

genomeW.fa

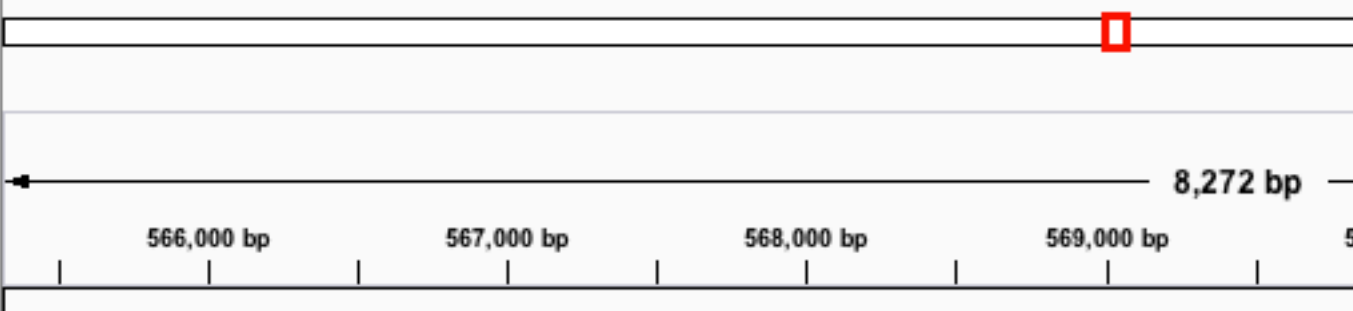
W

W:565,306-573,594

Go



NAME  
DATA TYPE  
DATA FILE



bamvasn\_W.sorted.bam Coverage



bamvasn\_W.sorted.bam

