

# 100X coverage, run 6

filter criterion

GATK<sub>Best Practice</sub>

raw MVs

QUAL

GQ

GATK<sub>Best Practice</sub>

DP<sub>max</sub>

DP<sub>min</sub>

AD

AB<sub>max</sub>

AB<sub>min</sub>



DNMs  
miscalled variants  
sequencing errors

0.00 0.25 0.50 0.75 1.00  
proportion