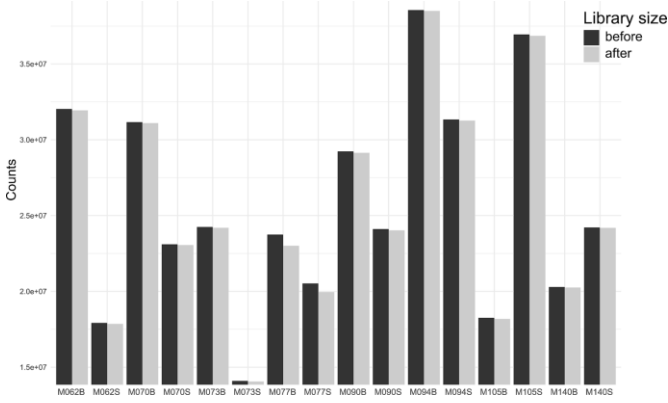
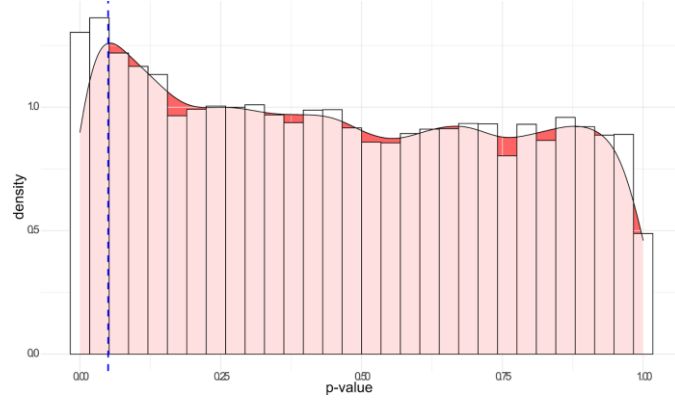


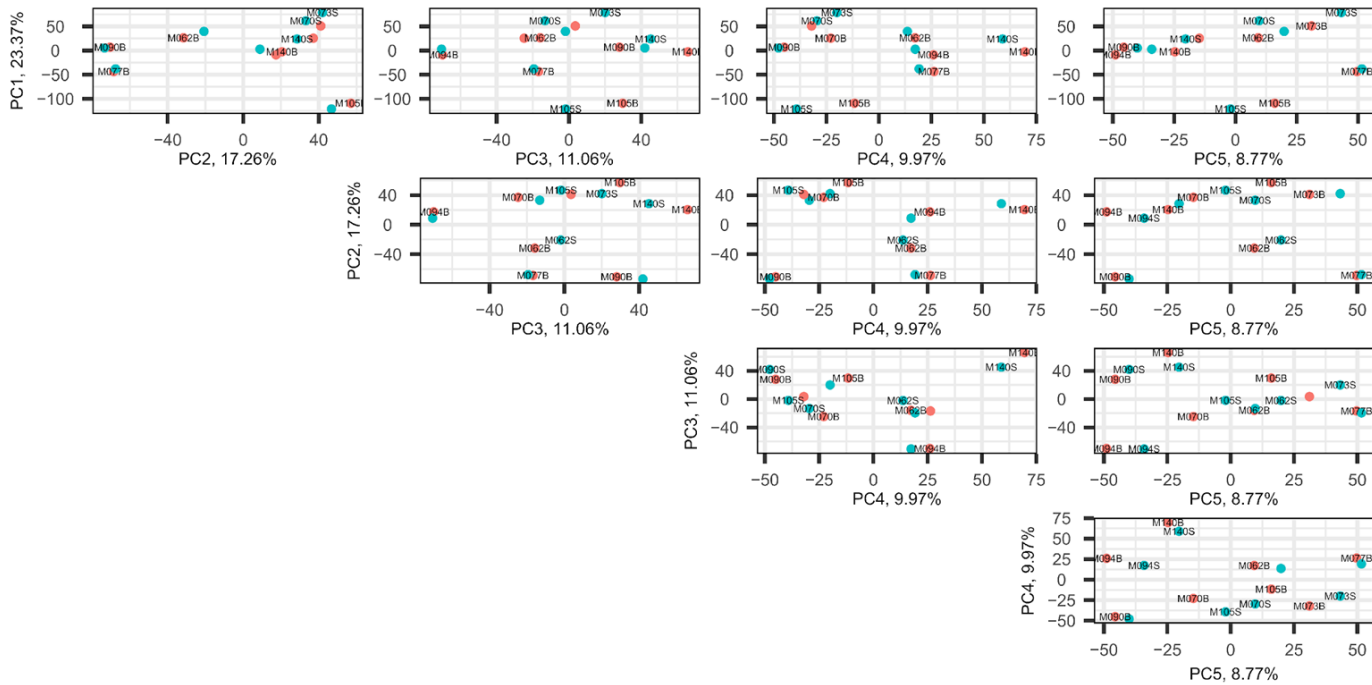
**A**



**B**



**C**



**Supplementary Figure S3. A**, Library size before and after low count filtering. **B**, P-value histogram distribution observed in the differential expression analysis. **C**, Pairwise scatter plots of the first five principal components from the principal component analysis of RNA-Seq data.