GFI1_R1.mLb.clN.sorted.bam 0.240 11.1 11.1 11.1 11.1 11.1 11.1 11.1 11.1 11.1 11.1 cross-correlation 11.1 11.1 0.238 11.1 0.236 -500 1500

500

strand-shift (140,155,175) NSC=1.02315,RSC=1.06712,Qtag=1

1000