

p-arm

q-arm

H9 hap2  
H9 hap1  
RPE1 hap2  
RPE1 hap1  
YAO pat  
YAO mat  
I002C hap2  
I002C hap1  
HG002 pat  
HG002 mat  
CHM13  
GRCh38

0.00 0.01 0.02 0.03 0.04 0.05  
Distance to end (%chr)

99.95 99.96 99.97 99.98 99.99 100.00  
Distance to end (%chr)

