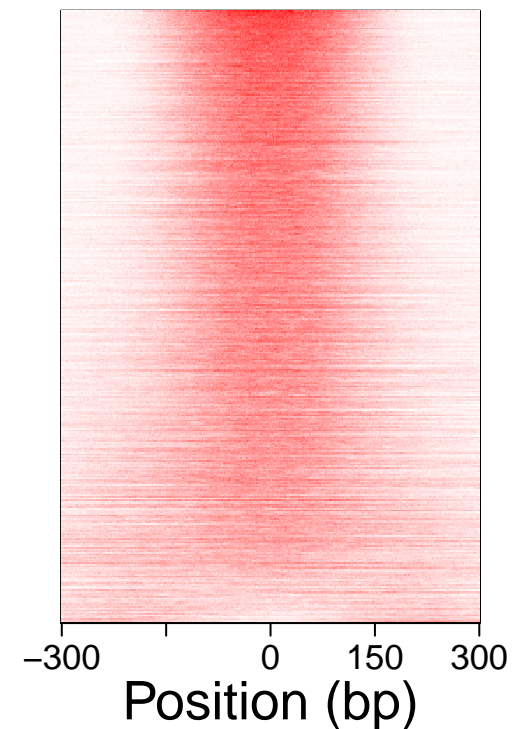
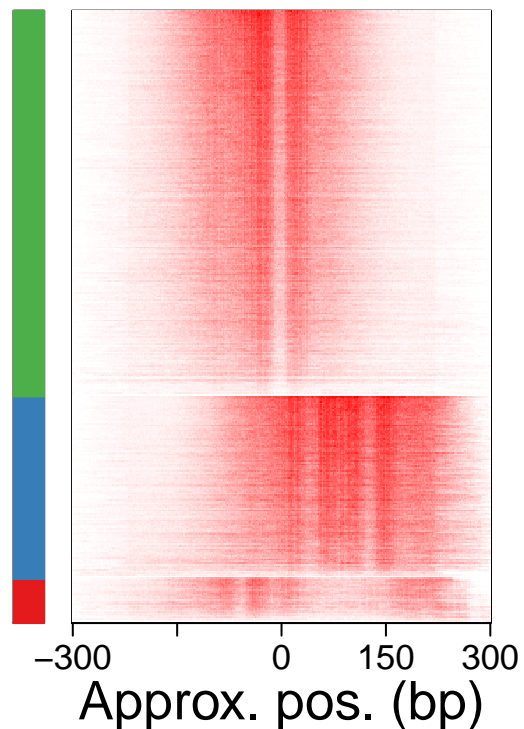
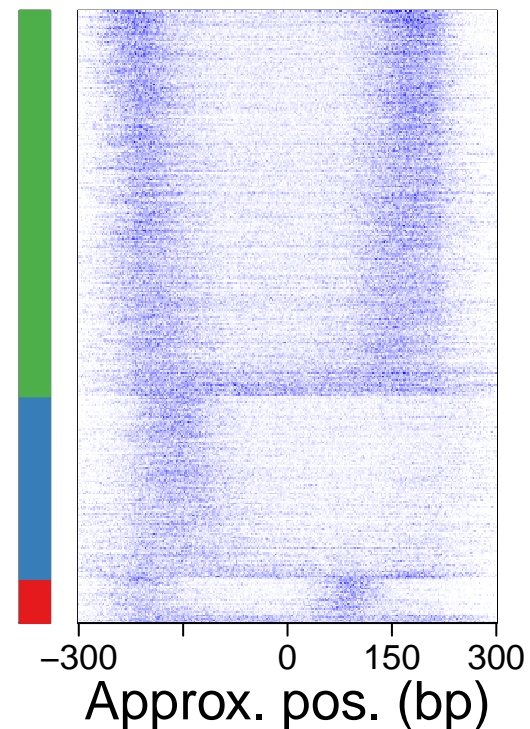
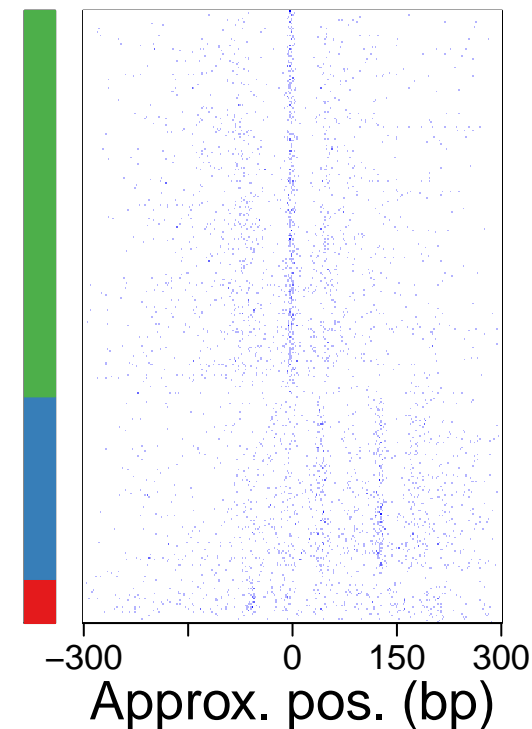
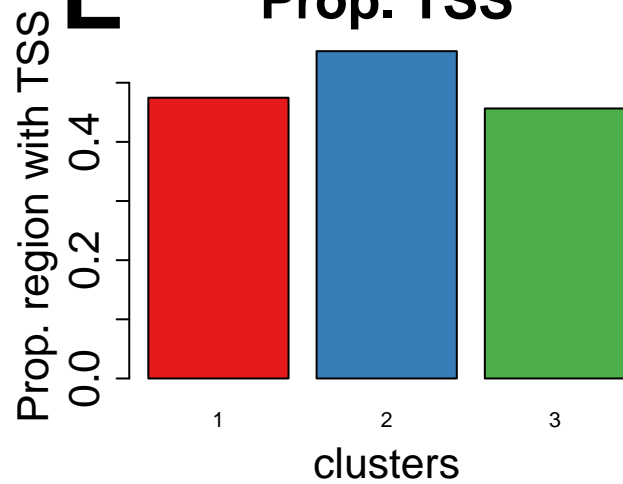
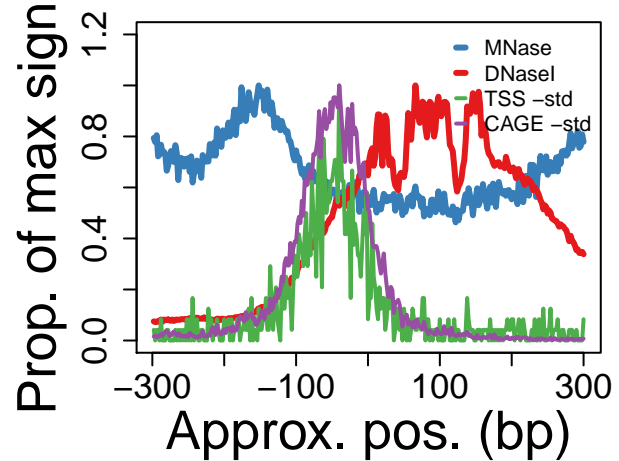


**A DNaseI data****B Aligned DNaseI****C Aligned MNase****D Aligned motifs****E Prop. TSS****F Cluster 2****G De novo discovered motifs**

- Cluster 1**  
 \*NFYA / NFYB  
 \*SP related
- Cluster 2 left**  
 \*SP related  
 \*NFYA / NFYB  
 GATA6 / GATA3  
 SFPI1 / SPIC  
 FOX related  
 ARNTL / BHLHE41
- Cluster 2 right**  
 \*SP related  
 \*NFYA / NFYB  
 ARNTL / BHLHE41
- Cluster 3**  
 \*SP related (c)  
 \*NFYA / NFYB  
 \*GATA related  
 ETS related  
 ATF1  
 \*AP1 related  
 \*NRF1 (c)