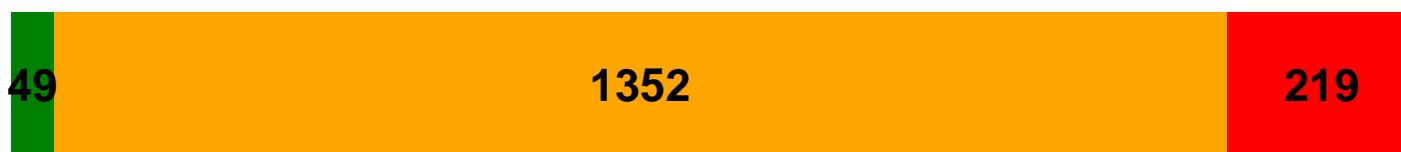


# 100X coverage, run 6

filter criterion

DNM candidates



AB<sub>min</sub>



DP<sub>min</sub>



GQ



AD



reassembly



DNMs  
miscalled variants  
sequencing errors

0.00 0.25 0.50 0.75 1.00  
proportion