

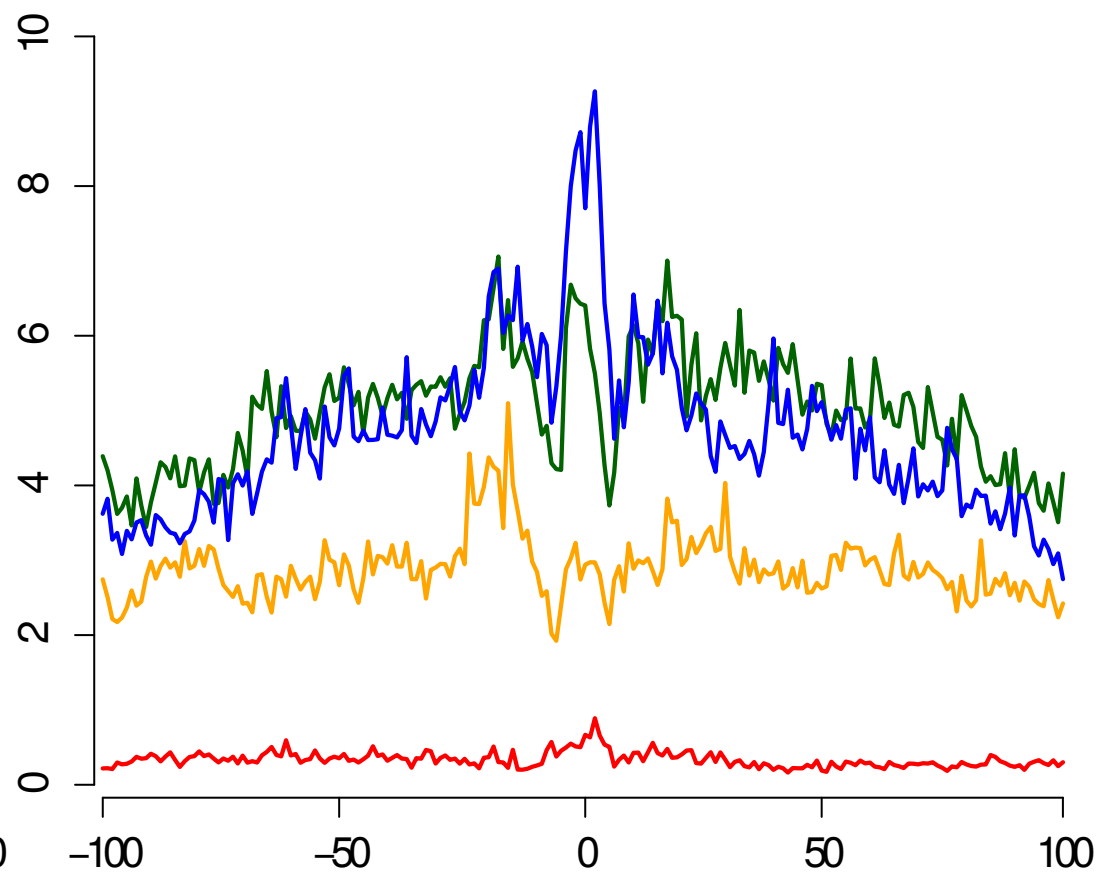
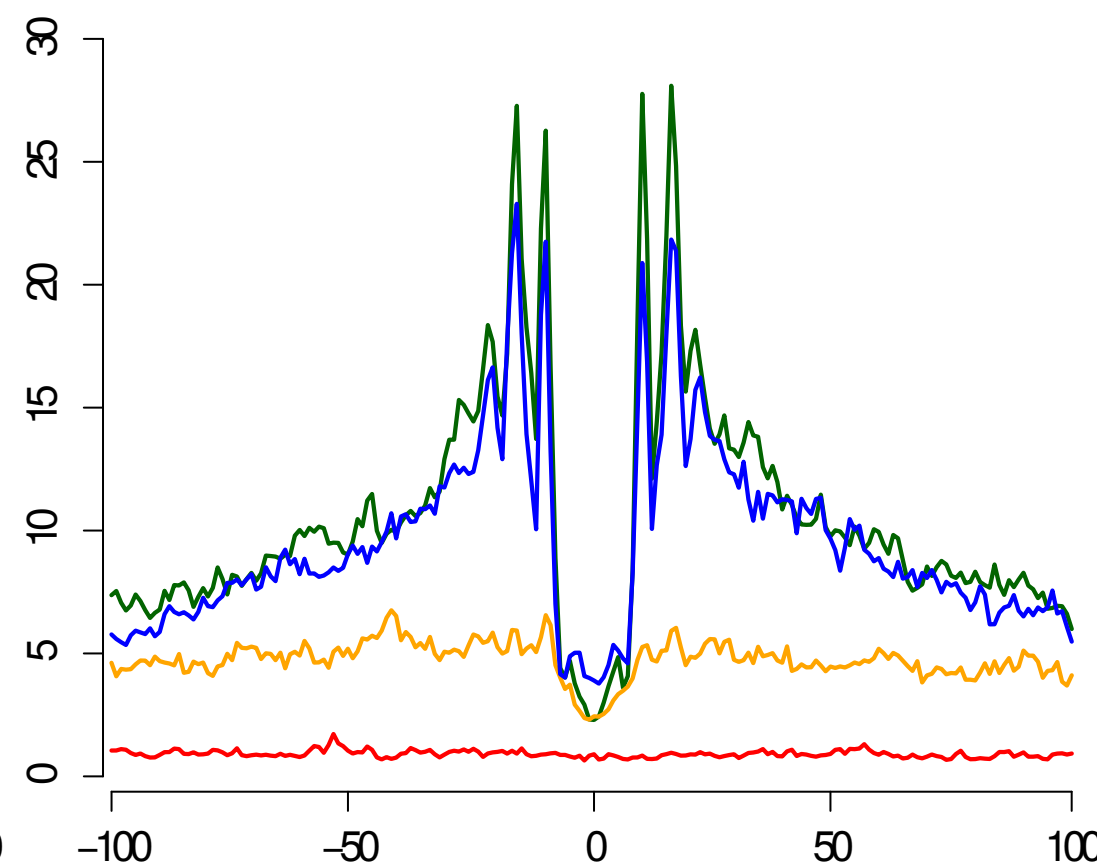
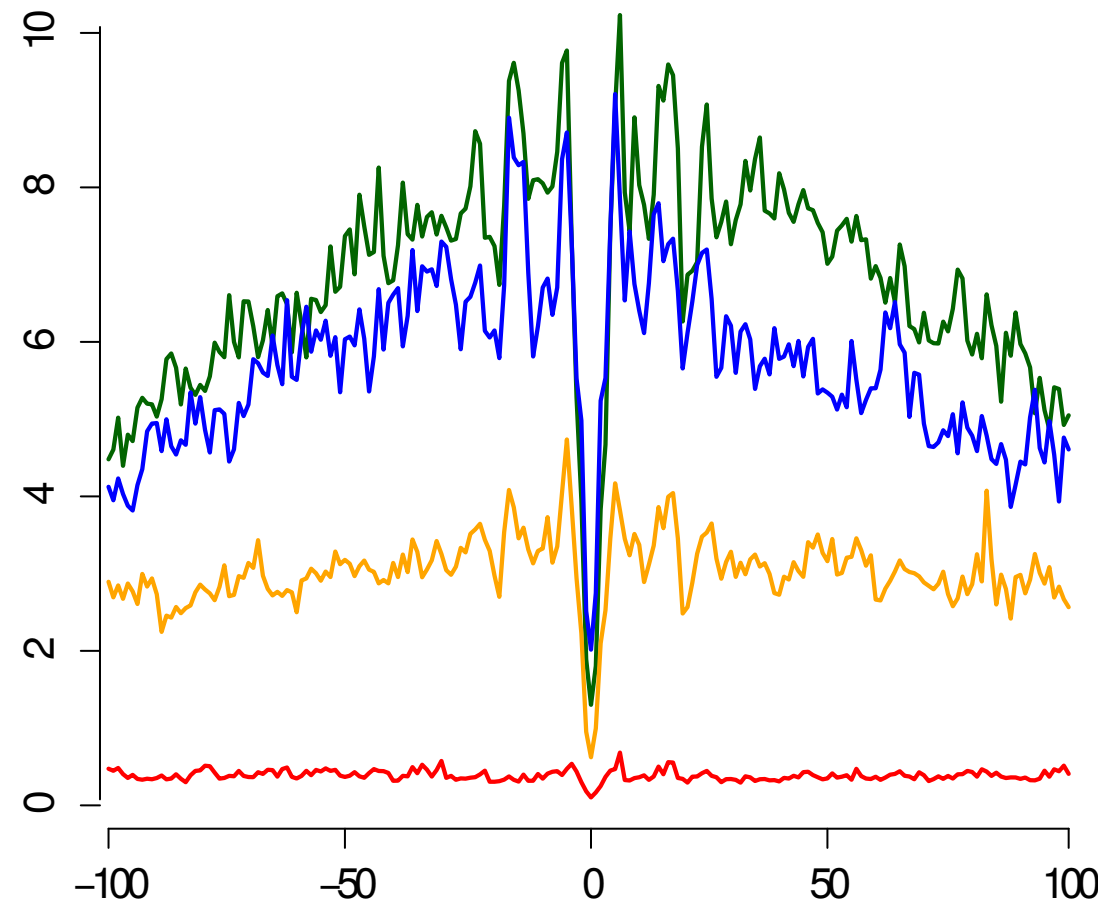
**a**

GABP

NRF1

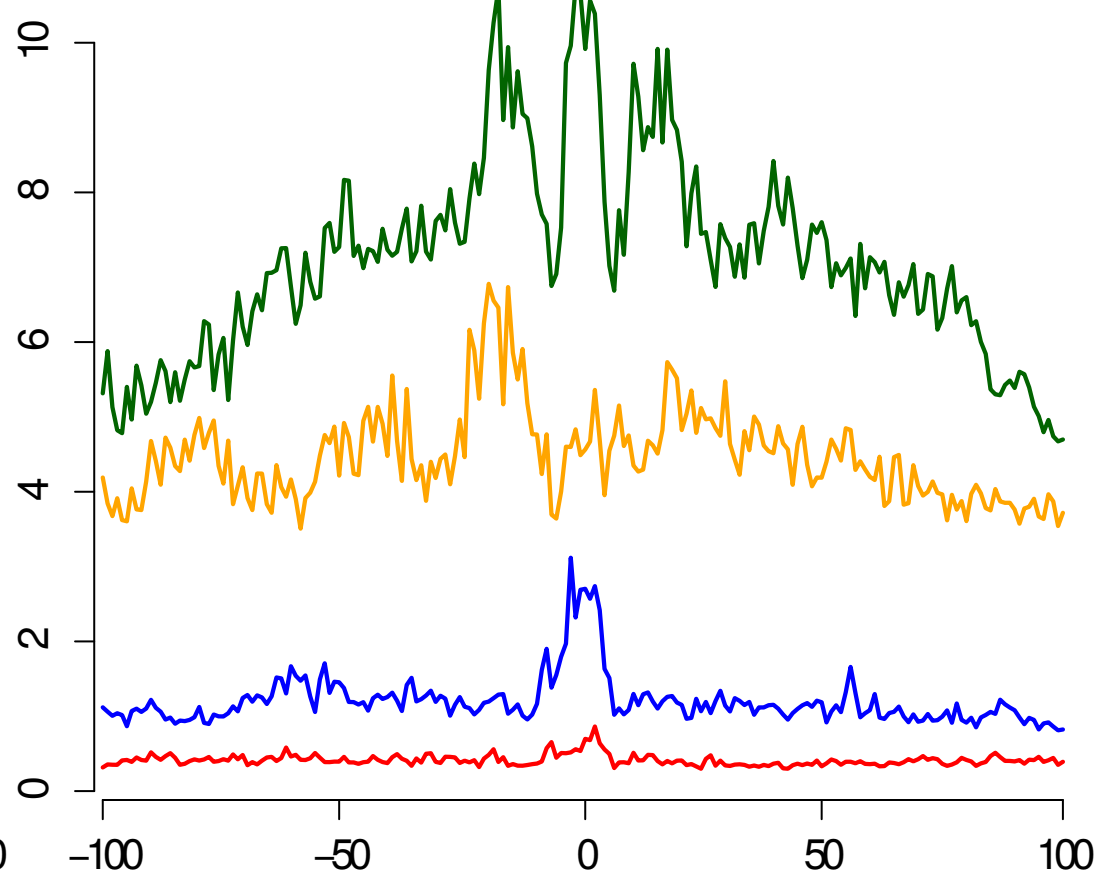
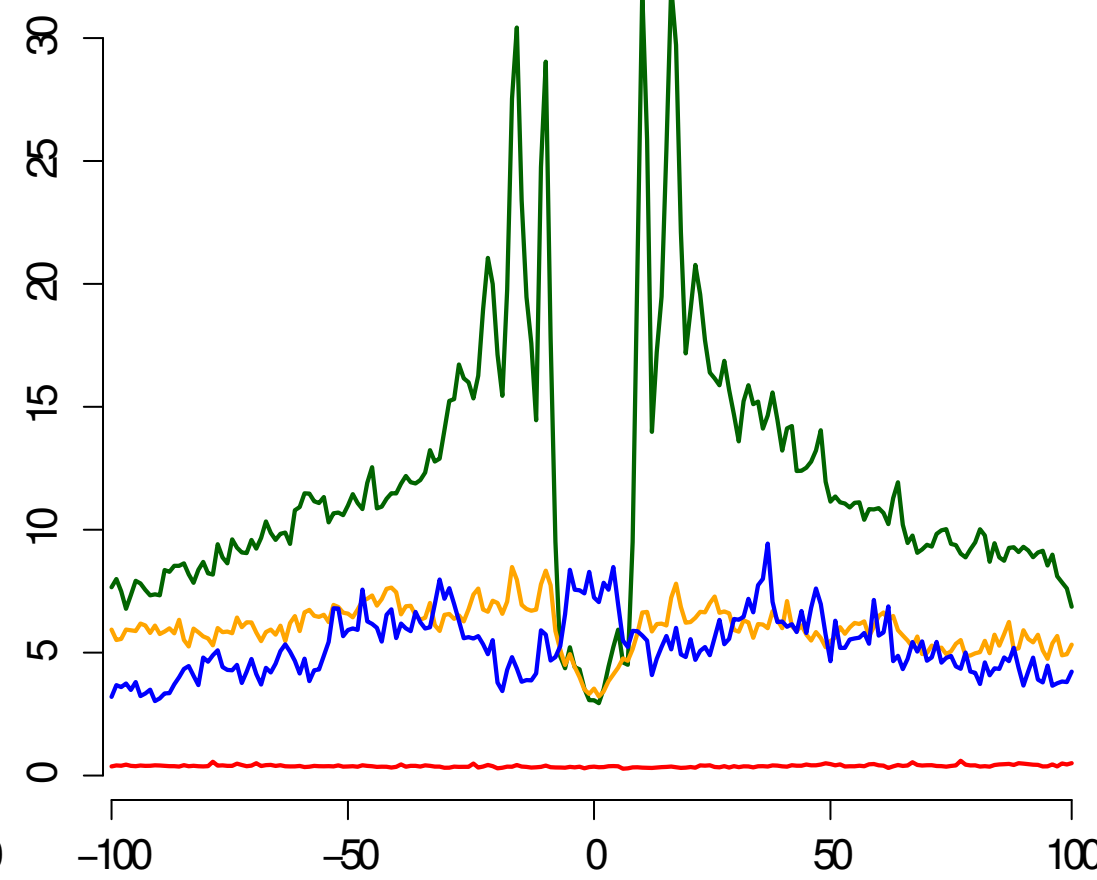
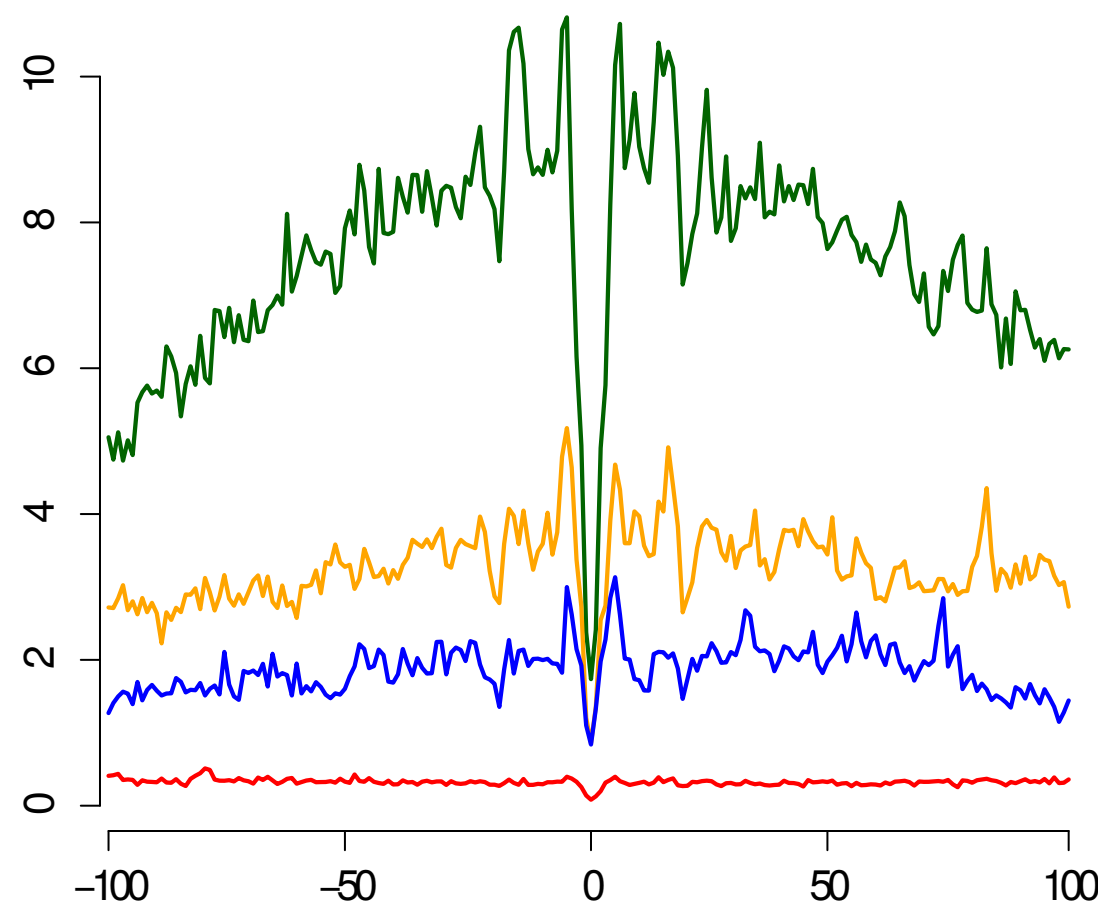
EGR1

Average Uncorrected  
DNase-seq Signal



**b**

Average Bias-Corrected  
DNase-seq Signal



Distance from Motif Center (bp)