

# 80X coverage, run 10

filter criterion

DNM candidates



DP<sub>min</sub>



AB<sub>min</sub>



GQ



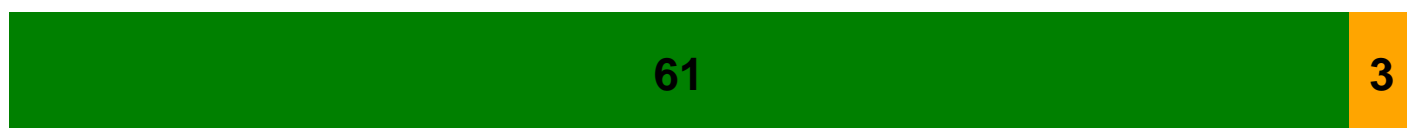
AB<sub>max</sub>



QUAL



reassembly



DNMs  
miscalled variants  
sequencing errors

0.00 0.25 0.50 0.75 1.00  
proportion