

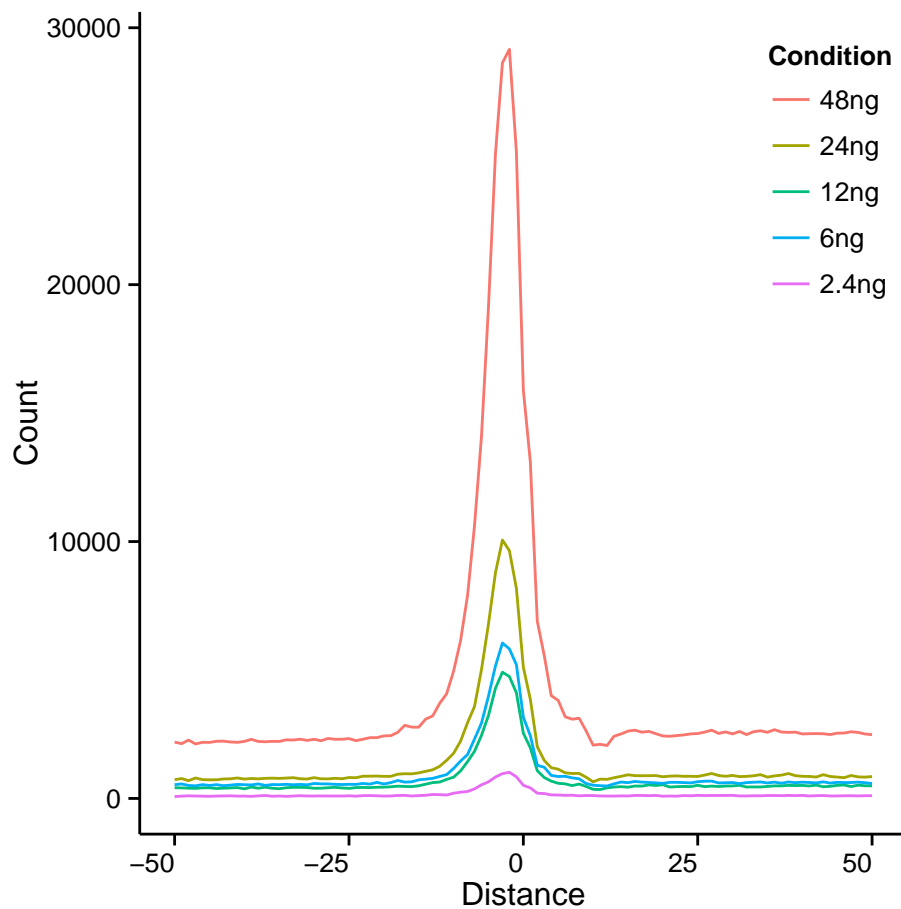
plot Jump-seq reads distribution

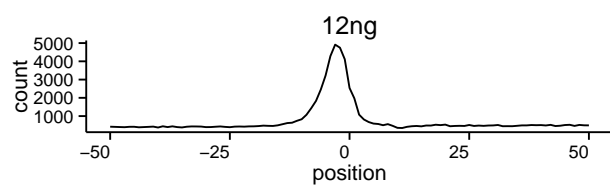
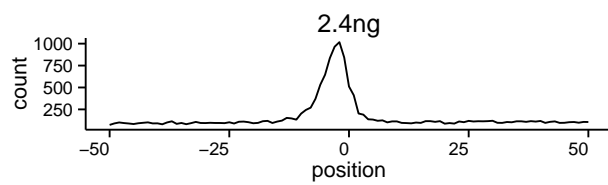
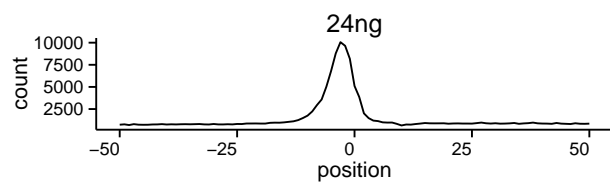
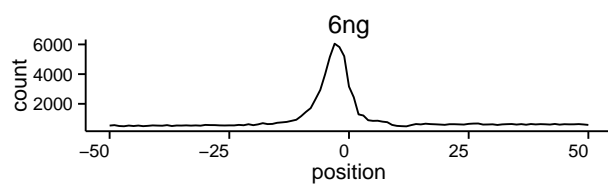
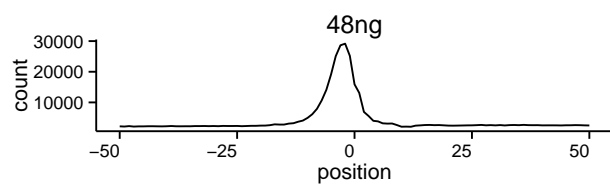
Use Tab-Seq FDR 0.484 peaks for hmc sites Use GSM1173118 strand-unspecific sites (signal 1, confidence 1)
plus reads and minus sites

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## Loading required package: lattice
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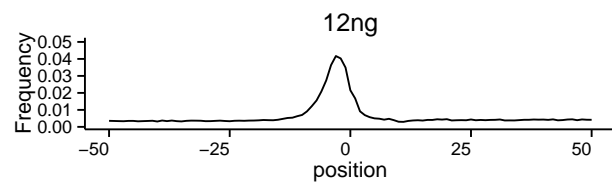
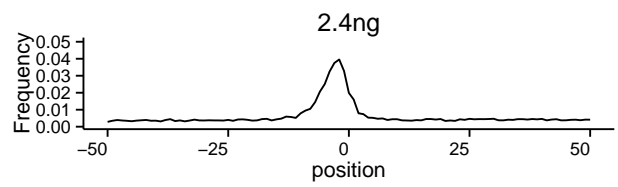
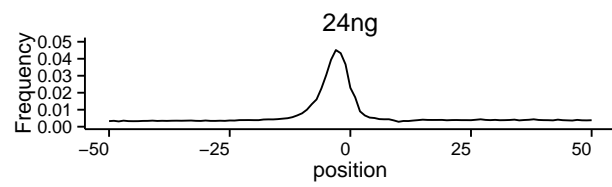
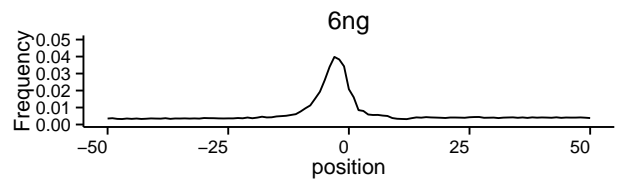
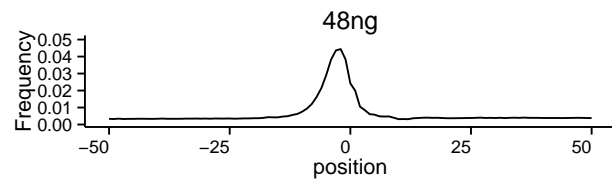
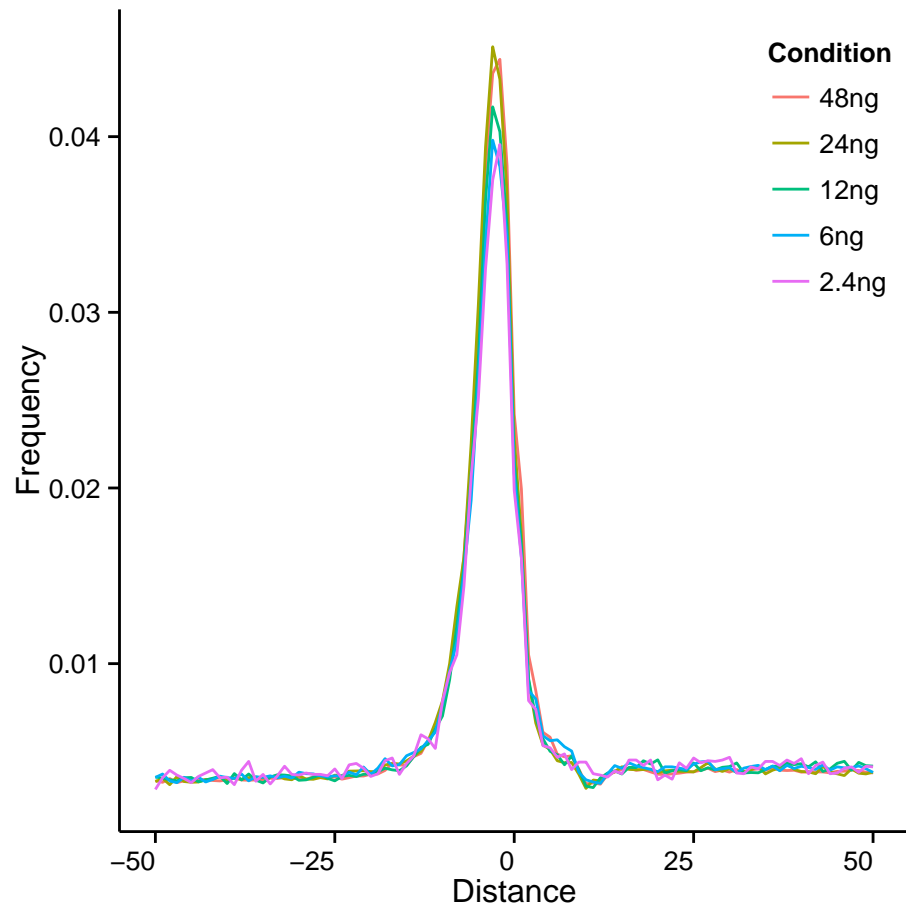
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## Loading required package: plyr
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- 5mc plus sites, unnormalized

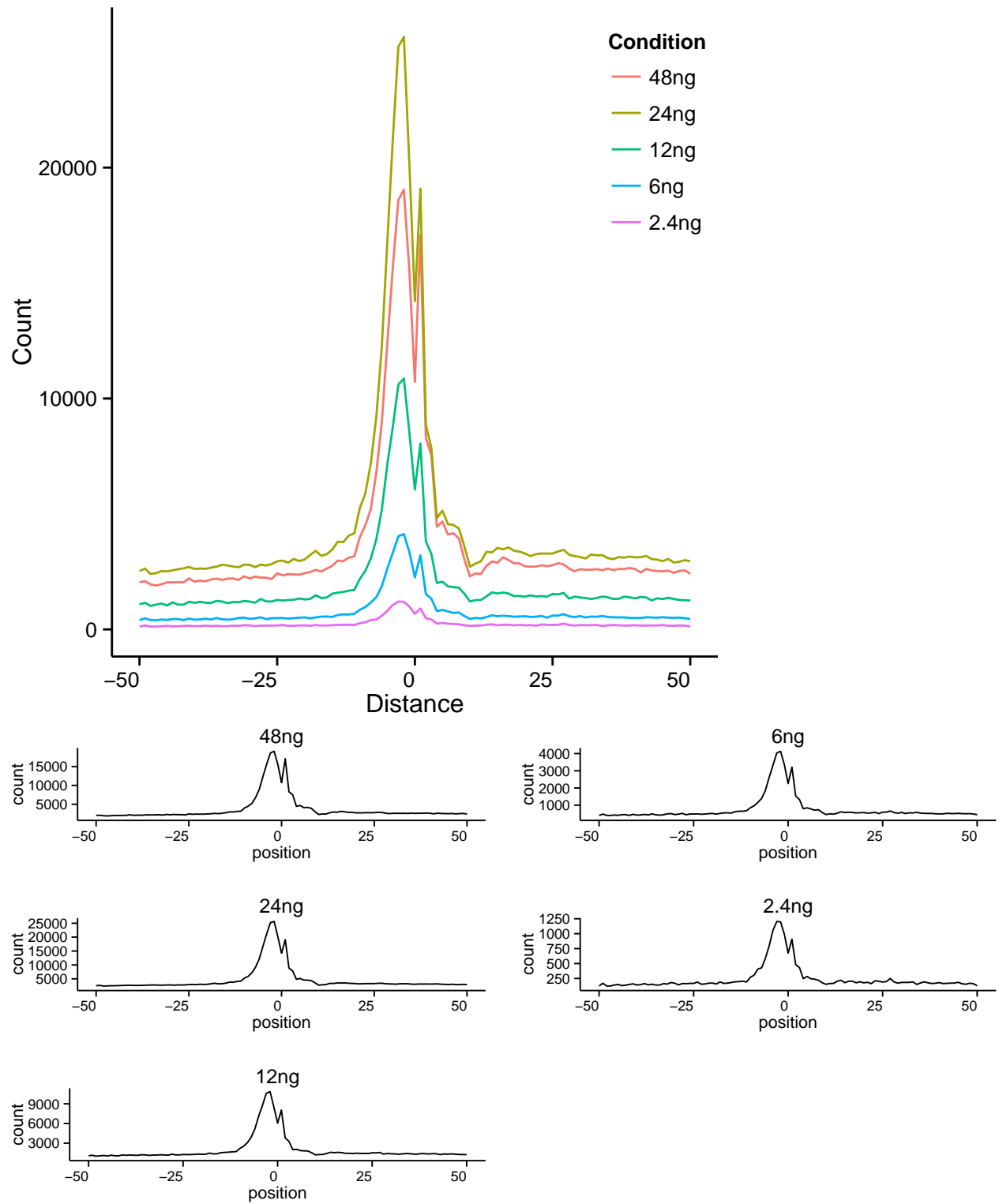




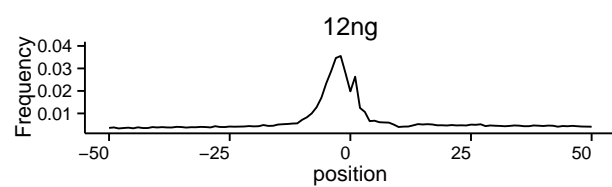
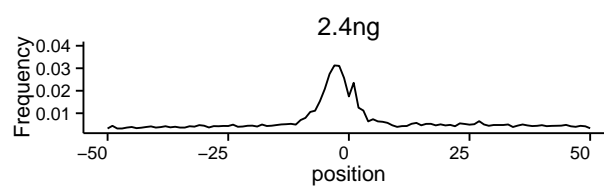
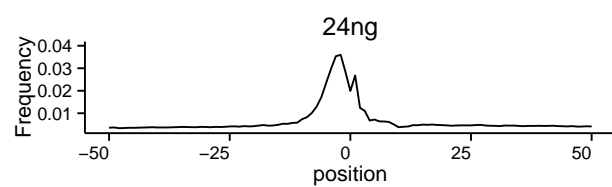
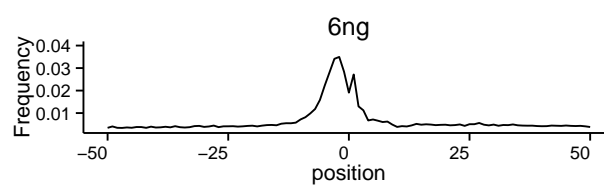
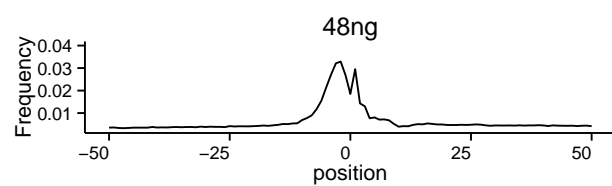
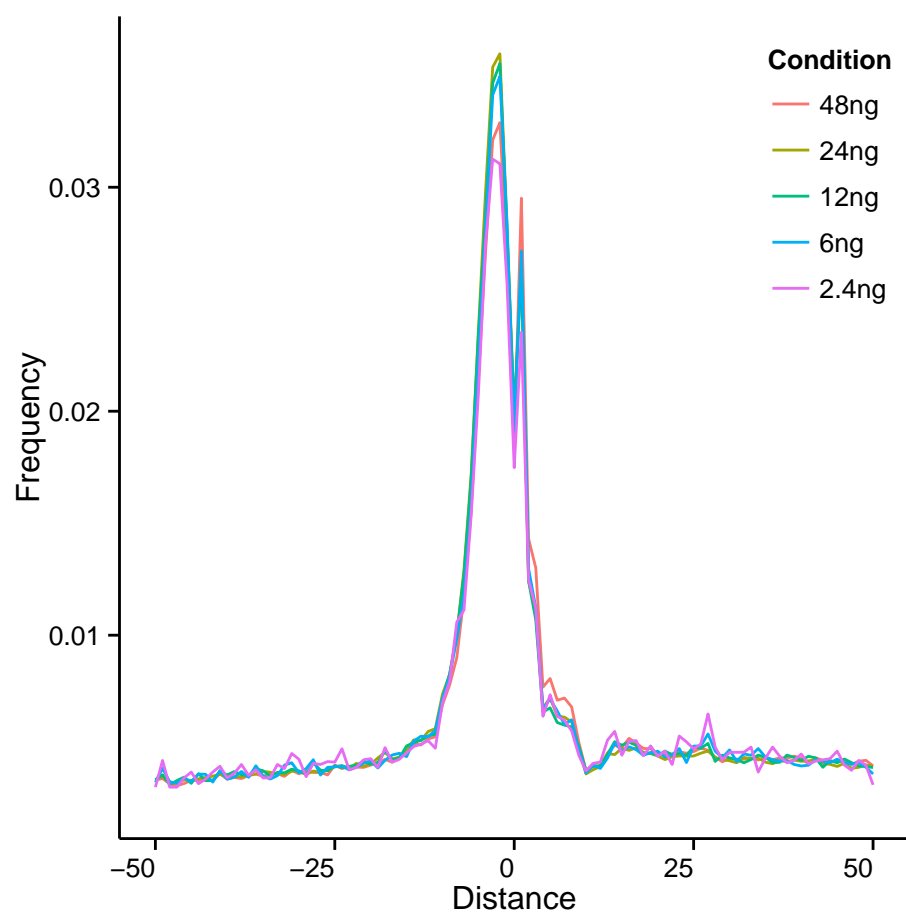
* 5mc plus sites, normalized



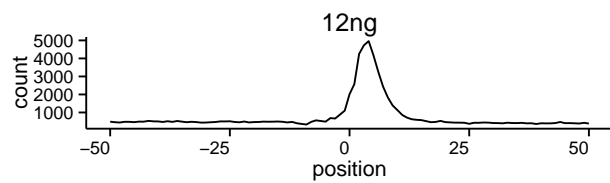
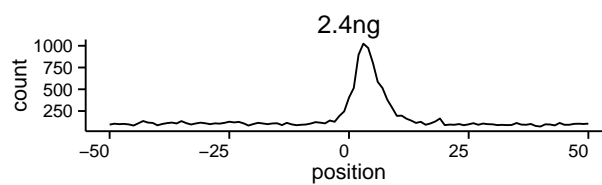
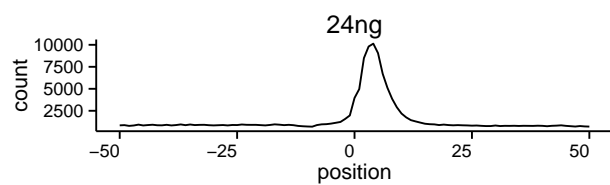
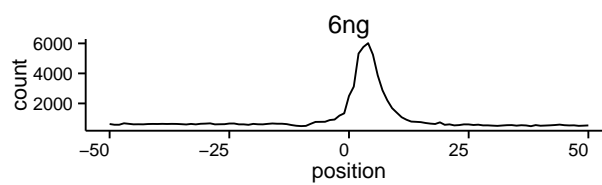
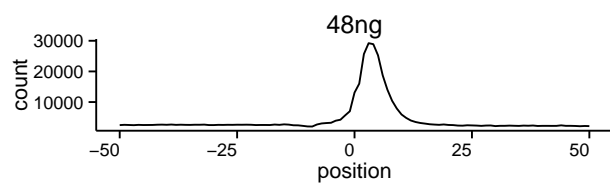
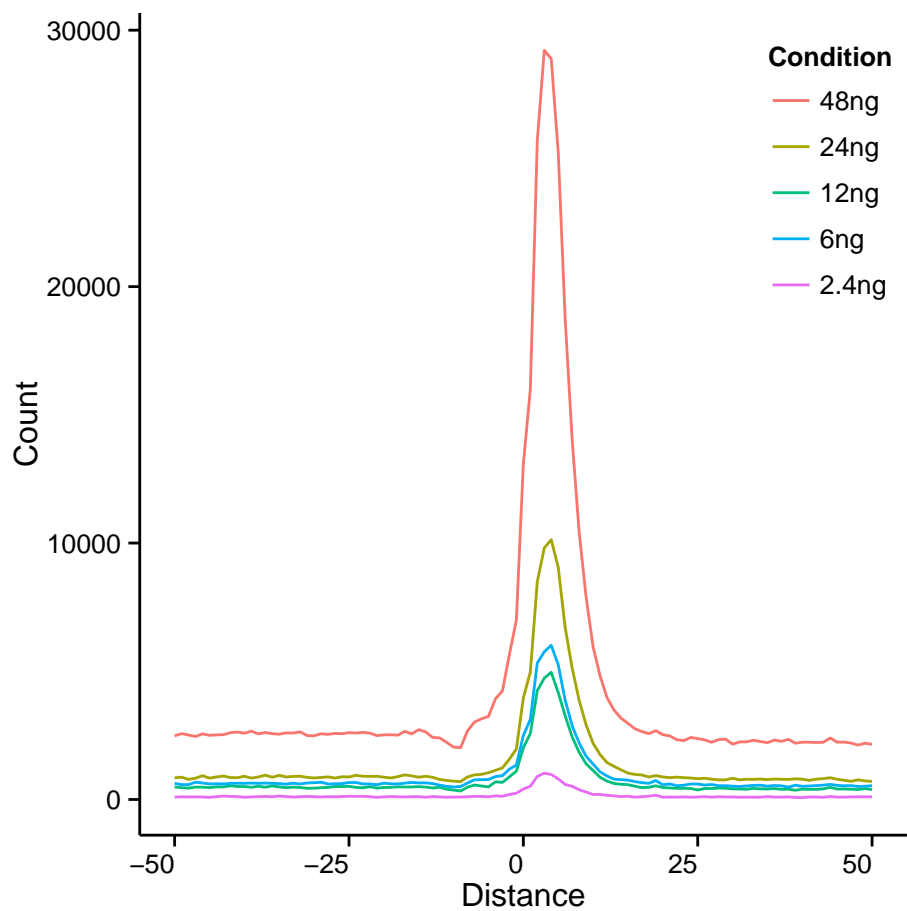
* 5hmc plus sites, unnormalized



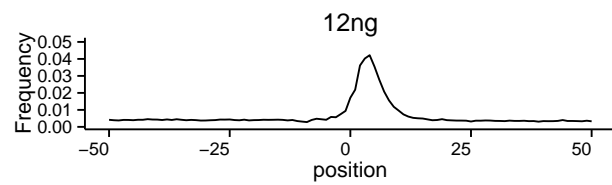
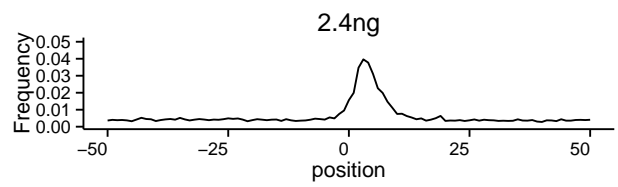
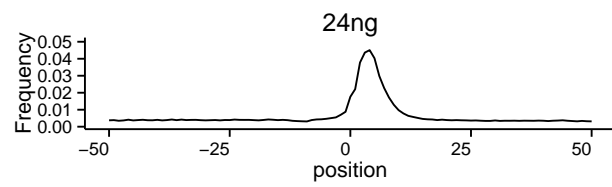
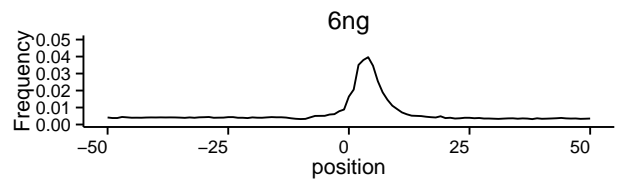
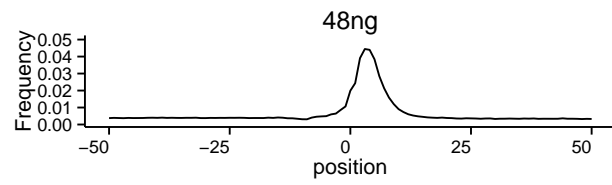
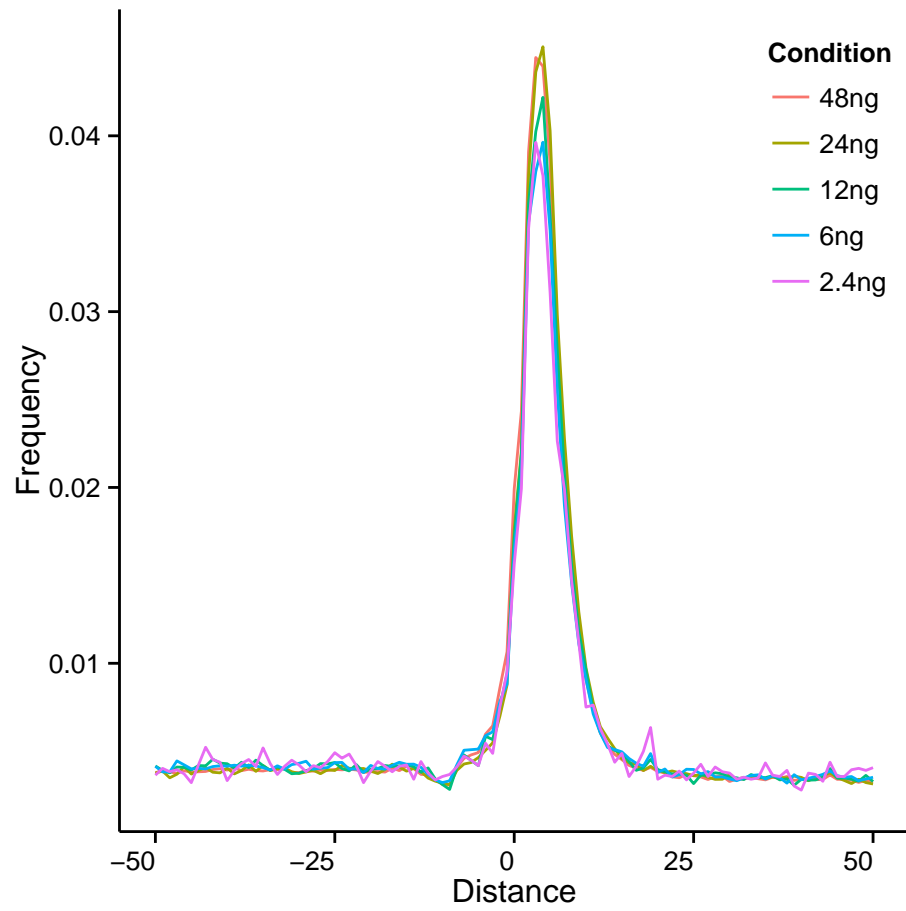
* 5hmc plus sites, normalized



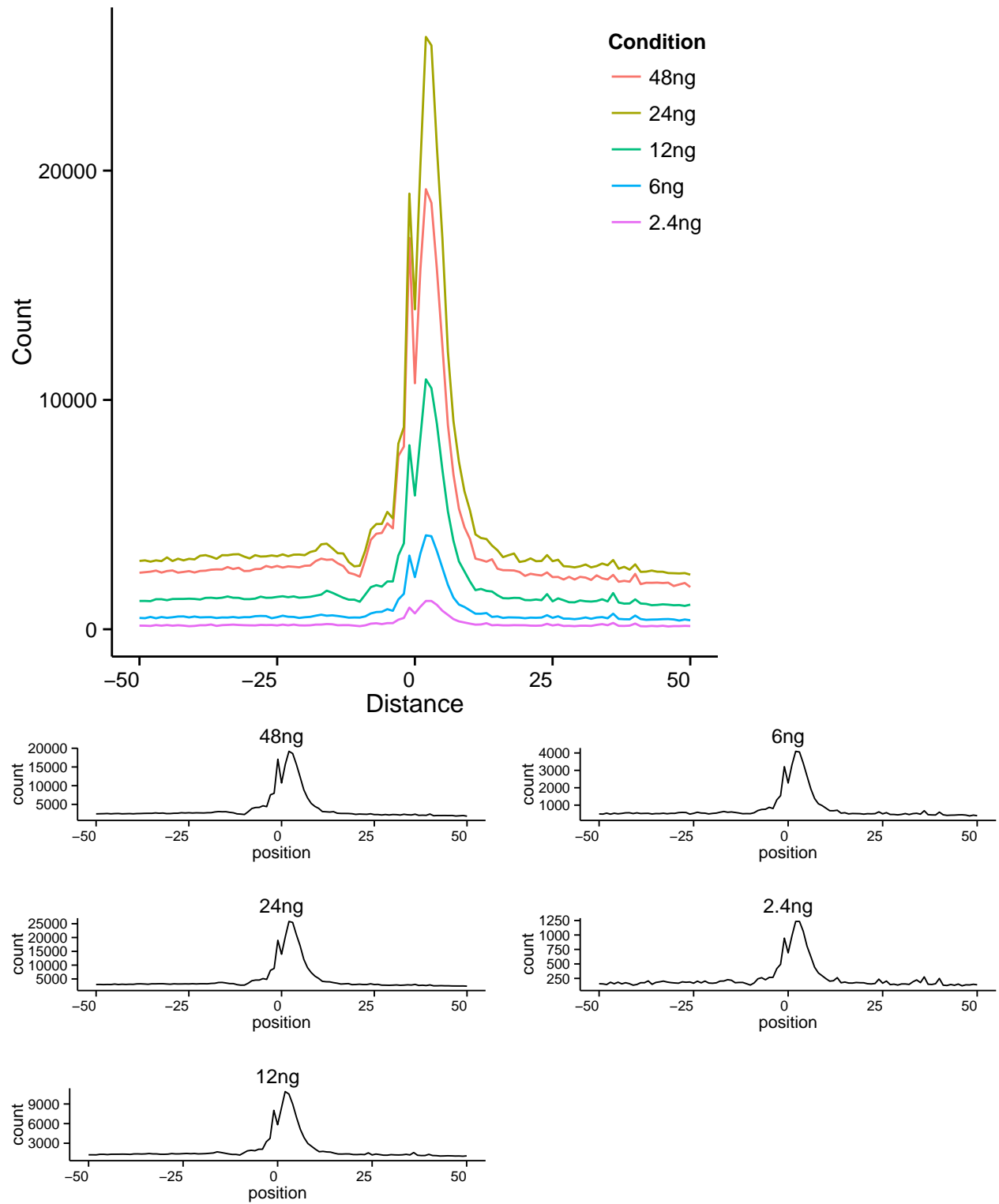
- 5mc minus sites, unnormalized



* 5mc minus sites, normalized



* 5hmc minus sites, unnormalized



* hmc minus sites, normalized

