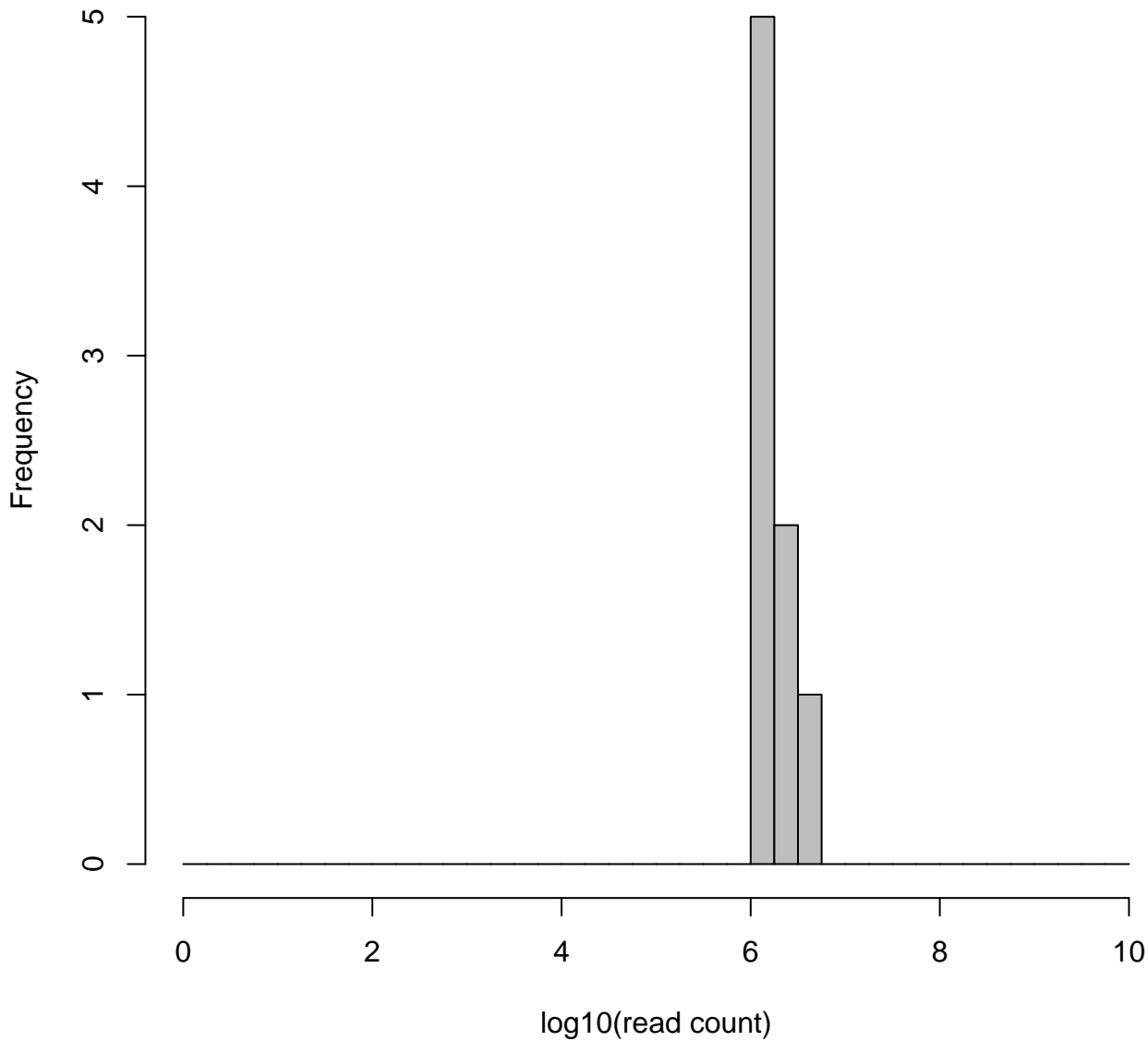
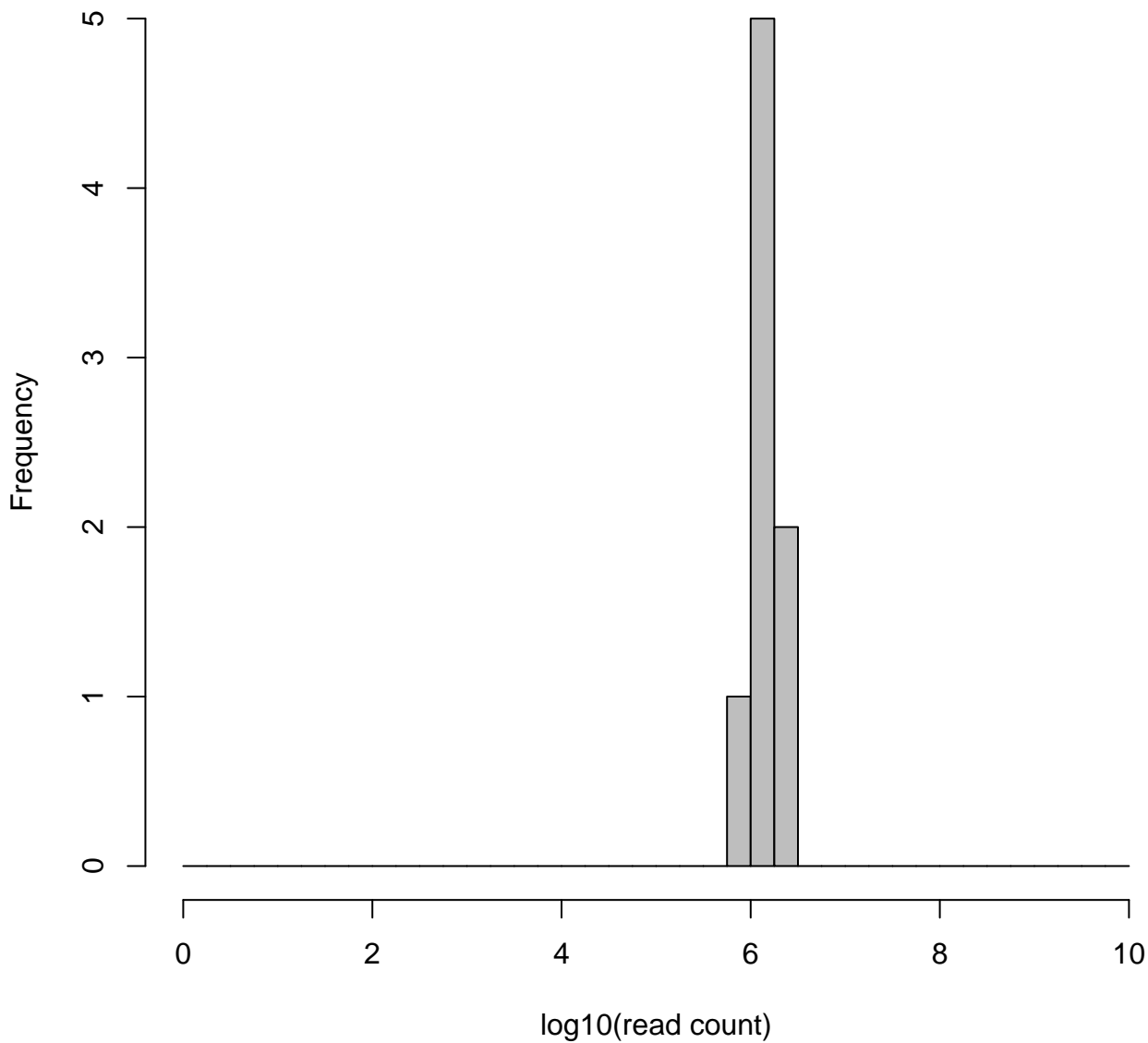


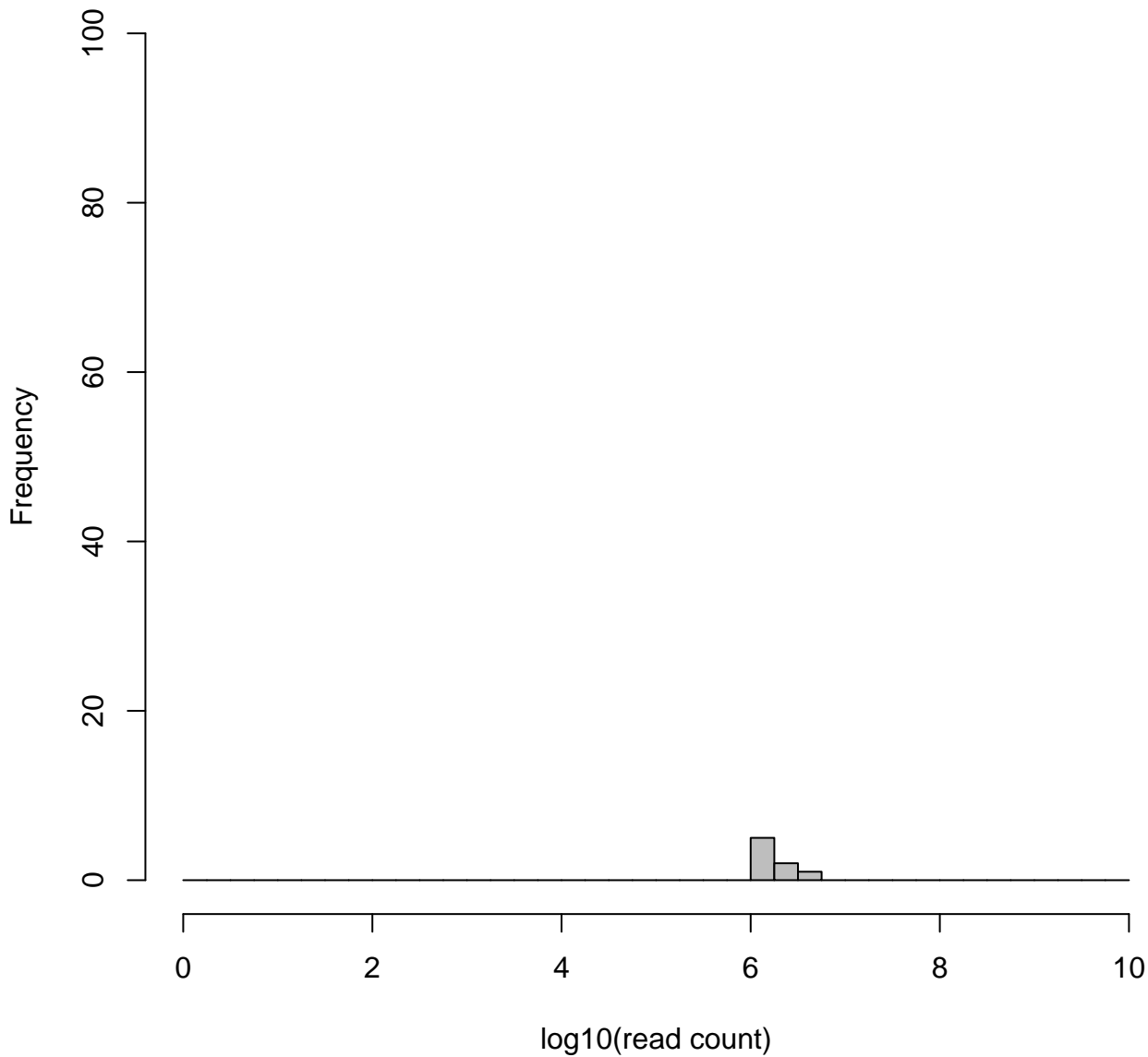
# Osa readcount RNA-Seq



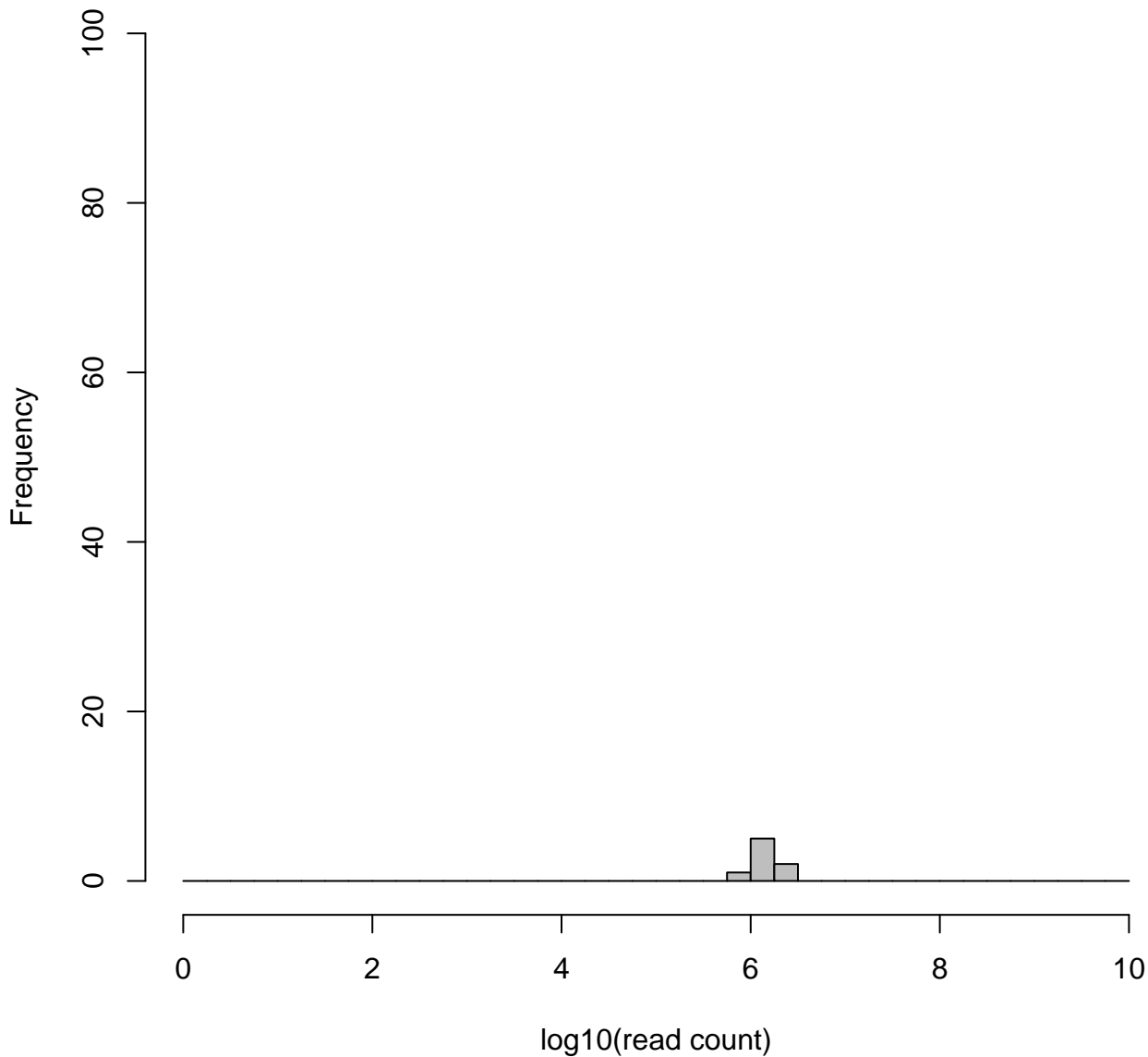
# Osa readcount RNA-Seq: data read

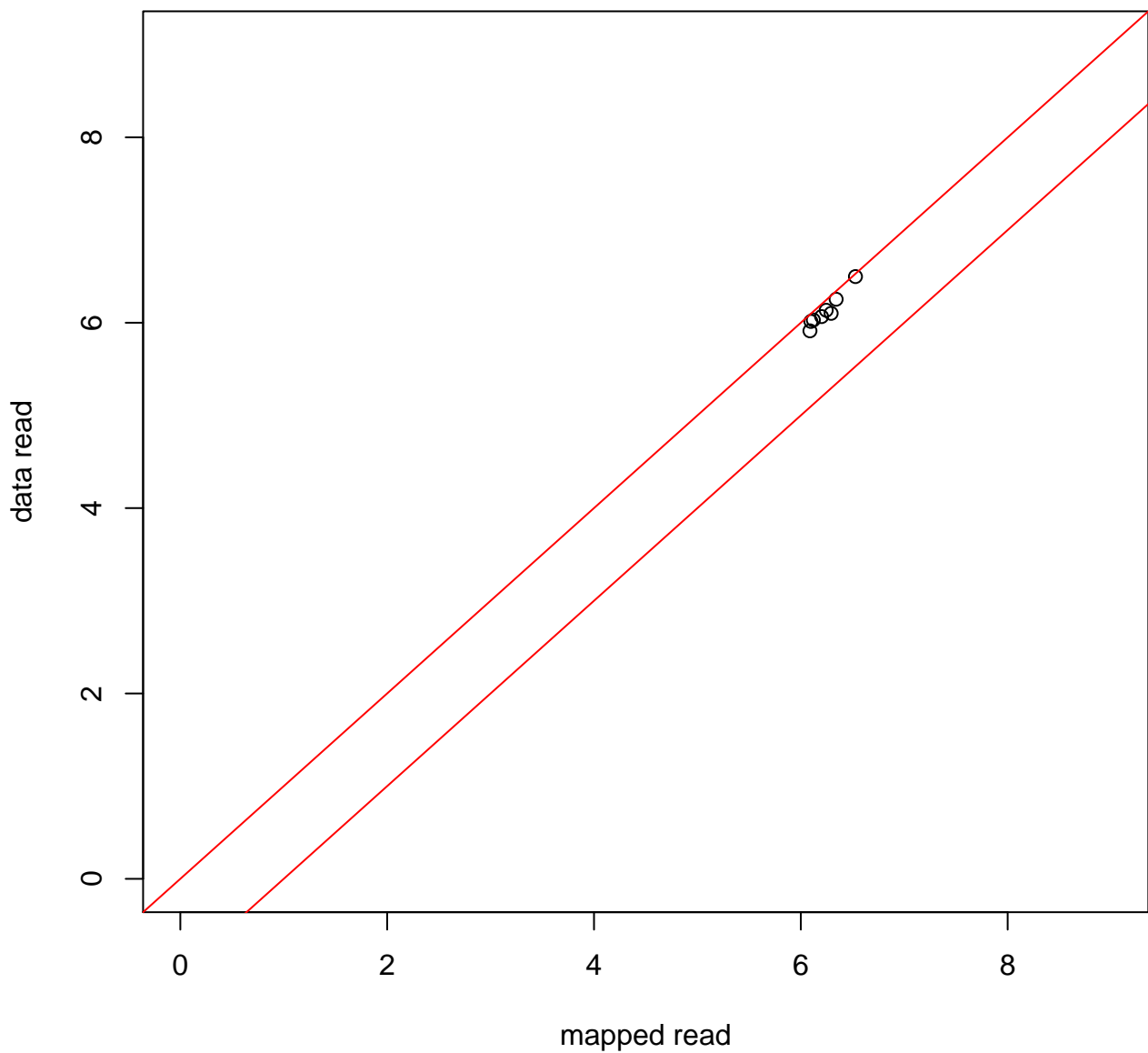


# Osa readcount RNA-Seq

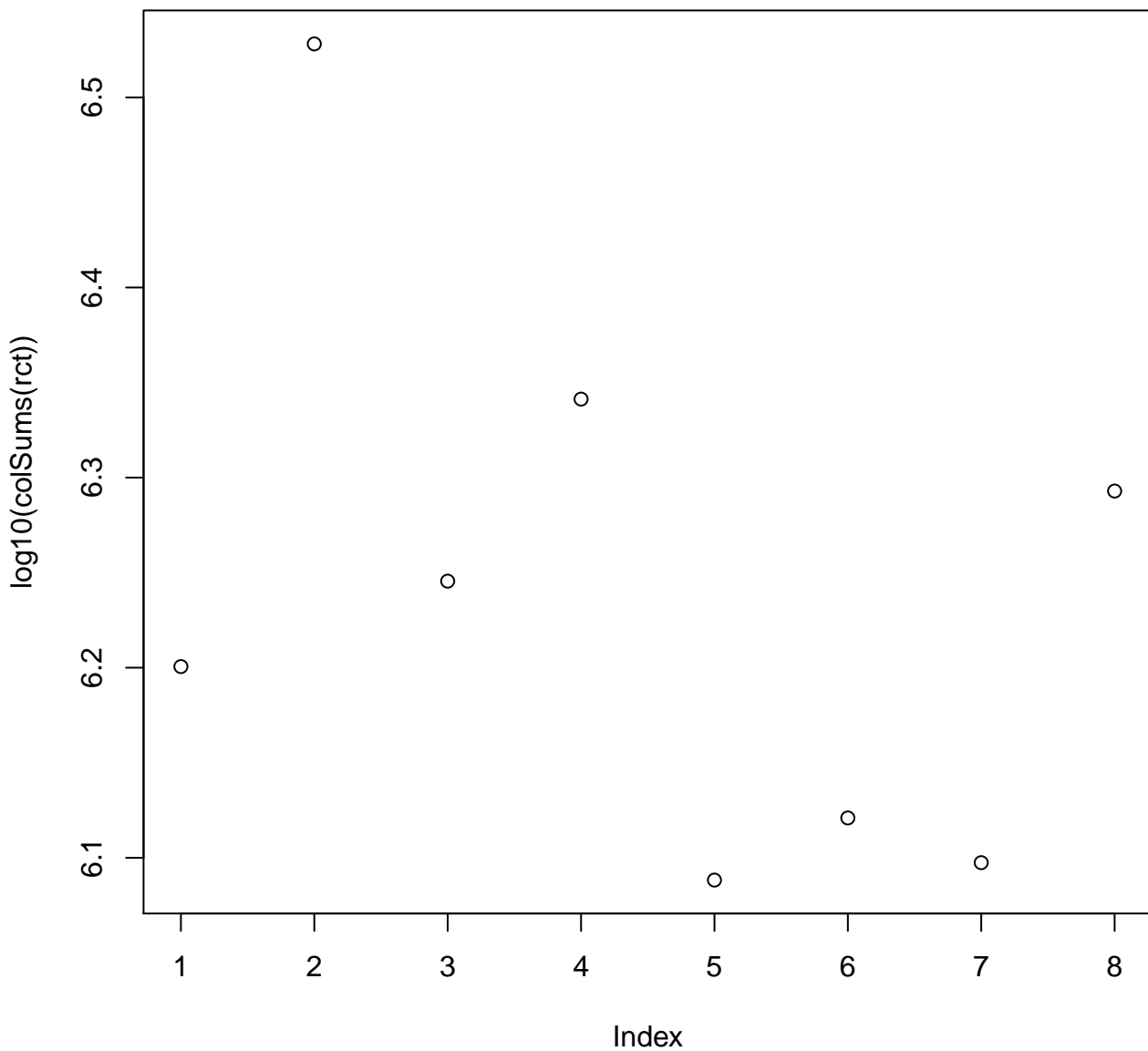


# Osa readcount RNA-Seq: data read

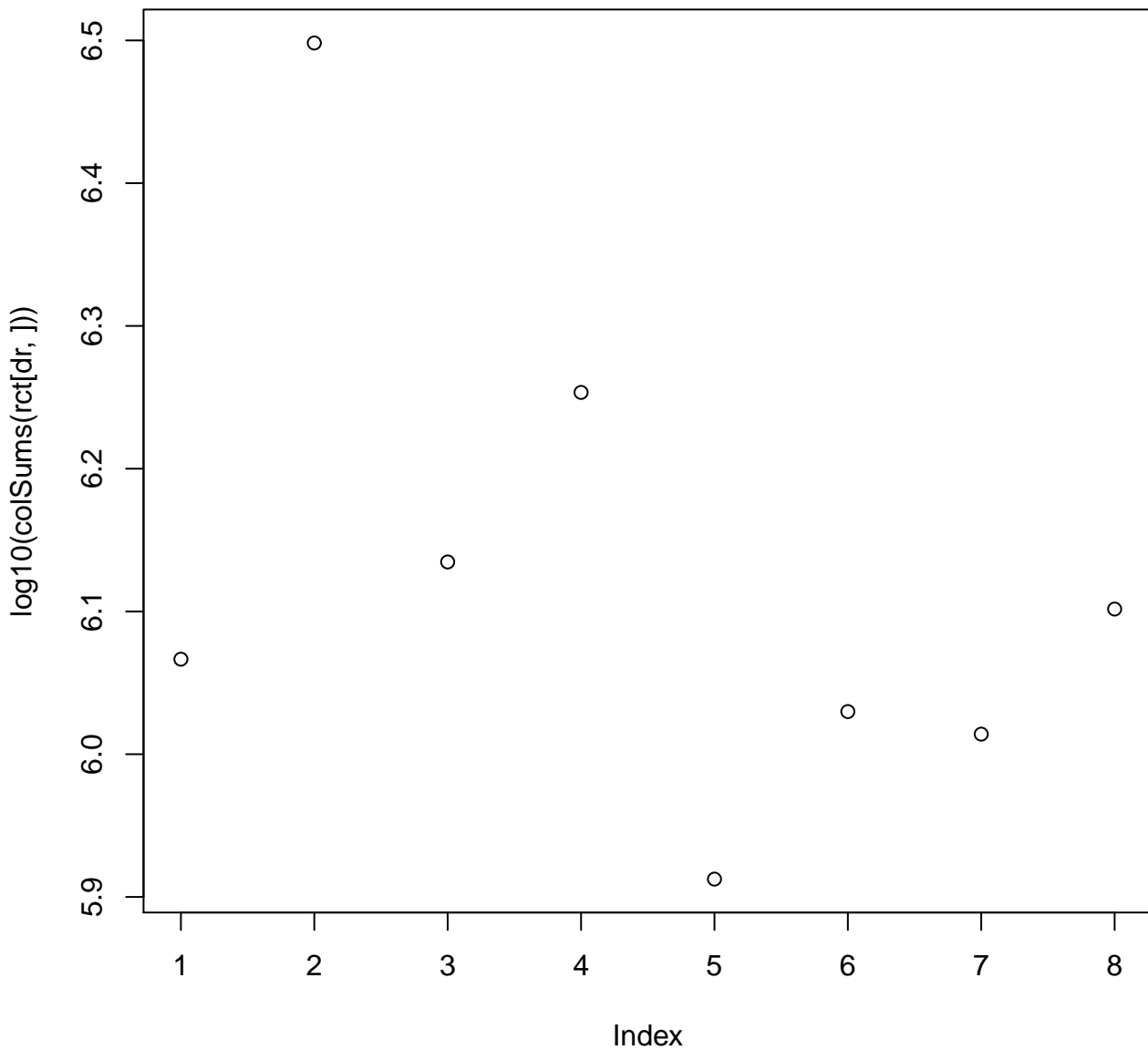




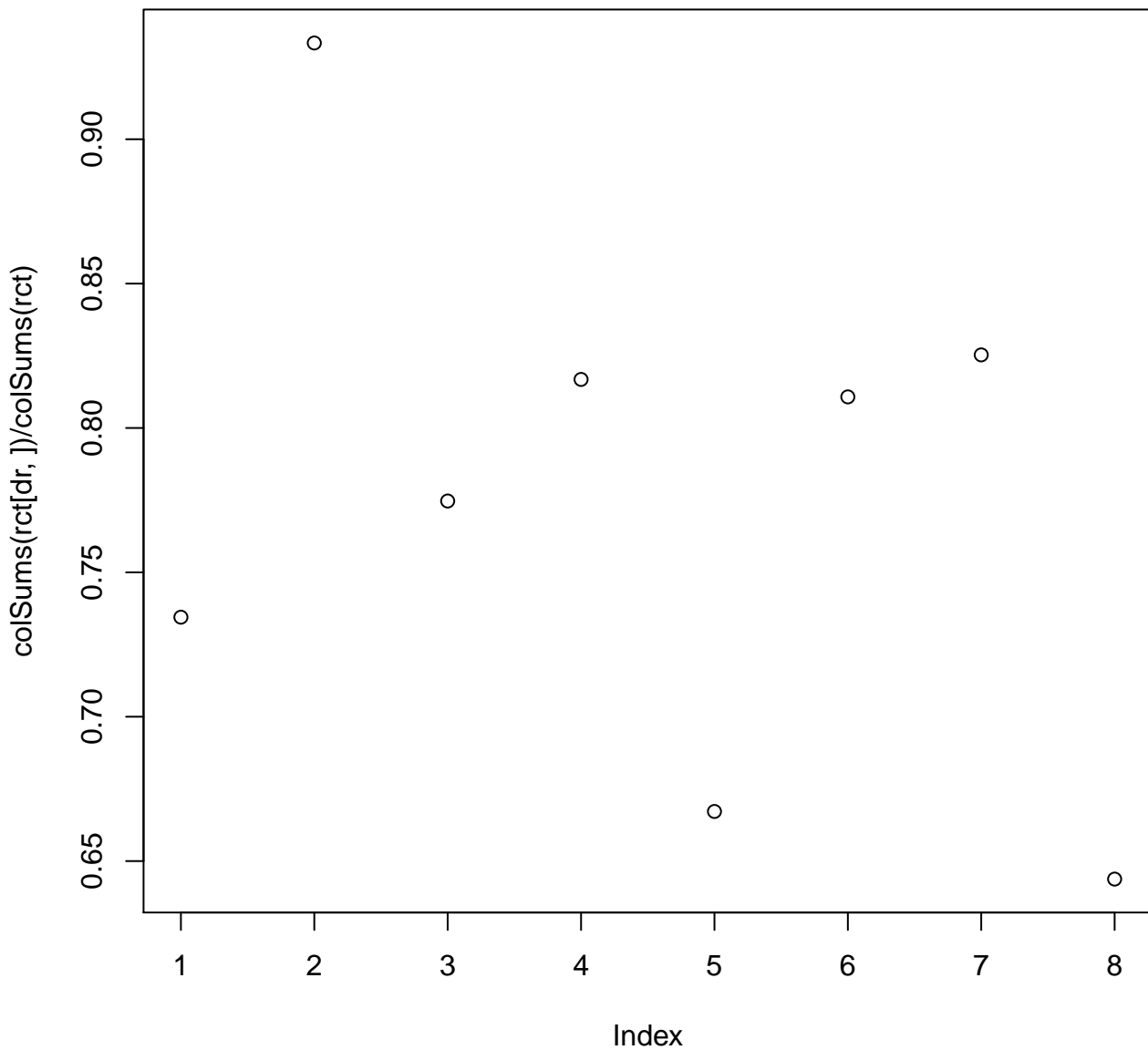
# mapped read



# data read

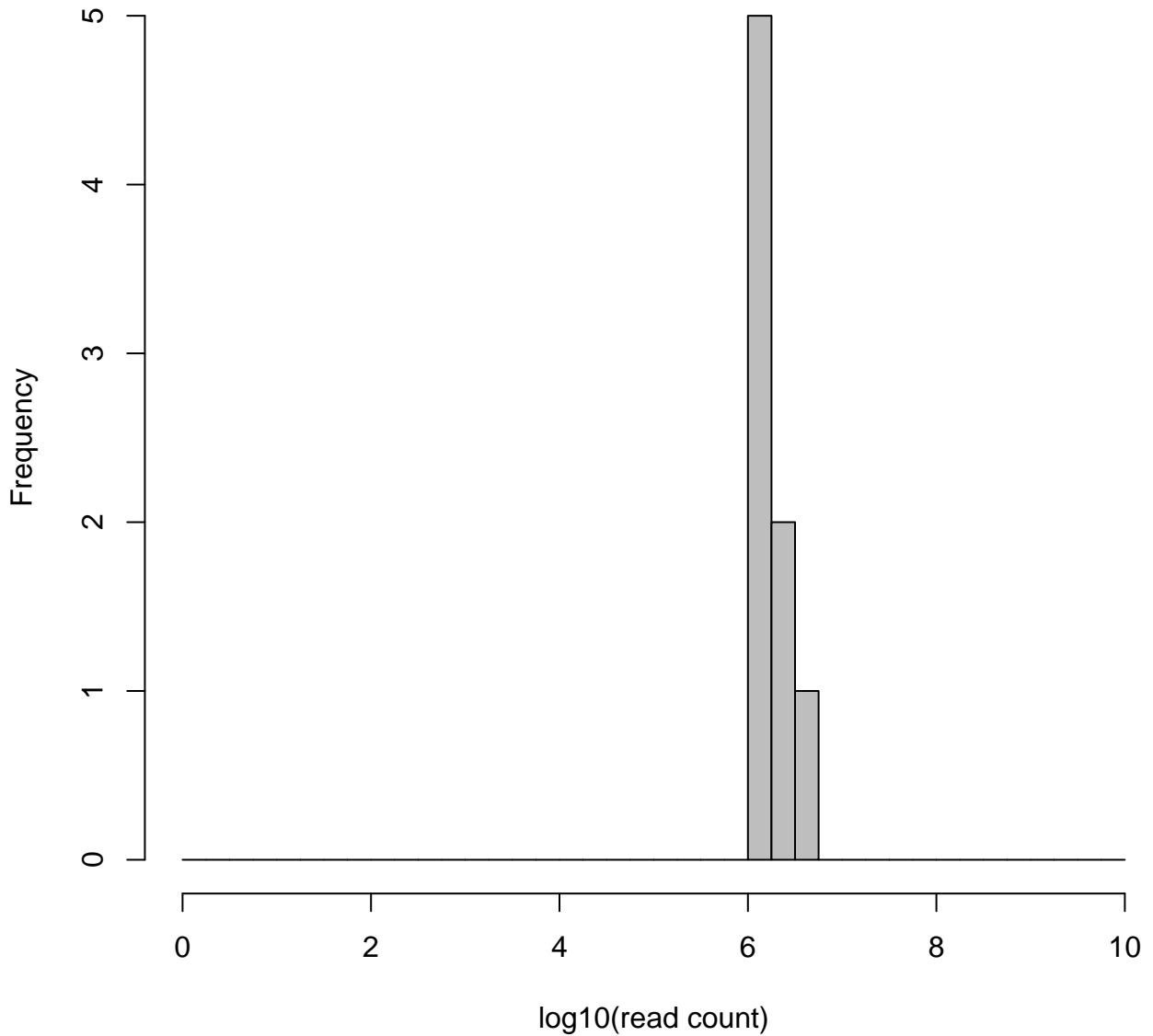


# data read / mapped read

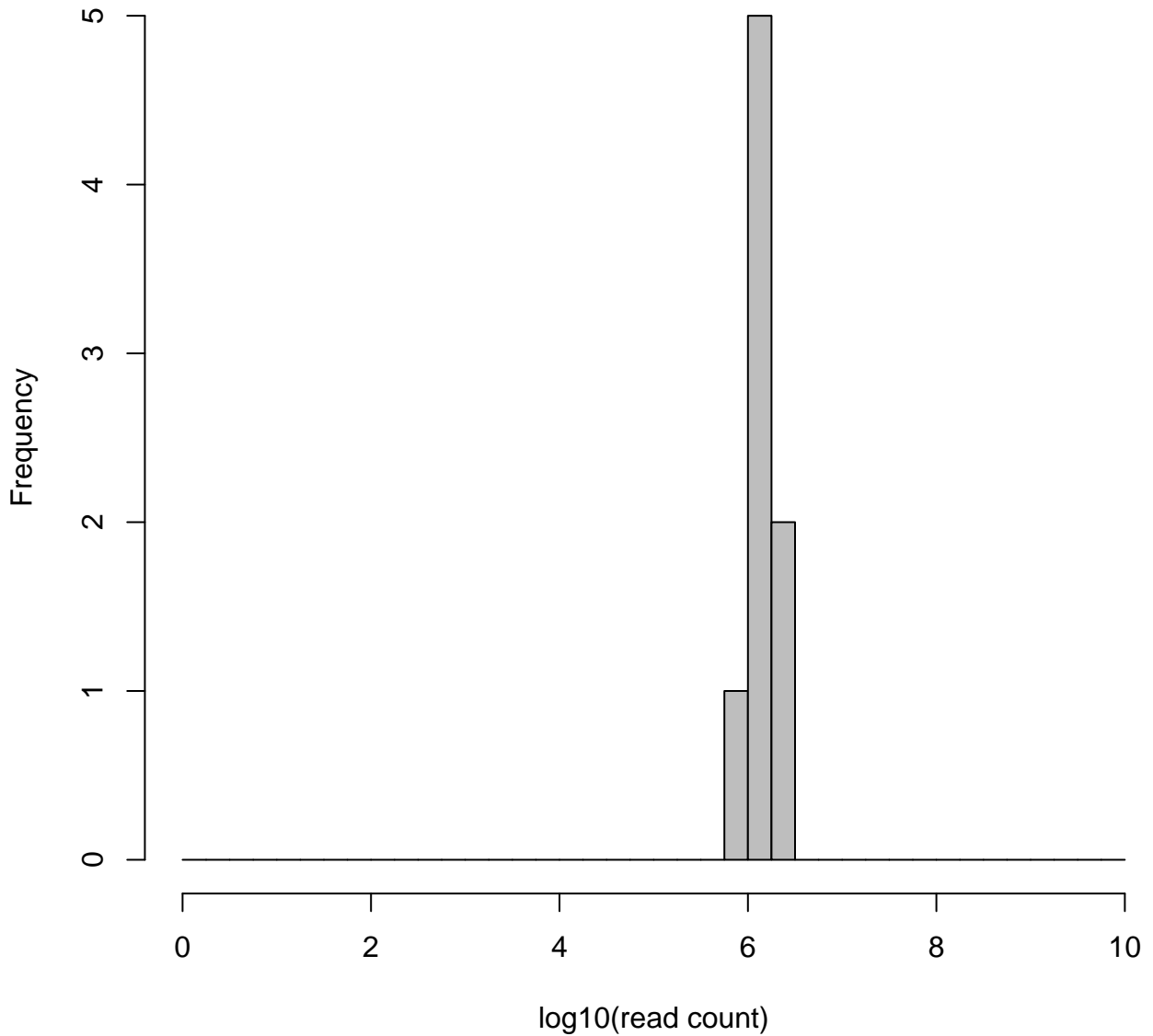




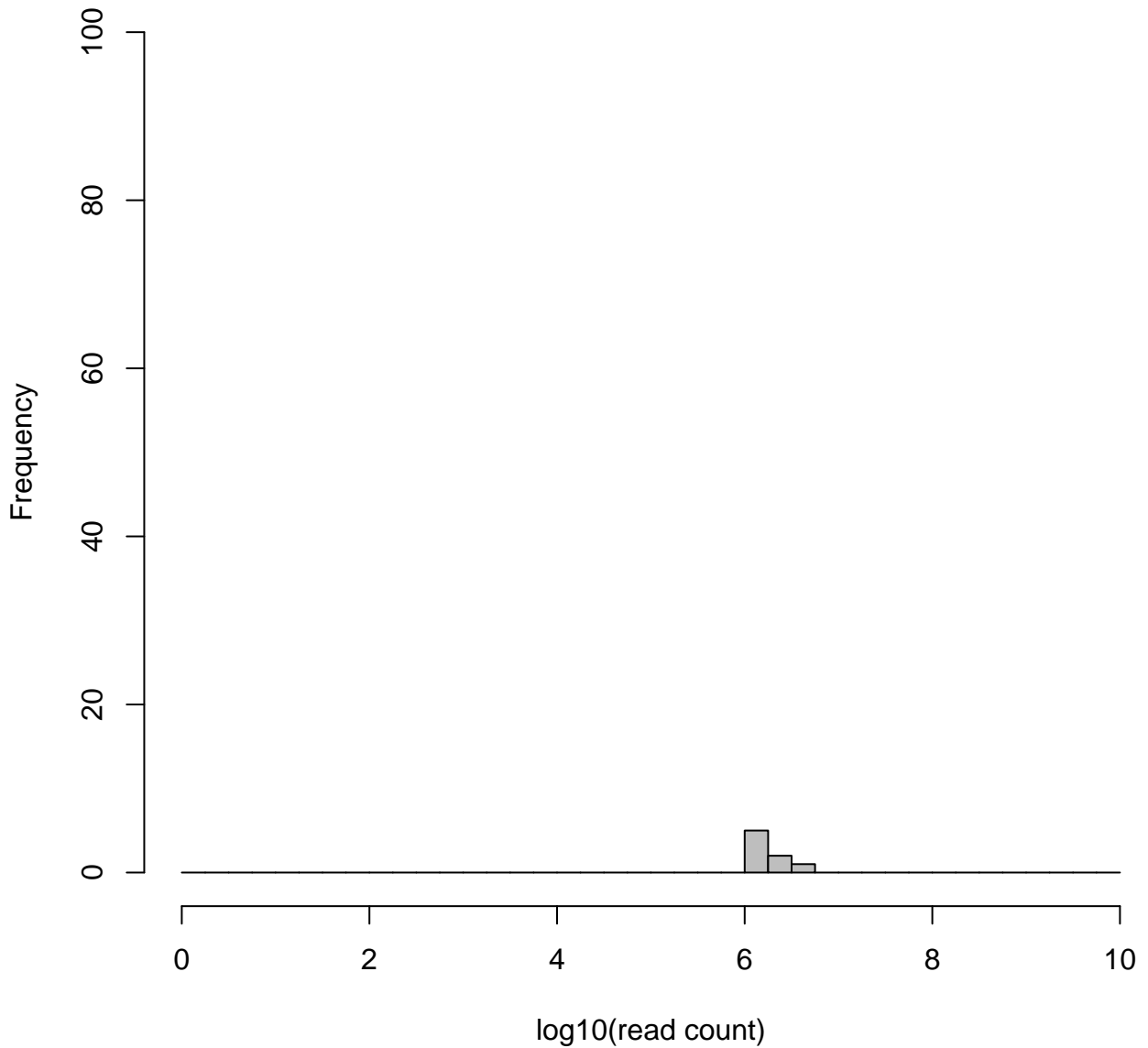
# ***Arabidopsis halleri* (2017) RNA-Seq**



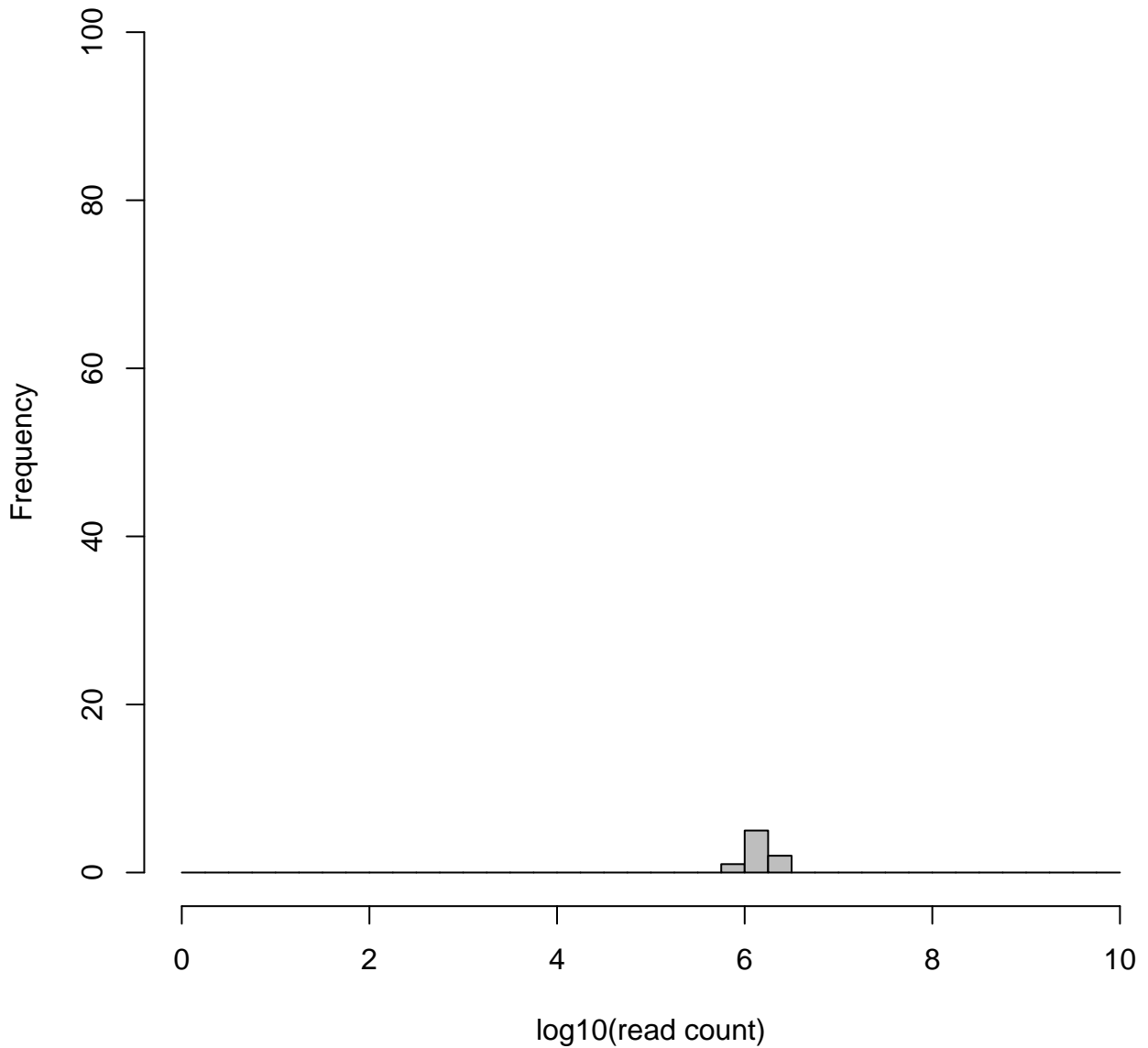
# ***Arabidopsis halleri* (2017) RNA-Seq: data read**

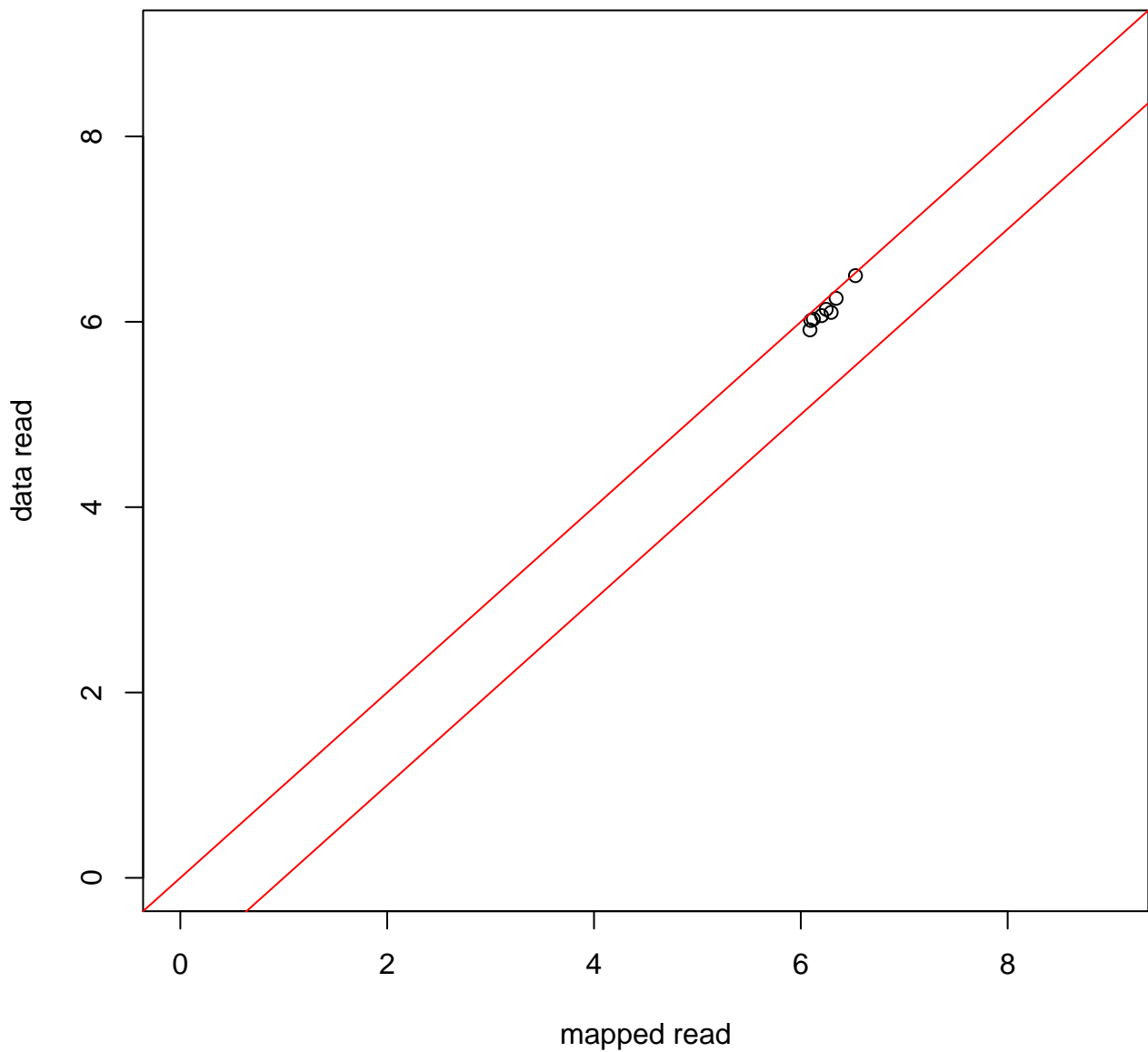


# ***Arabidopsis halleri* (2017) RNA-Seq**

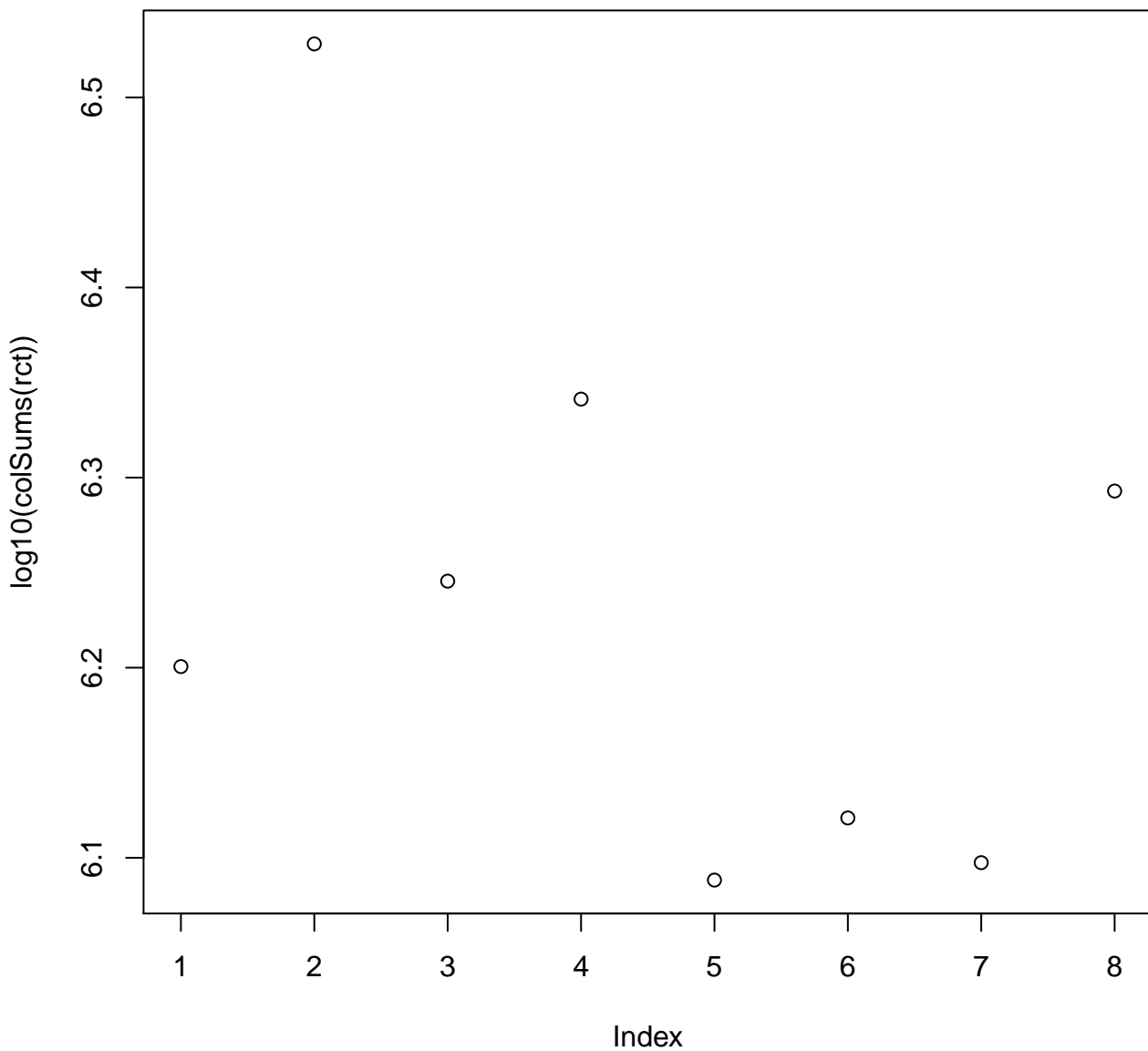


# ***Arabidopsis halleri* (2017) RNA-Seq: data read**

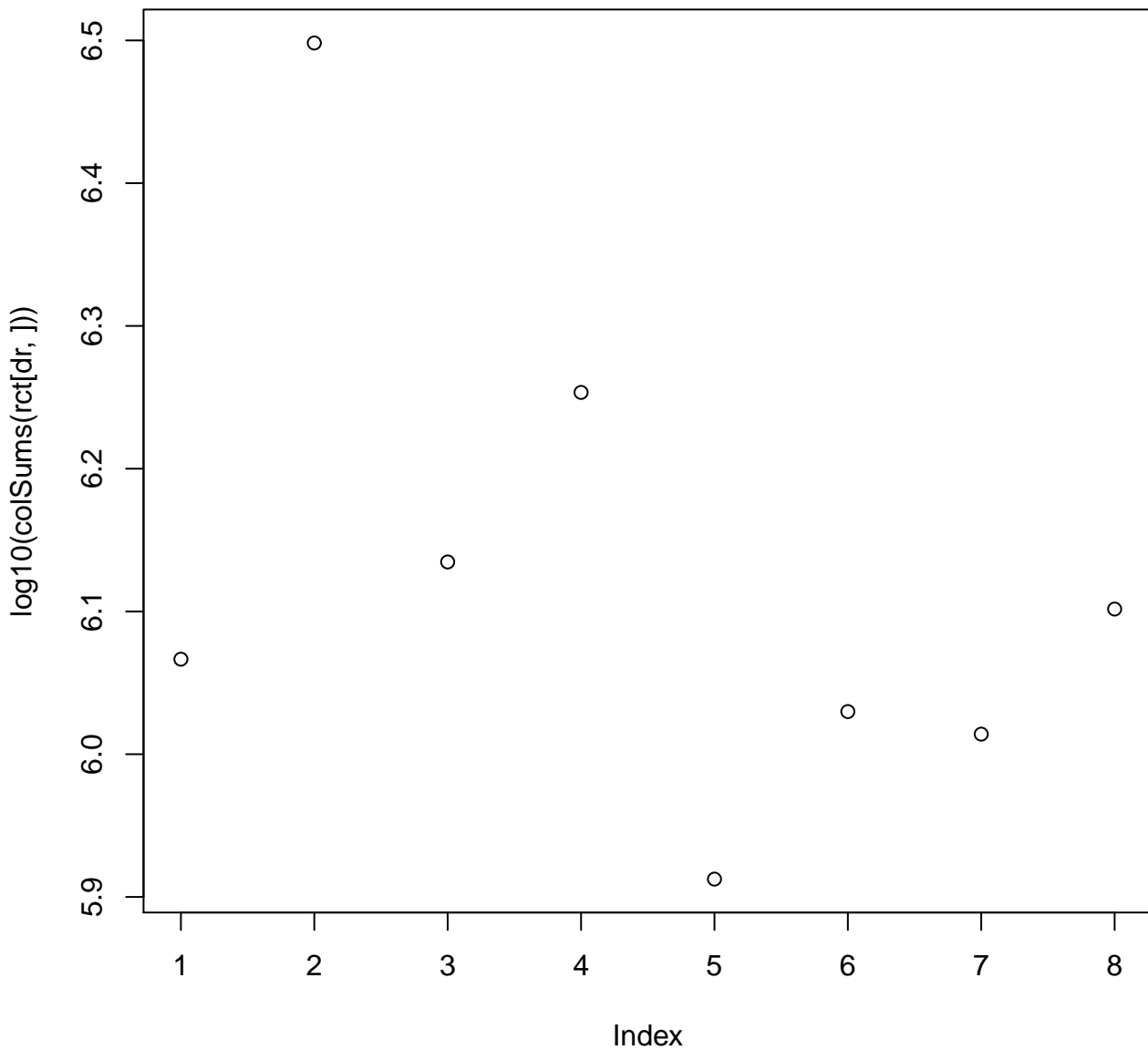




# mapped read



# data read



# data read / mapped read

