

# General Statistics

42 samples

bowtie2  
% Aligned



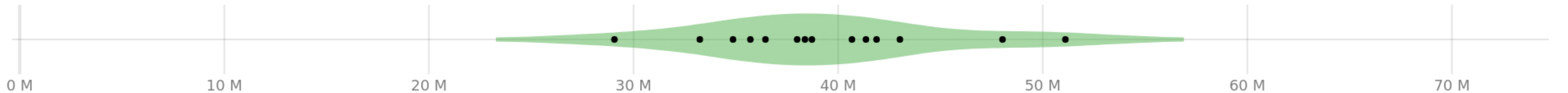
Samtools Stats: stats  
Error rate



Samtools Stats: stats  
Non-primary



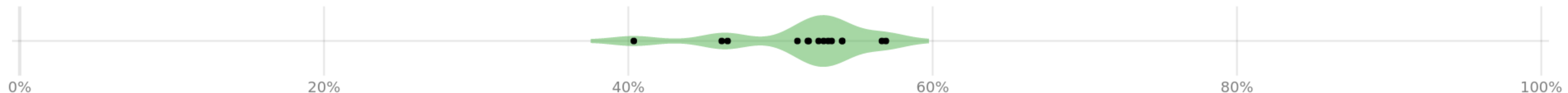
Samtools Stats: stats  
Reads mapped



Samtools Stats: stats  
% Mapped



Samtools Stats: stats  
% Proper pairs



Samtools Stats: stats  
Total seqs

