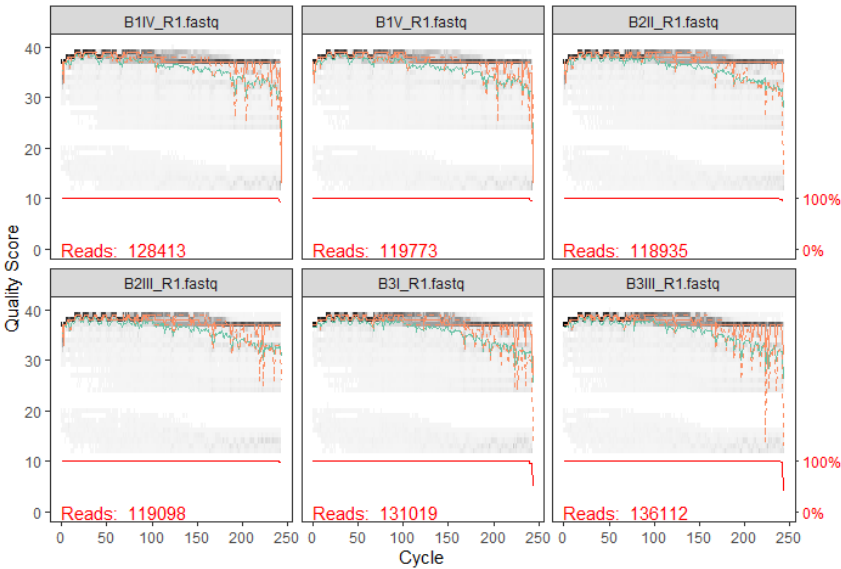
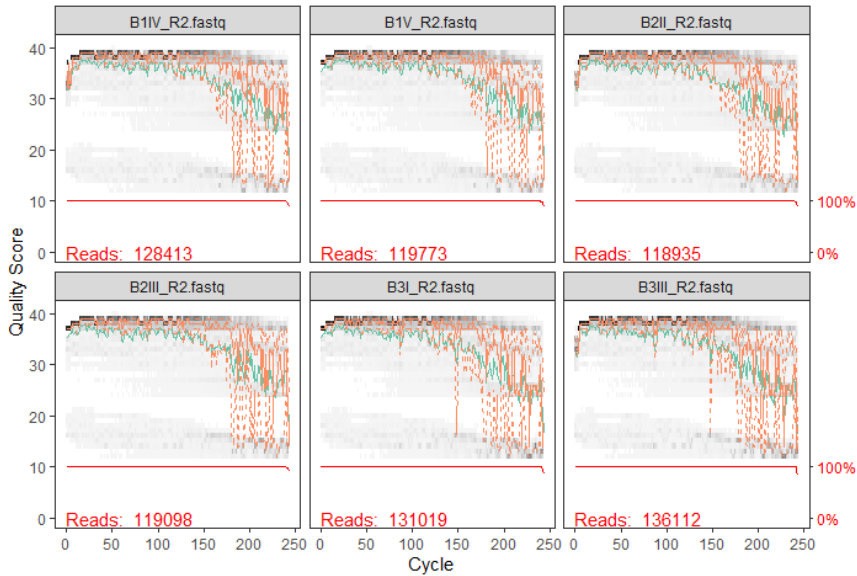


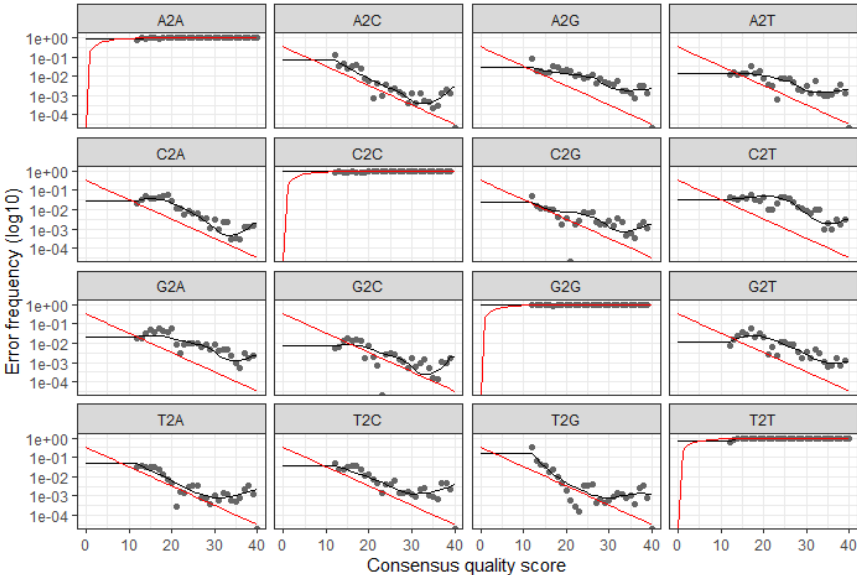
Rplot 1



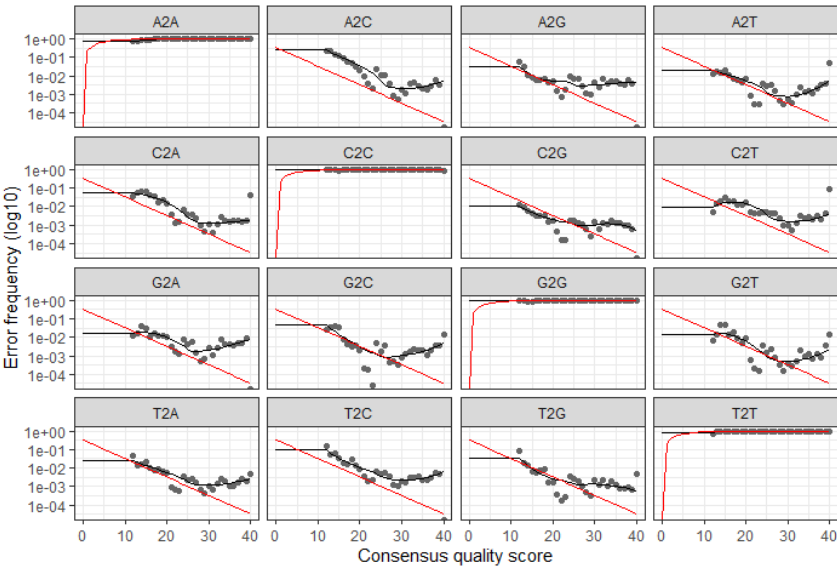
Rplot 2



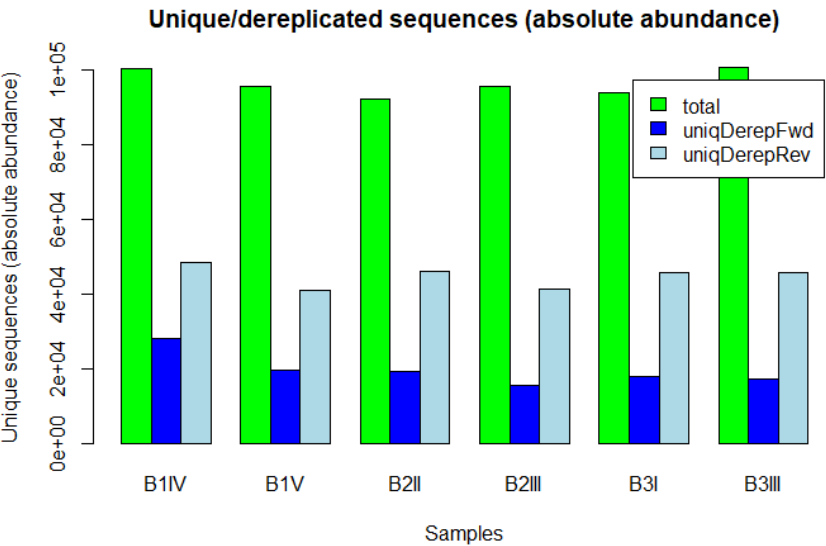
Rplot 3



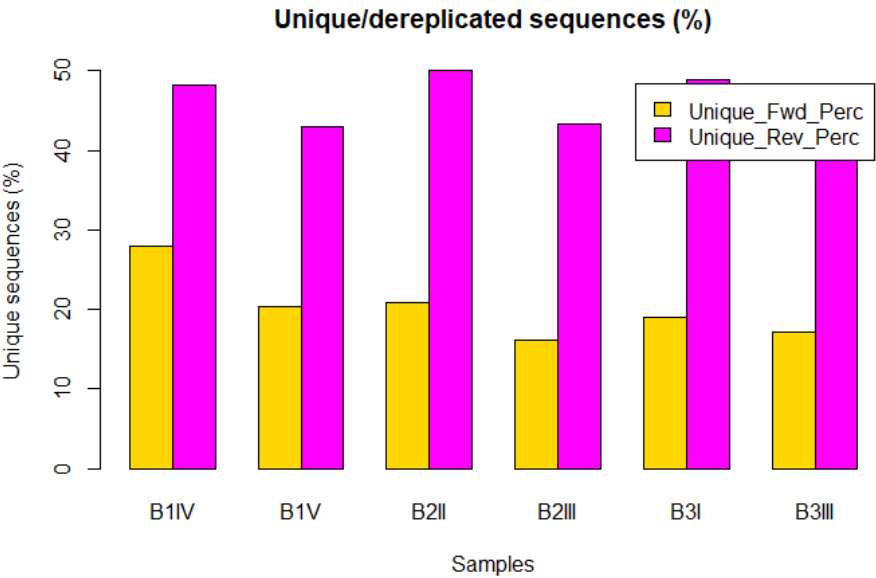
Rplot 4



Rplot 5

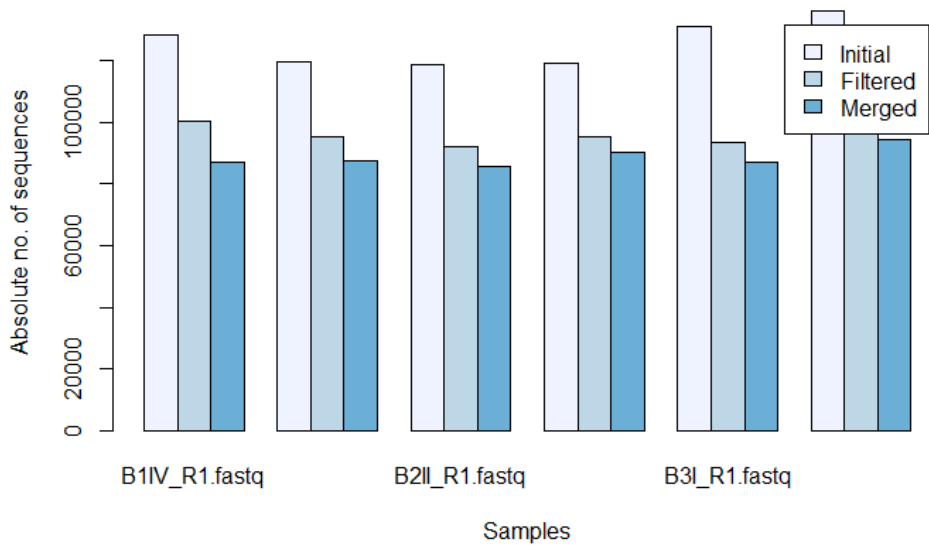


Rplot 6



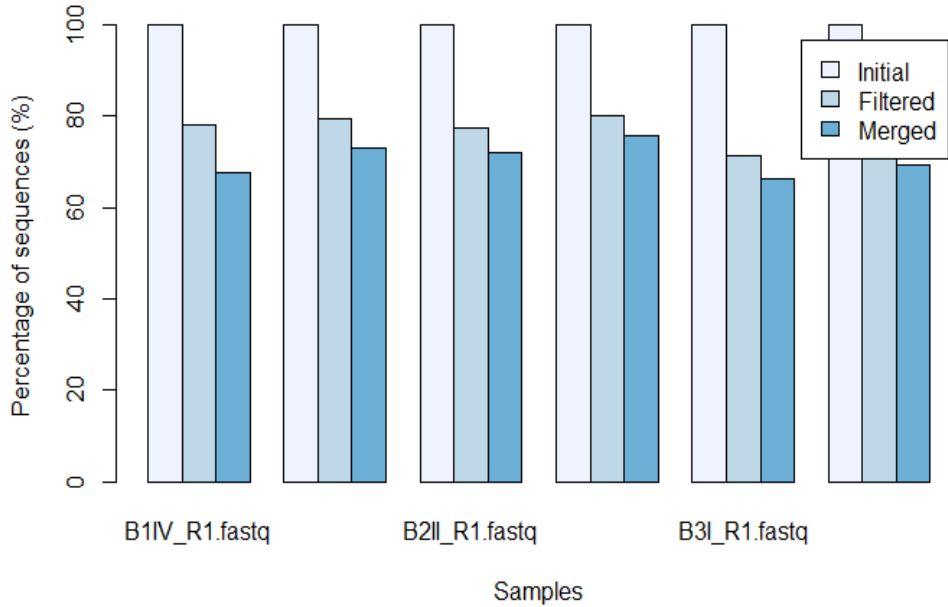
Rplot 7

Comparison of no. of sequences kept across pipeline steps: initial, filtered and merged

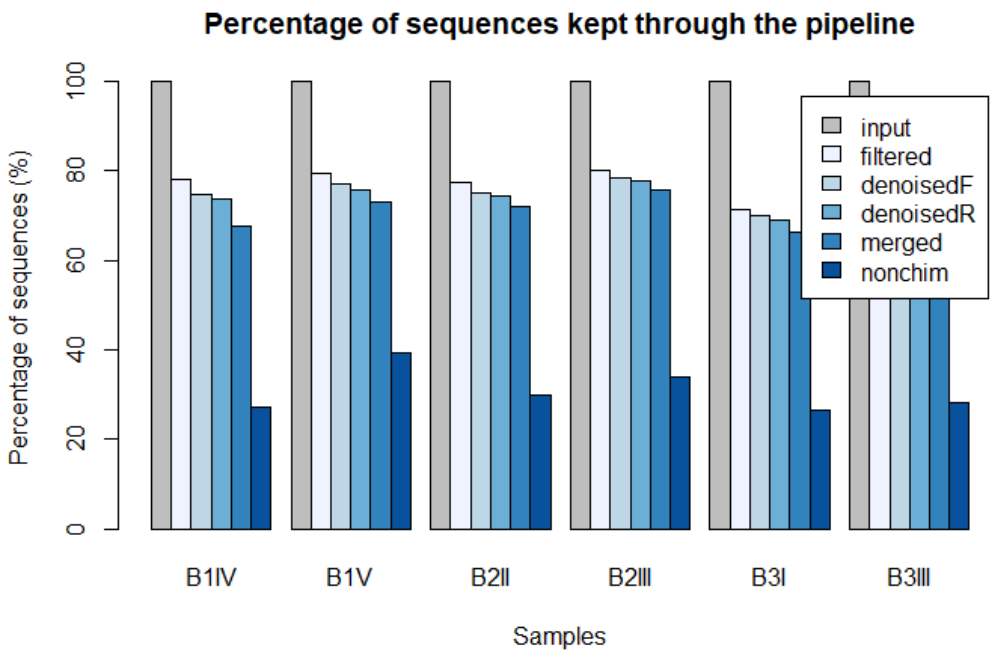


Rplot 8

Comparison of proportion of sequences kept across pipeline steps: initial, filtered and merged



Rplot 9



Rplot 10

