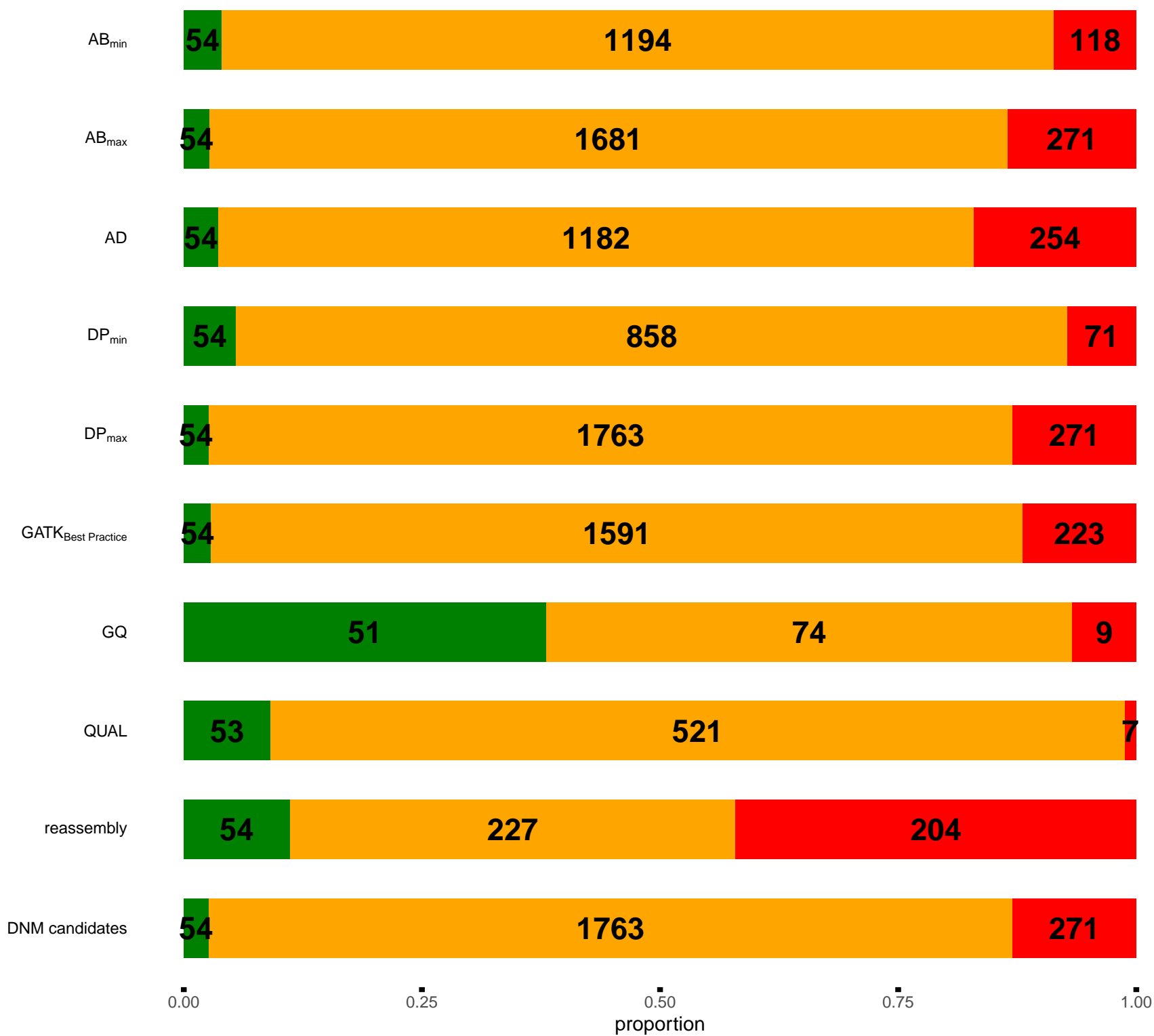


# 50X coverage, run 4

filter criterion



DNMs  
miscalled variants  
sequencing errors

0.00 0.25 0.50 0.75 1.00  
proportion