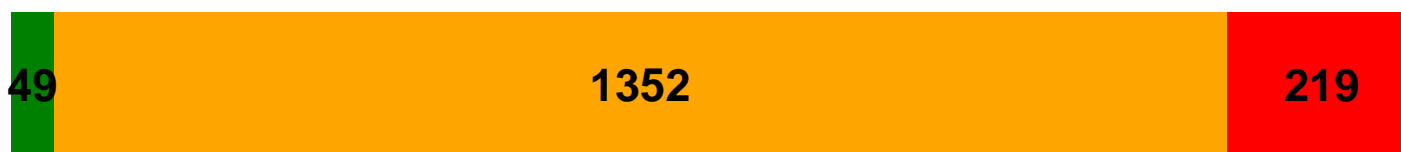


100X coverage, run 6

DNM candidates



GQ



QUAL



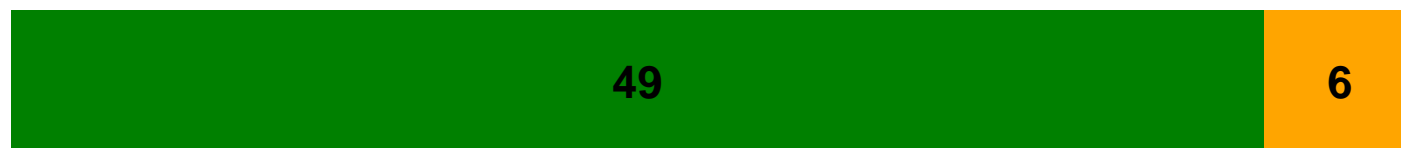
AD



AB_{min}



reassembly



DNMs
miscalled variants
sequencing errors

0.00 0.25 0.50 0.75 1.00
proportion