

90X coverage, run 7

filter criterion

DNM candidates



AB_{min}



DP_{min}



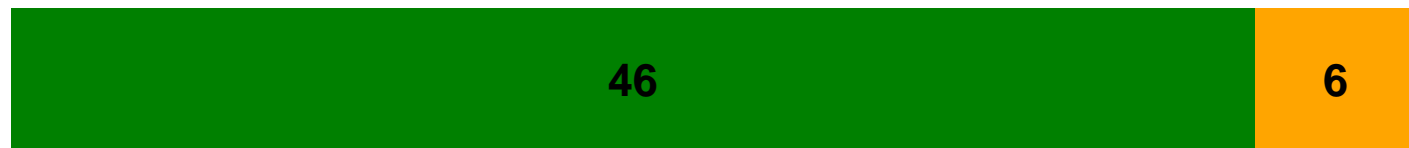
GQ



AD



reassembly



■ DNMs
■ miscalled variants
■ sequencing errors

0.00

0.25

0.50

0.75

1.00

proportion