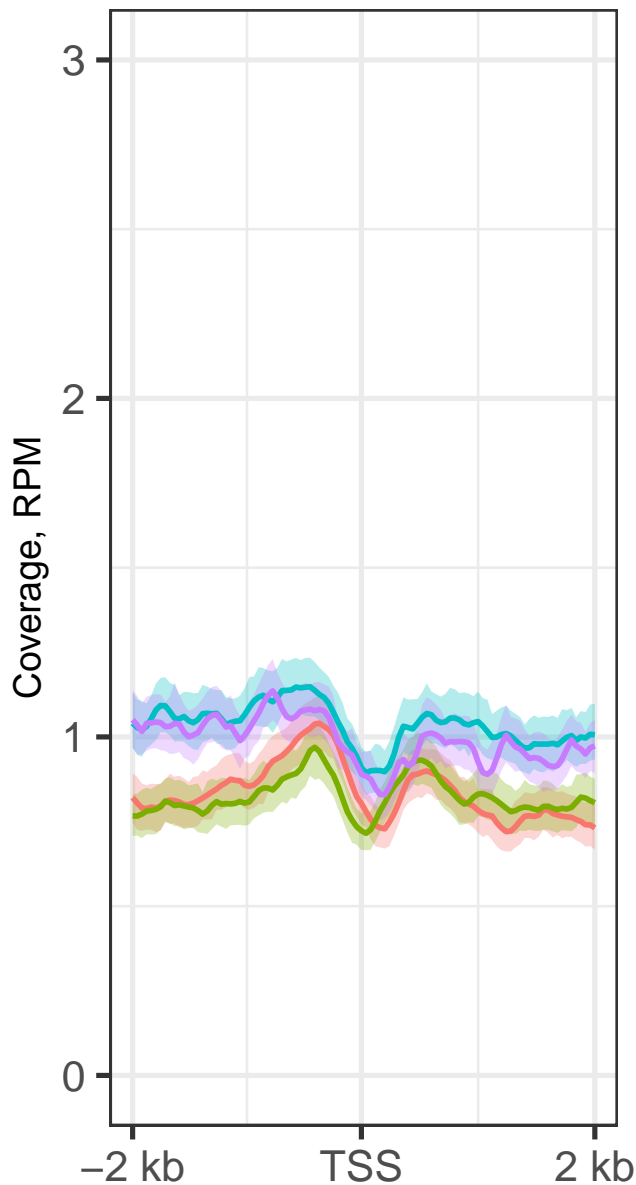


# input\_TPR1\_eds1



## group

- cluster2\_overexpr\_24h\_t3
- TAIR\_2000
- TPR1\_Col\_bound\_genes
- TPR1\_eds1\_bound\_genes