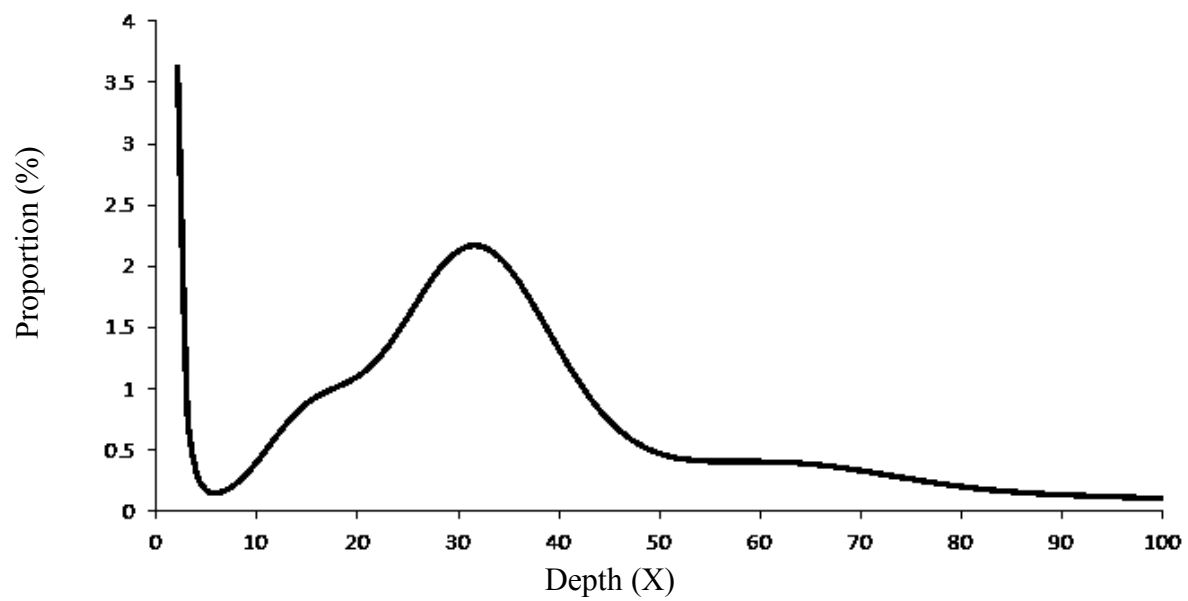




Supplementary Figure 1: 17-mer coverage distribution. The distribution was used to estimate the *S. caffer* genome size. The x-axis is 17-mercoverage depth and the y-axis is the percentage of the total number of 17-mers.