

20X coverage, run 10

filter criterion

DNM candidates

62

8009

1624

GQ

61

6186

69

DP_{min}

61

4868

59

AD

61

2838

53

AB_{min}

61

2837

50

reassembly

59

119

27

DNMs
miscalled variants
sequencing errors

0.00

0.25

0.50

0.75

1.00

proportion