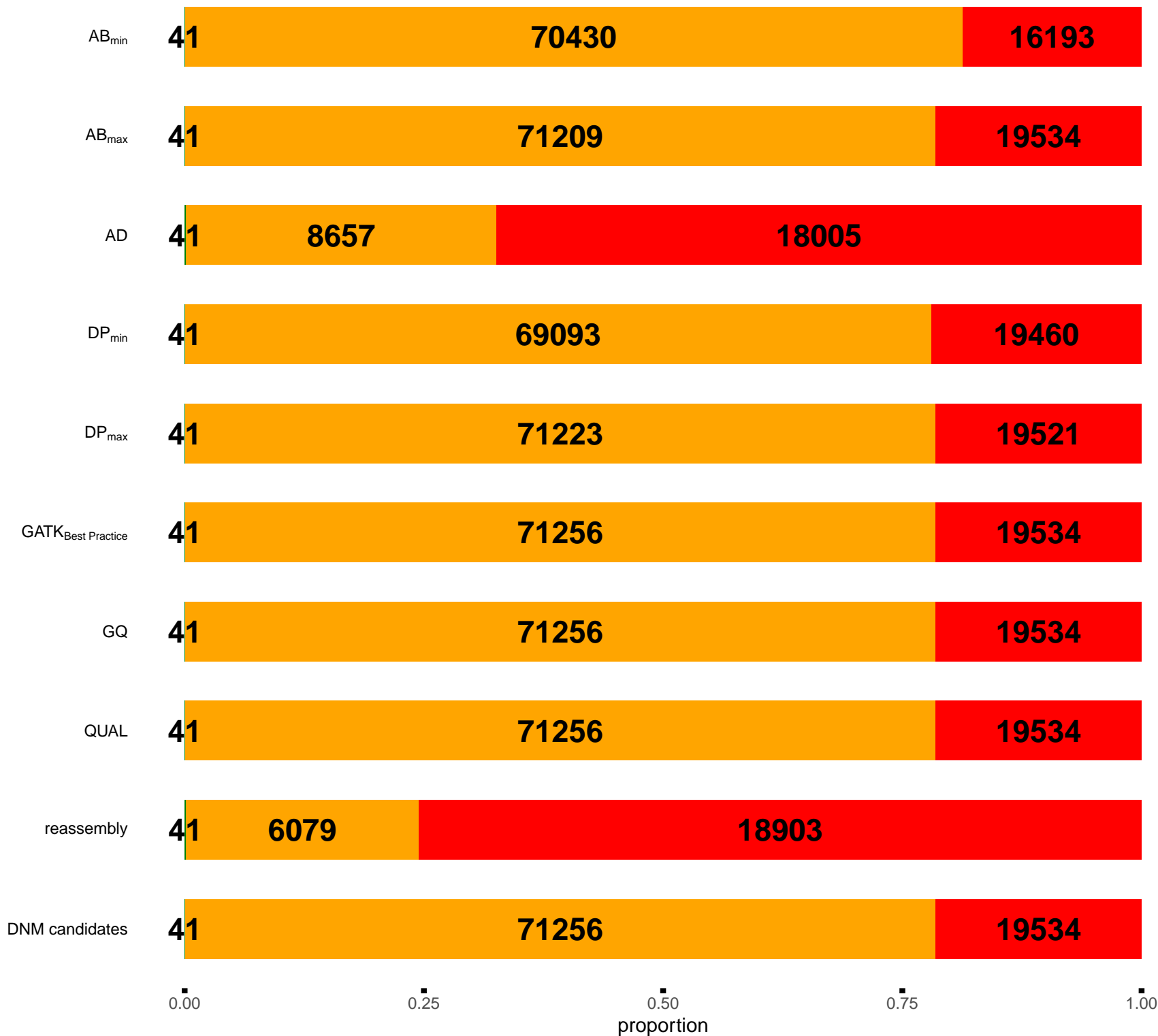


10X coverage, run 3

filter criterion



DNMs
miscalled variants
sequencing errors

0.00 0.25 0.50 0.75 1.00
proportion