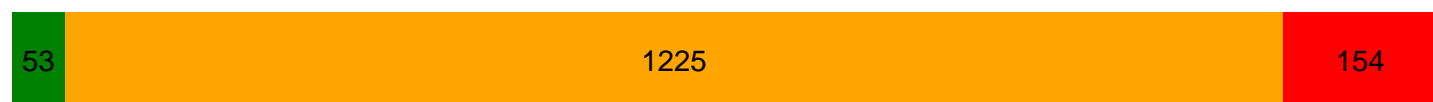


90X coverage, run 9

filter criterion

GATK_{Best Practice}



GQ



QUAL



raw MVs



AD



DP_{min}



DP_{max}



AB_{max}



AB_{min}



DNMs
miscalled variants
sequencing errors

0.00 0.25 0.50 0.75 1.00

proportion