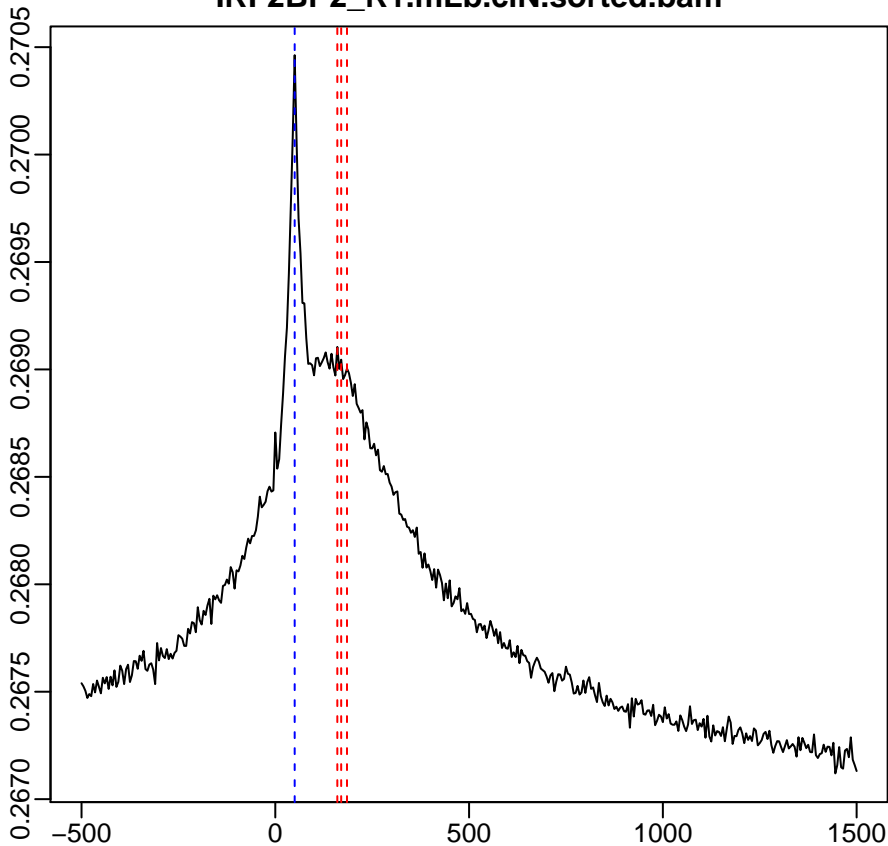


# IRF2BP2\_R1.mLb.cIN.sorted.bam

cross-correlation



strand-shift (160,170,185)  
NSC=1.00739,RSC=0.592468,Qtag=0