

# 60X coverage, run 3

filter criterion

DNM candidates



GQ



DP<sub>min</sub>



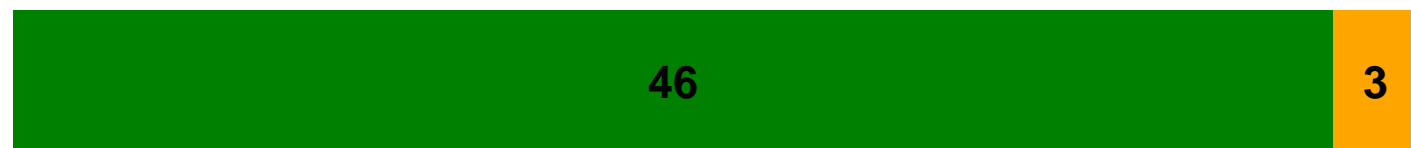
AB<sub>min</sub>



DP<sub>max</sub>



reassembly



■ DNMs  
■ miscalled variants  
■ sequencing errors

0.00

0.25

0.50

0.75

1.00

proportion