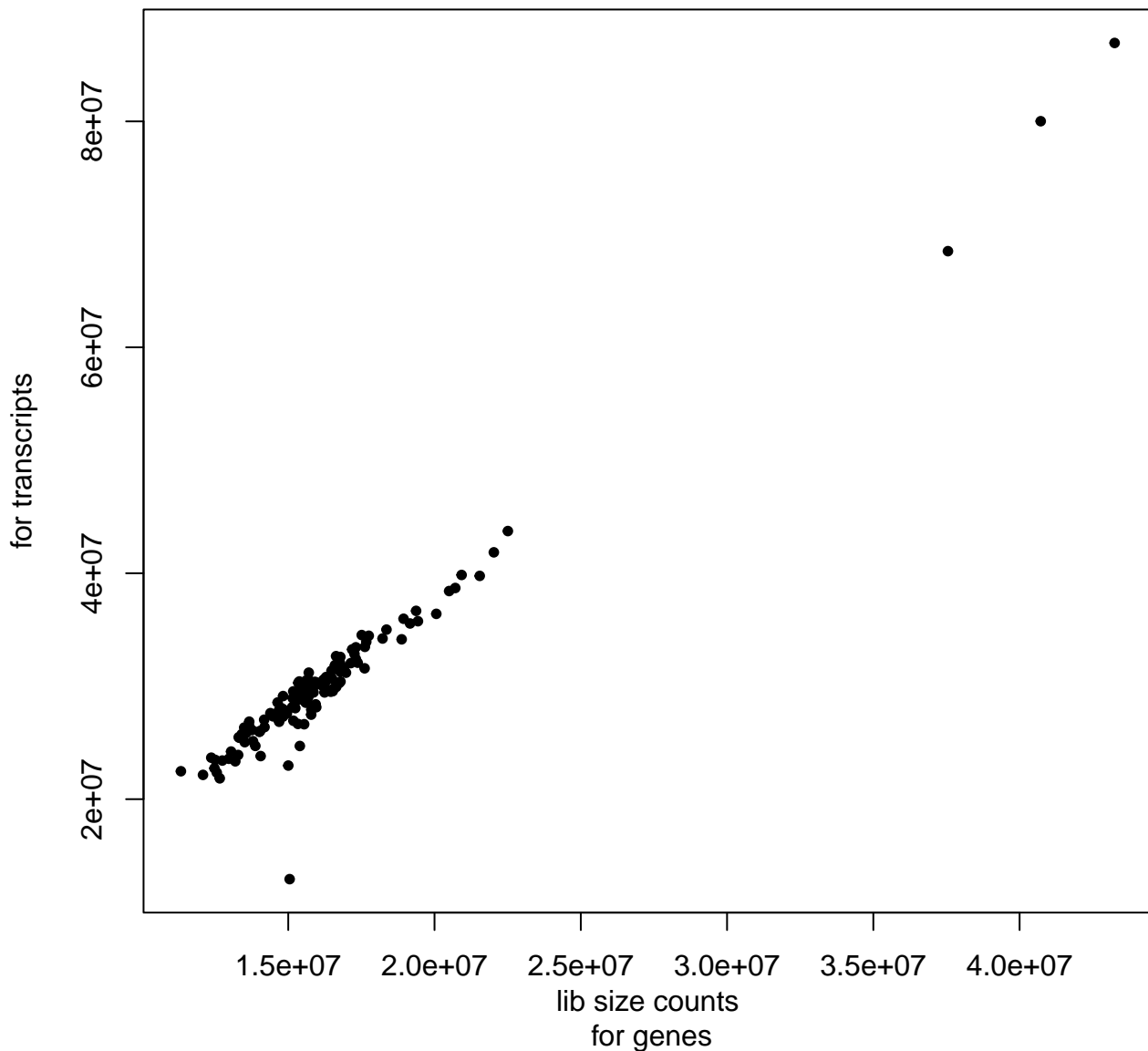
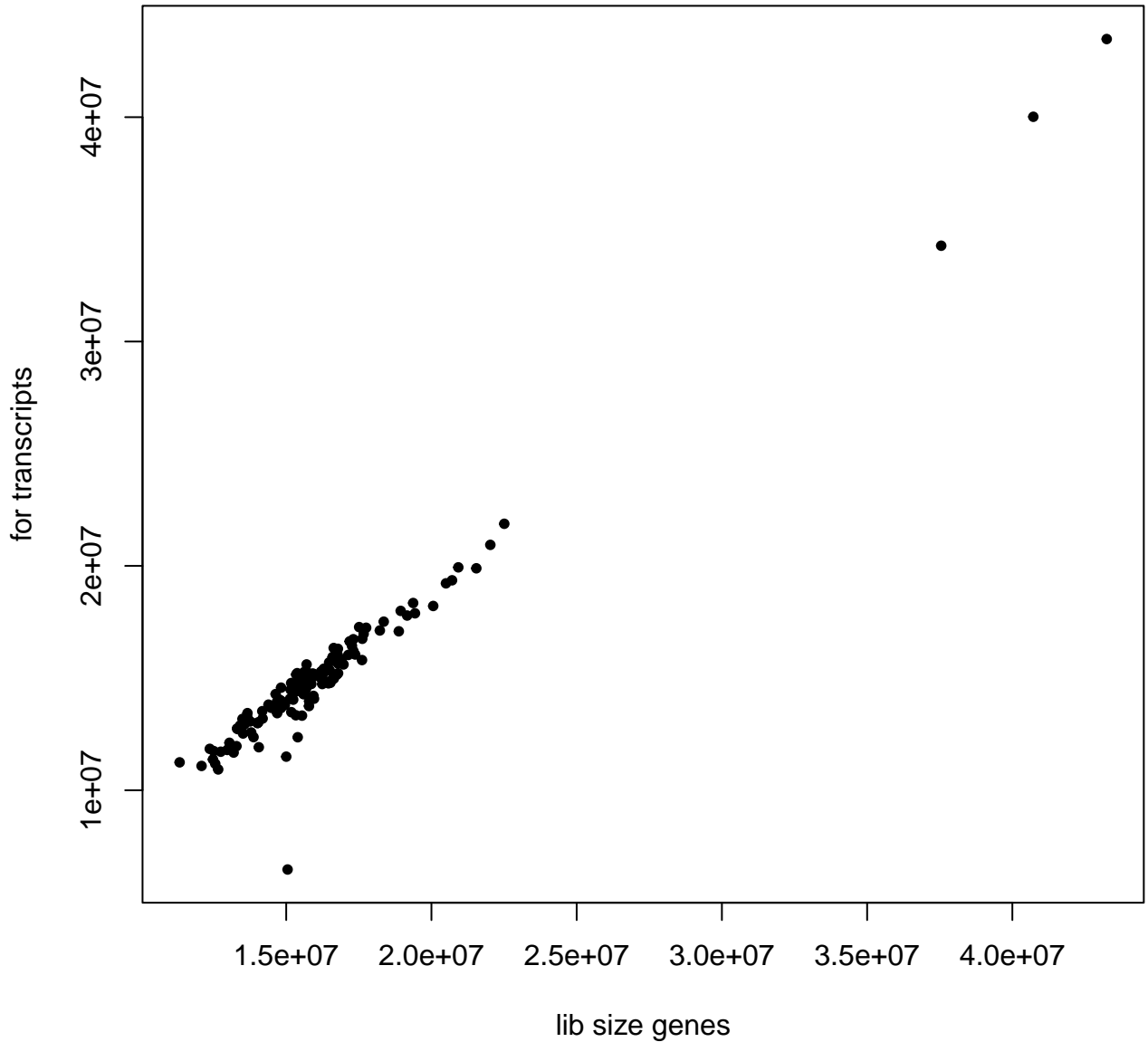


**Converted counts with read  
length = 75  
cor= 0.979061932083825**



Converted counts with read  
length = 150  
cor= 0.979071995373833



# Transcripts and genes with more than 1 CPM

