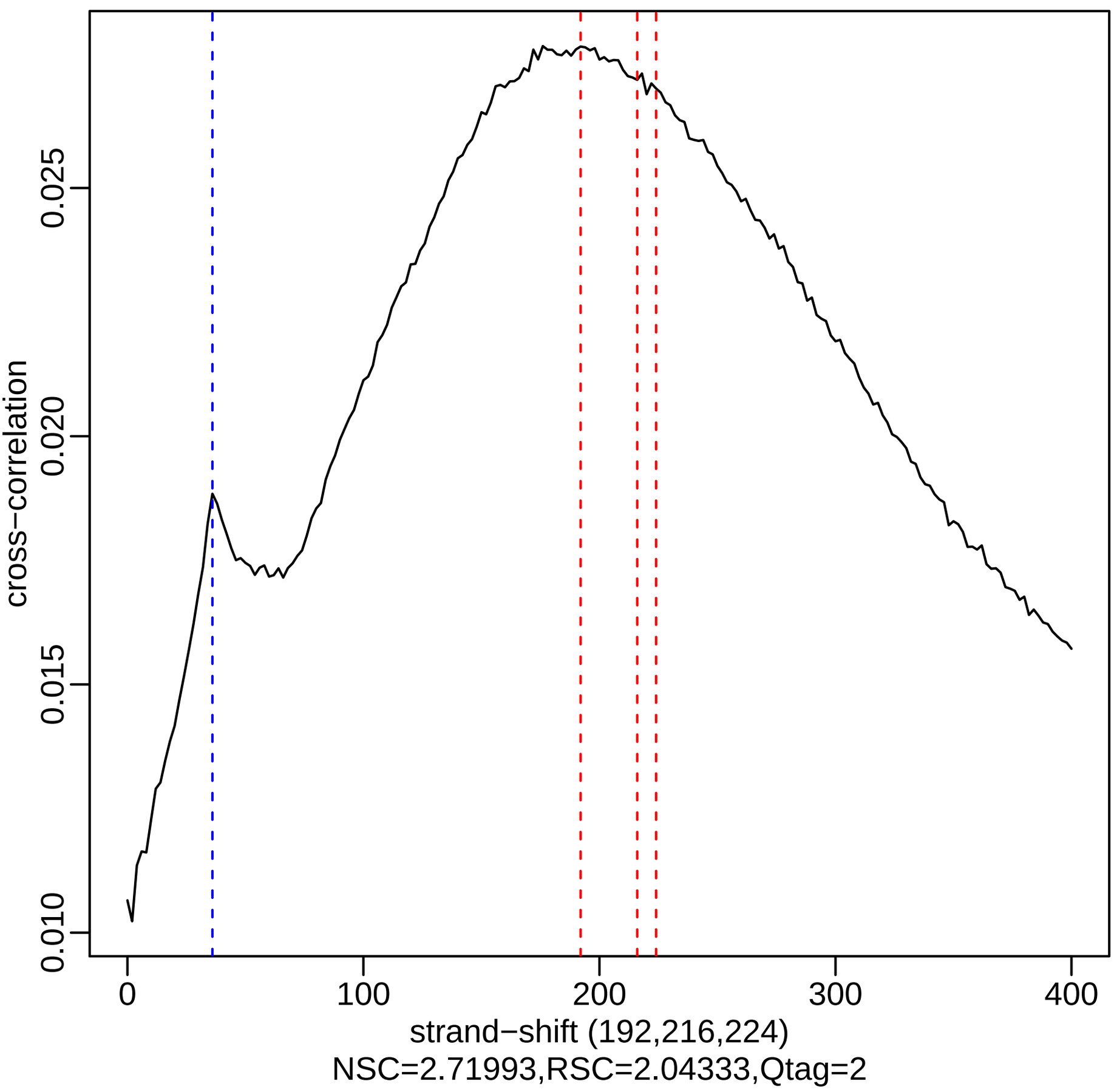
GRF2_B73.1x36mers.unique.bam
cross−correlation
0.010
0.015
0.020
0.025
0
100
200
300
400
strand−shift (192,216,224)
NSC=2.71993,RSC=2.04333,Qtag=2