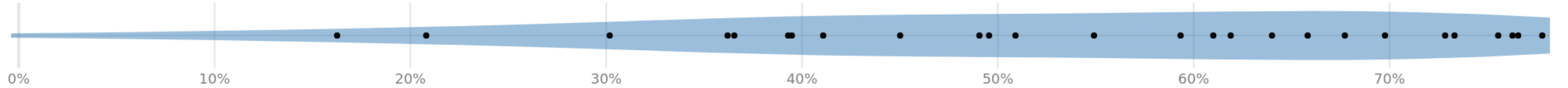


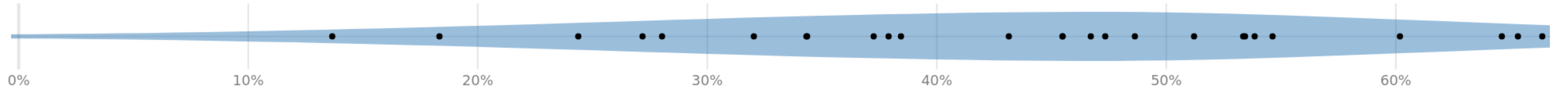
General Statistics

98 samples

STAR
Aligned



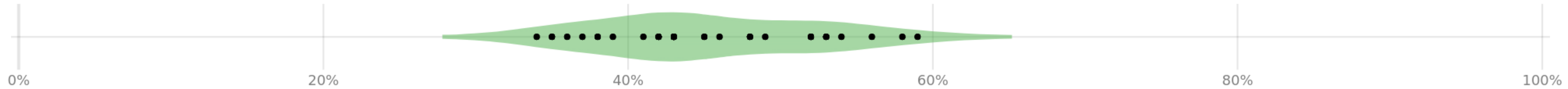
STAR
Uniq aligned



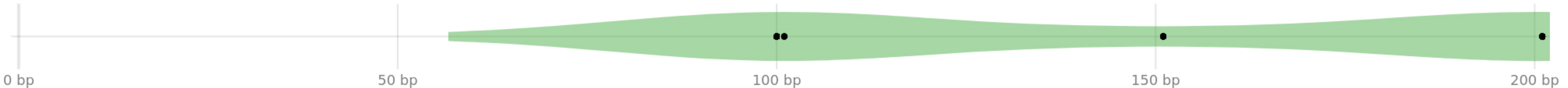
FastQC
Dups



FastQC
GC



FastQC
Median len



FastQC
Seqs

