

100X coverage, run 3

filter criterion

GATK_{Best Practice}

raw MVs

QUAL

GQ

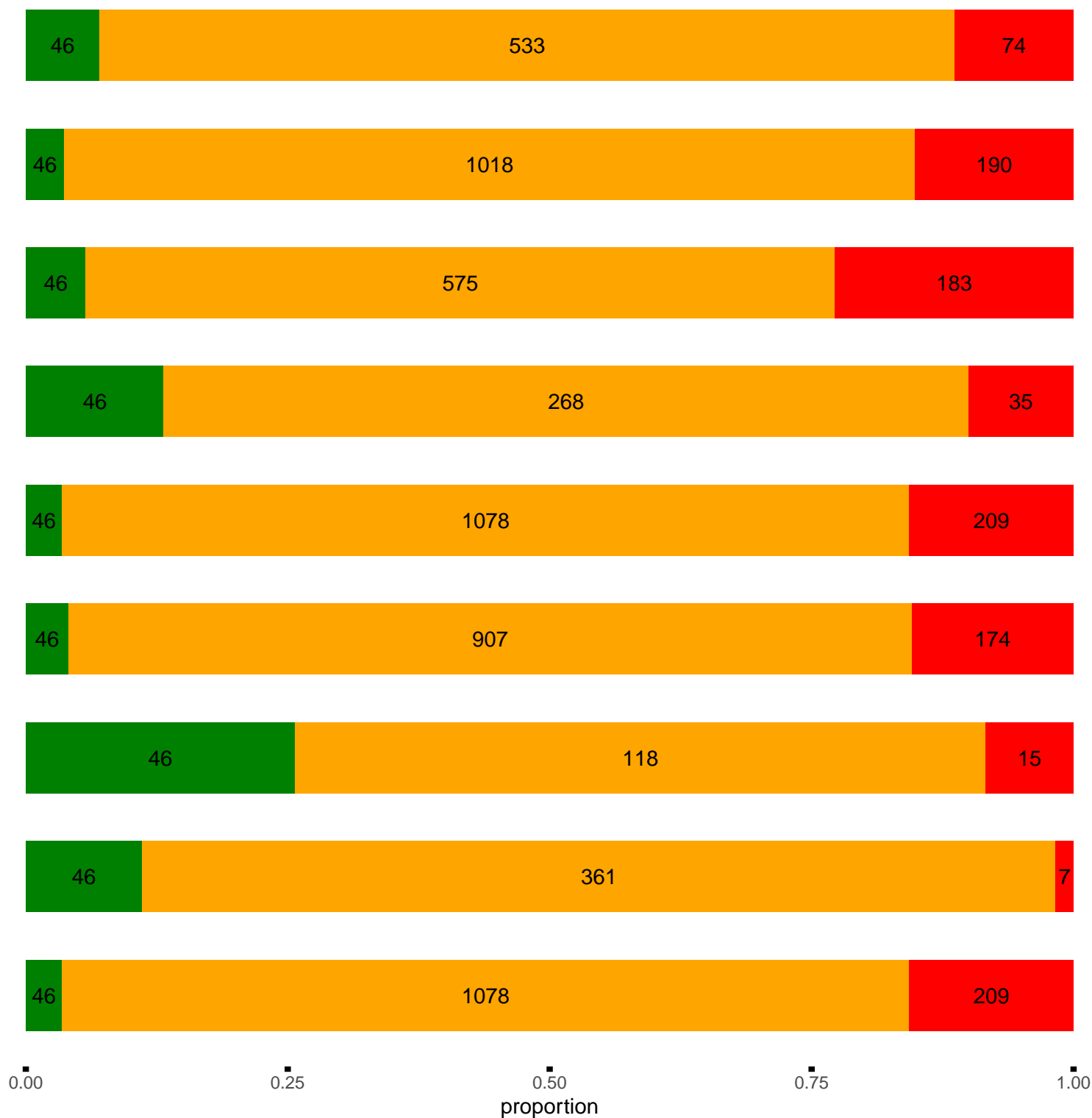
DP_{max}

DP_{min}

AD

AB_{max}

AB_{min}



DNMs
miscalled variants
sequencing errors

0.00 0.25 0.50 0.75 1.00
proportion