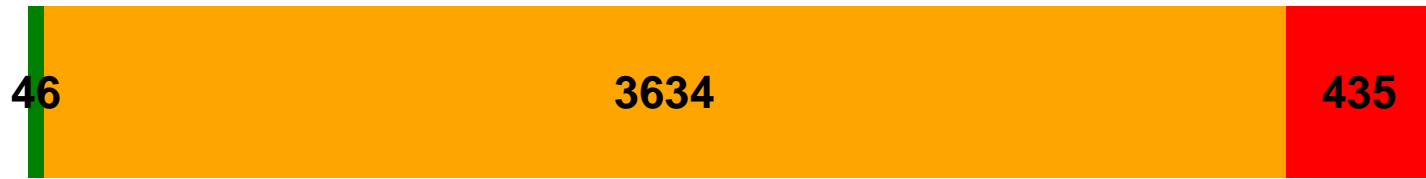


30X coverage, run 7

filter criterion

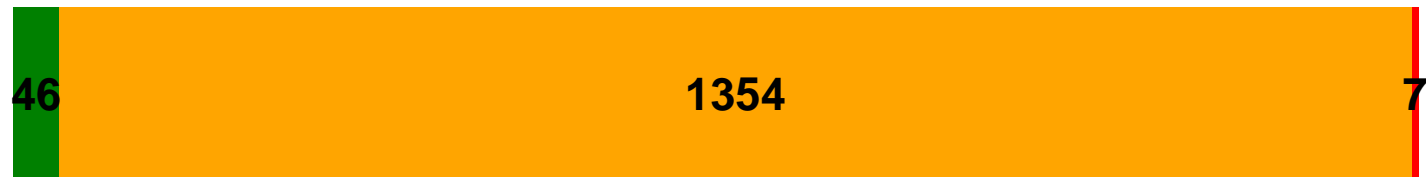
DNM candidates



DP_{min}



AB_{min}



DNMs
miscalled variants
sequencing errors

GQ



reassembly



0.00

0.25

0.50

0.75

1.00

proportion