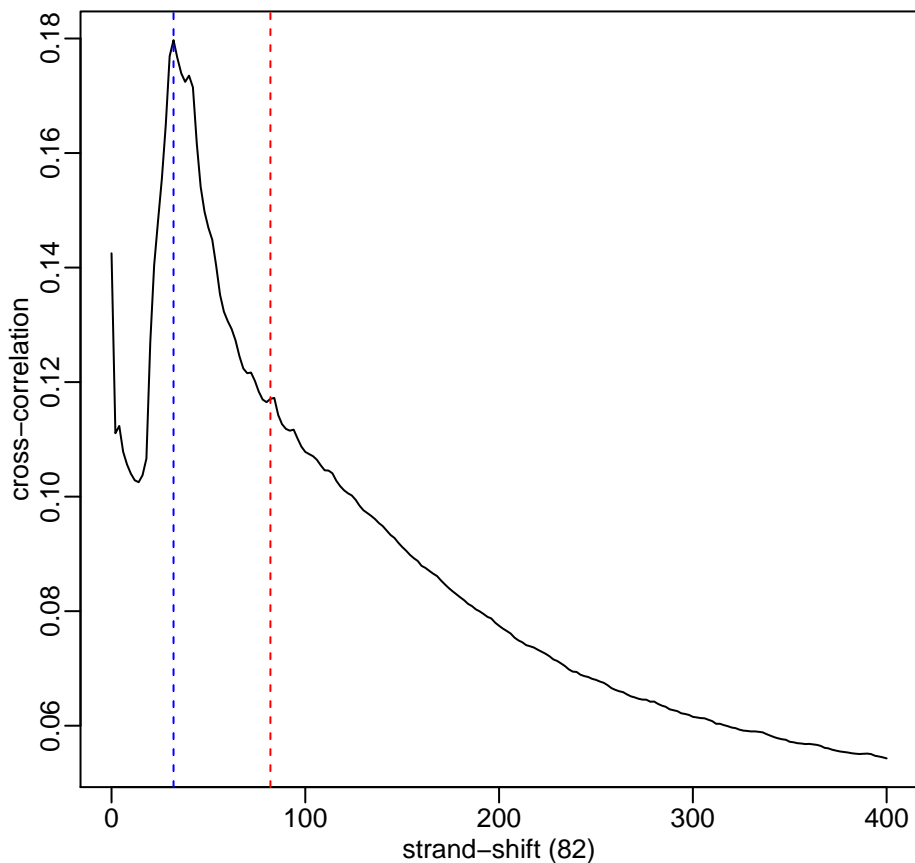


# DNaseSeq\_Lung\_B.bam



NSC=2.15376,RSC=0.499491,Qttag=-1