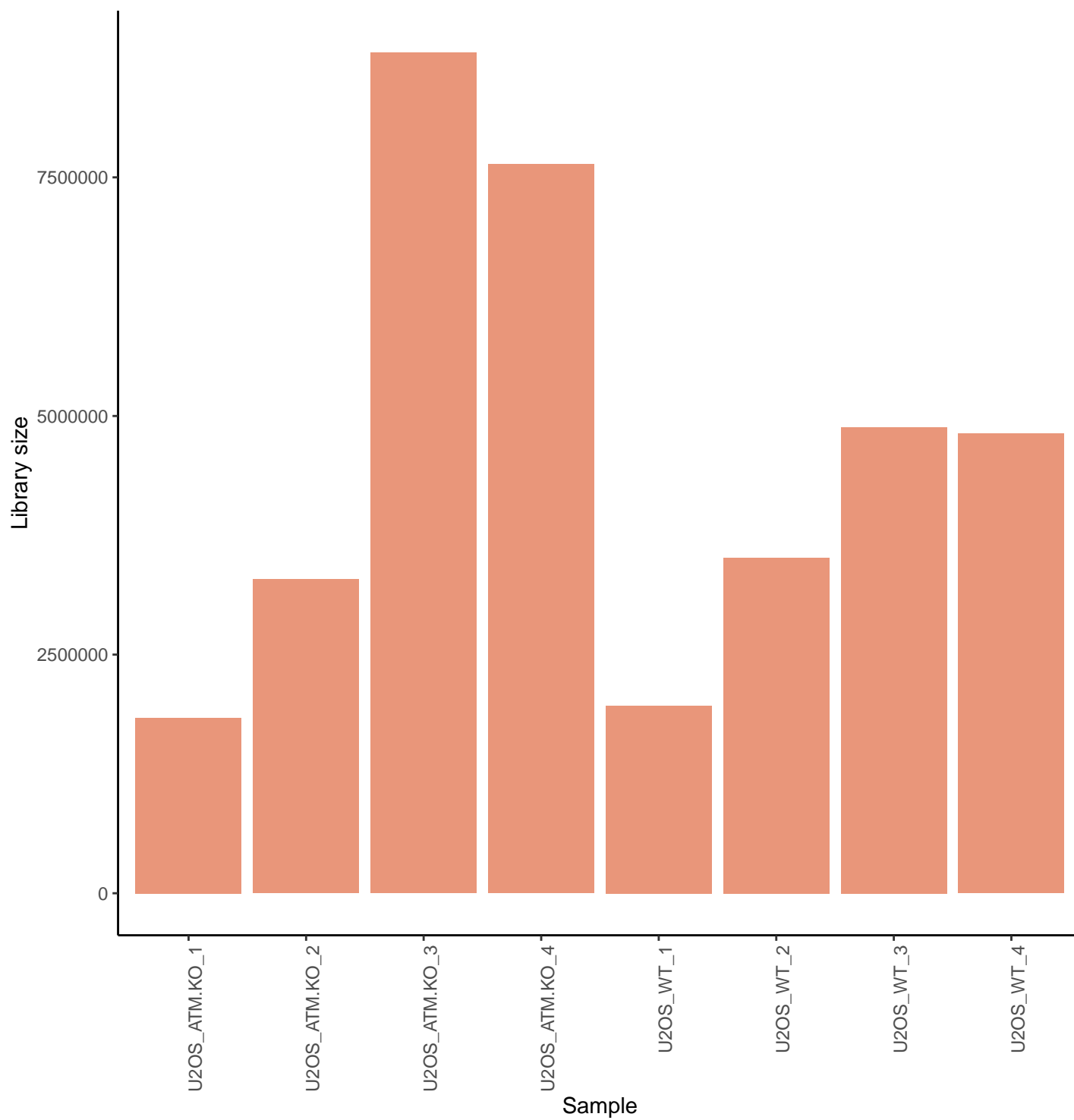


Barplot of library sizes



Number of genes with count > 0

