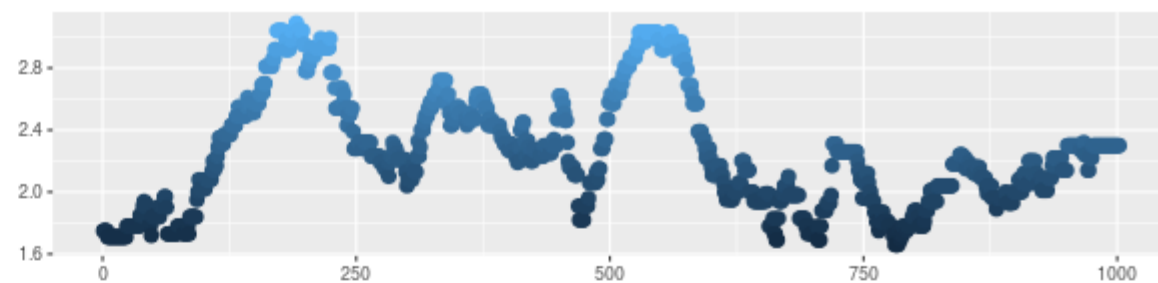
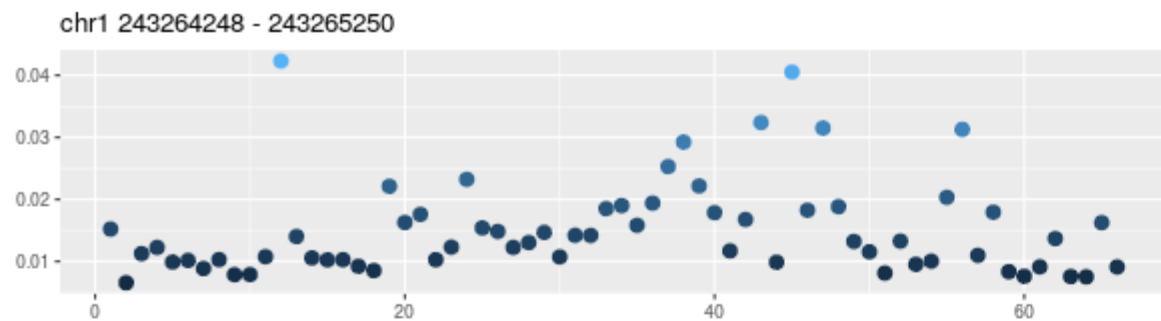


Attention weights

ChIP-seq score

Single attention module

E2F1/K562

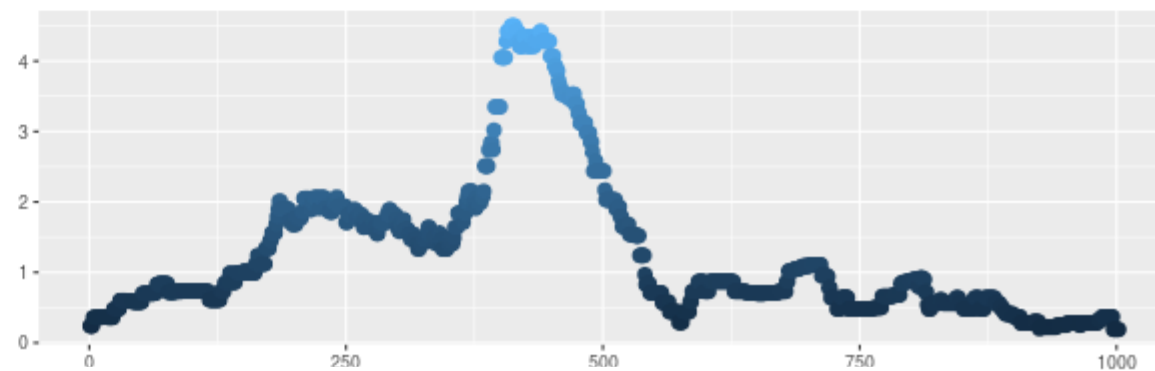
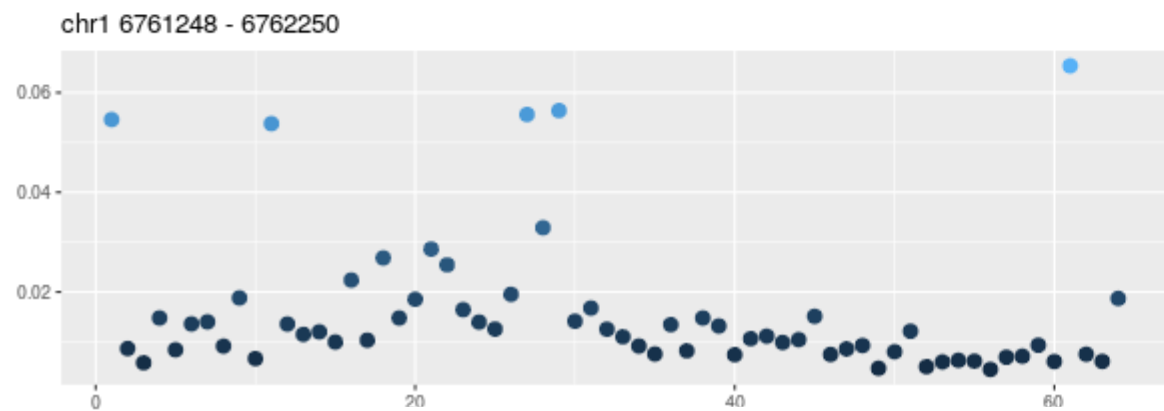
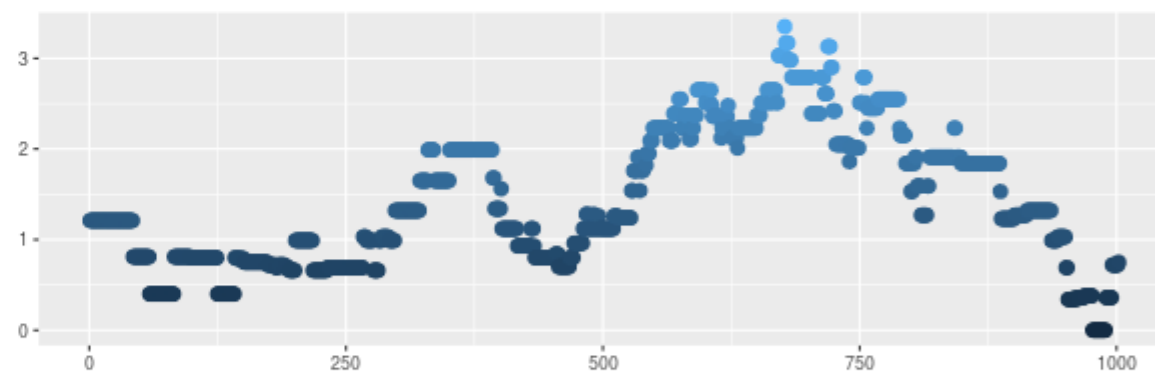
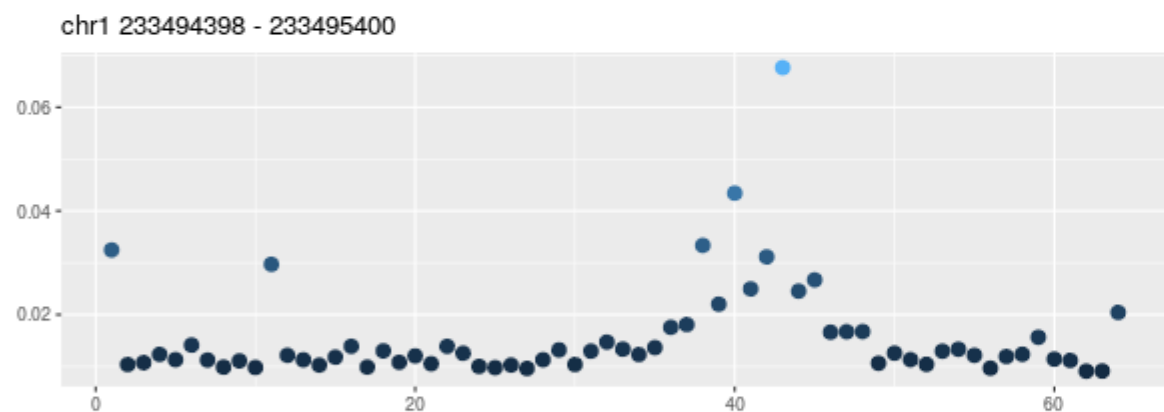
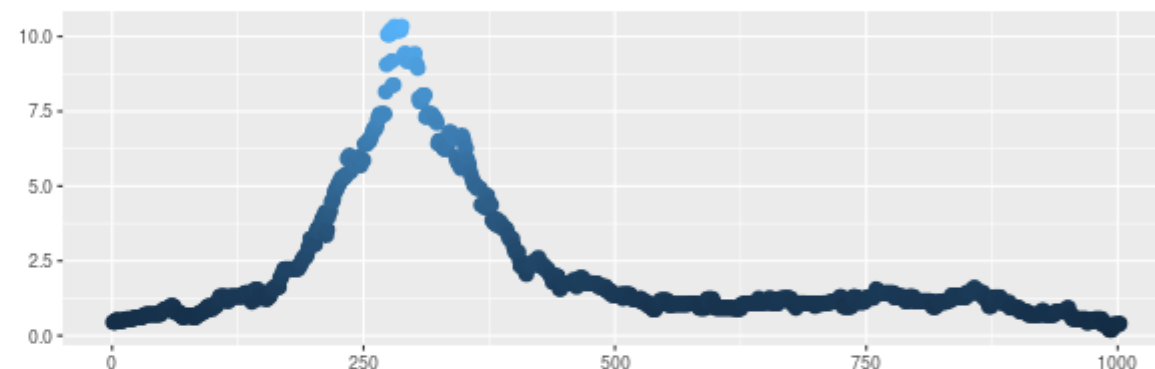
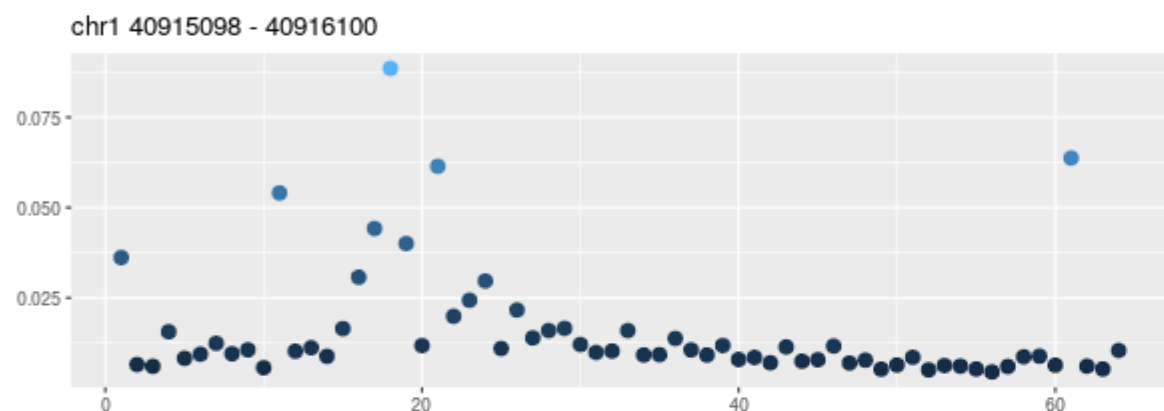
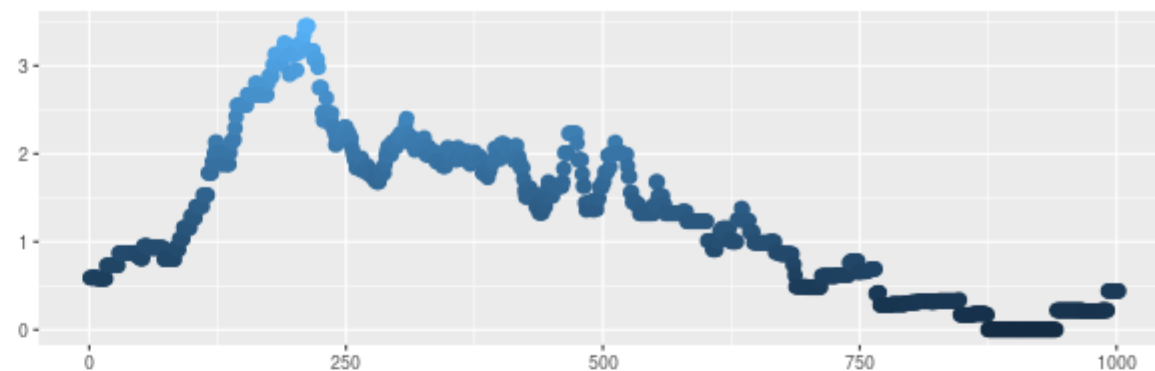
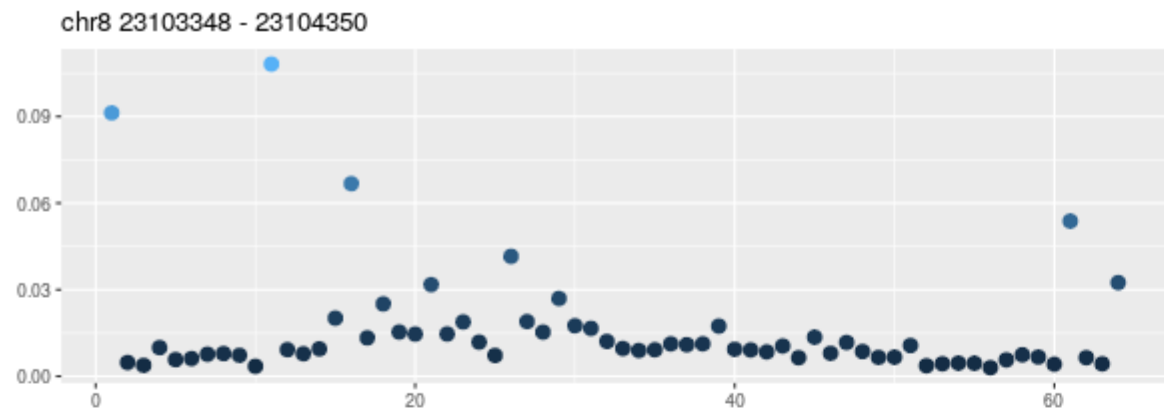


Attention weights

CHIP-seq score

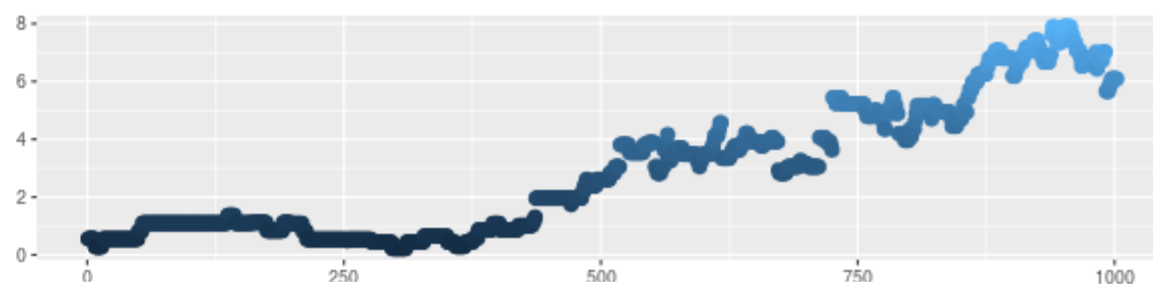
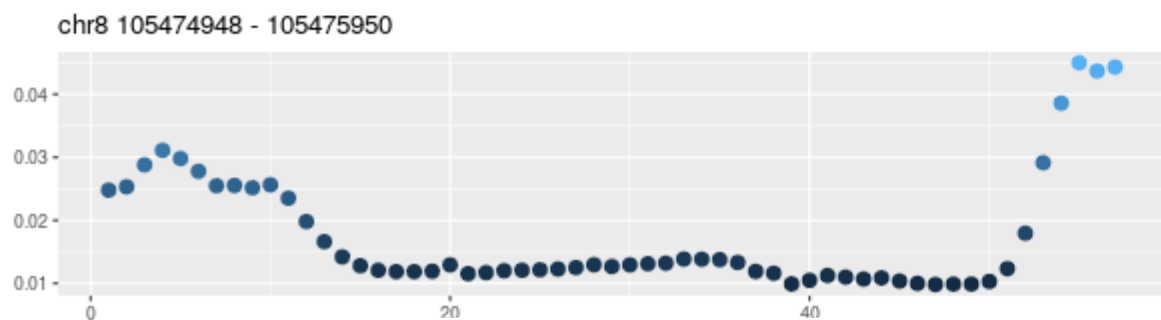
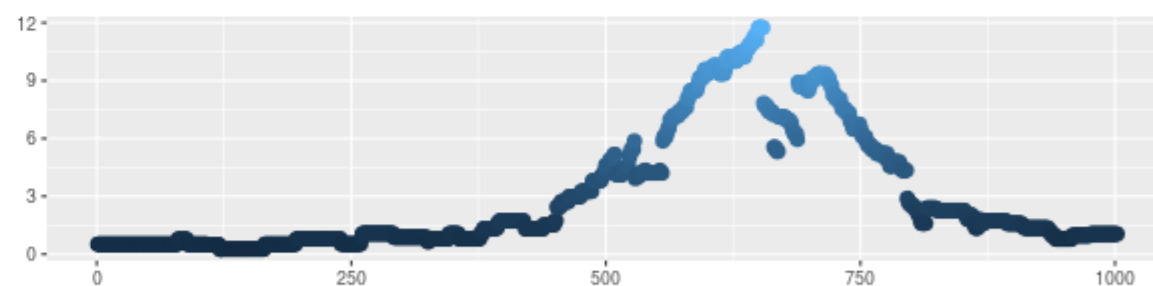
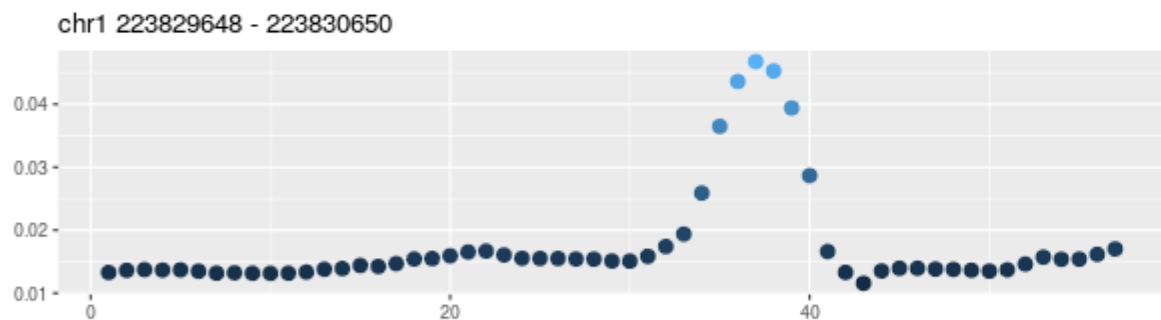
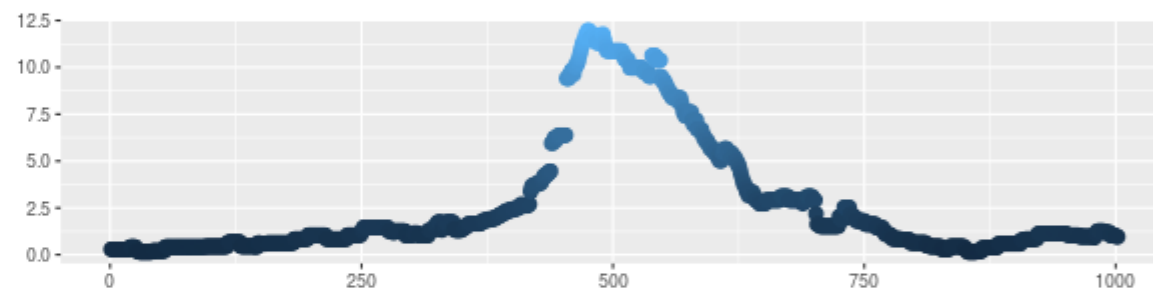
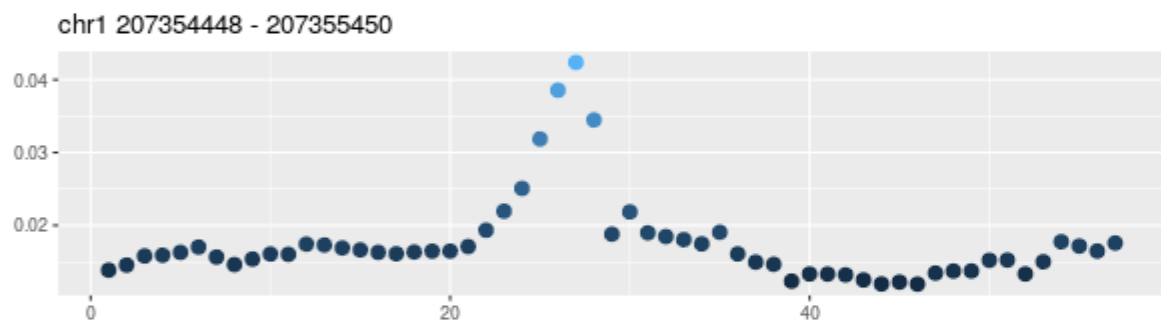
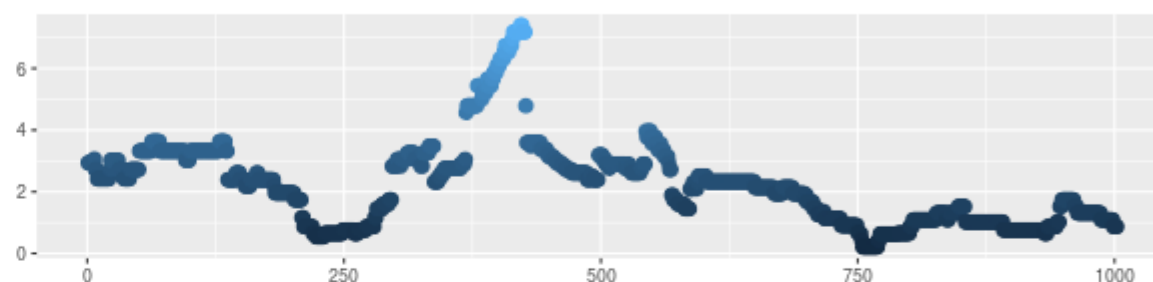
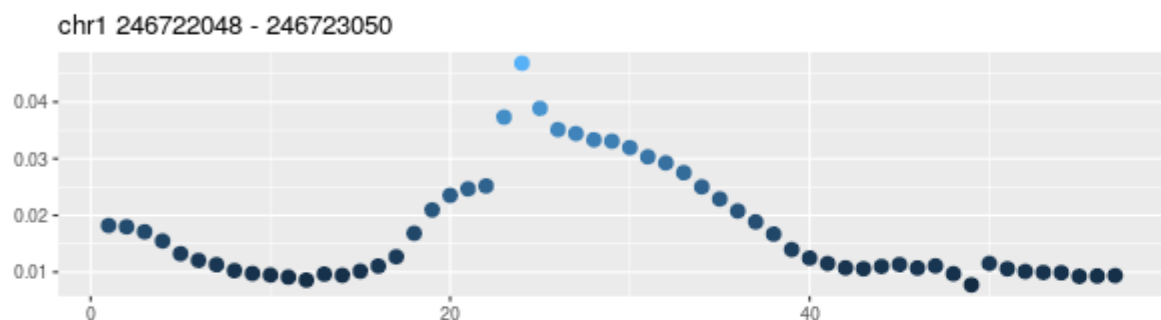
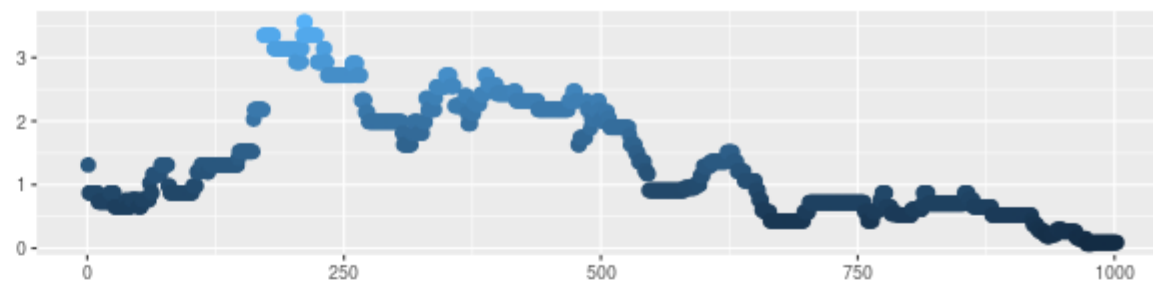
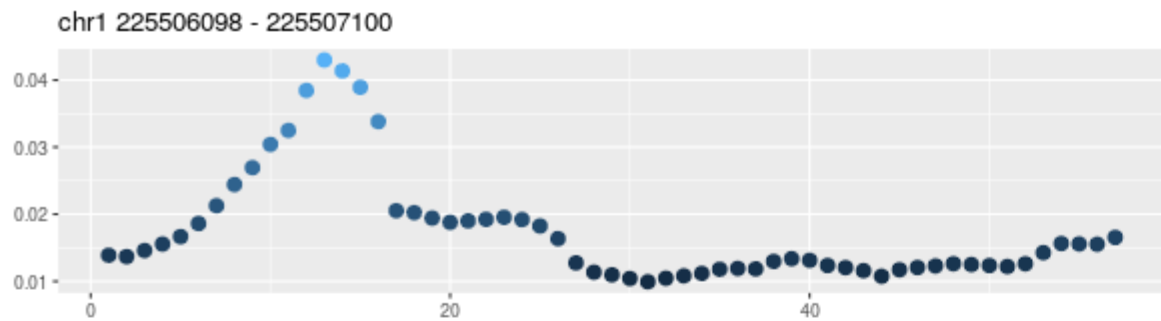
Single attention module

EGR1/liver



Single attention module

FOXA1/liver

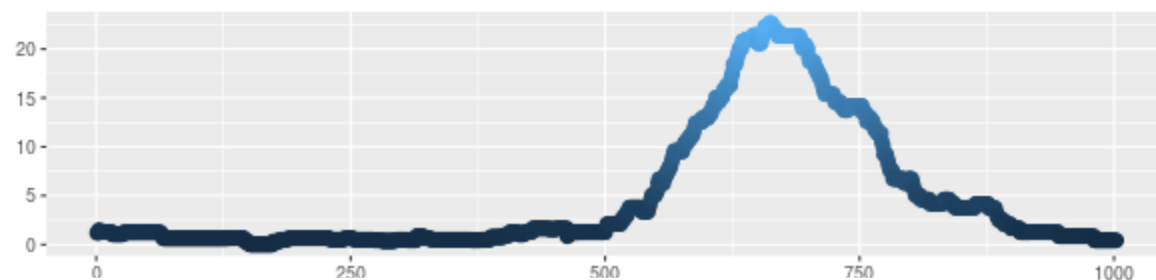
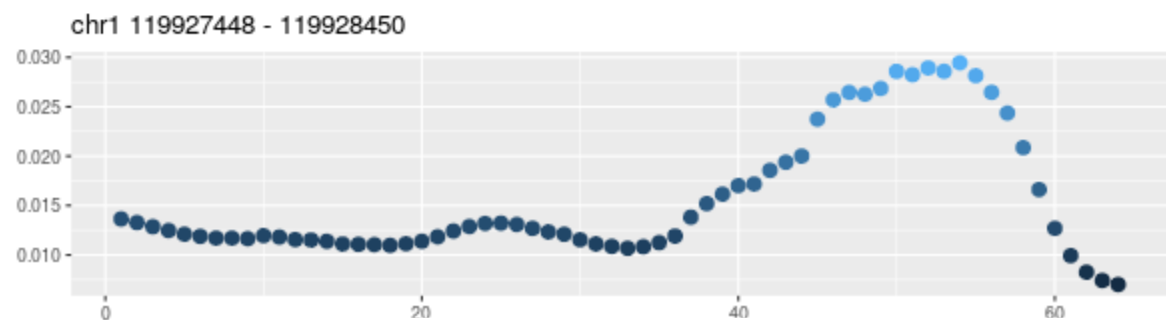
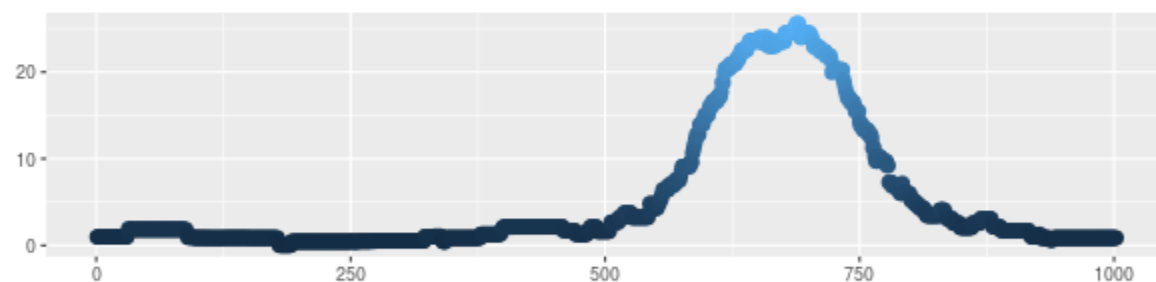
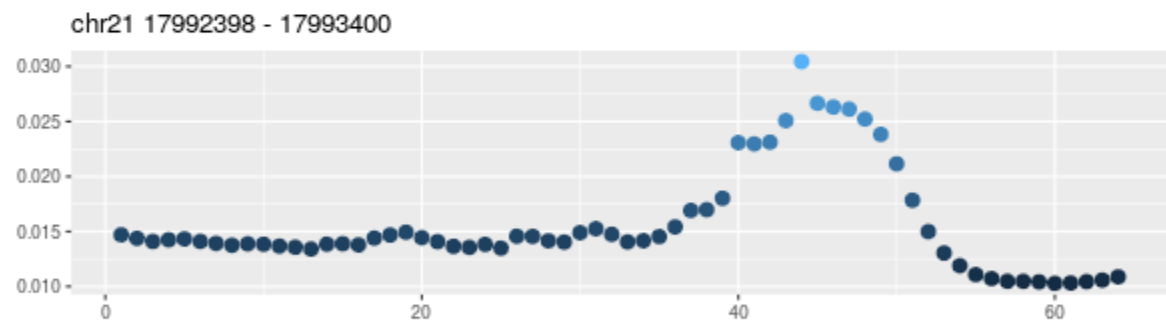
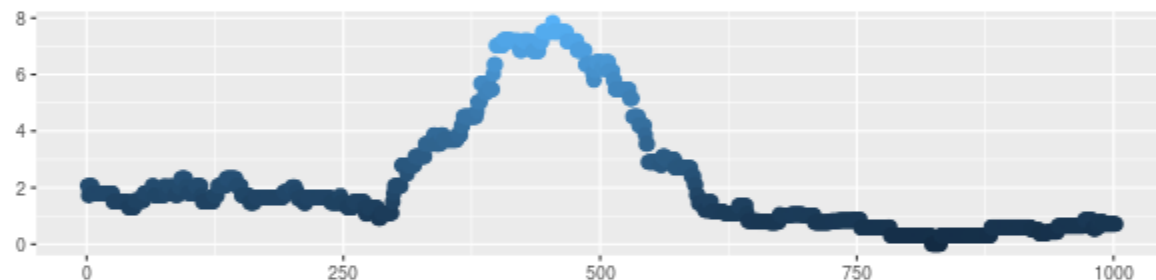
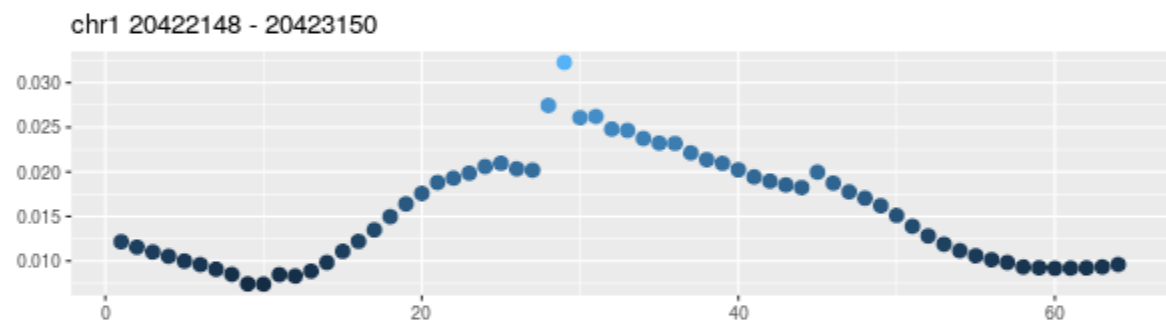
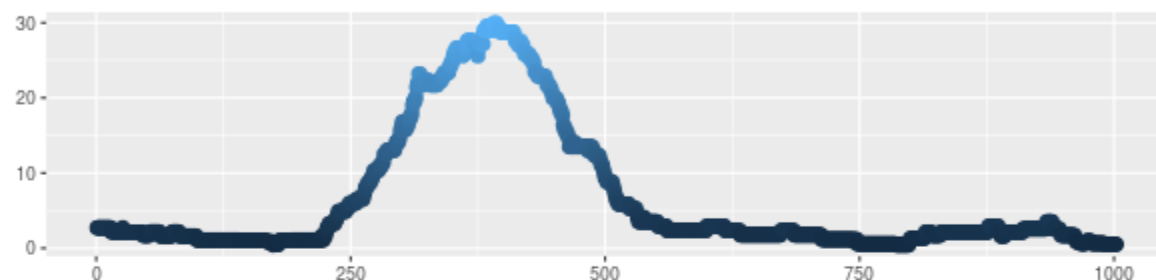
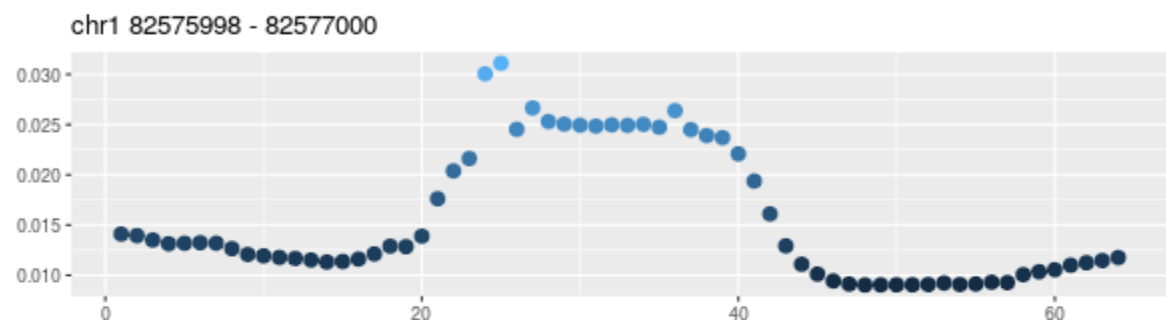
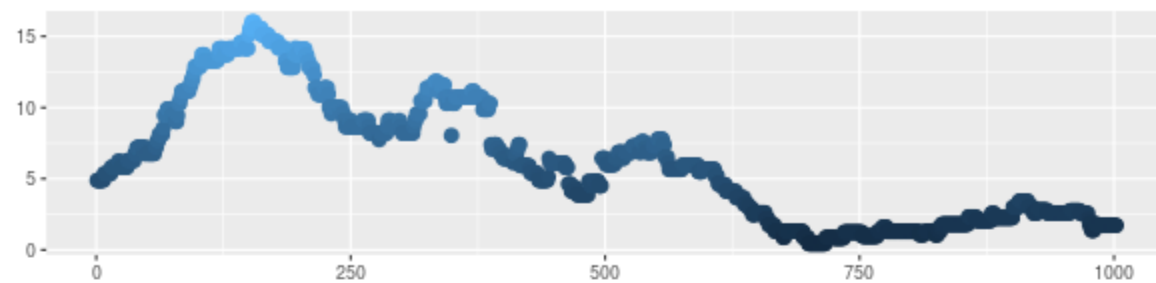
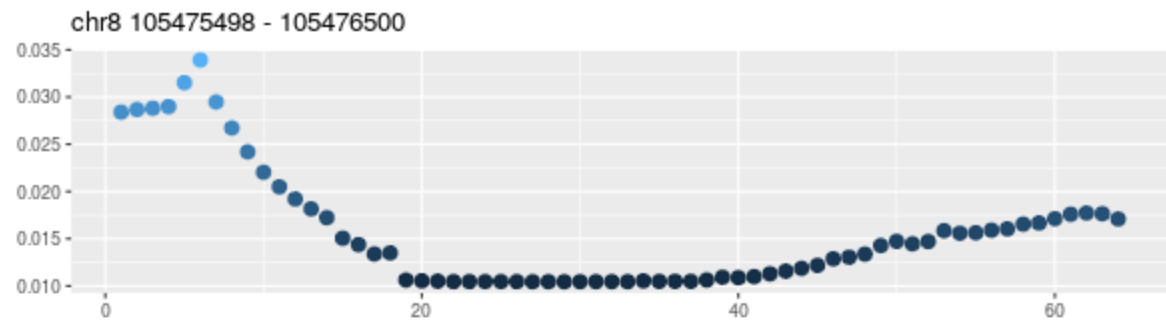


Attention weights

ChIP-seq score

Single attention module

FOXA2/liver

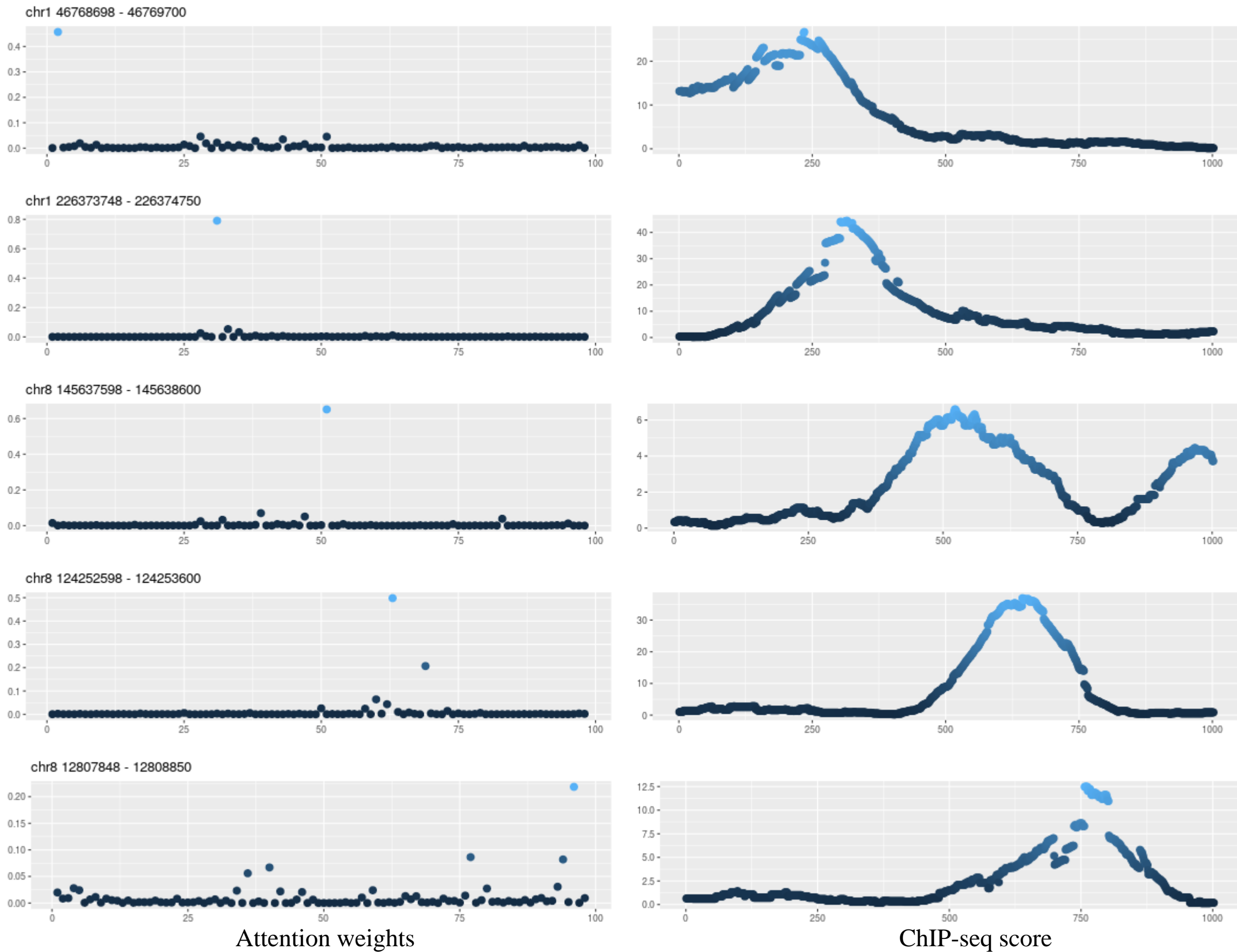


Attention weights

ChIP-seq score

Single attention module

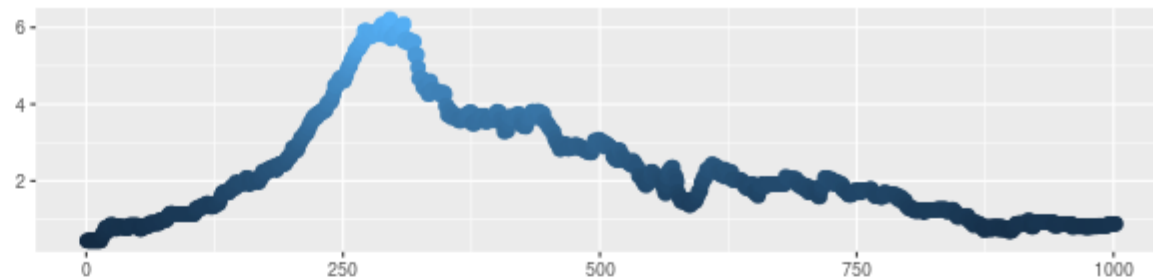
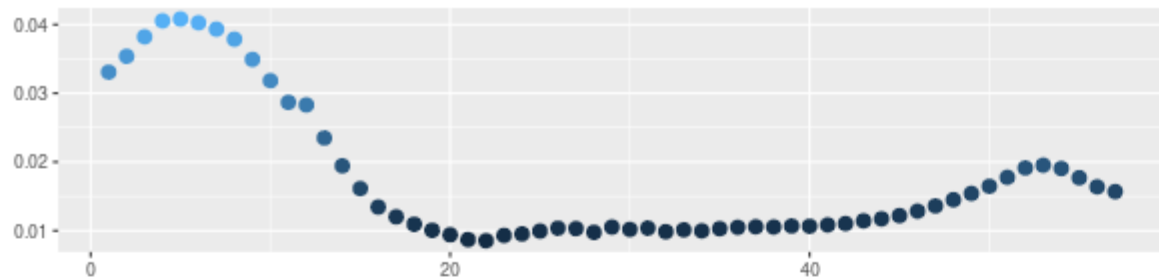
GABPA/liver



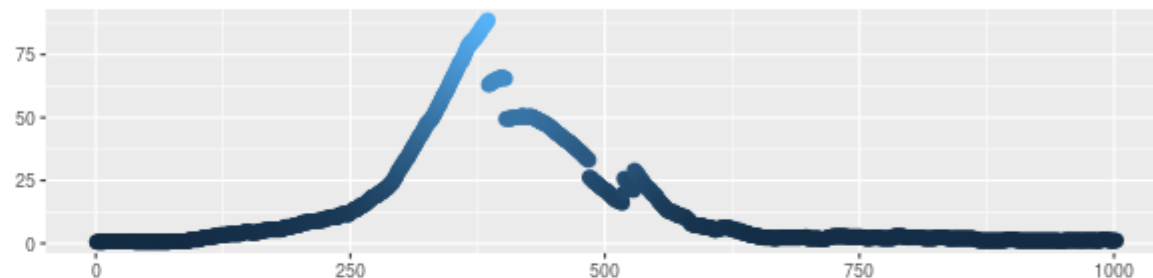
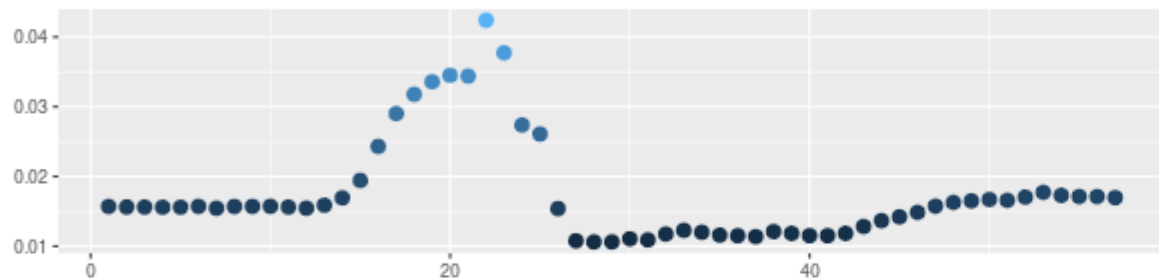
Single attention module

HNF4A/liver

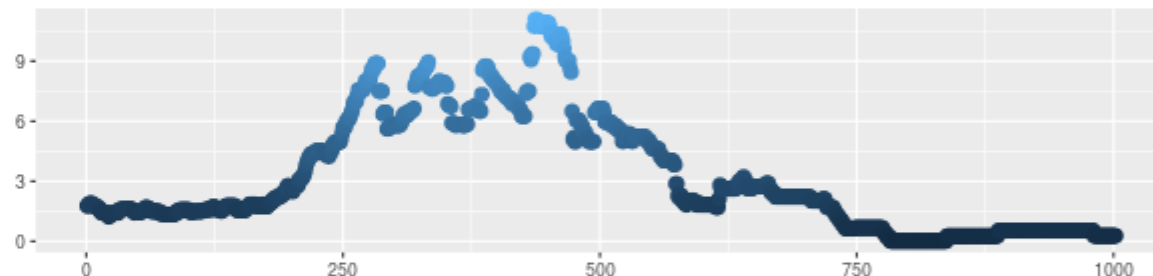
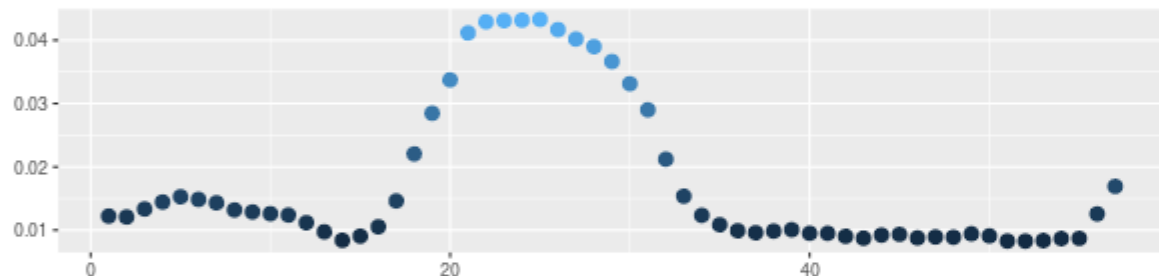
chr8 23103248 - 23104250



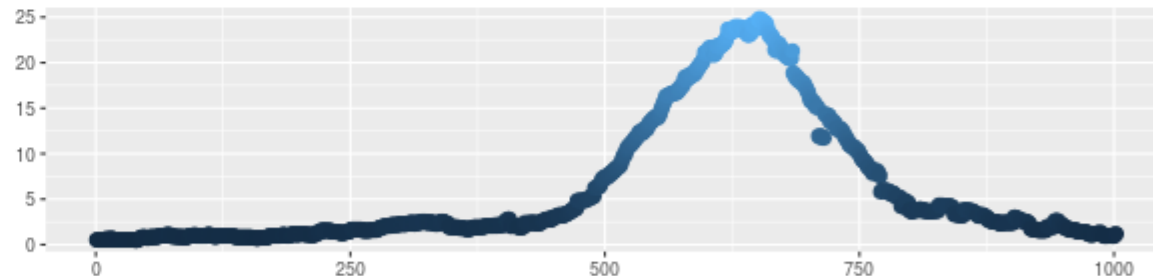
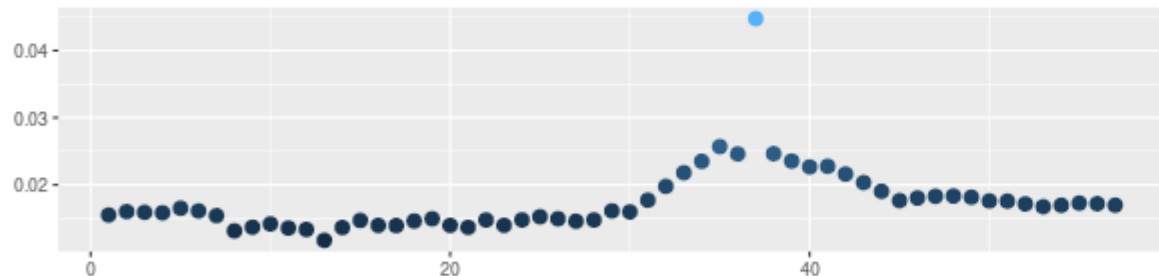
chr1 187724048 - 187725050



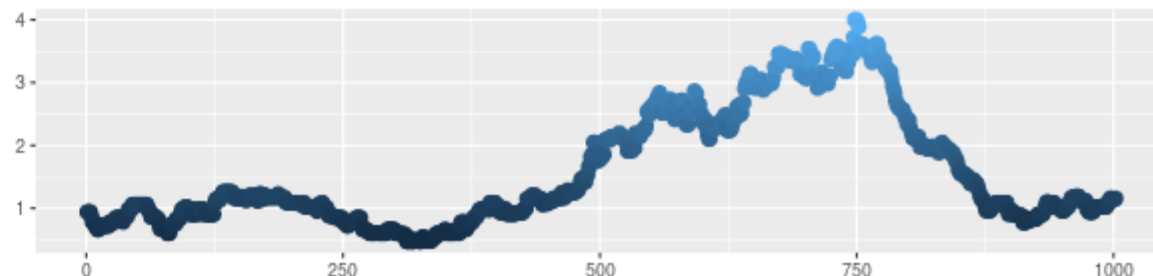
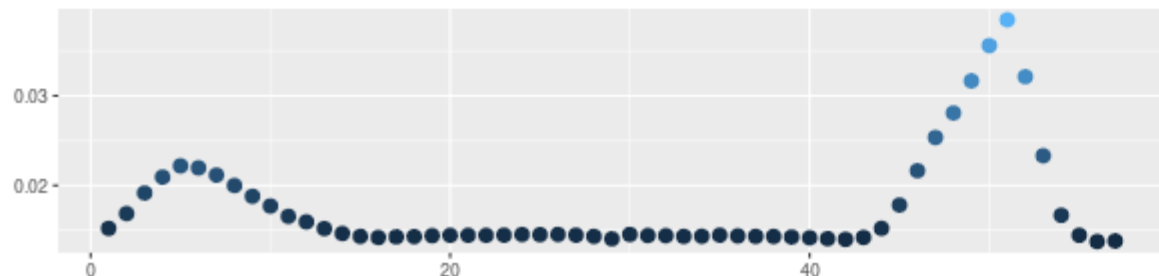
chr1 150382548 - 150383550



chr1 227281448 - 227282450



chr1 118471248 - 118472250



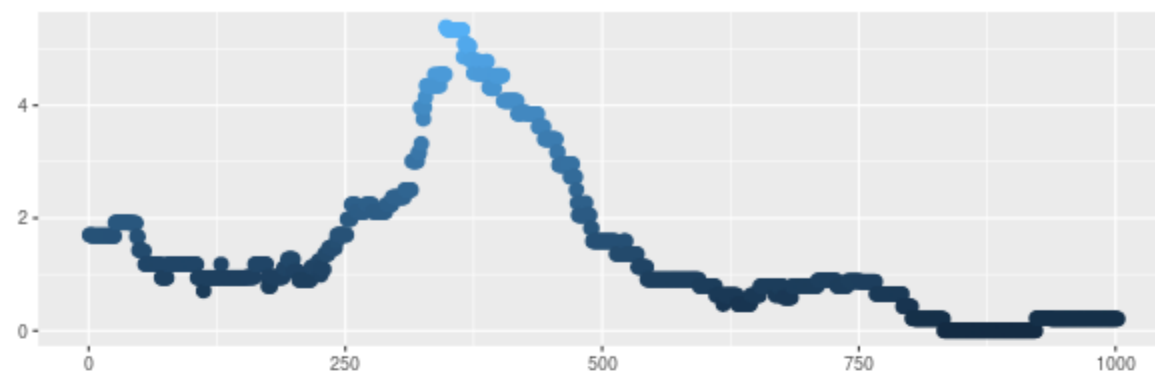
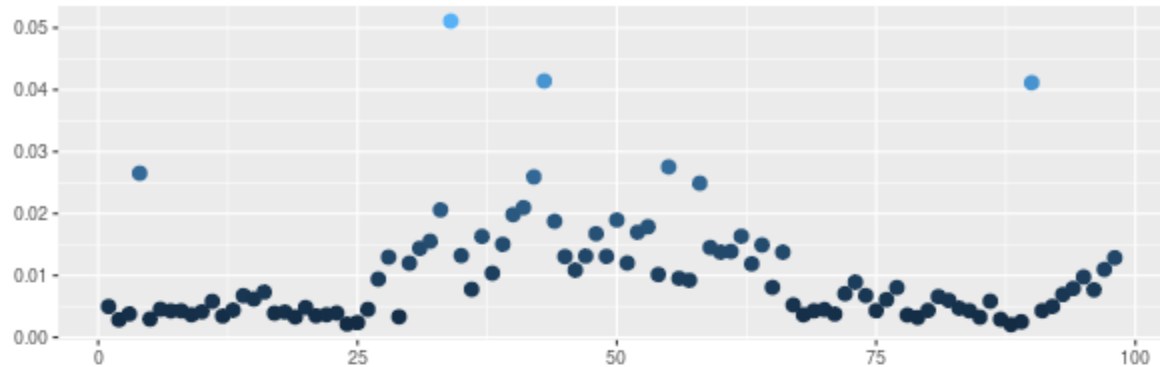
Attention weights

ChIP-seq score

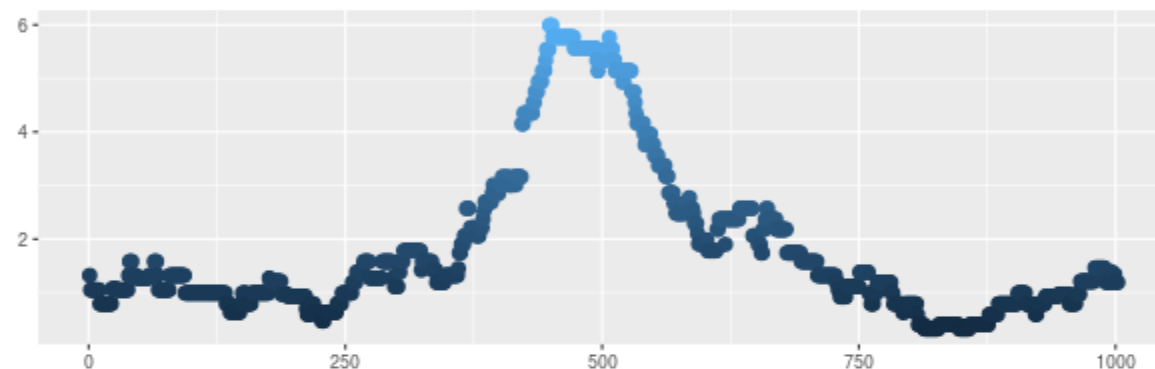
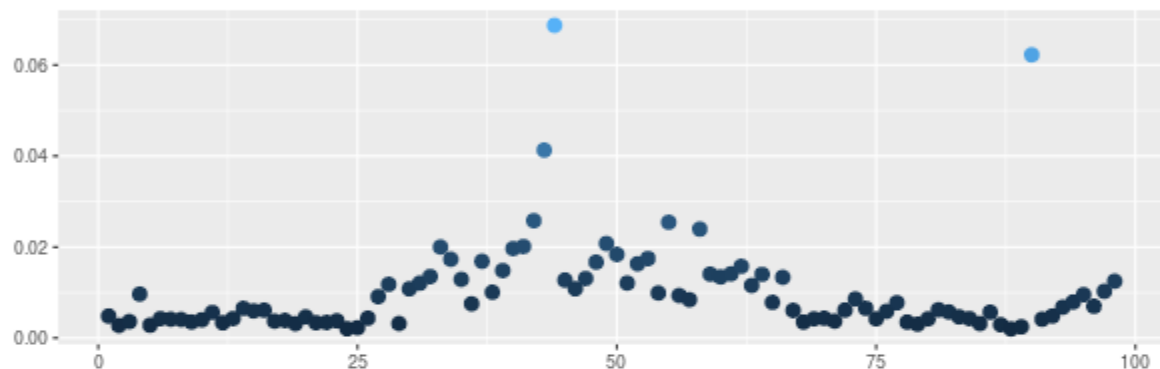
Single attention module

JUND/liver

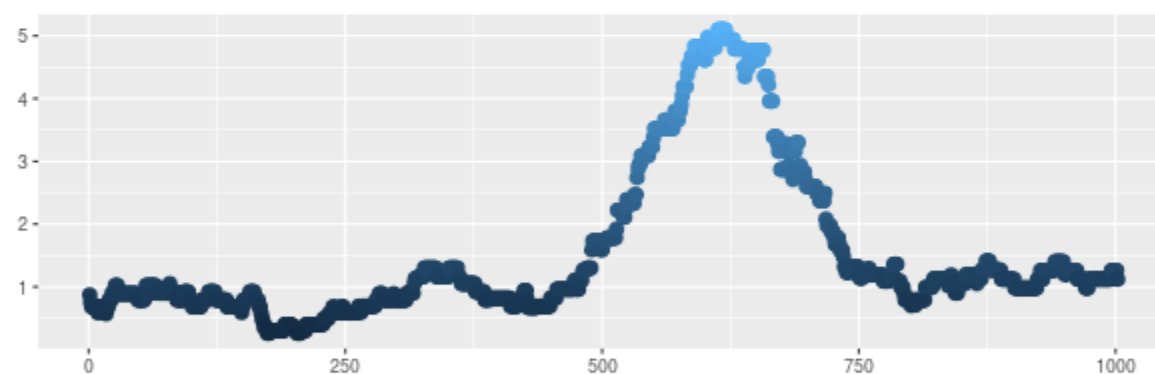
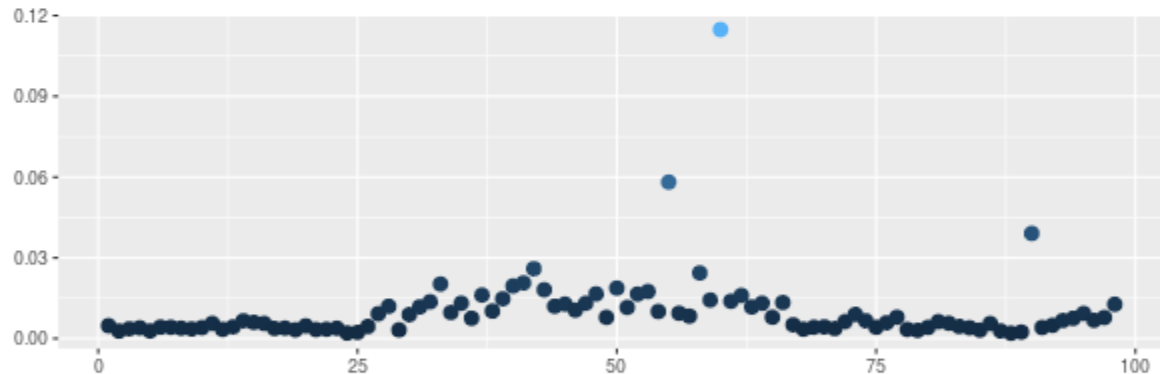
chr1 227052748 - 227053750



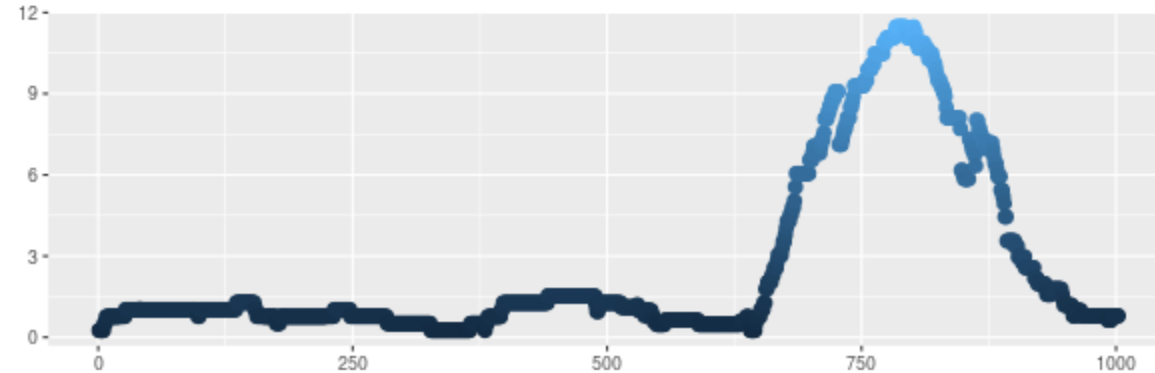
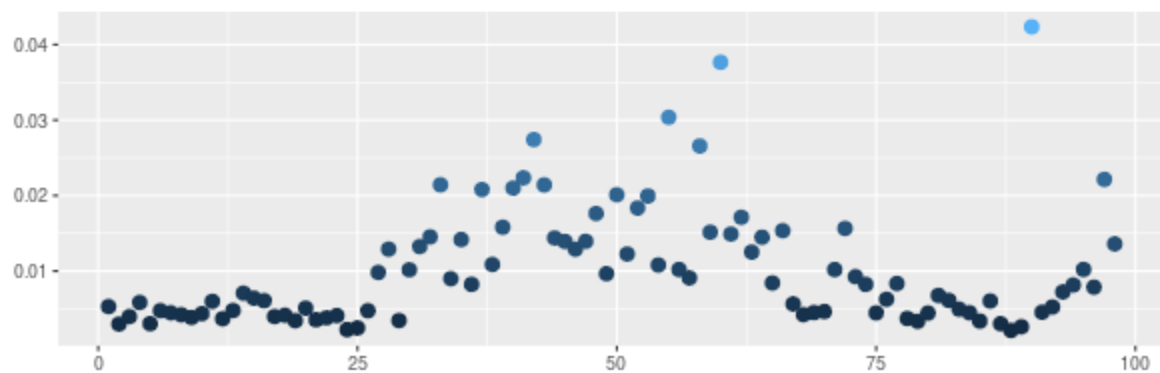
chr1 19528148 - 19529150



chr21 44153548 - 44154550



chr8 67683248 - 67684250

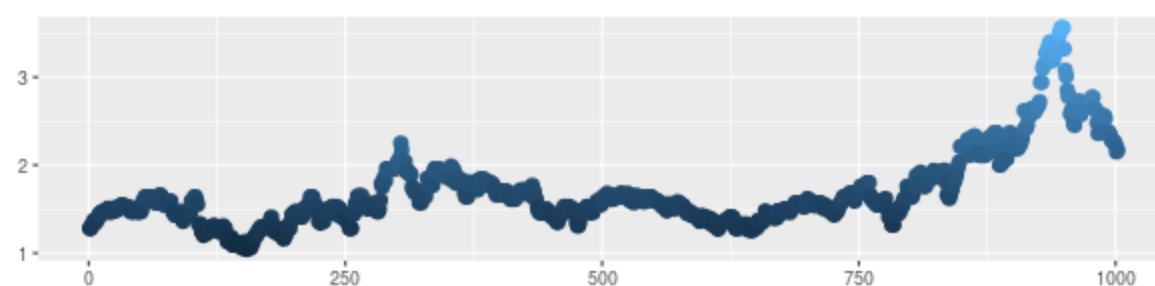
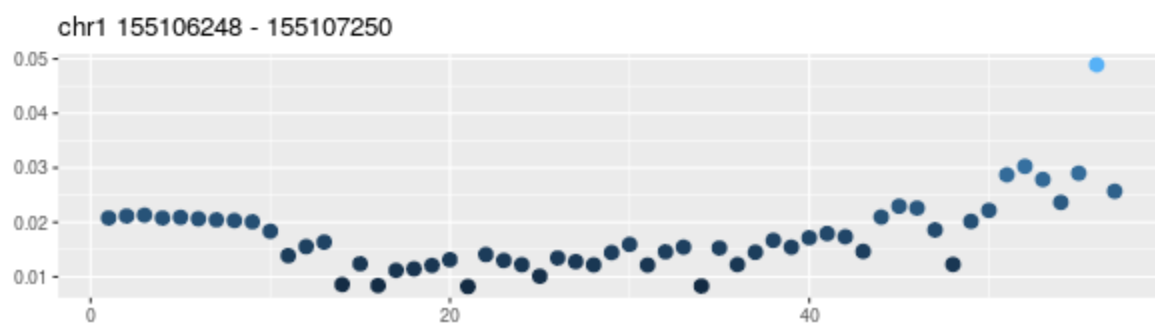
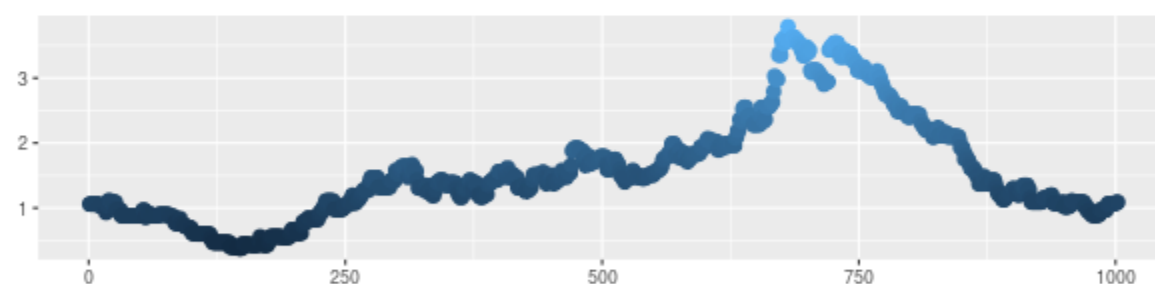
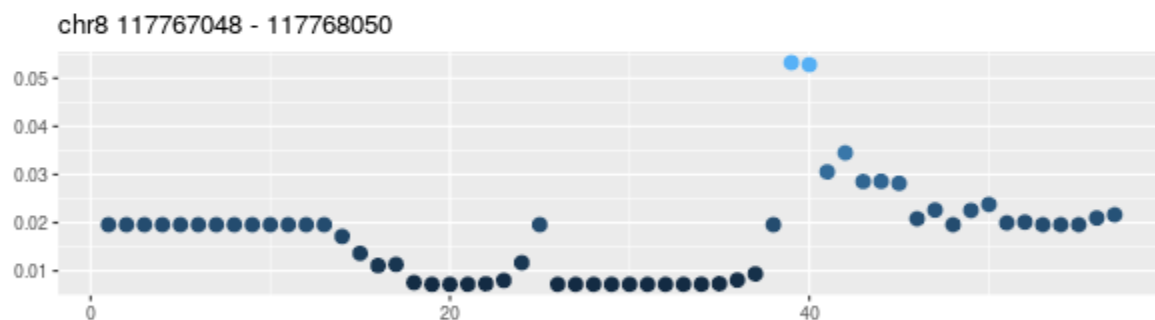
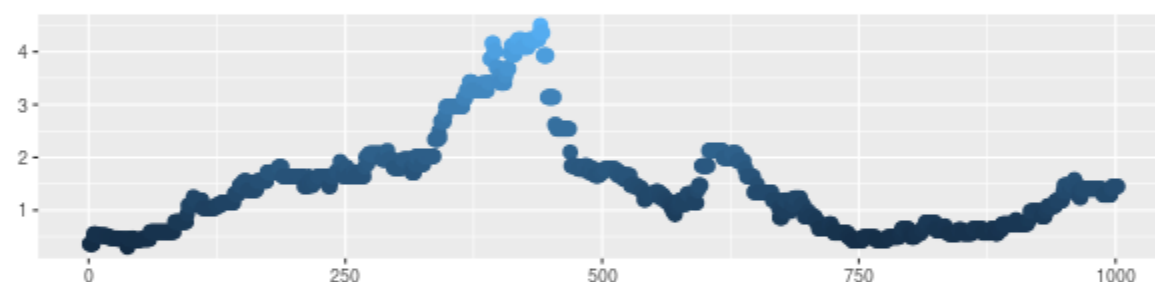
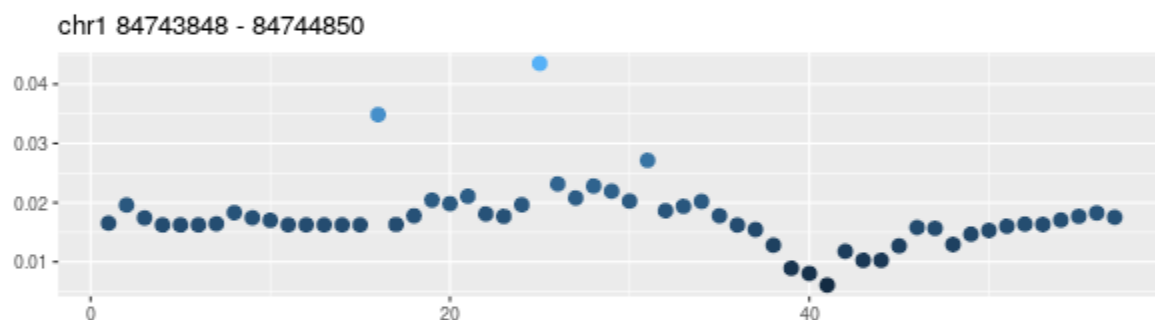
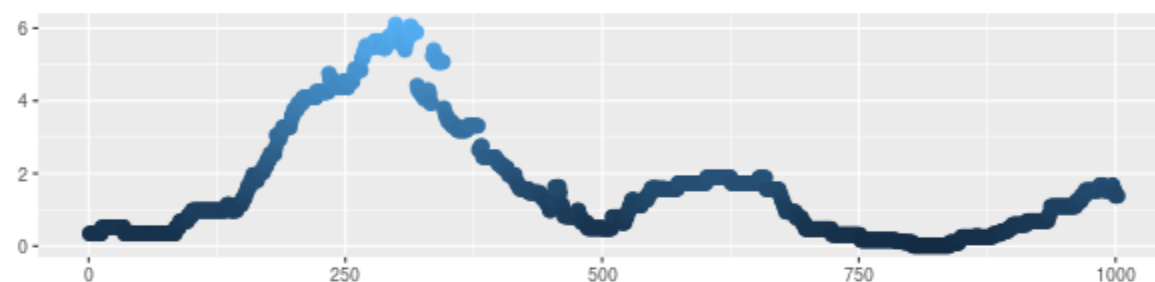
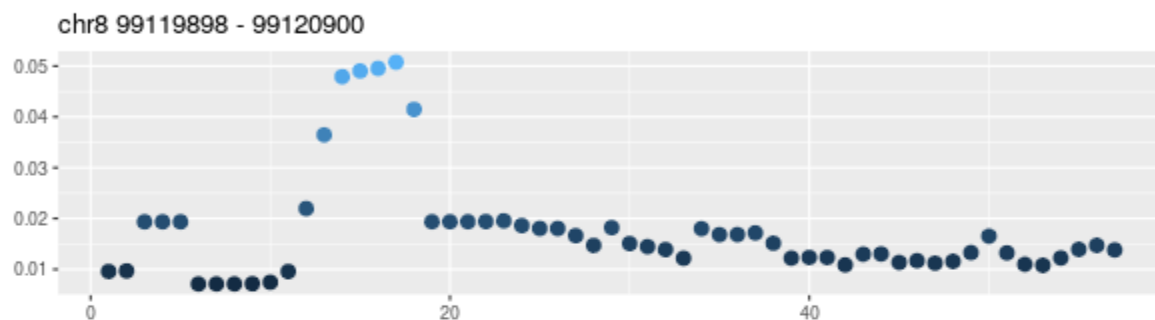
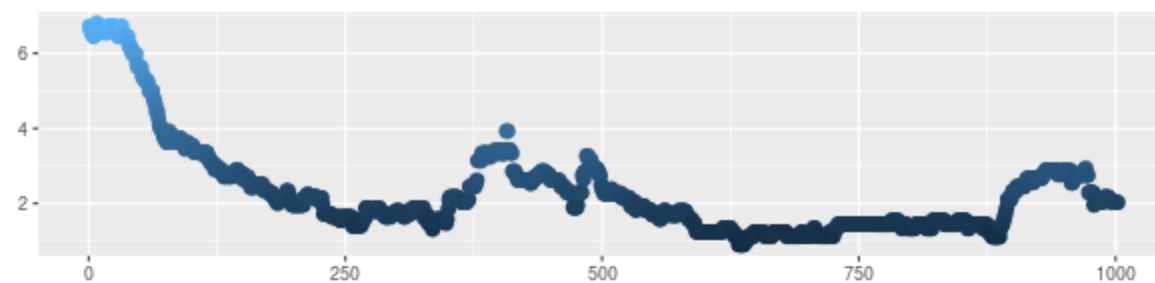
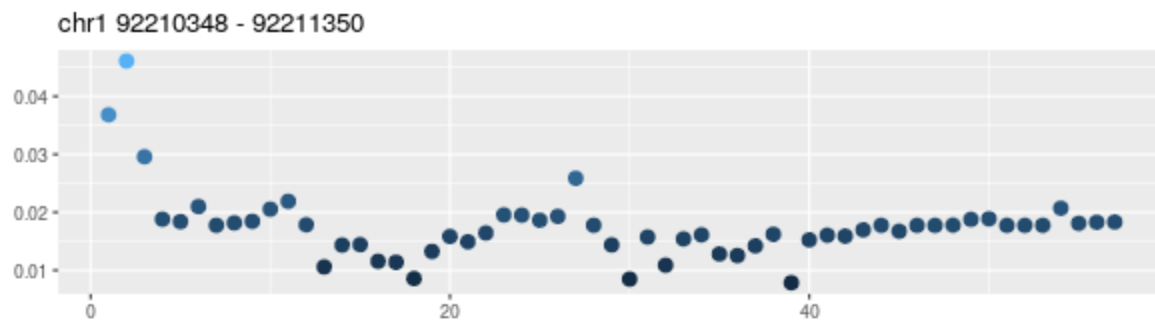


Attention weights

CHIP-seq score

Single attention module

MAX/liver

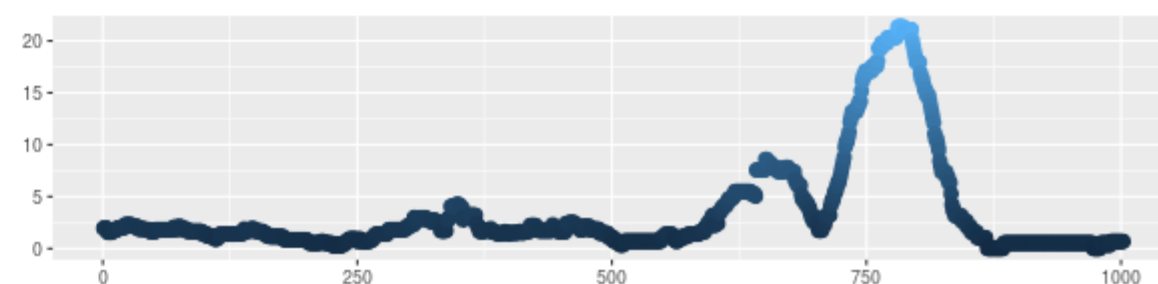
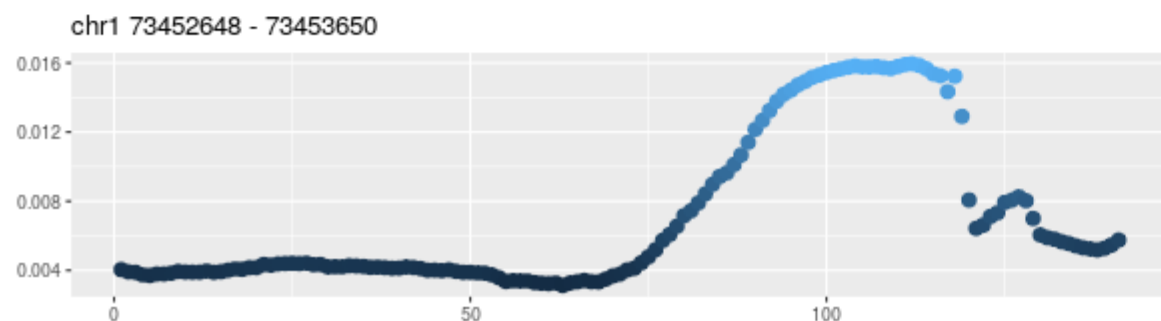
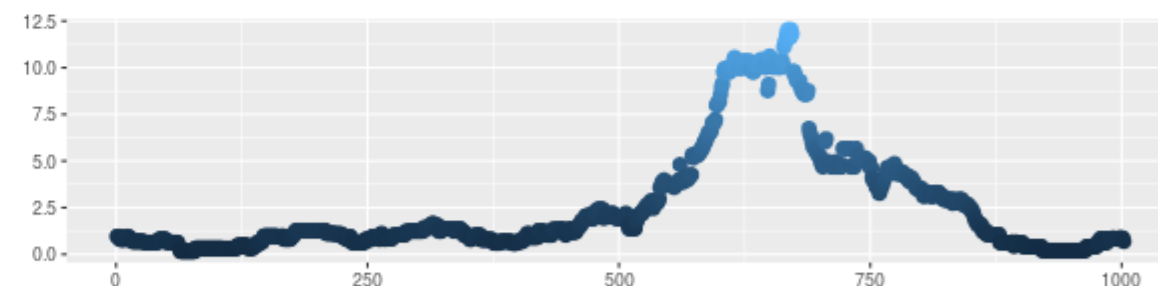
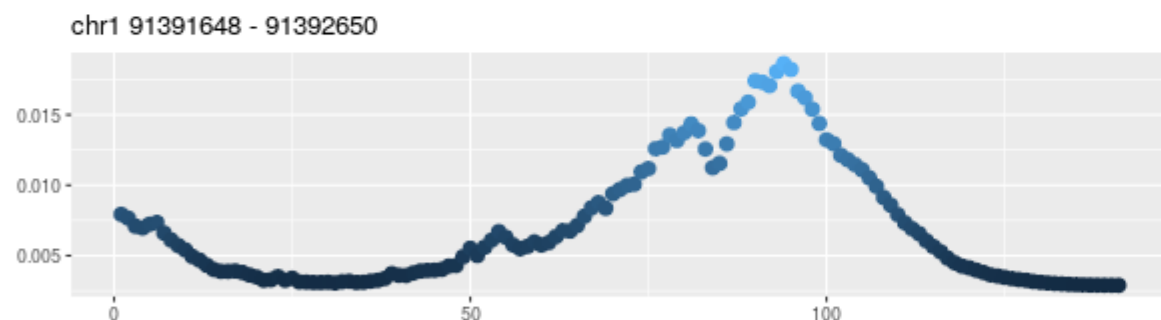
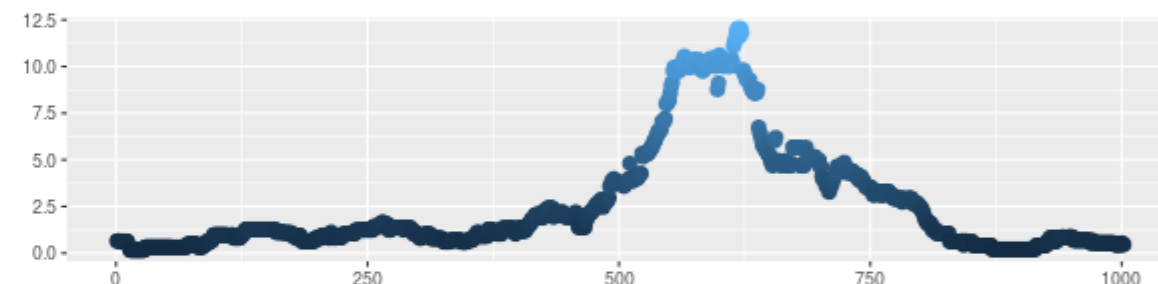
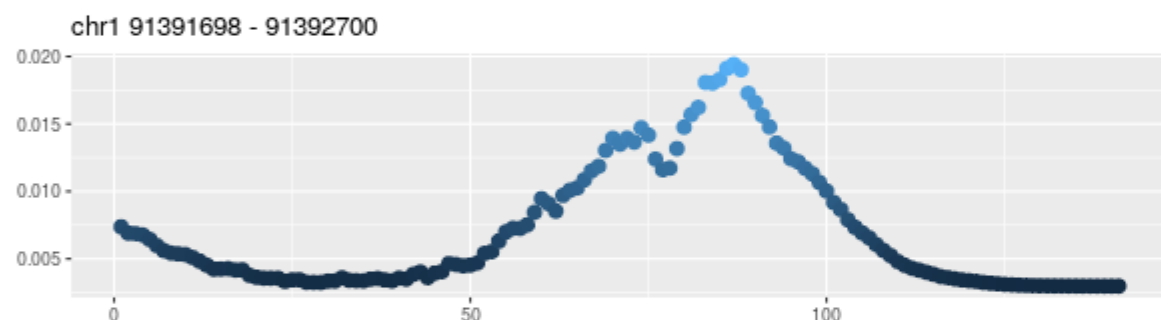
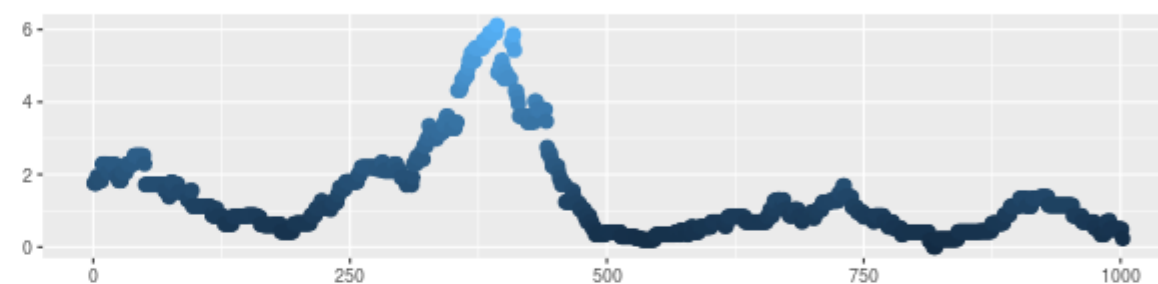
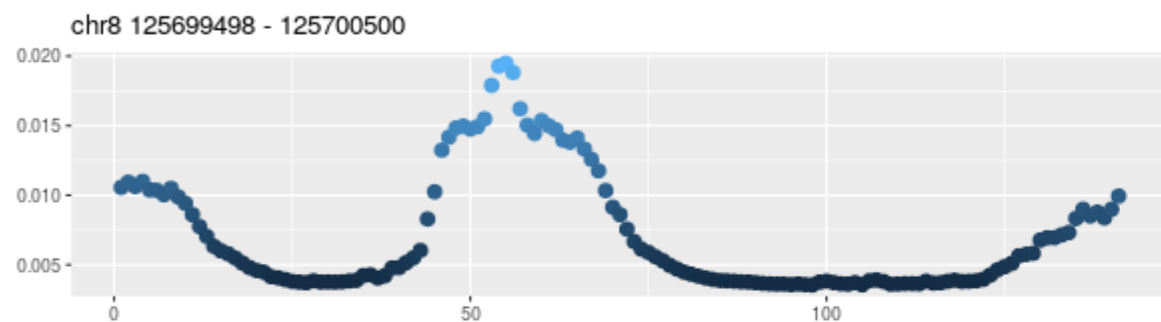
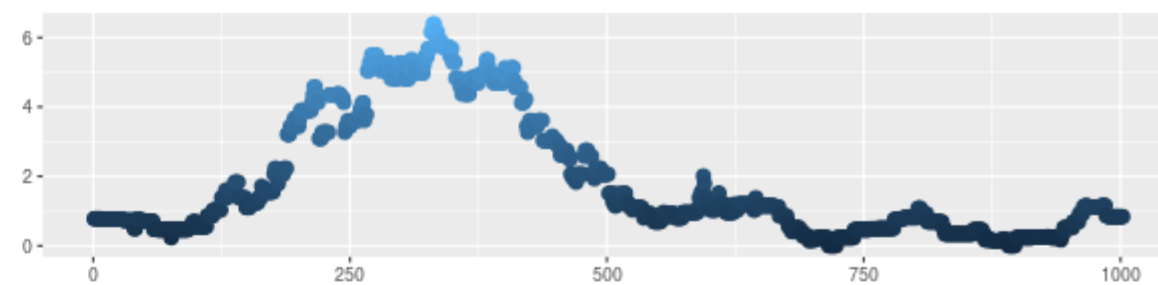
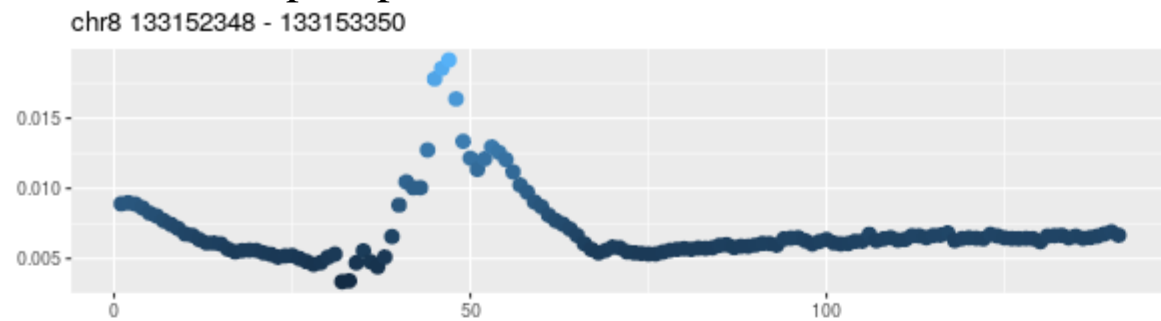


Attention weights

CHIP-seq score

Single attention module

NANOG/induced pluripotent stem cell

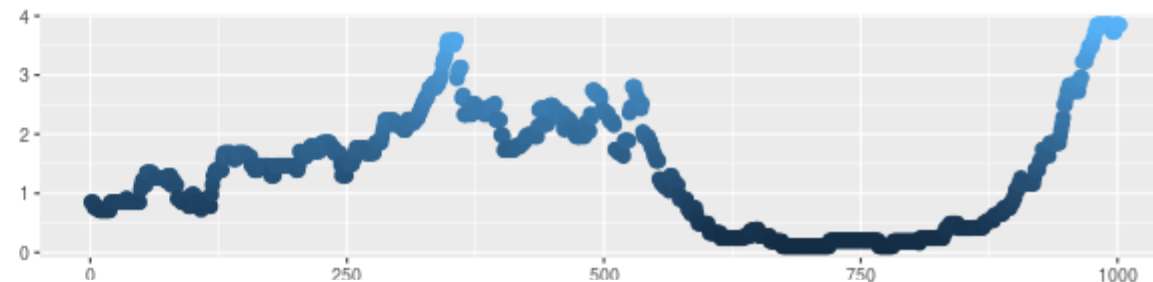
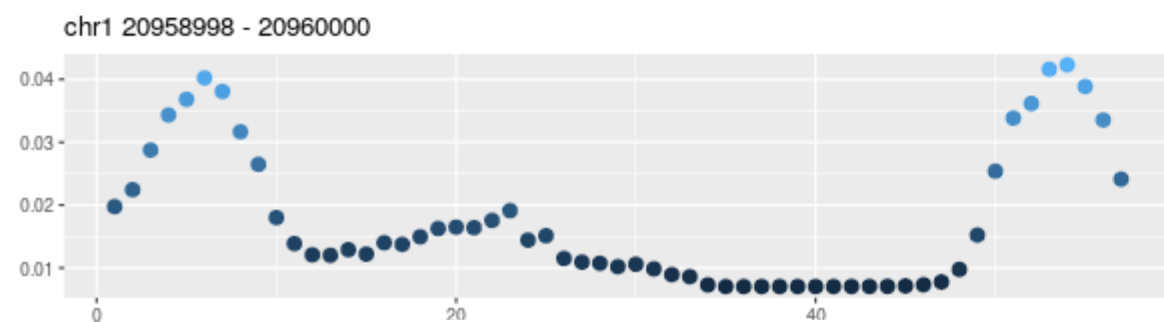
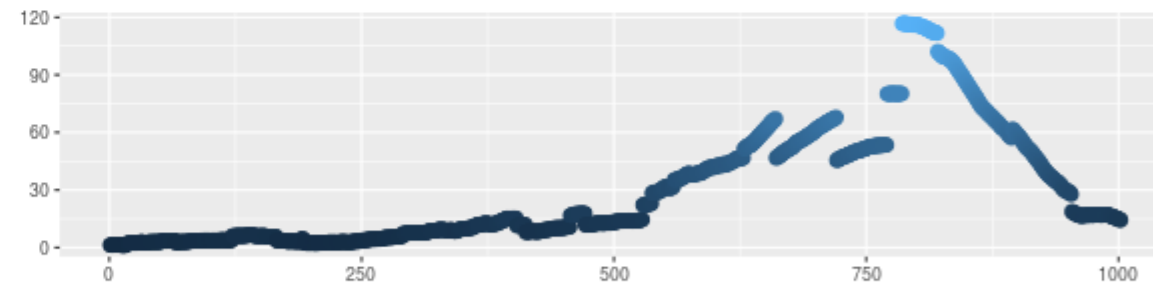
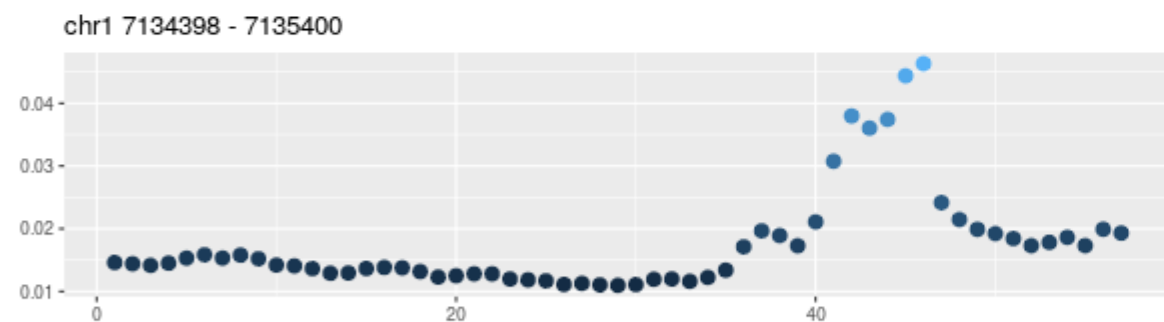
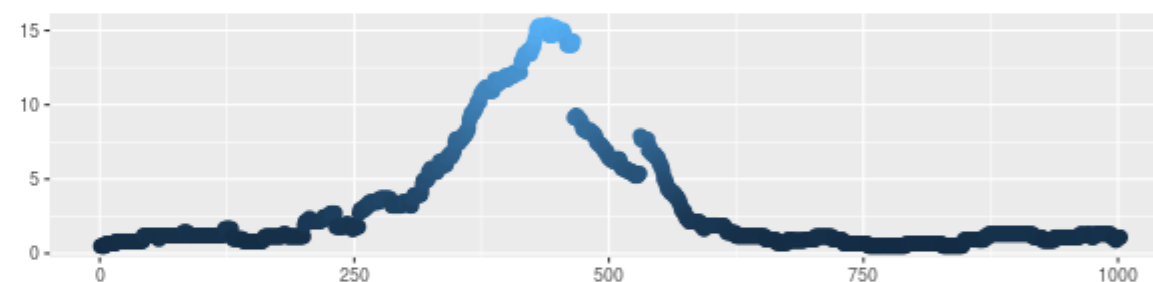
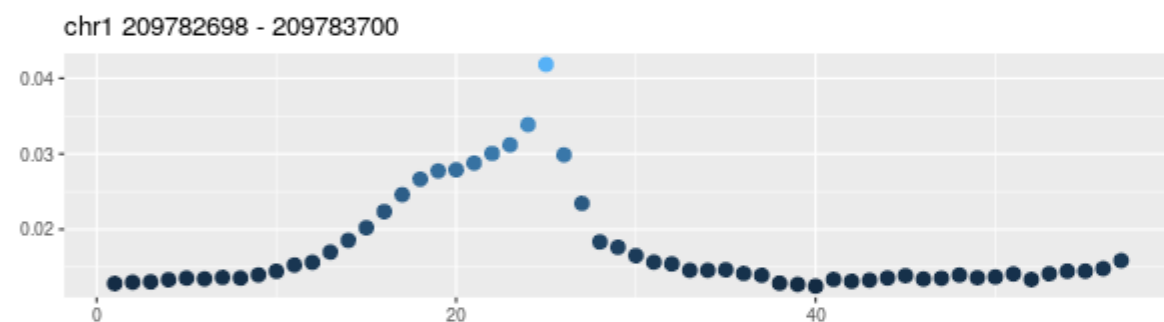
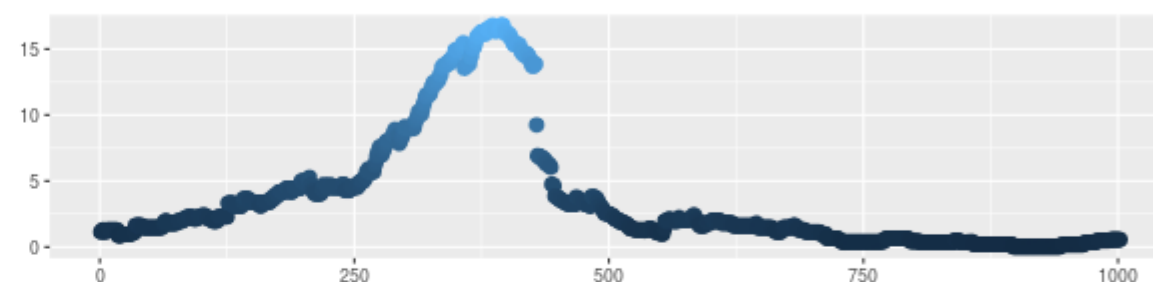
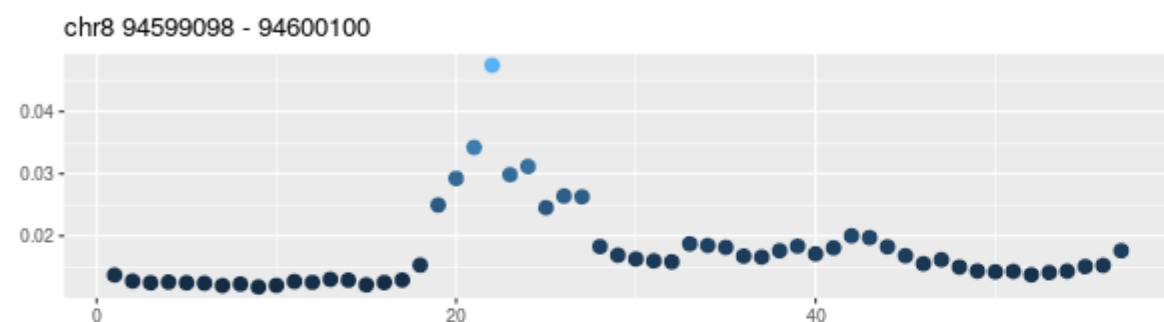
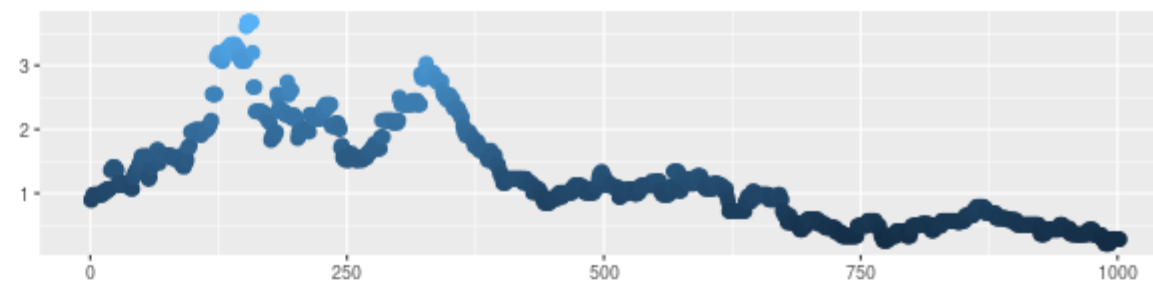
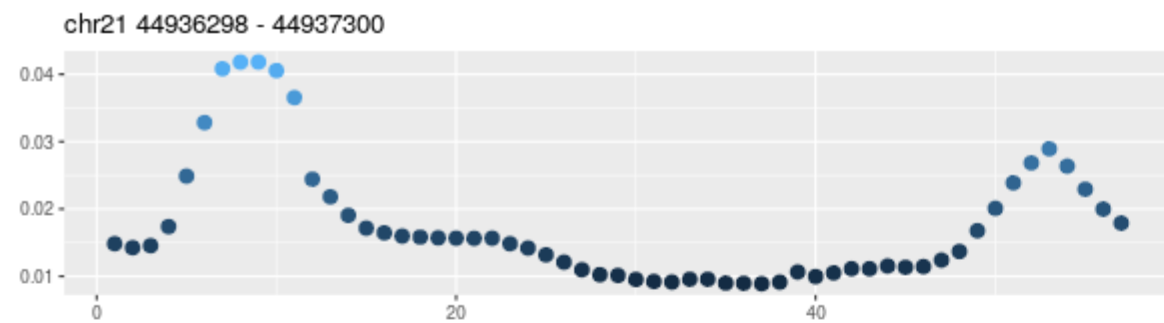


Attention weights

ChIP-seq score

Single attention module

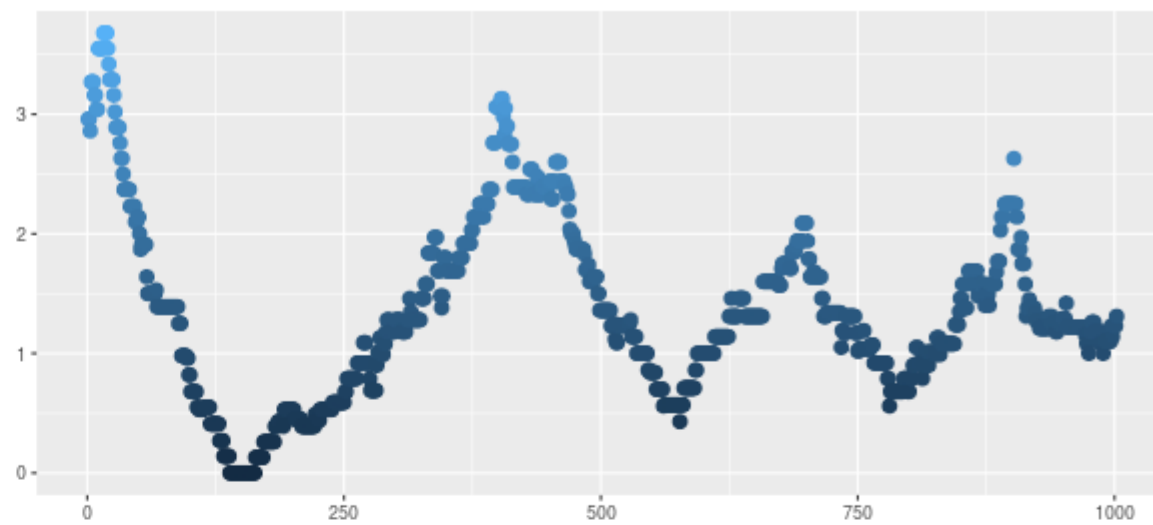
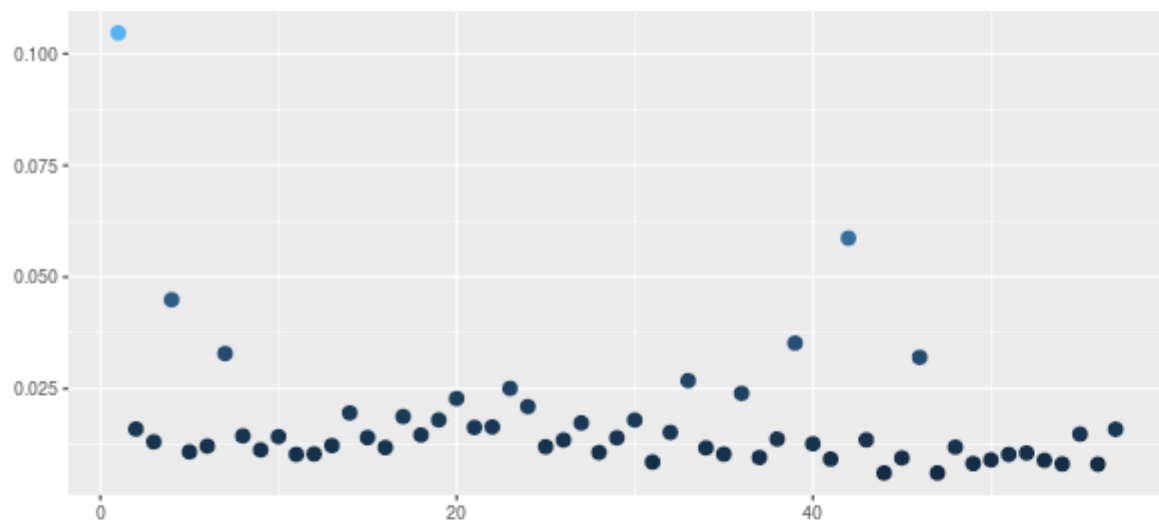
REST/liver



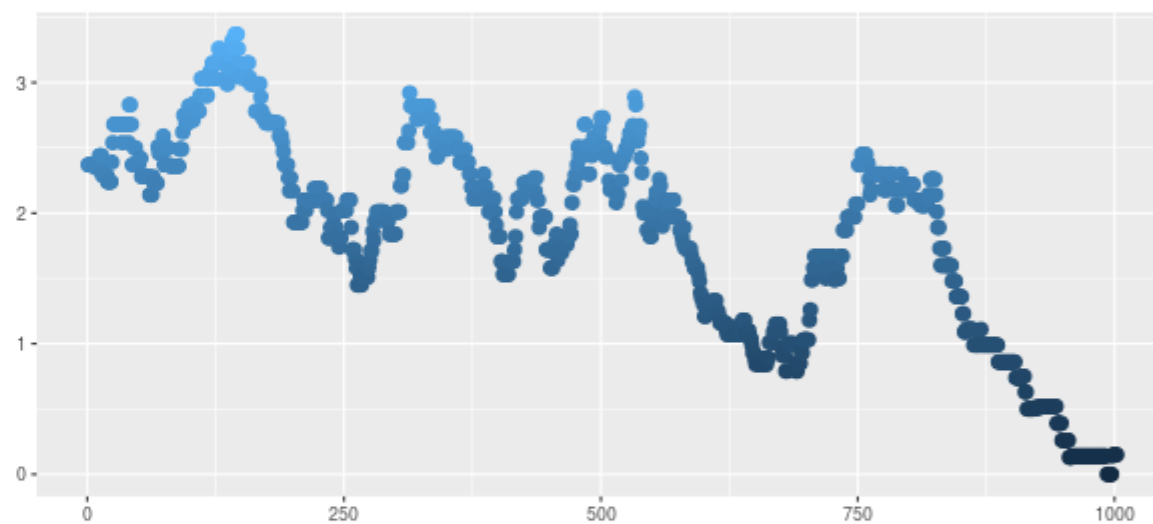
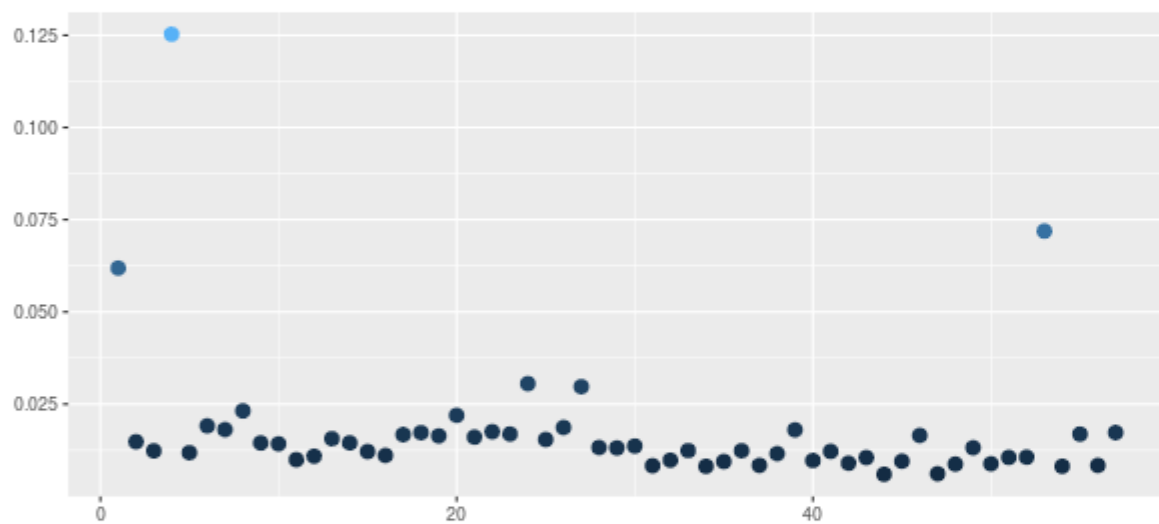
Single attention module

TAF1/liver

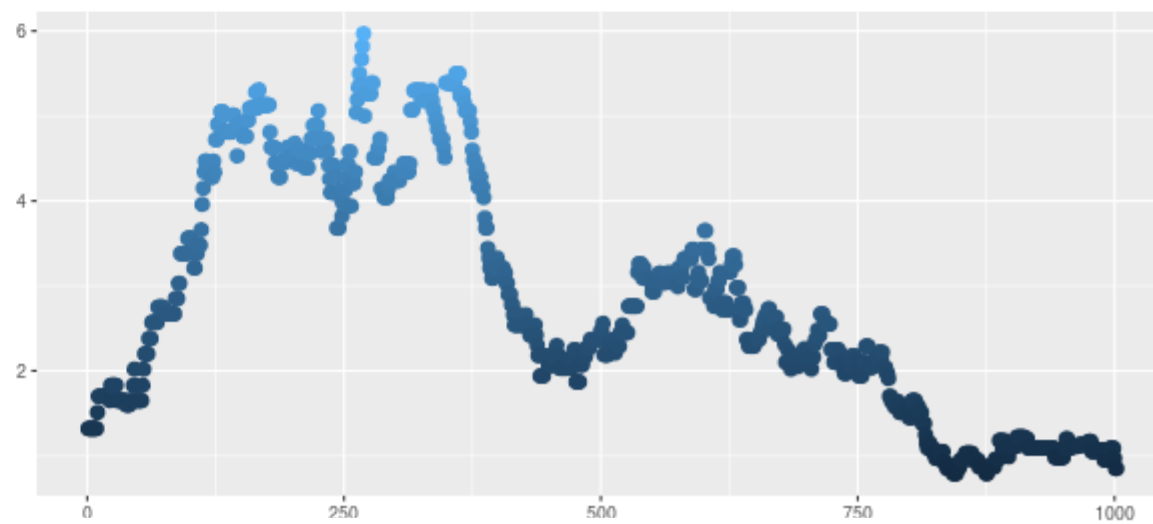
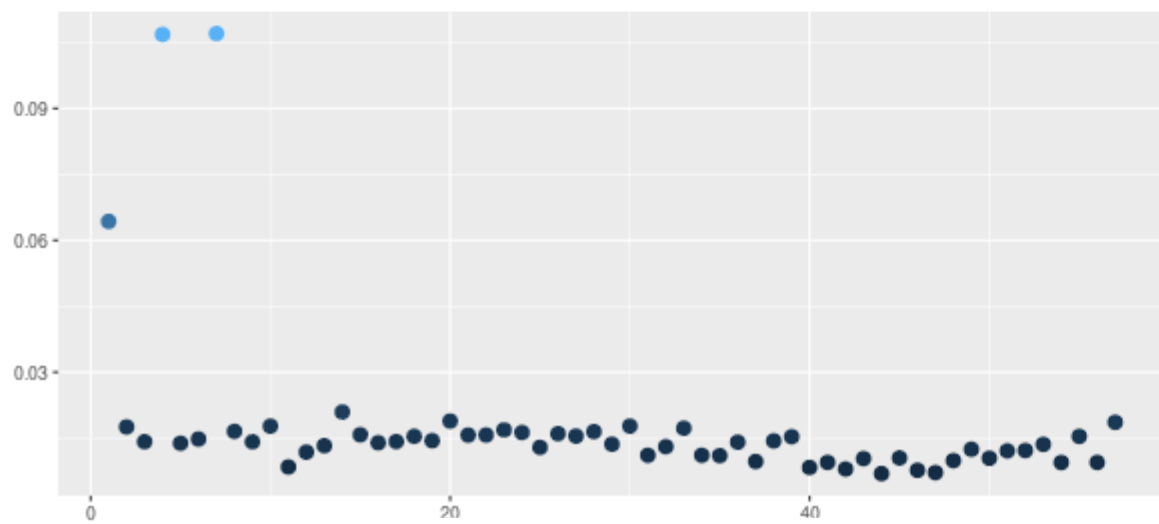
chr1 145476748 - 145477750



chr1 115322348 - 115323350



chr1 205179748 - 205180750

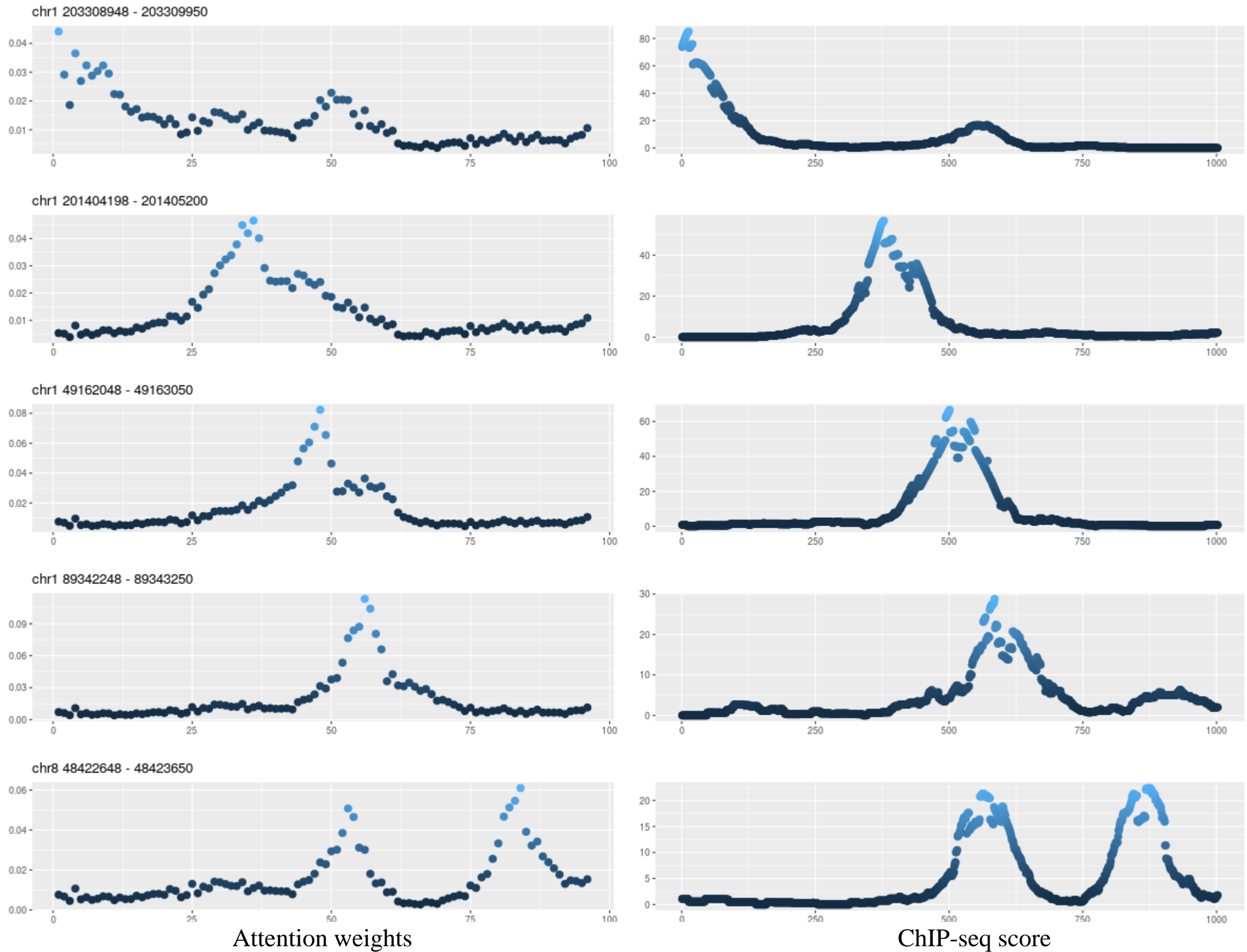


Attention weights

CHIP-seq score

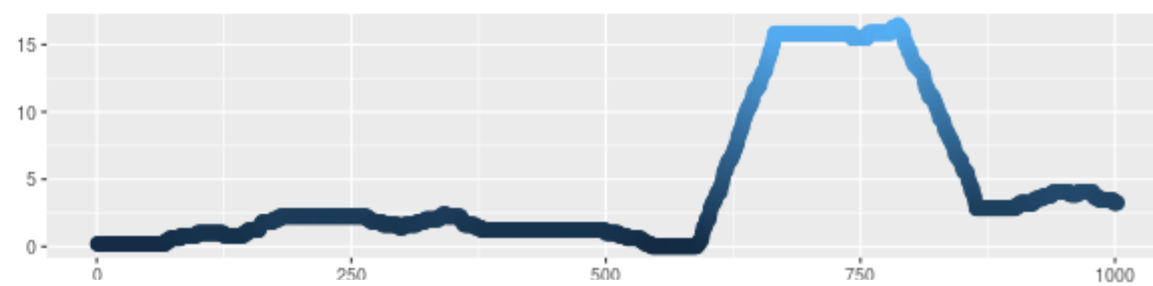
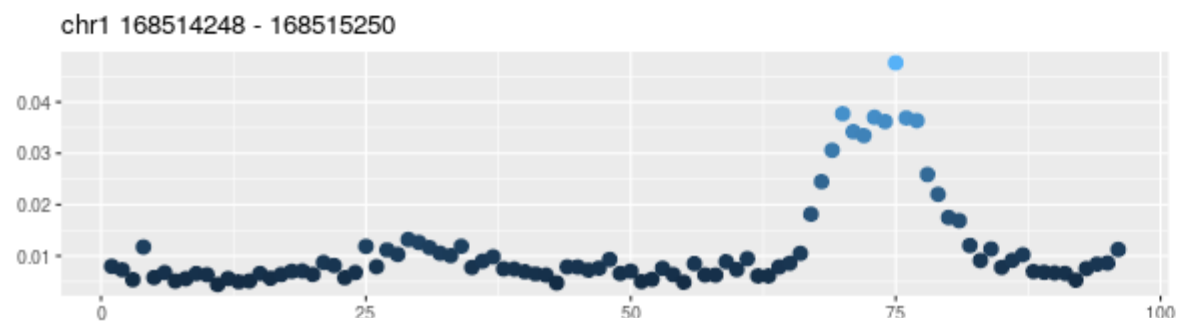
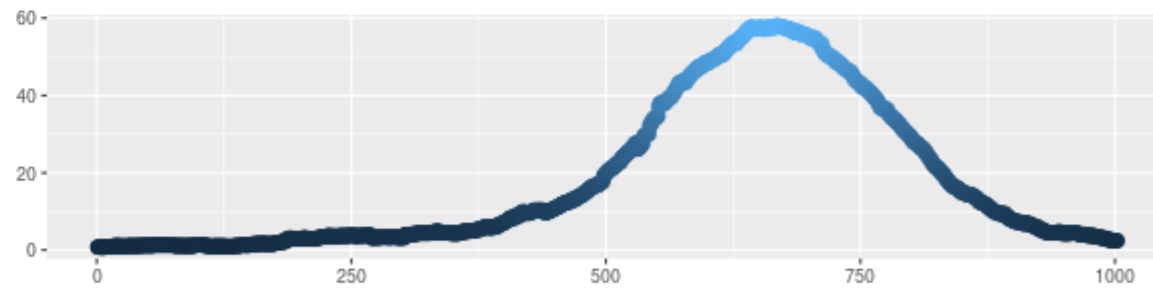
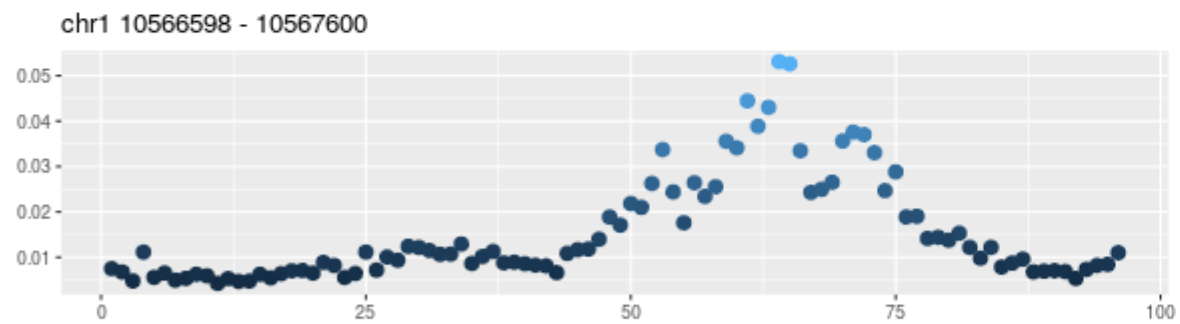
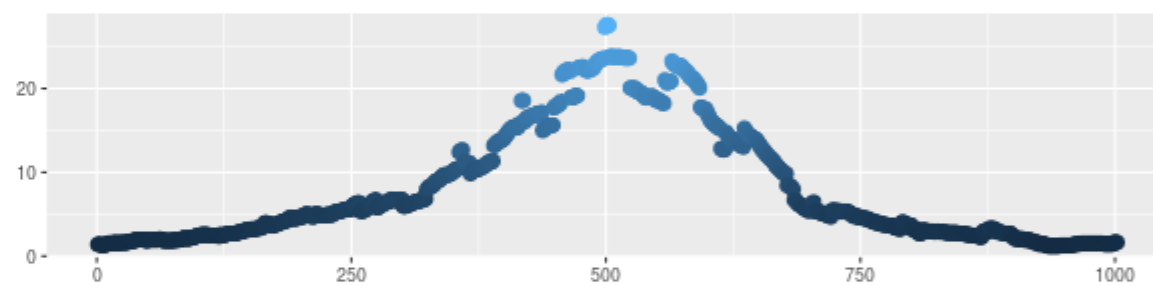
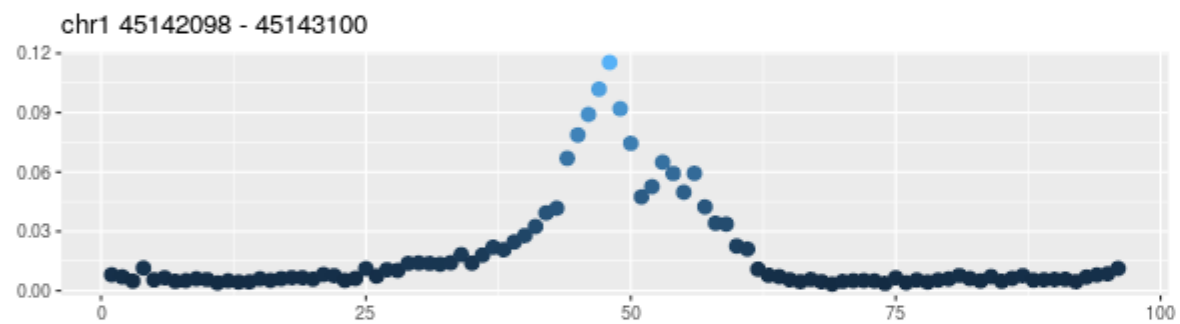
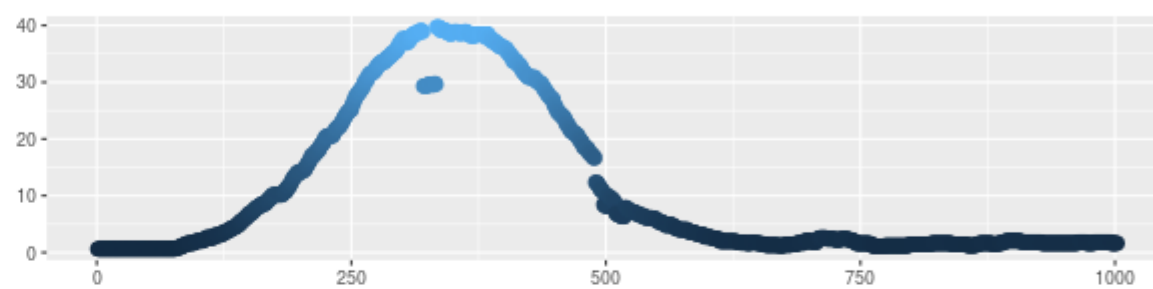
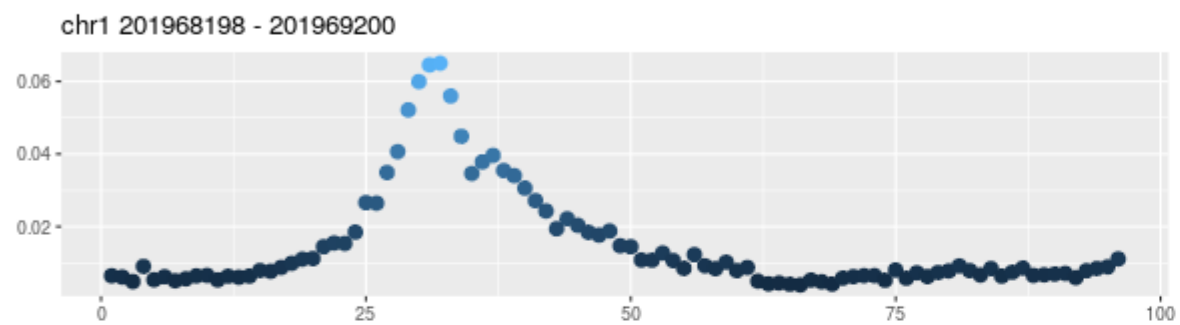
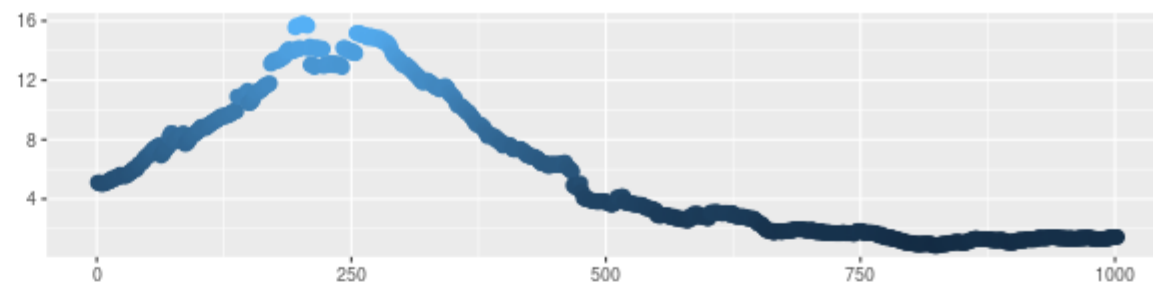
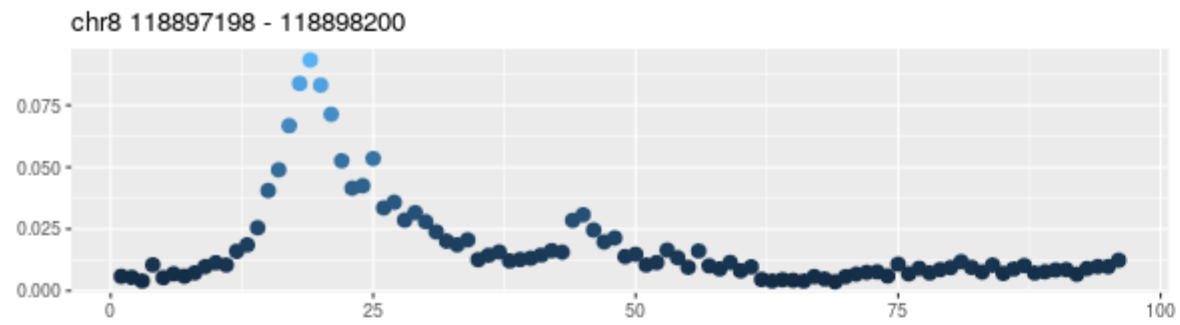
Pairwise attention module

CTCF/induced pluripotent stem cell



Pairwise attention module

CTCF/PC-3

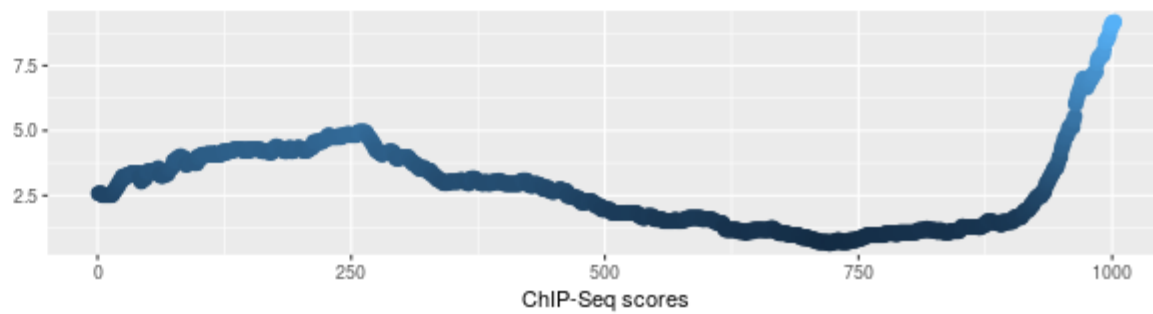
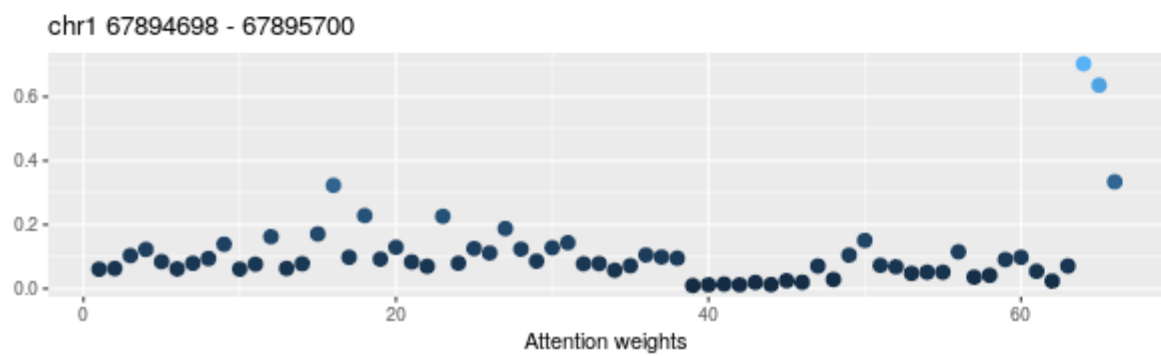
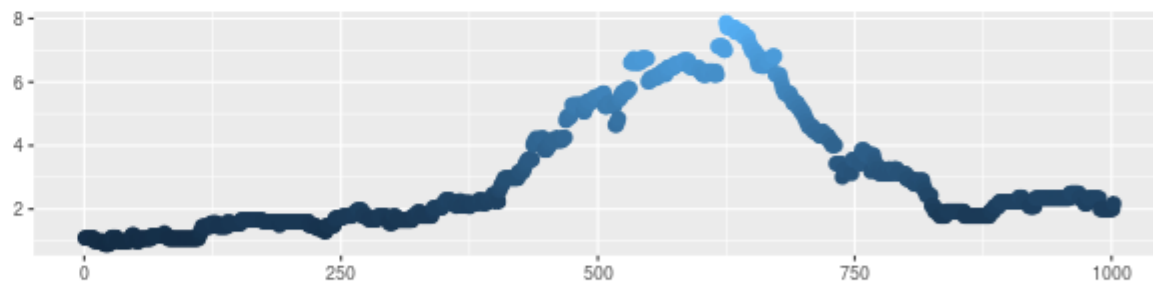
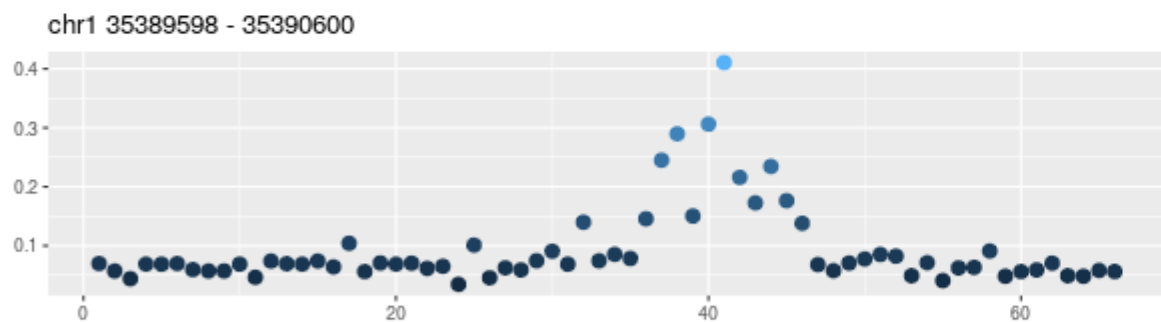
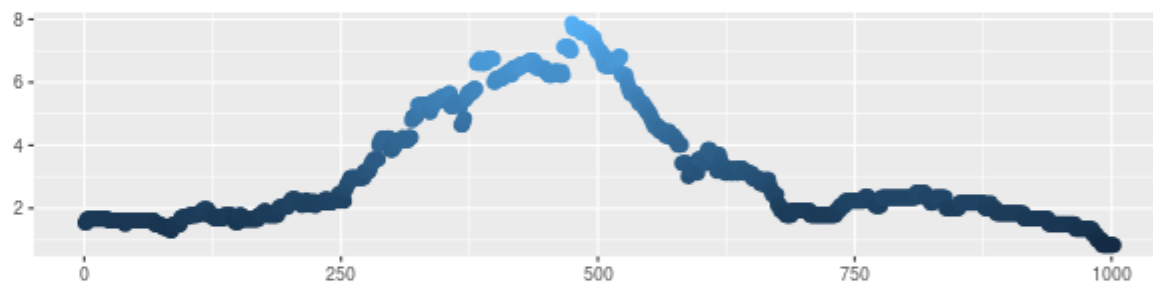
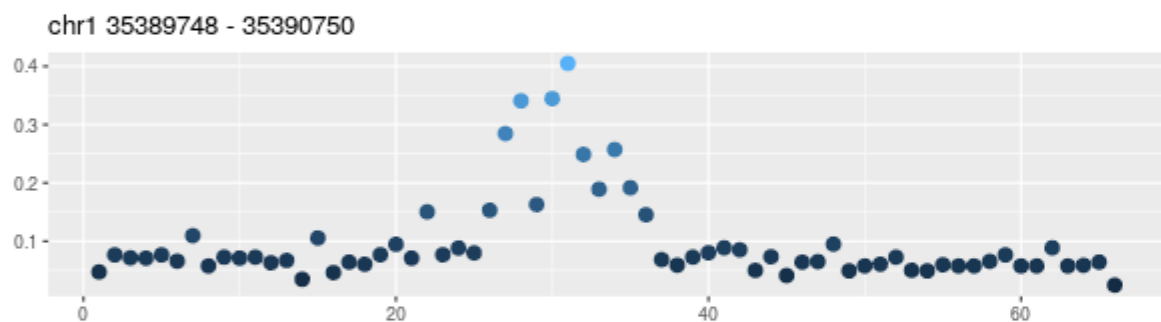
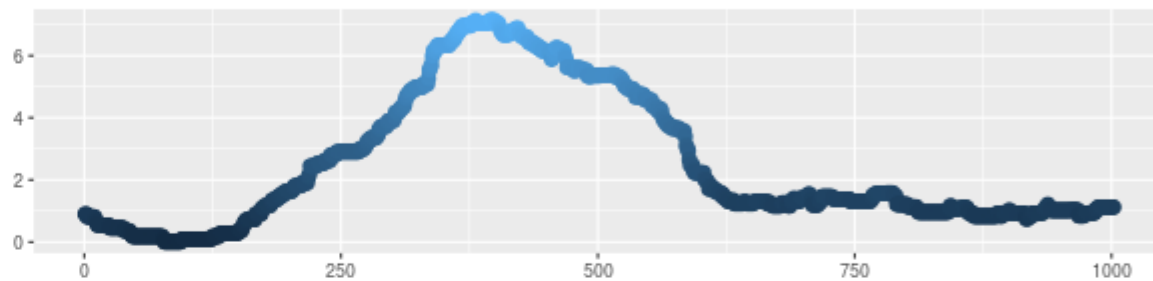
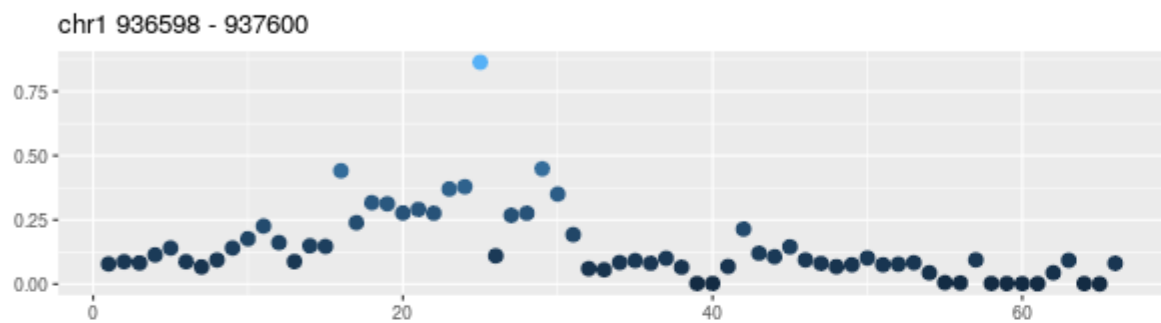
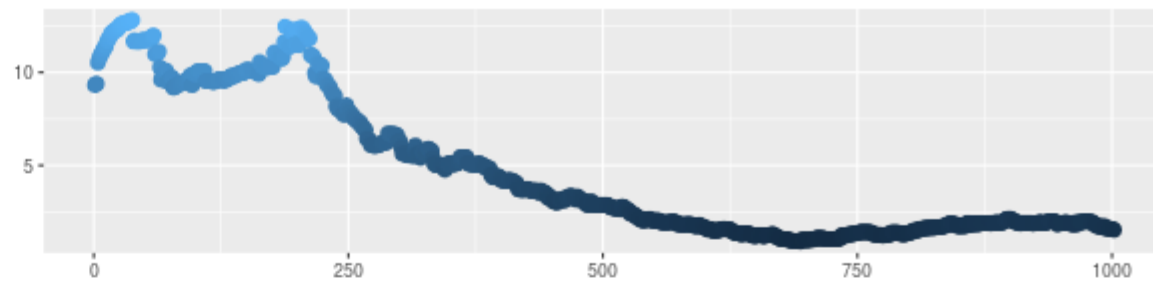
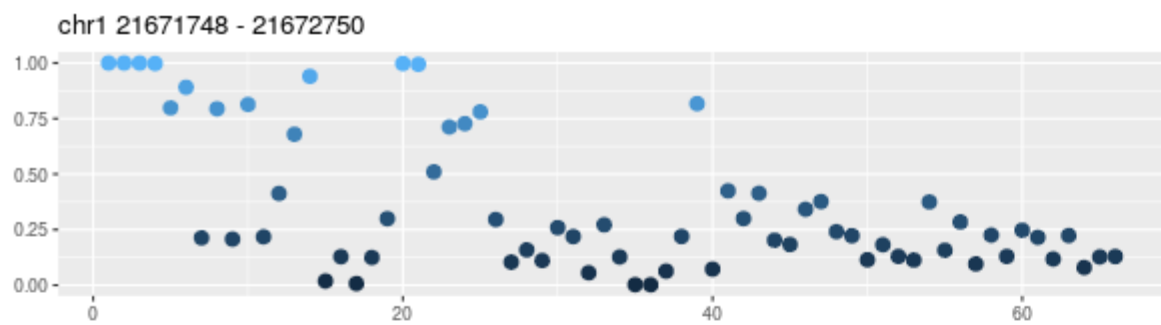


Attention weights

ChIP-seq score

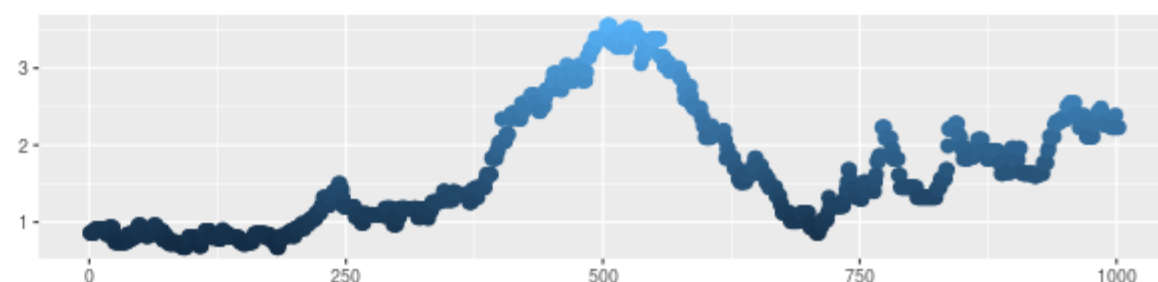
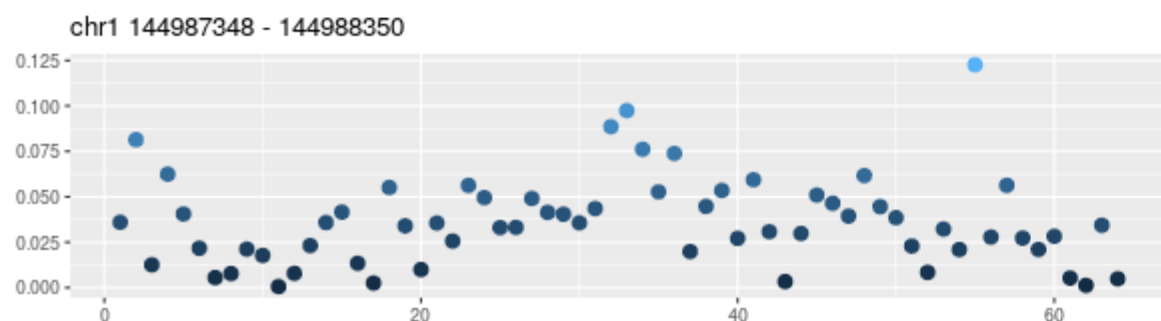
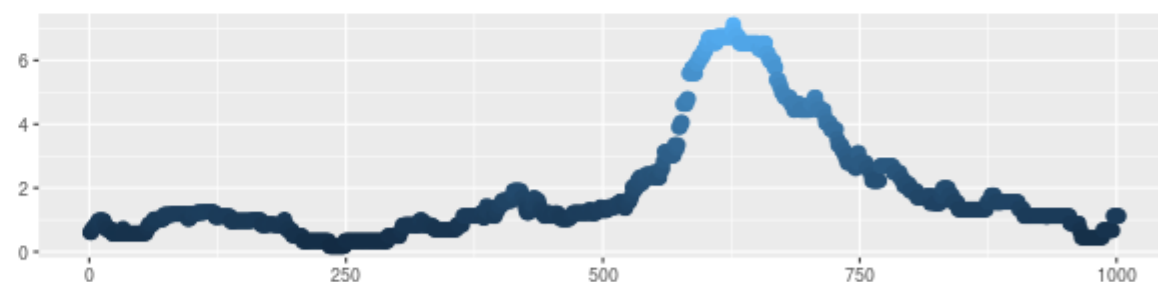
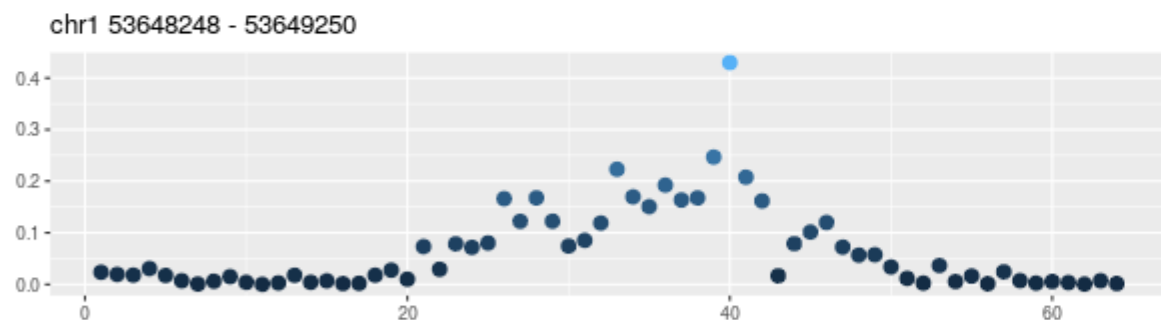
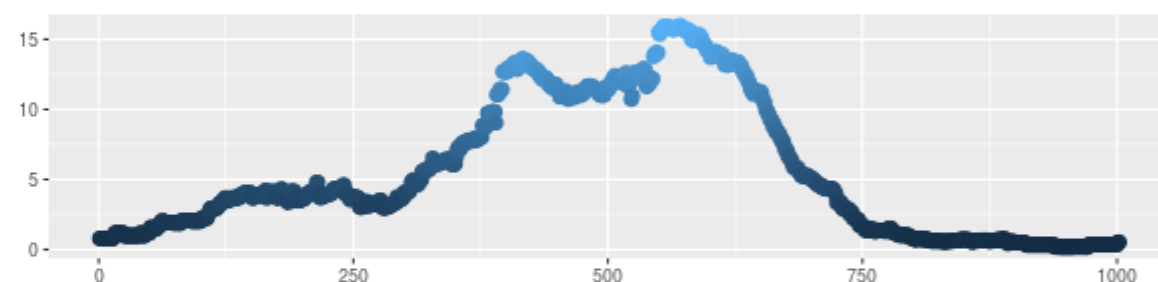
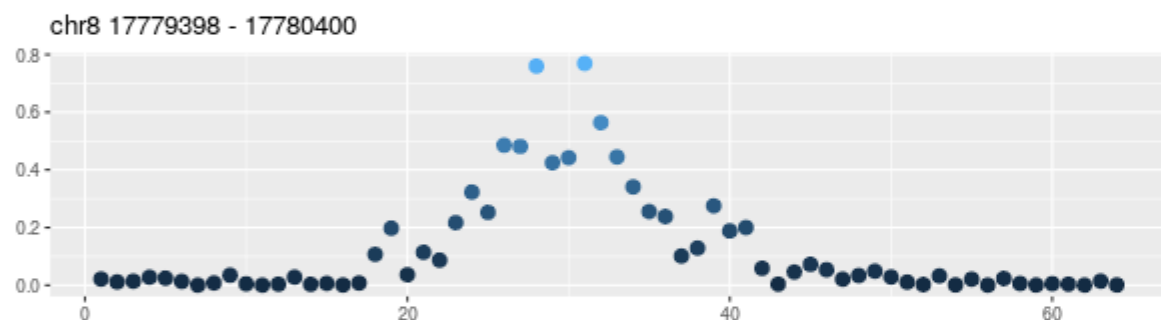
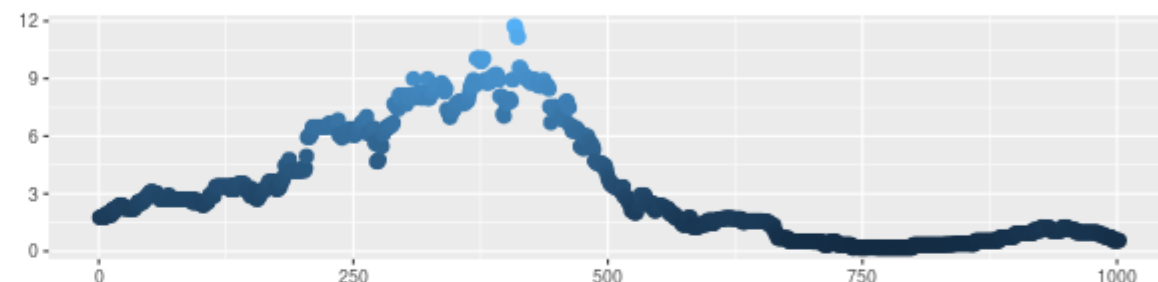
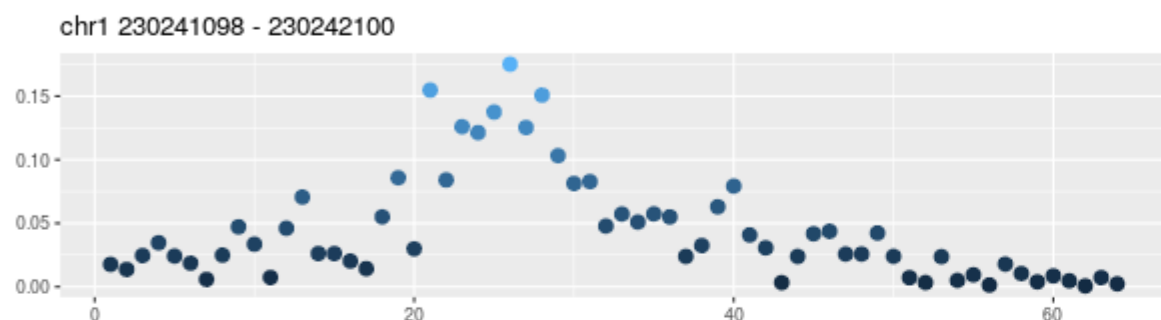
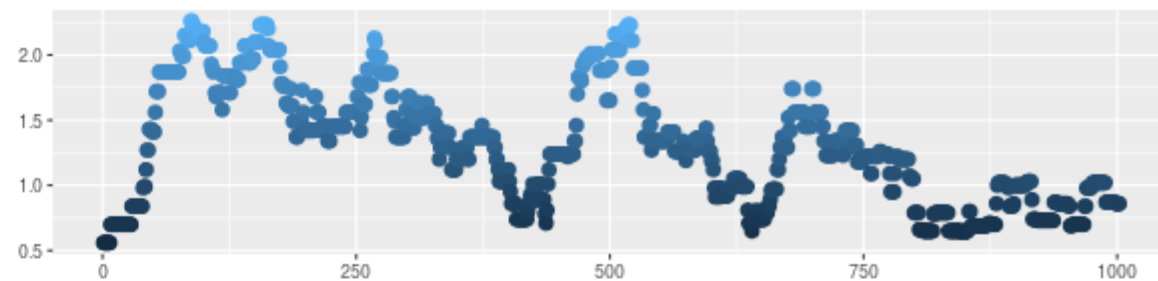
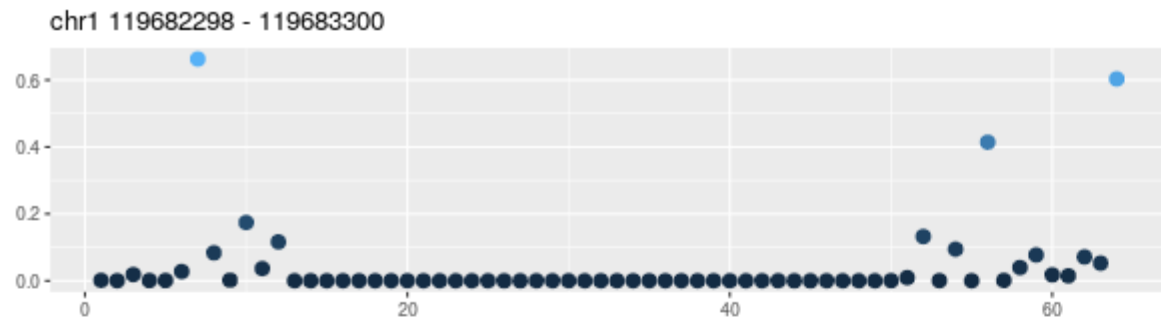
Pairwise attention module

E2F1/K562



Pairwise attention module

EGR1/liver

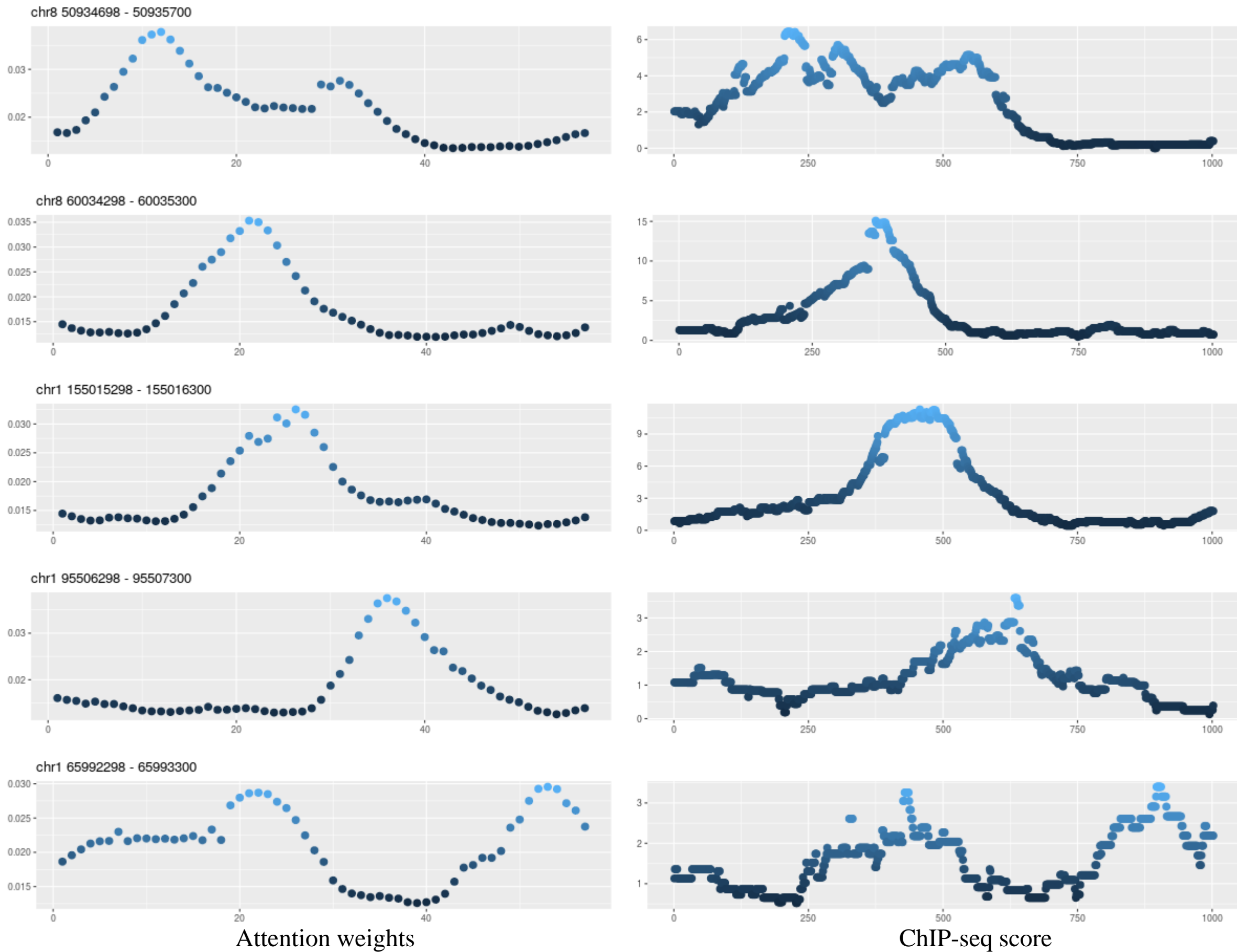


Attention weights

CHIP-seq score

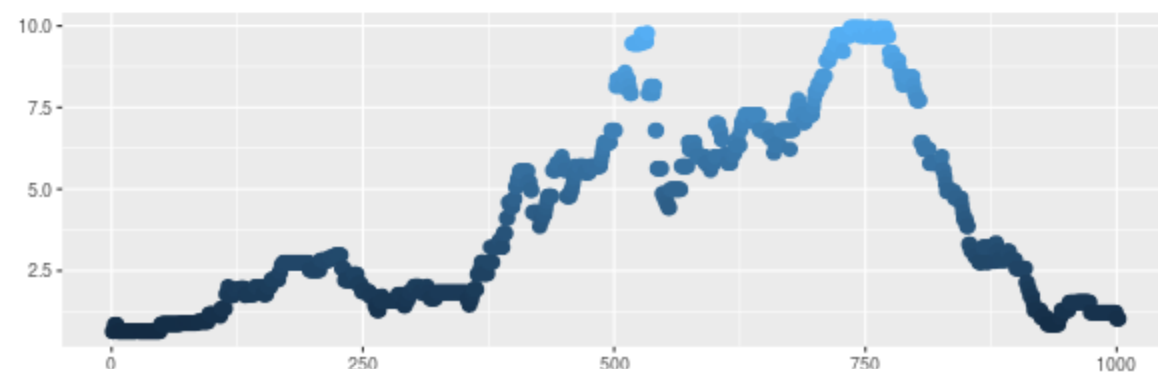
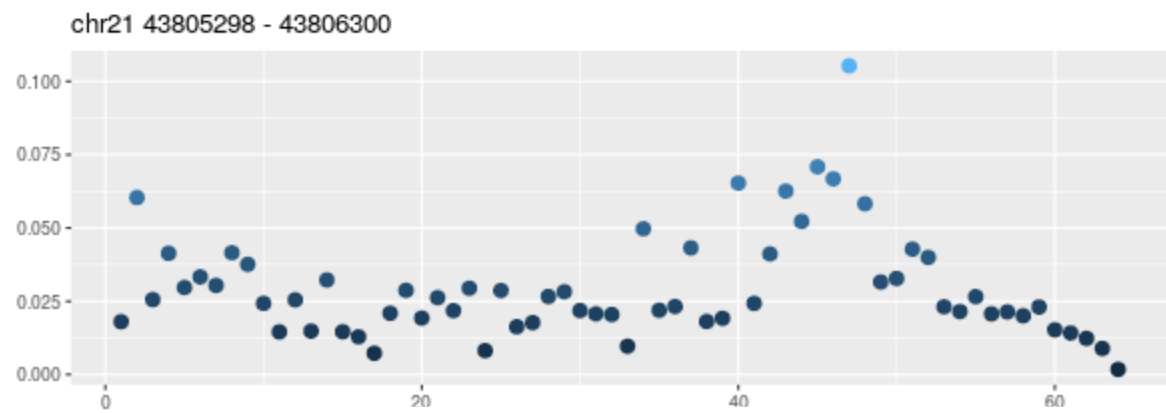
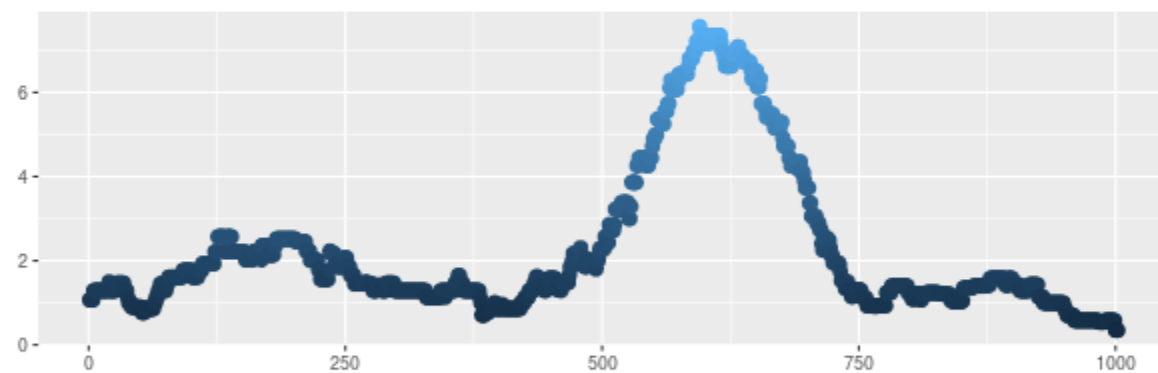
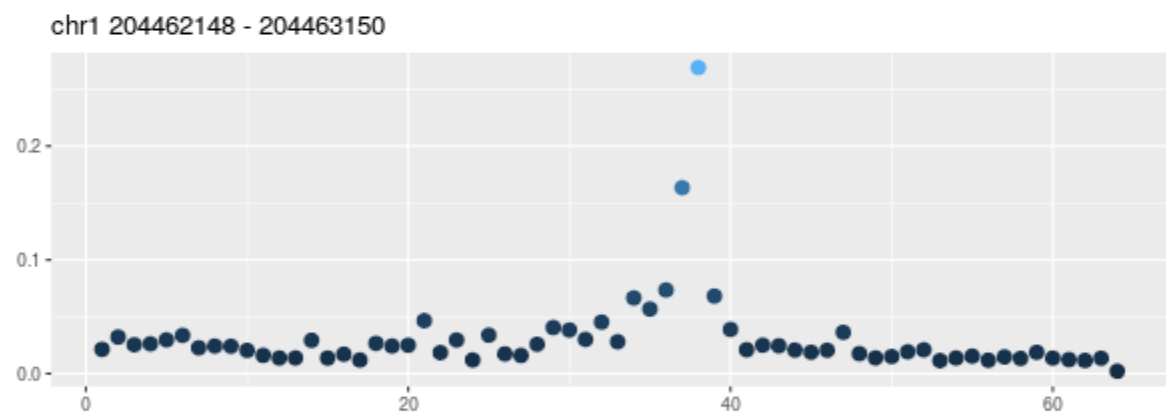
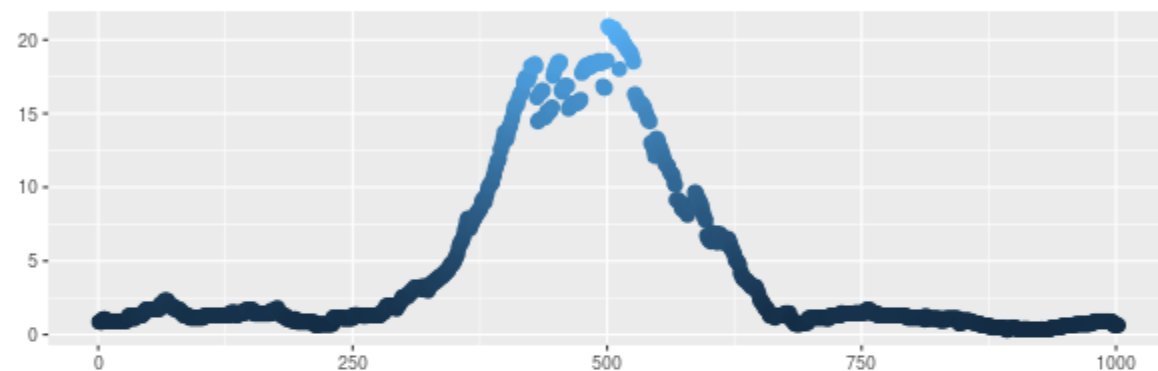
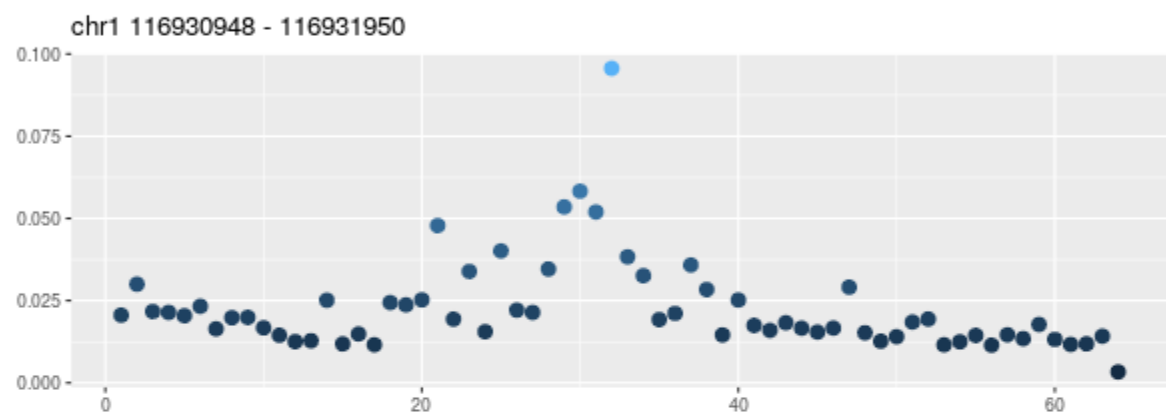
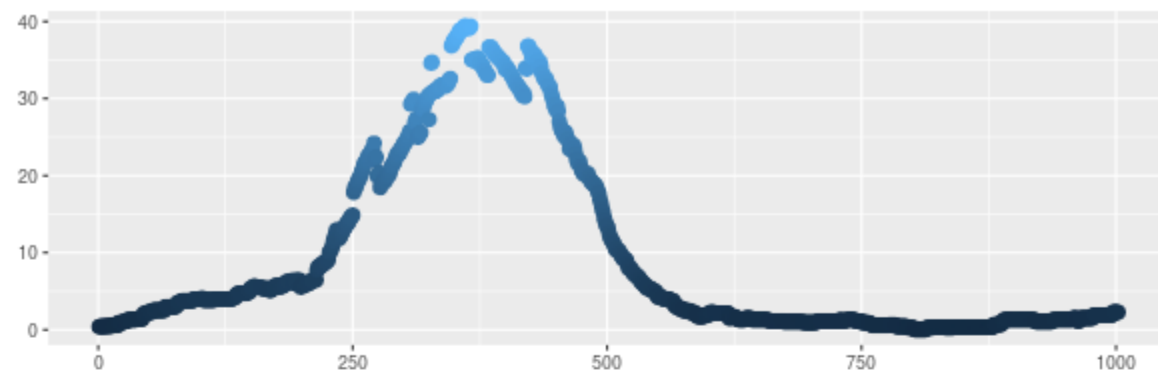
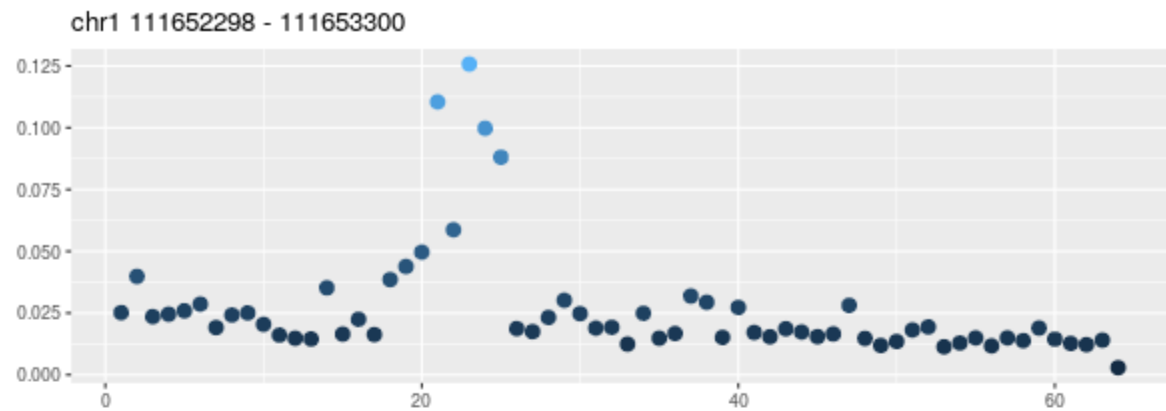
Pairwise attention module

FOXA1/liver



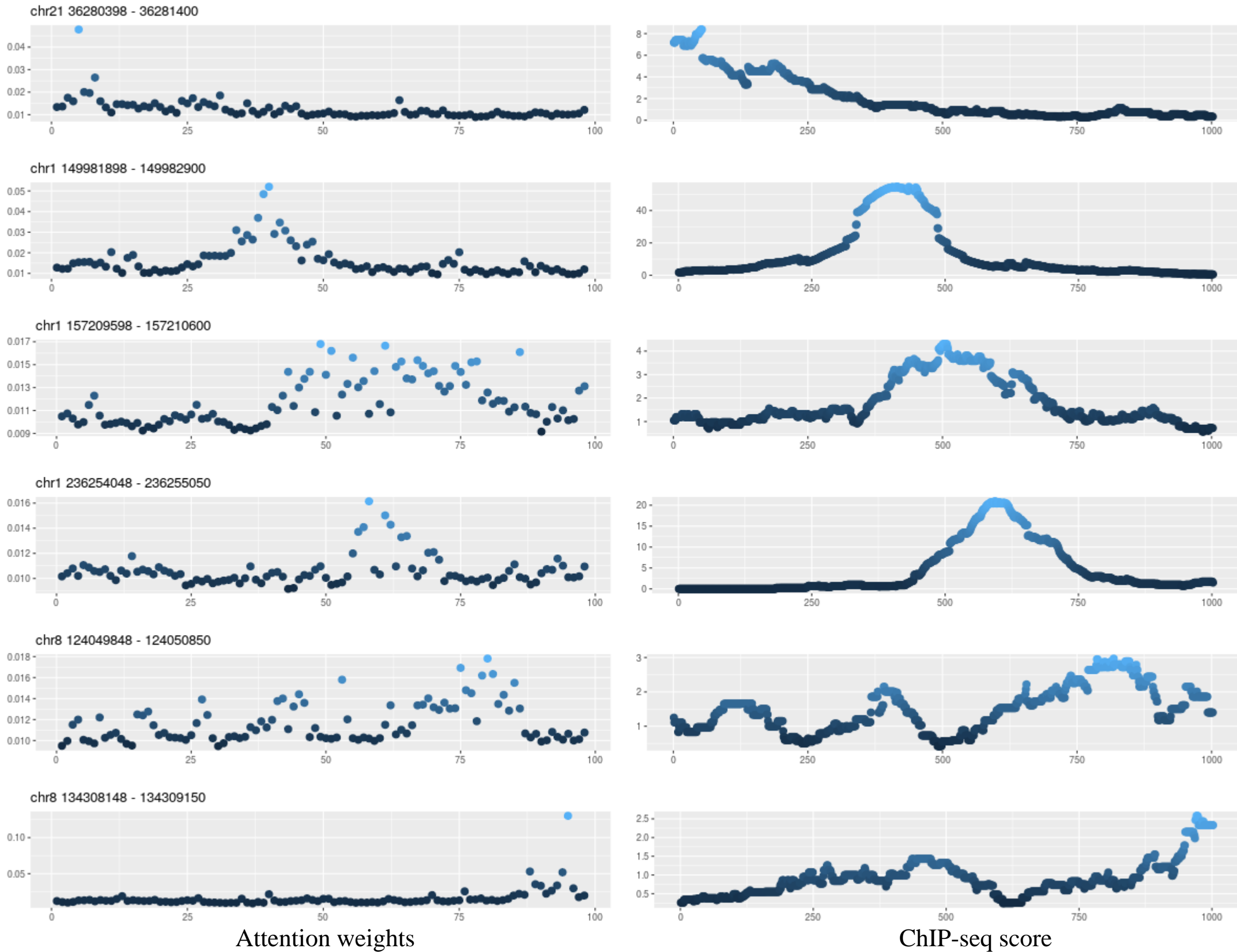
Pairwise attention module

FOXA2/liver



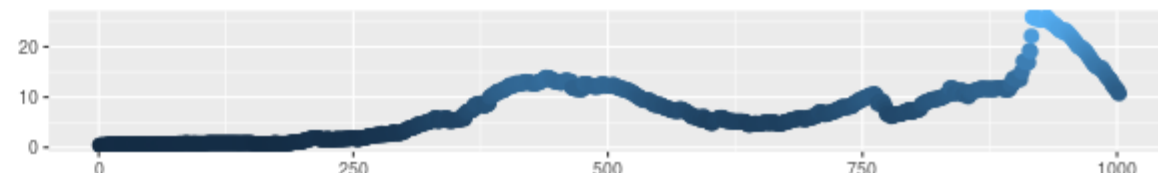
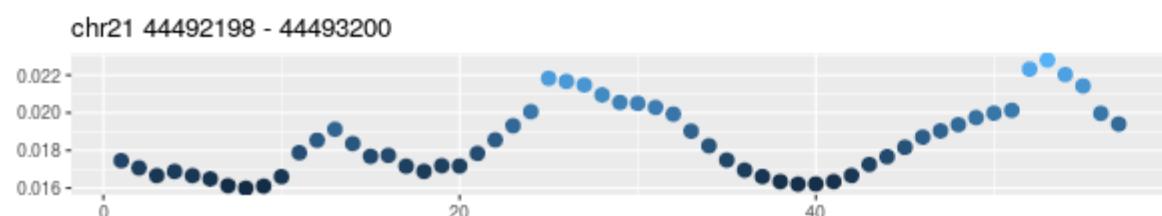
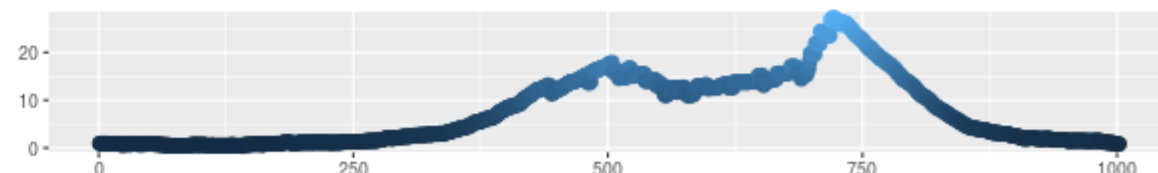
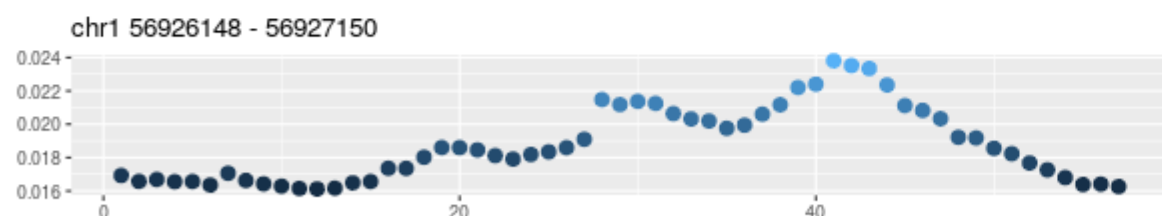
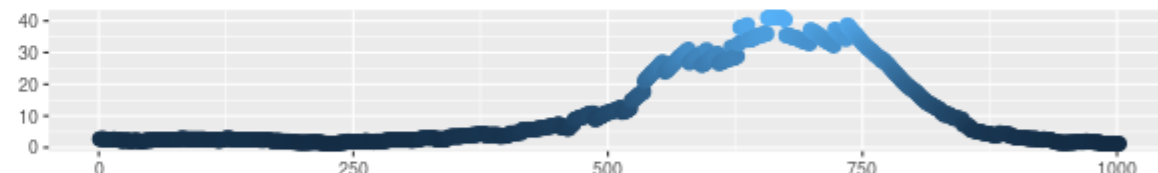
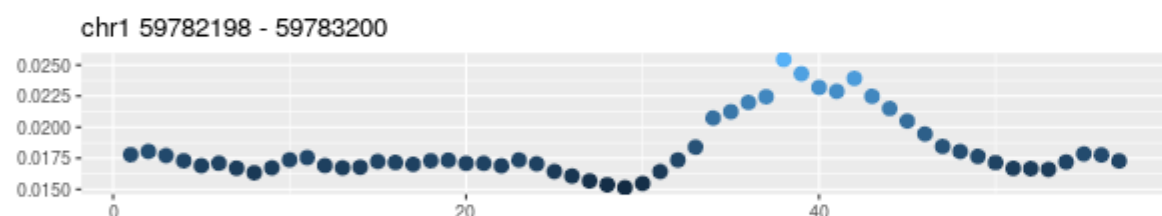
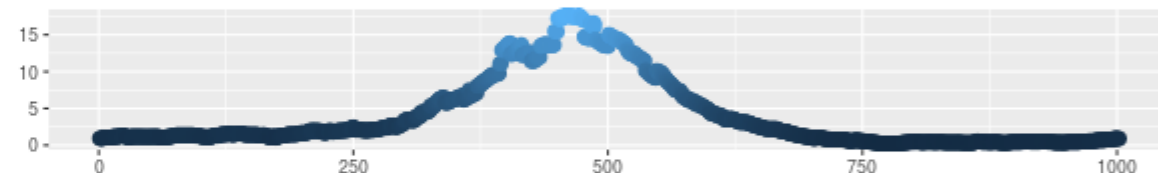
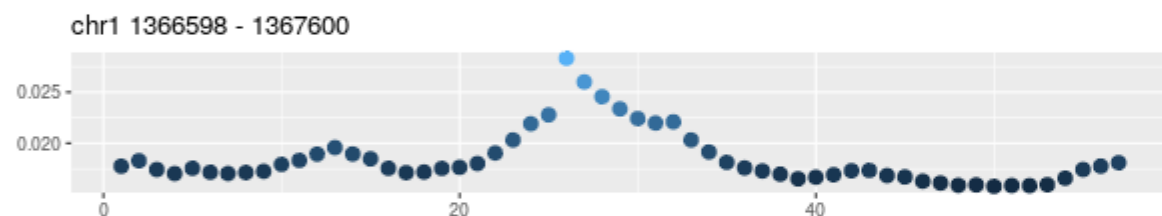
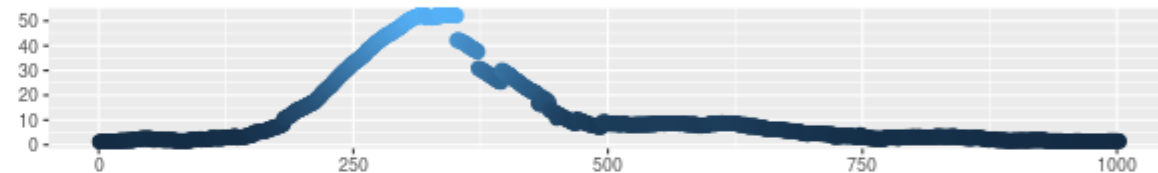
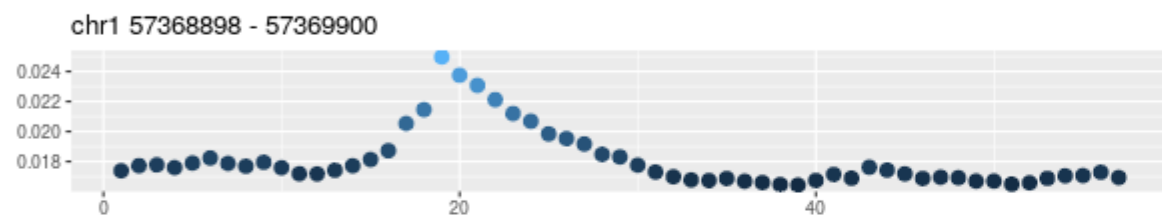
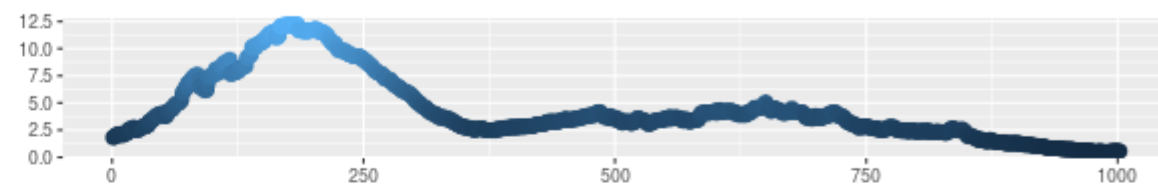
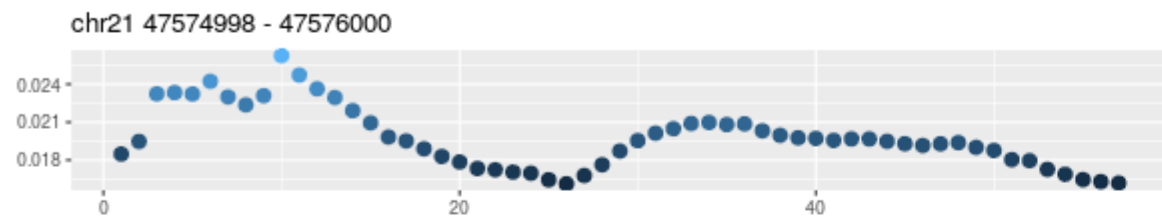
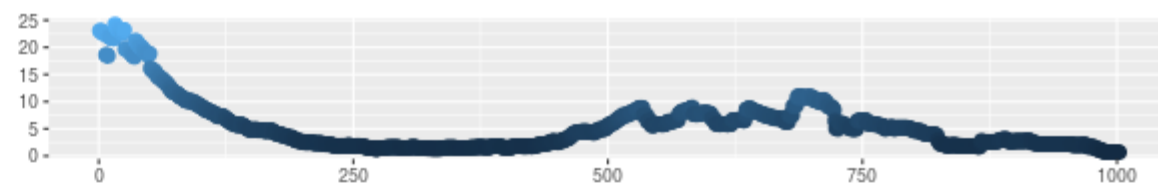
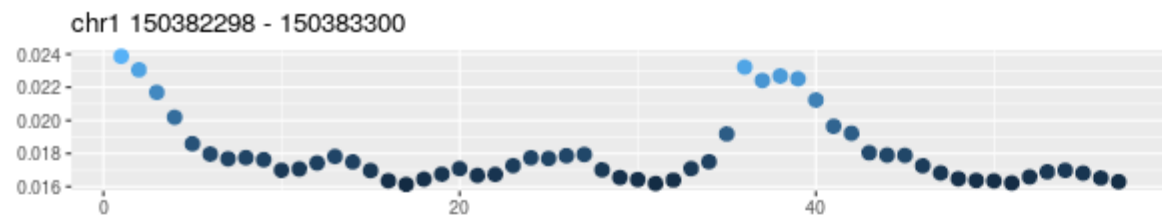
Pairwise attention module

GABPA/liver



Pairwise attention module

HNF4A/liver

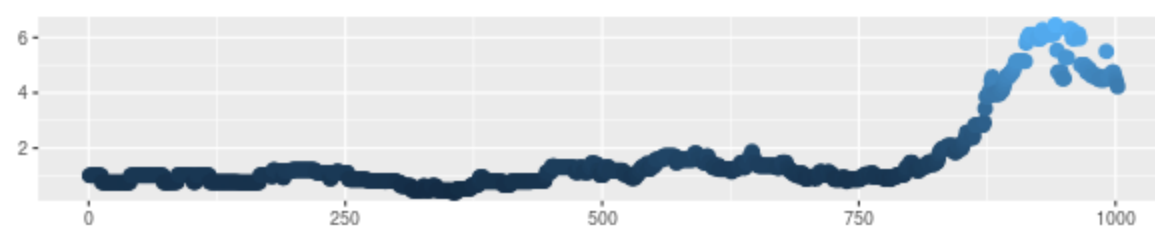
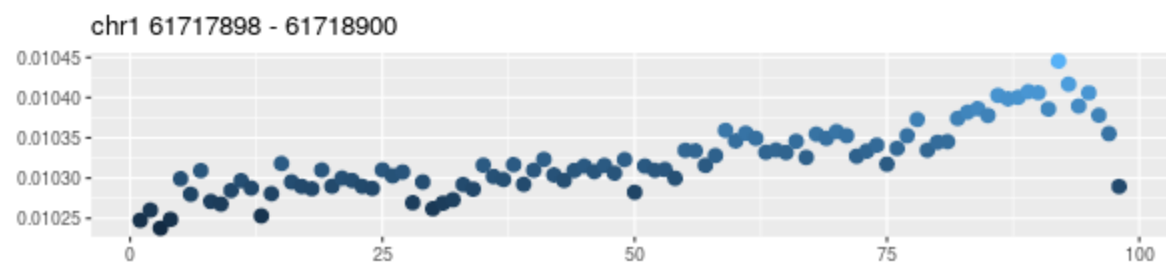
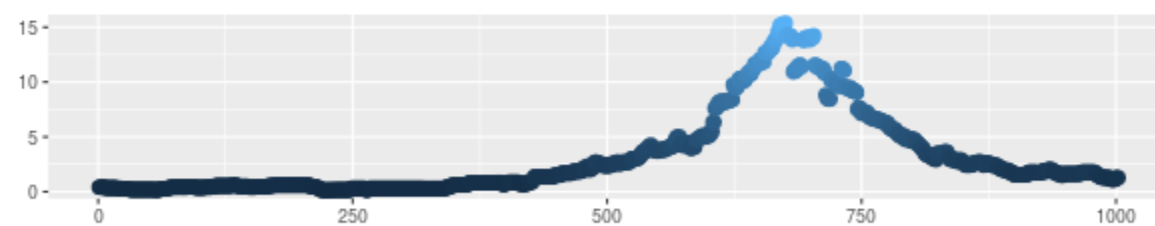
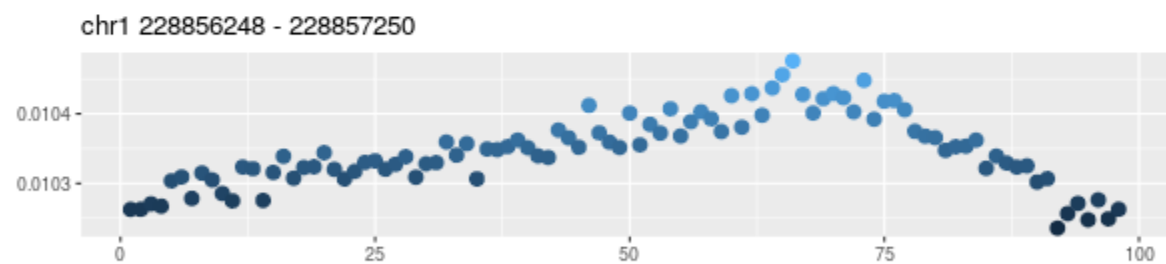
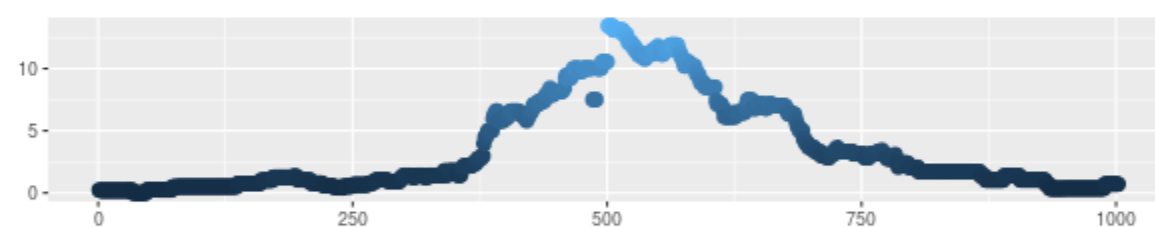
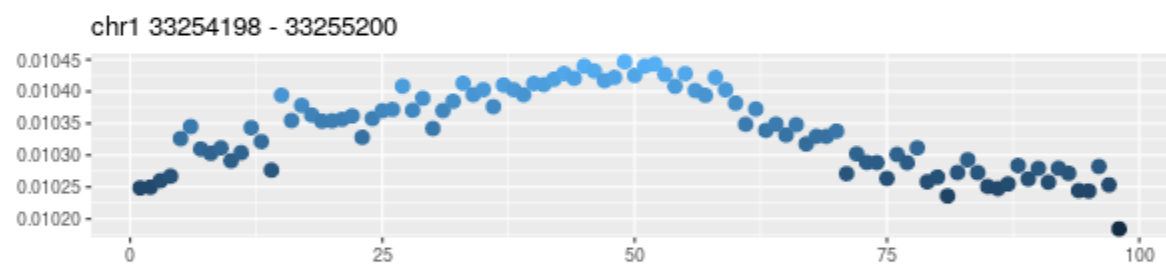
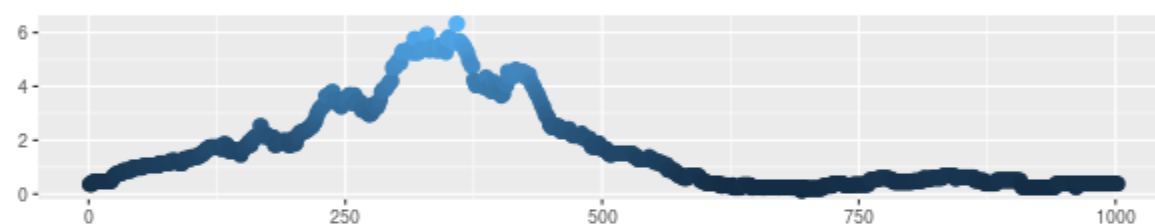
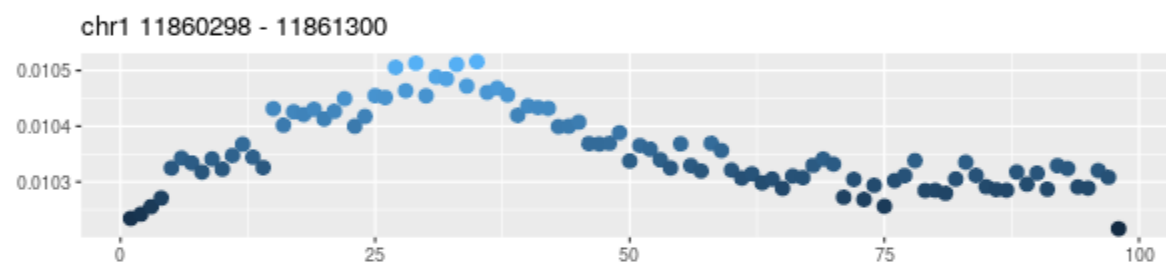
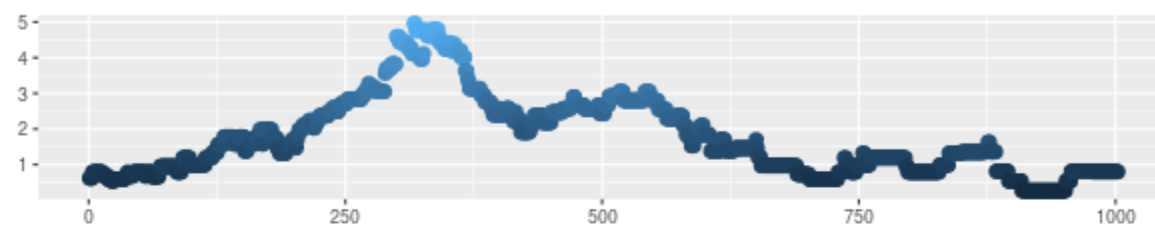
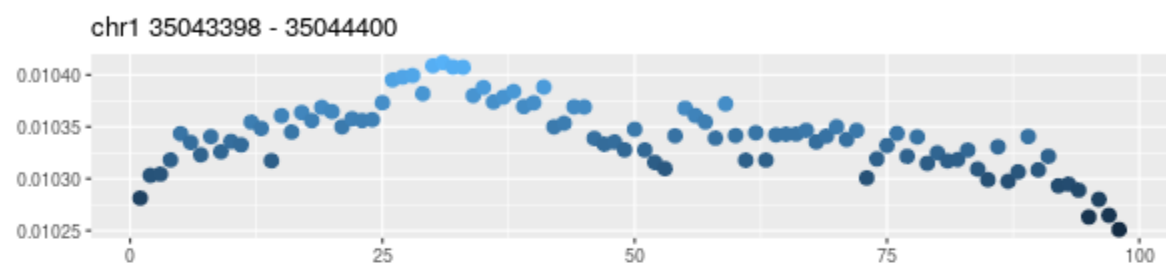
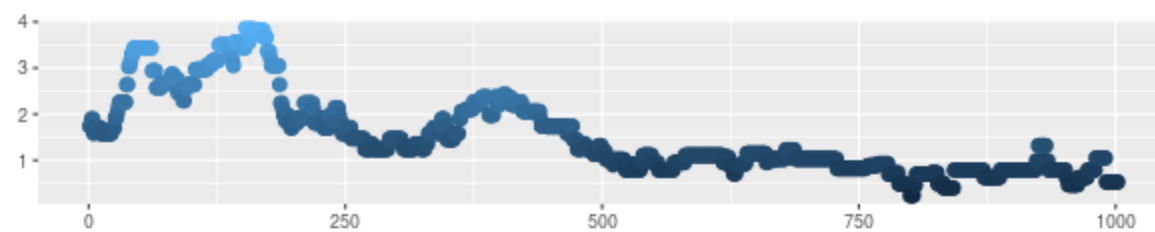
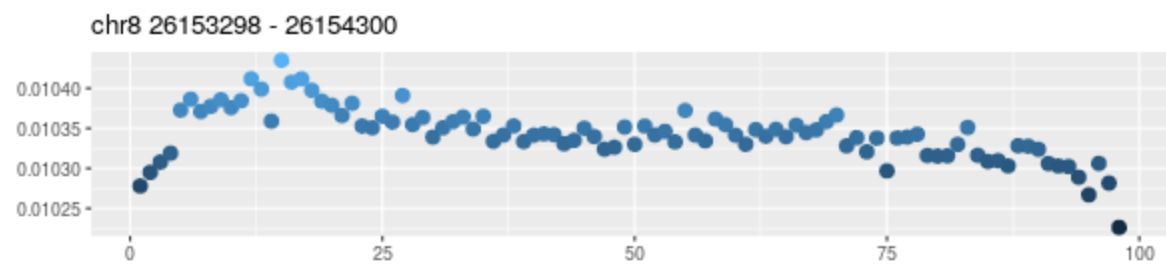


Attention weights

CHIP-seq score

Pairwise attention module

JUND/liver



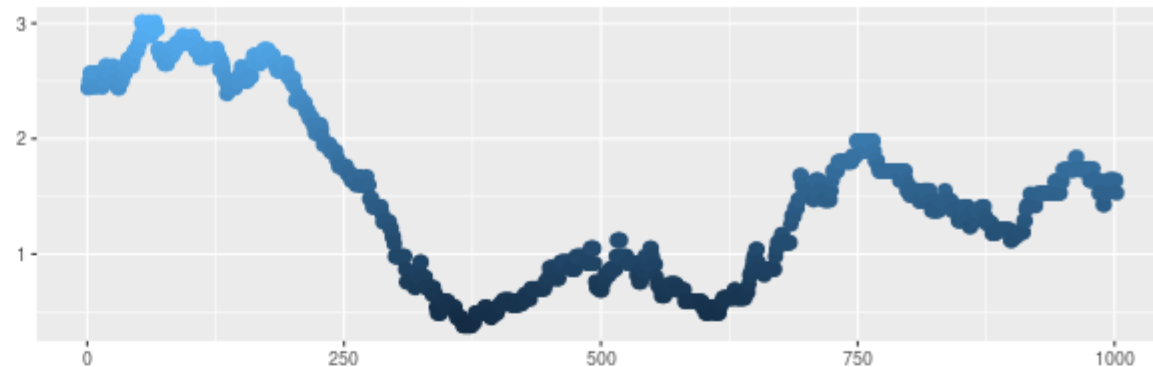
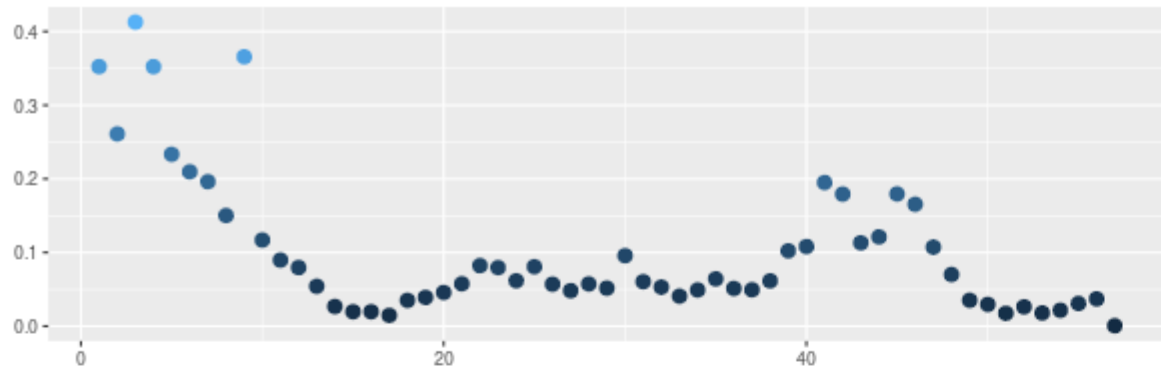
Attention weights

CHIP-seq score

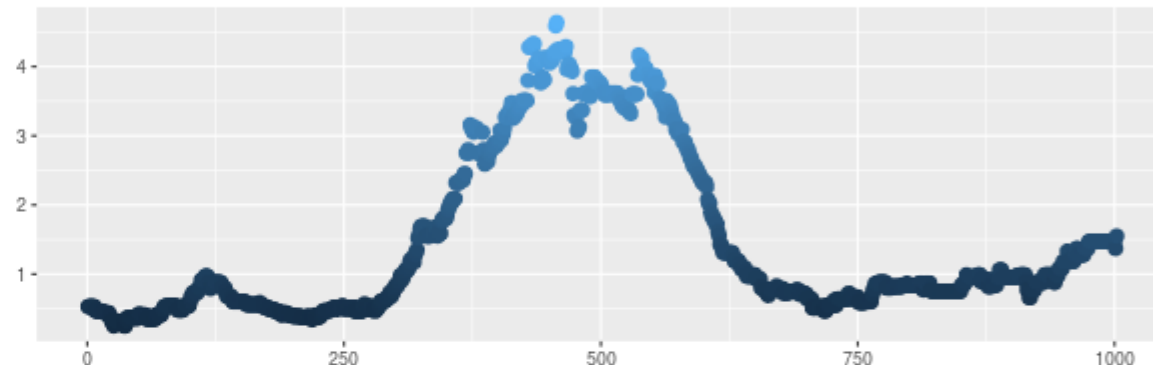
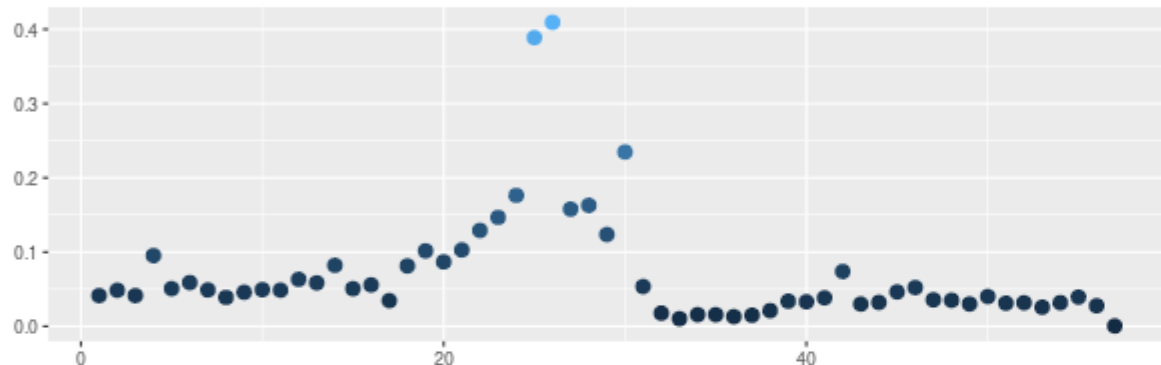
Pairwise attention module

MAX/liver

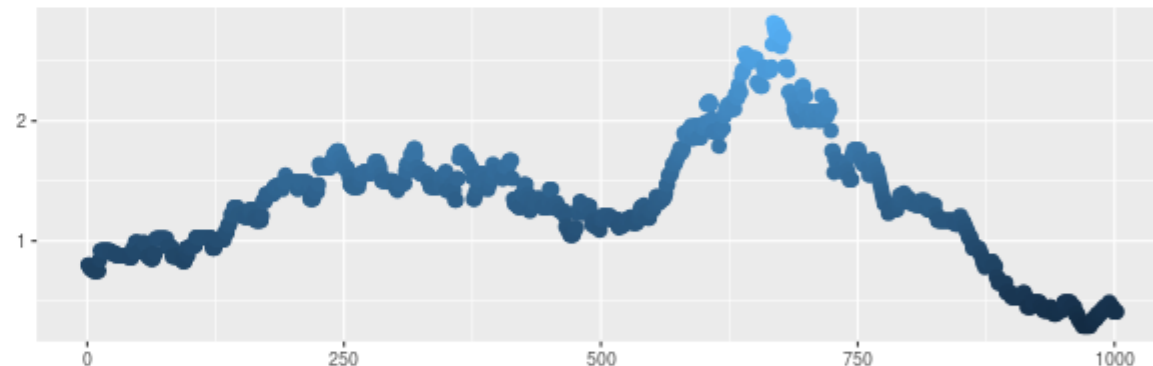
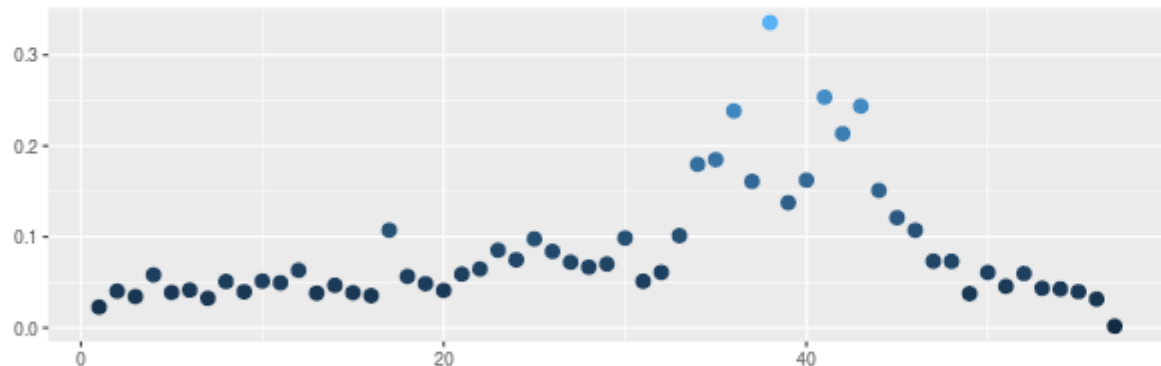
chr1 150980398 - 150981400



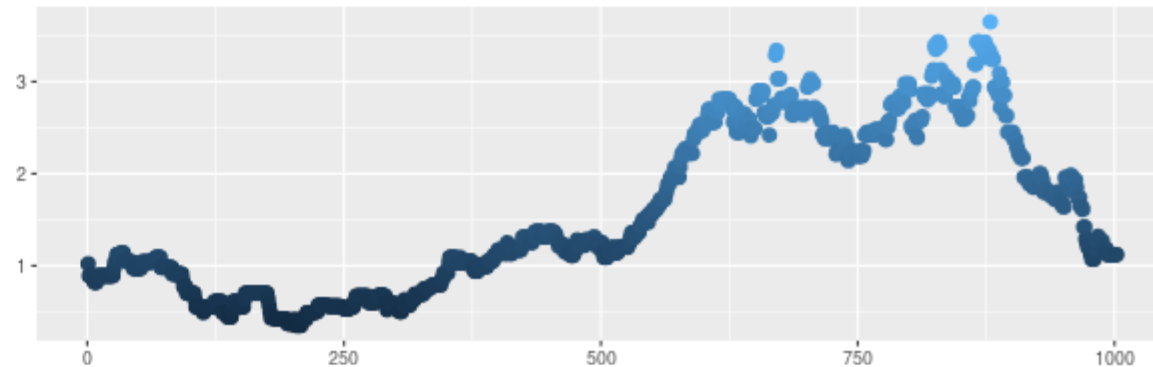
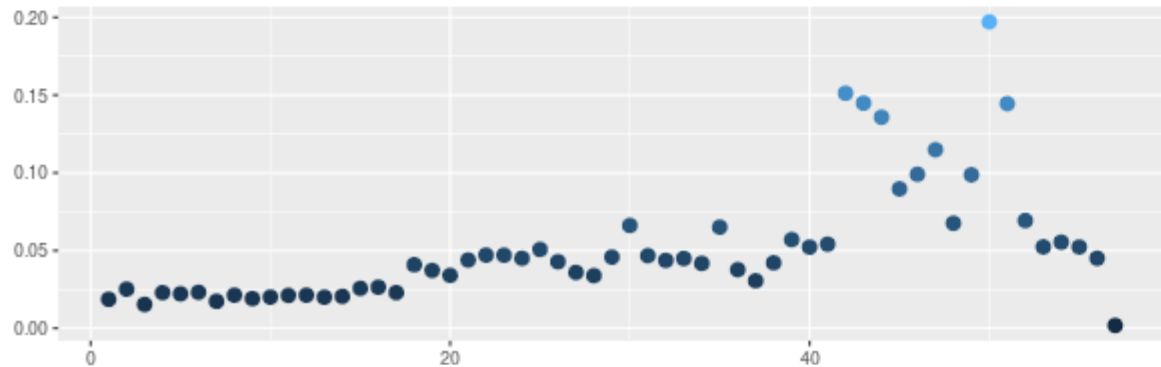
chr1 28558698 - 28559700



chr1 171453548 - 171454550



chr1 234667348 - 234668350

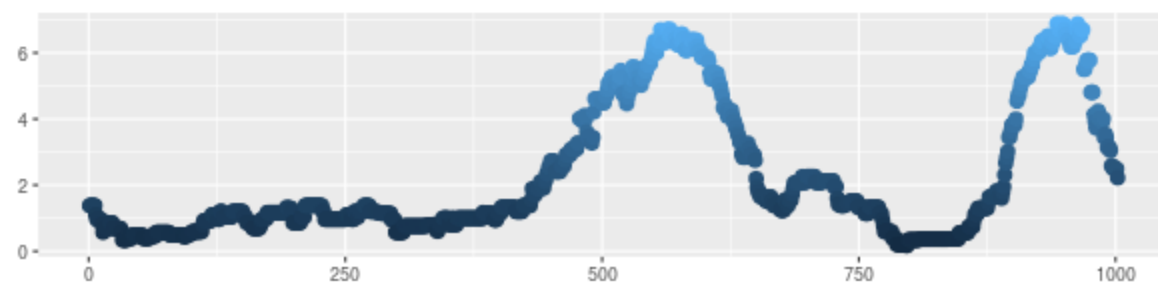
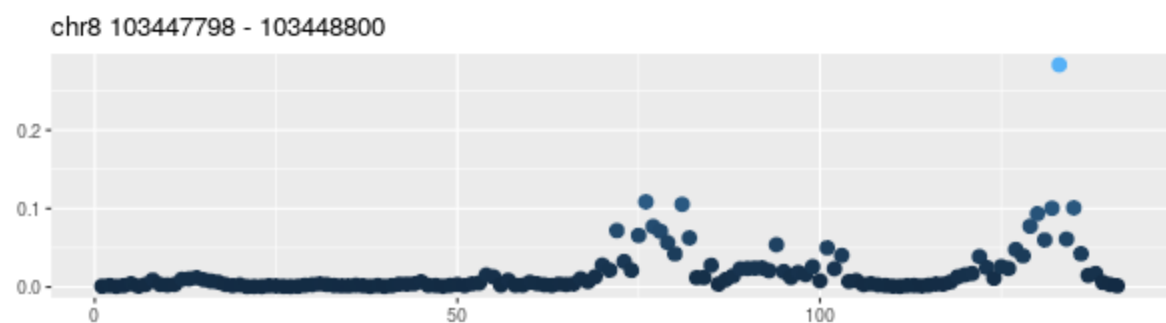
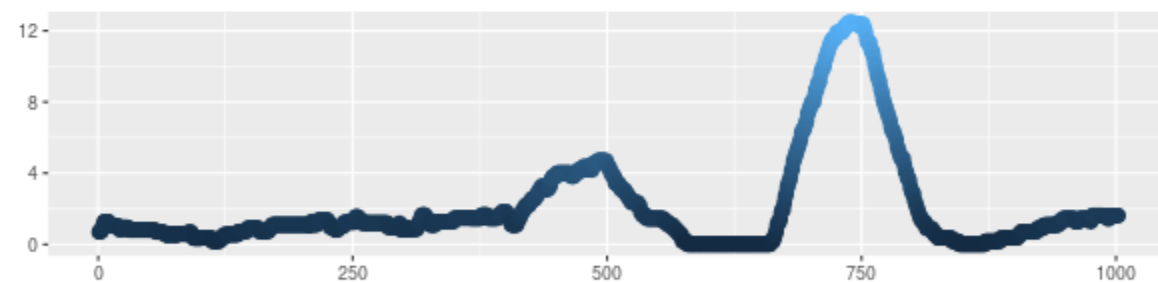
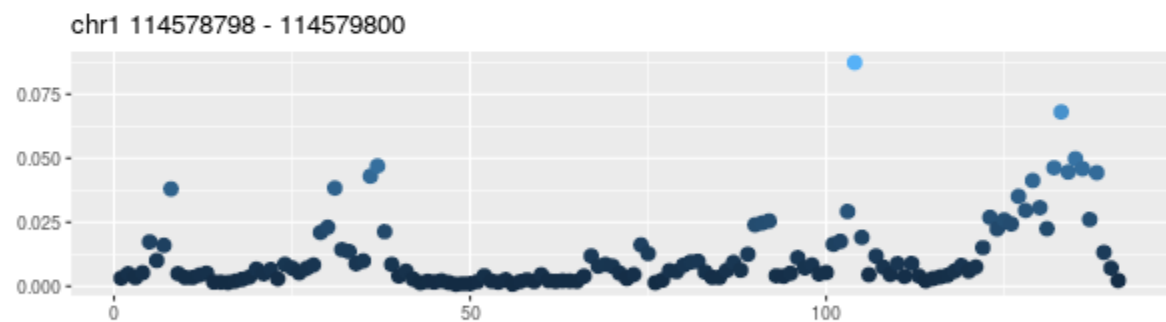
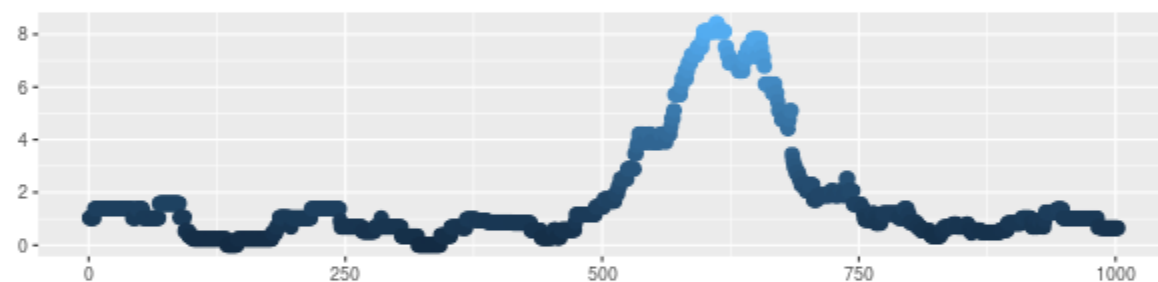
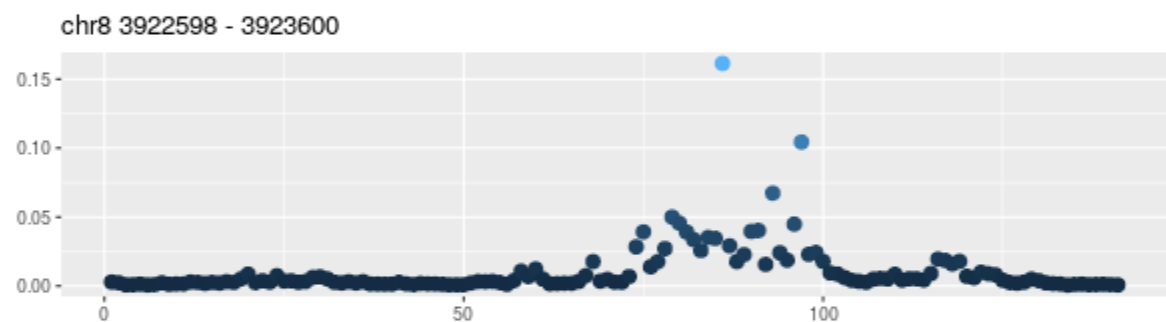
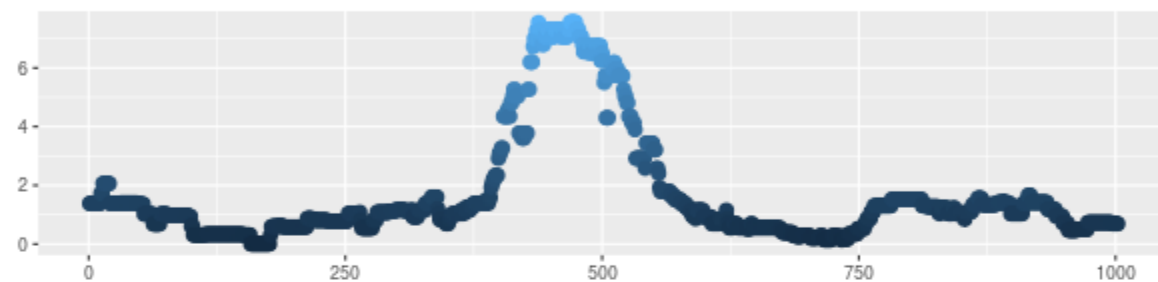
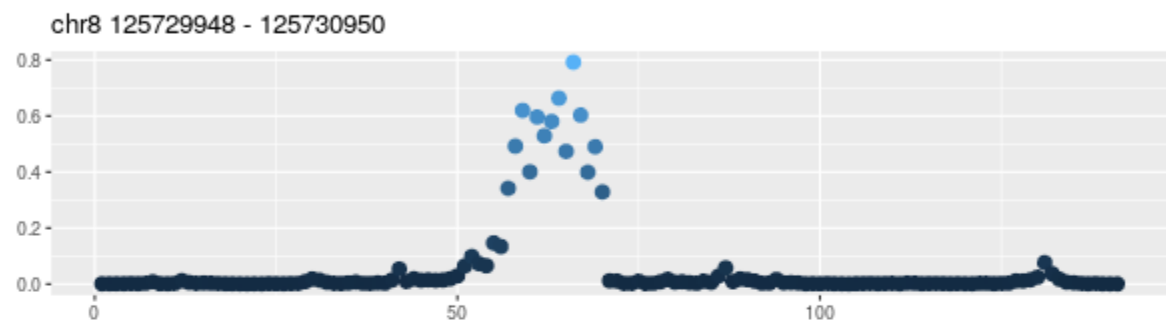
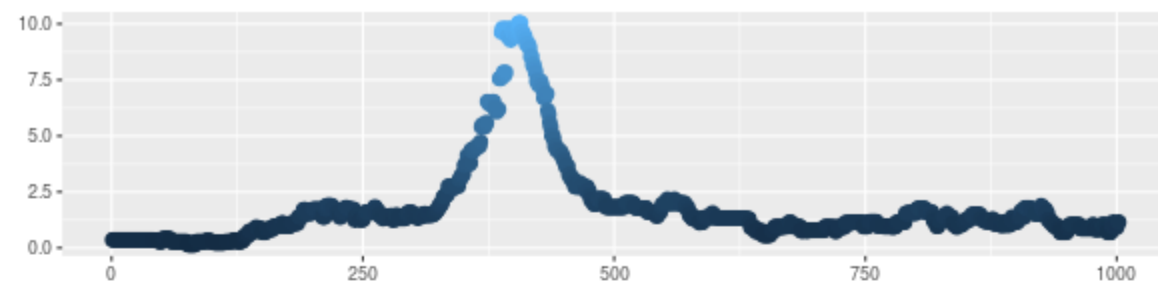
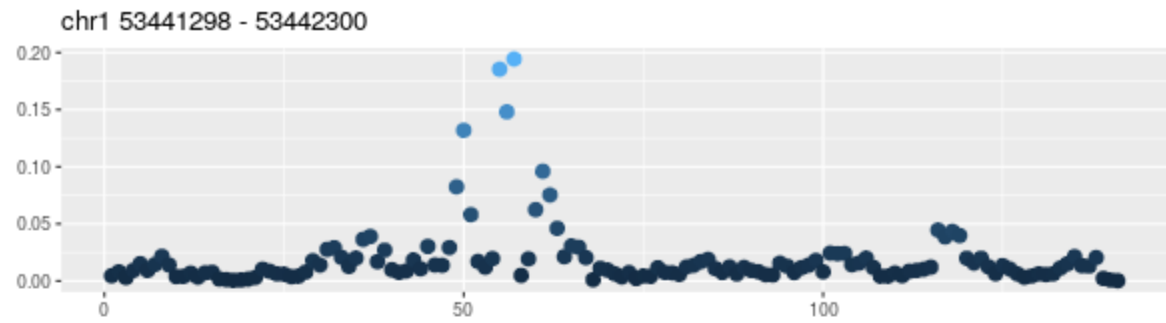


Attention weights

CHIP-seq score

Pairwise attention module

NANOG/induced pluripotent stem cell



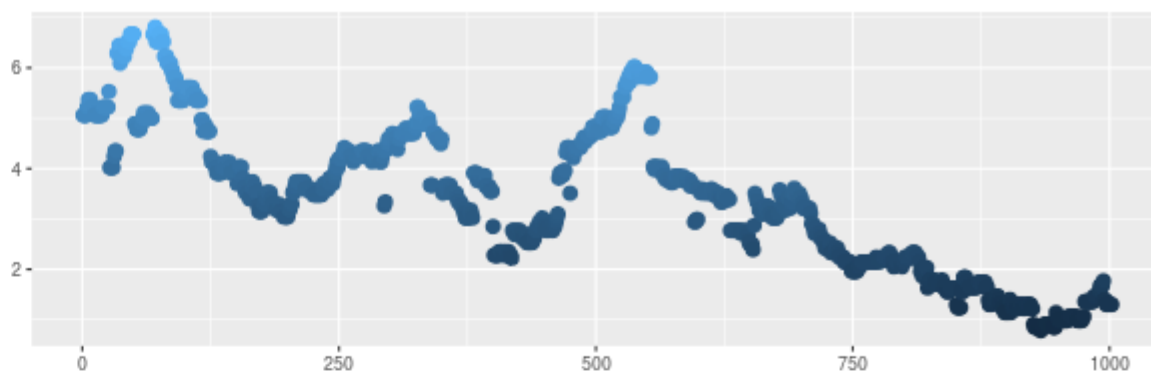
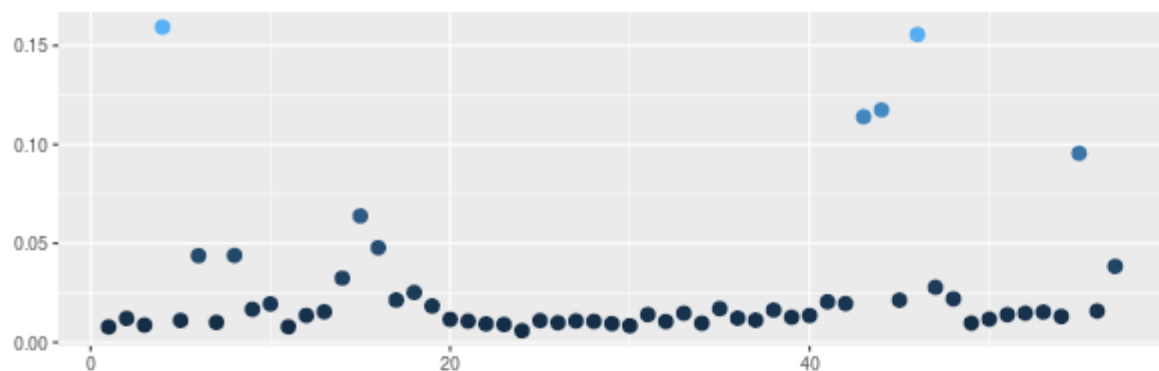
Attention weights

CHIP-seq score

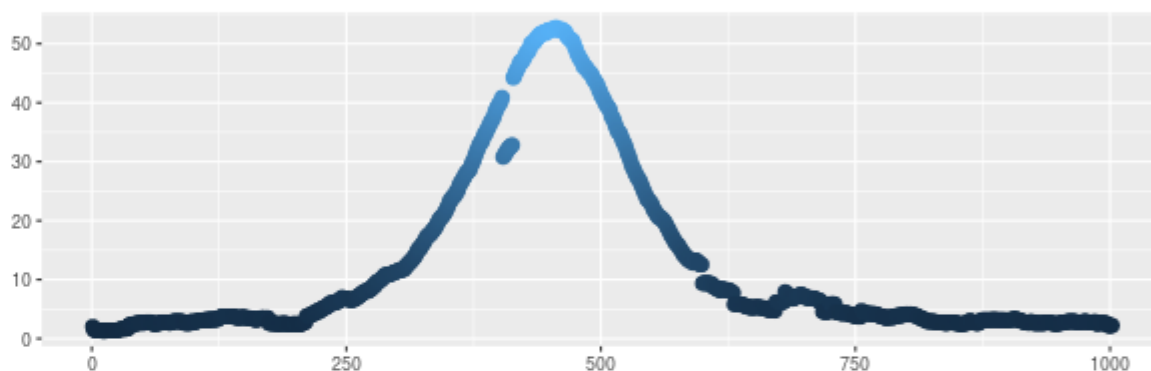
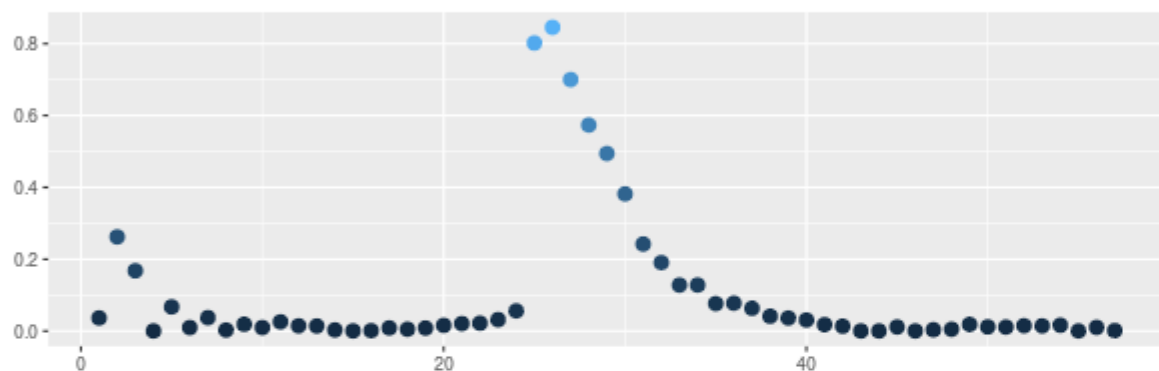
Pairwise attention module

REST/liver

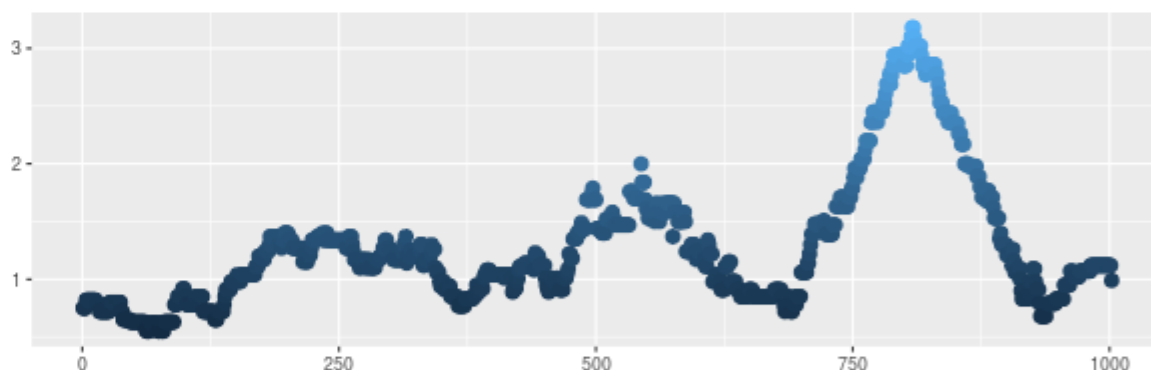
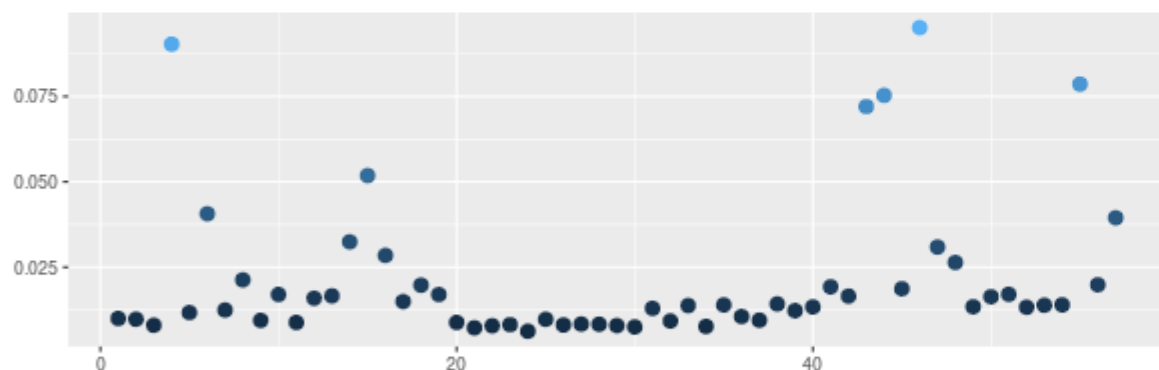
chr8 124428098 - 124429100



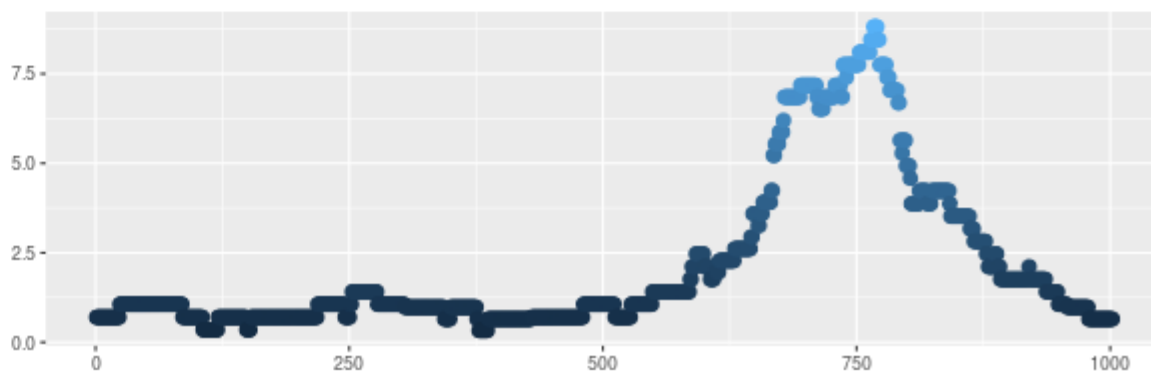
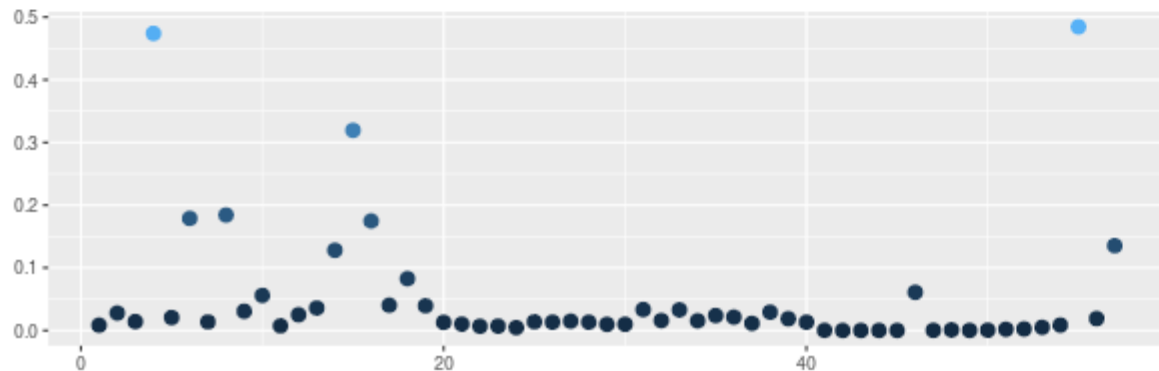
chr1 44433348 - 44434350



chr1 28051348 - 28052350



chr1 211326348 - 211327350



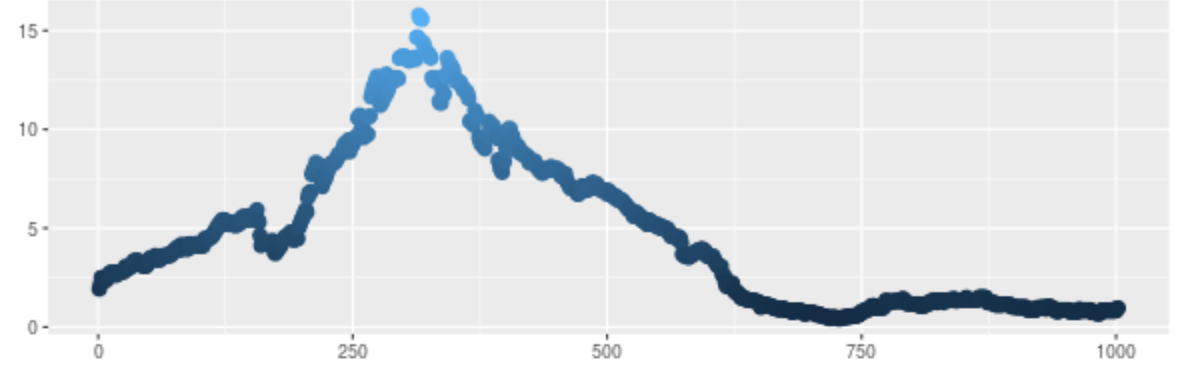
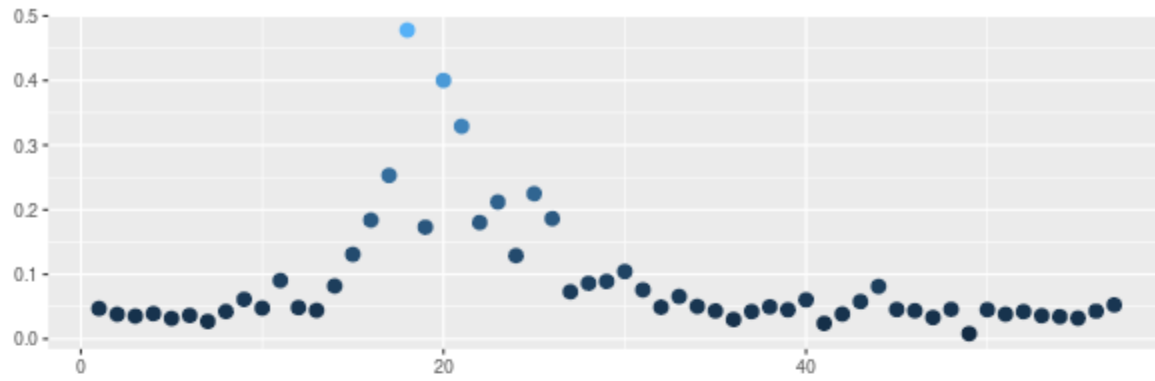
Attention weights

CHIP-seq score

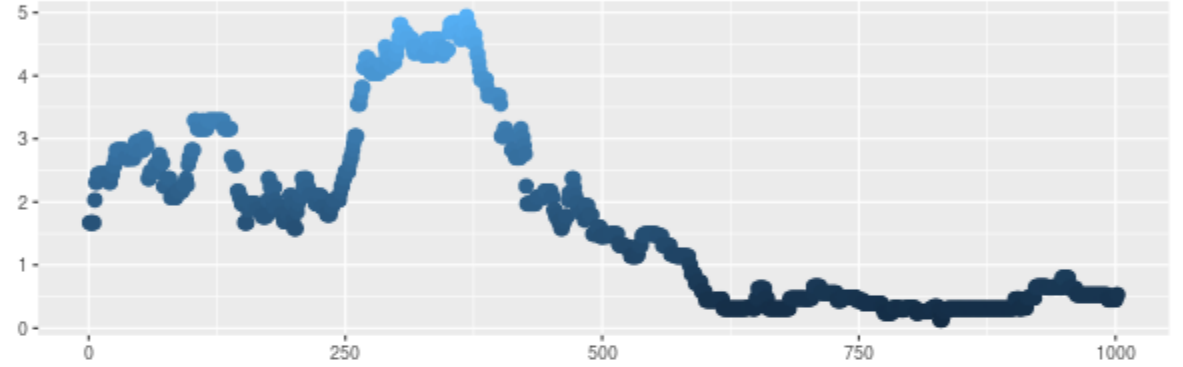
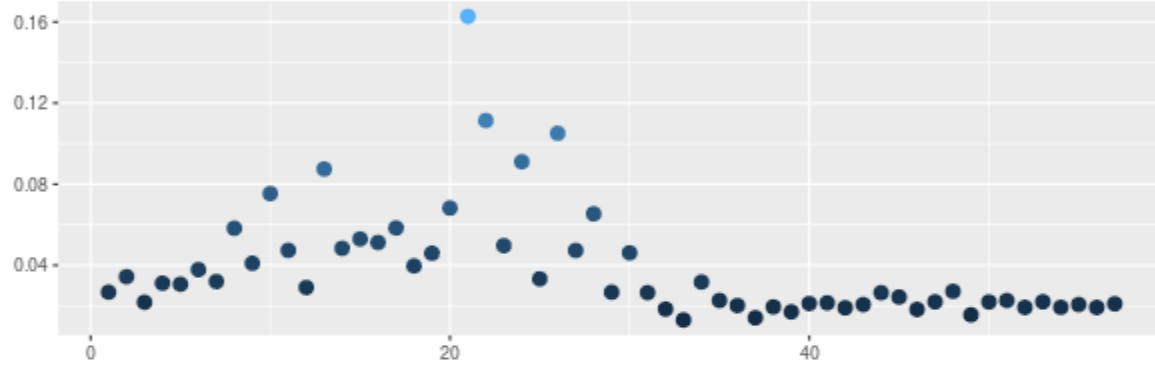
Pairwise attention module

TAF1/liver

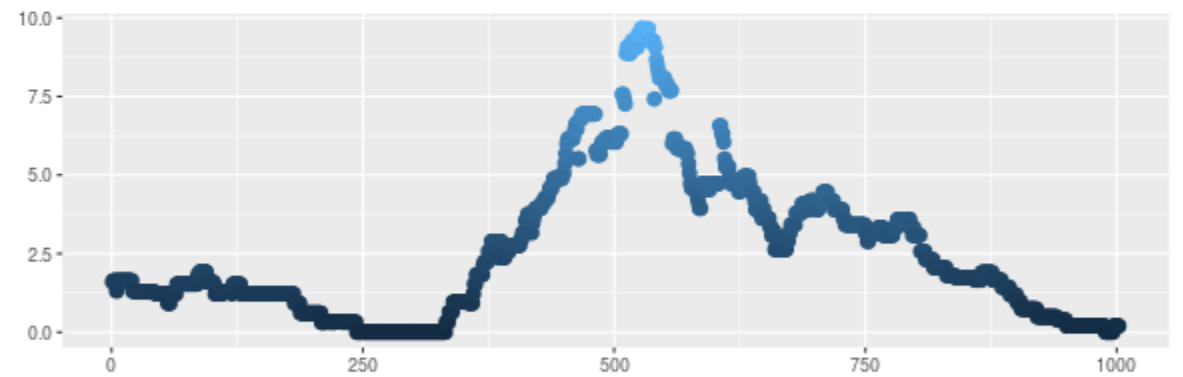
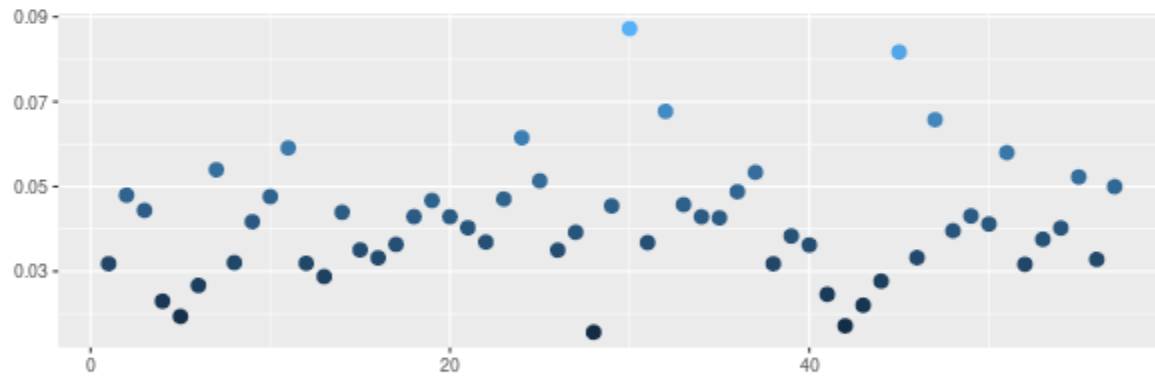
chr1 45986848 - 45987850



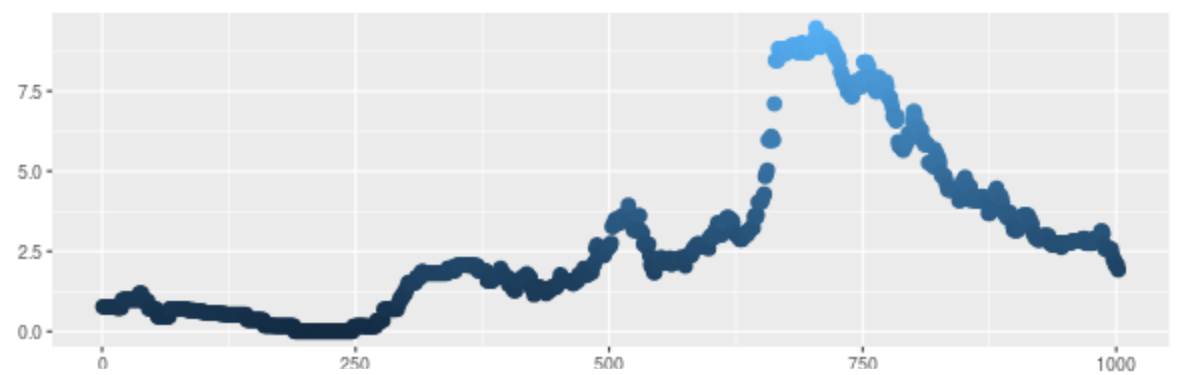
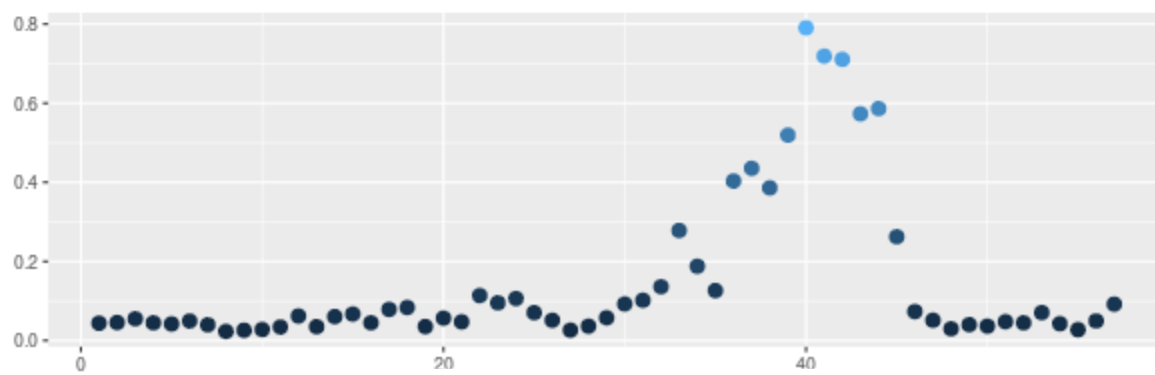
chr1 179050298 - 179051300



chr8 30241148 - 30242150



chr1 150458748 - 150459750



Attention weights

CHIP-seq score