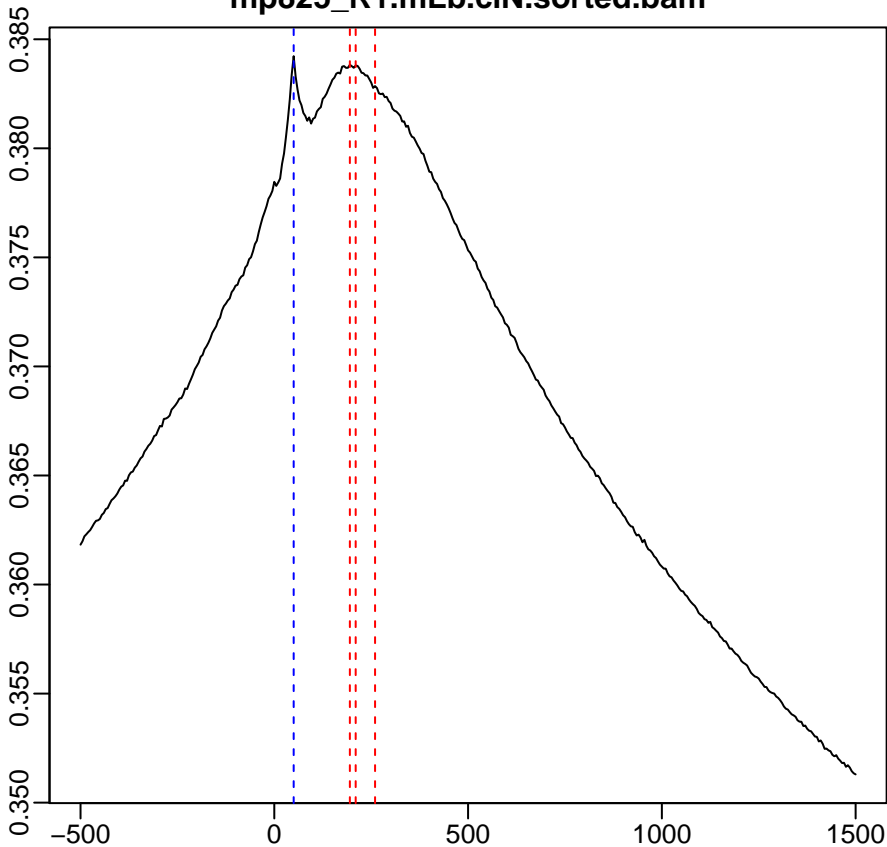


mp825_R1.mLb.cIN.sorted.bam

cross-correlation



strand-shift (195,210,260)

NSC=1.09262,RSC=0.988028,Qtag=0