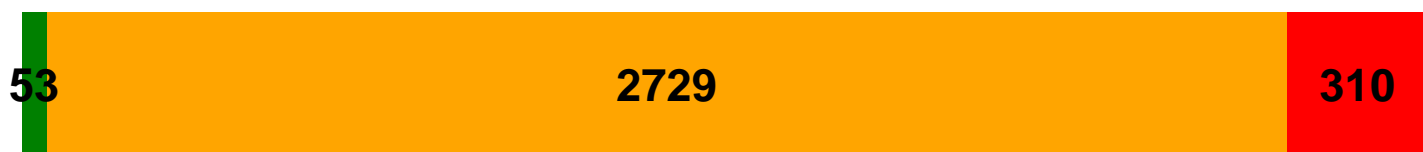


# 40X coverage, run 9

filter criterion

DNM candidates



DP<sub>min</sub>



GQ



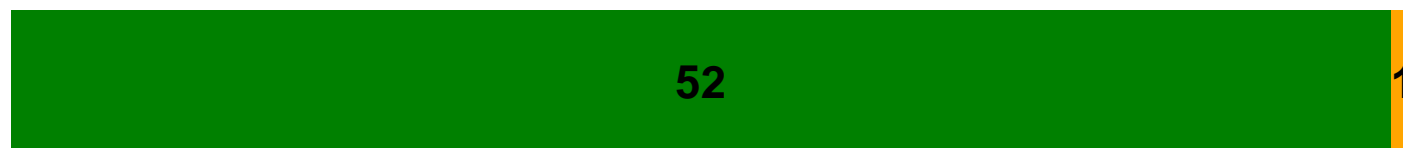
AB<sub>min</sub>



AD



reassembly



■ DNMs  
■ miscalled variants  
■ sequencing errors

0.00 0.25 0.50 0.75 1.00  
proportion