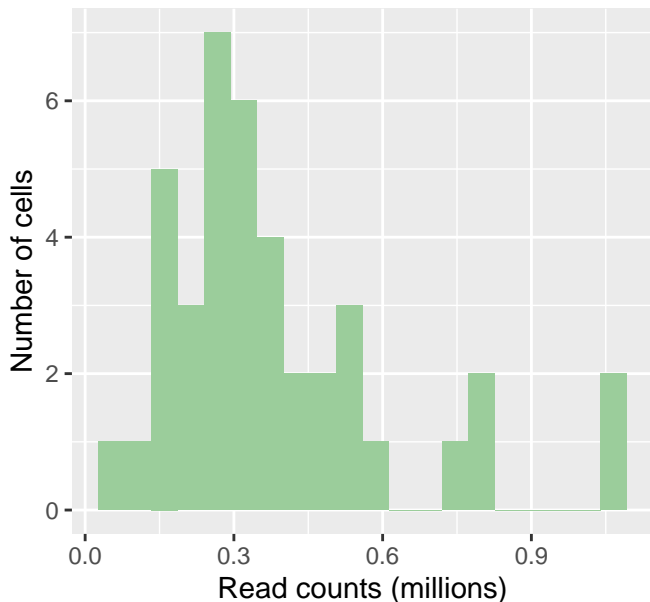
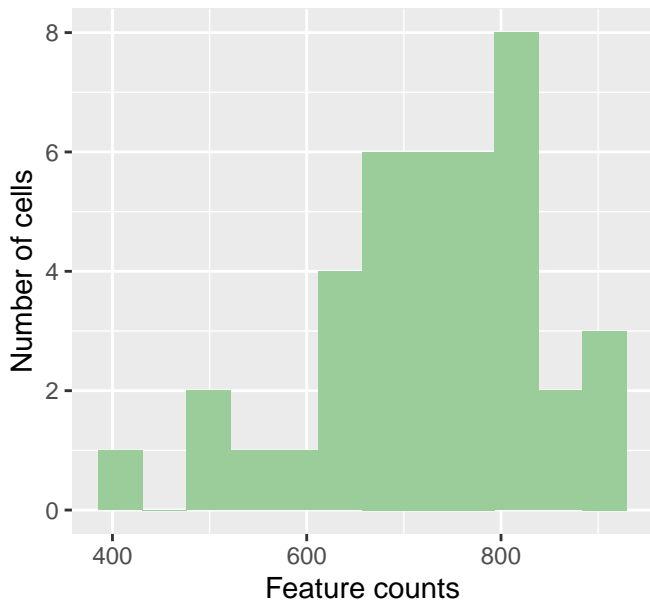


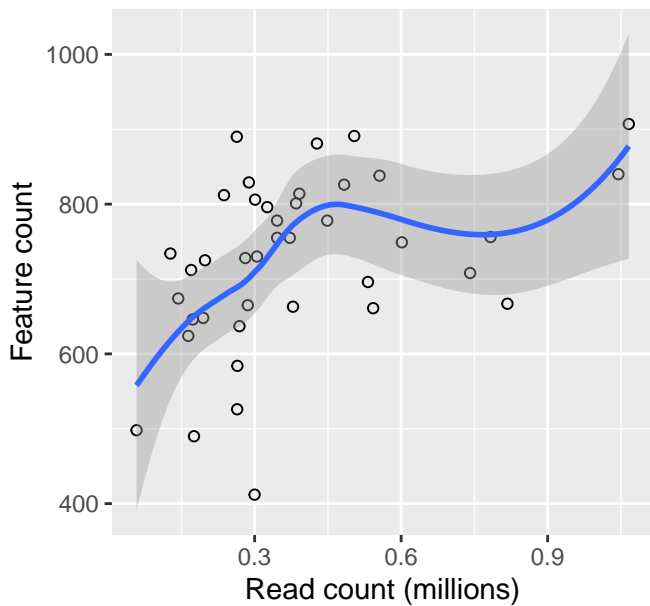
Read counts per cell



Feature counts per cell



Scatterplot of reads vs features



% MT genes

