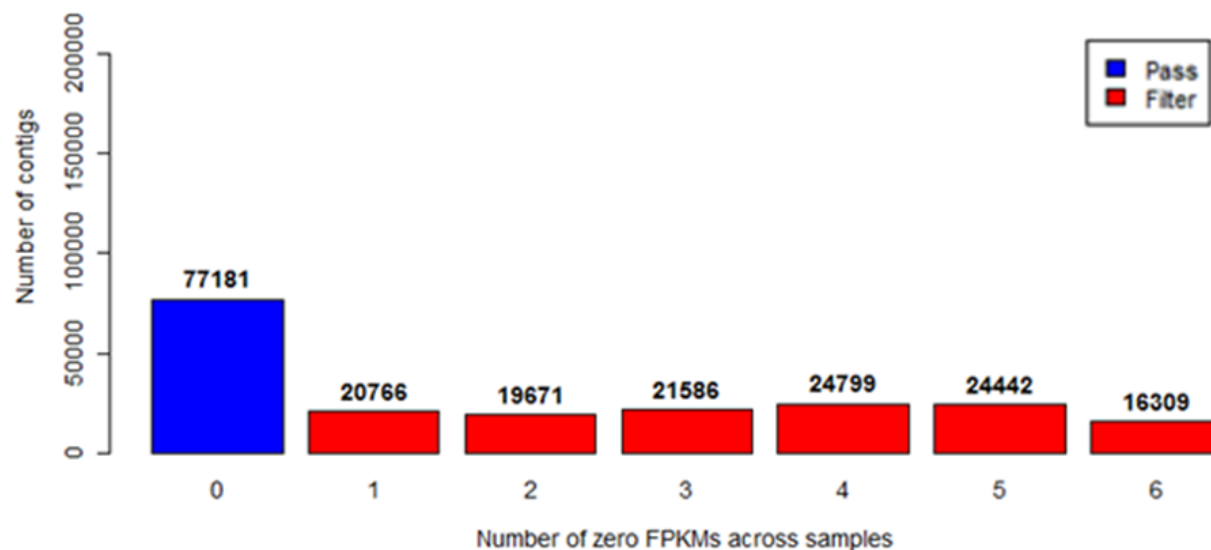


Supplementary Material



Supplementary Figure 1. Analysis of expression values of the contigs assembled from *F. chiloensis* libraries. The FPKM values of 127,573 contigs were compared within the different RNA-seq libraries. The contigs were classified according to the number of zero FPKM values in the samples. A total of 77,181 contigs continued in the analysis.