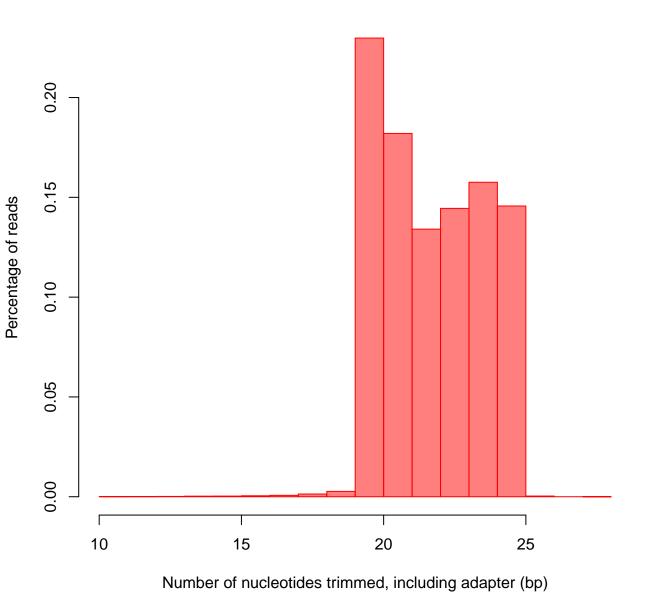
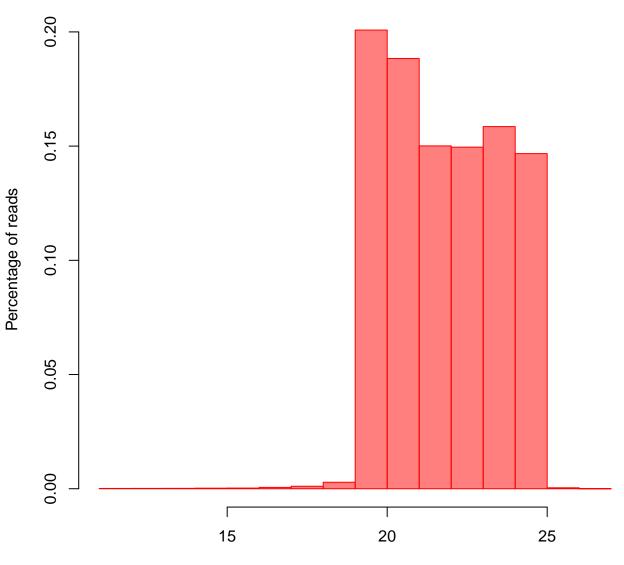
Trimming statistics, week0.1_forward

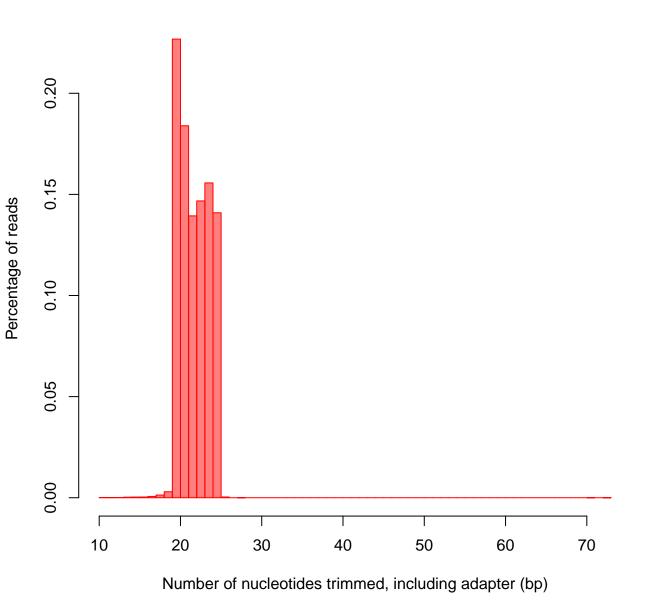


Trimming statistics, week0.2_forward

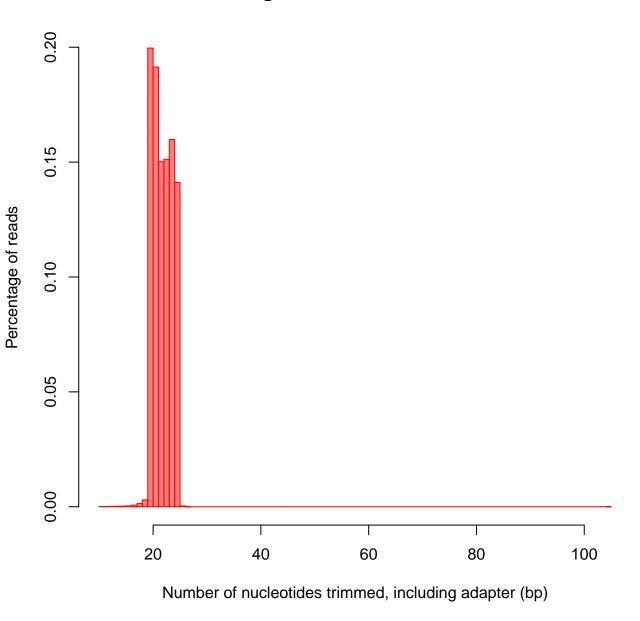


Number of nucleotides trimmed, including adapter (bp)

Trimming statistics, week4.1_forward



Trimming statistics, week4.2_forward



Trimmed reads

