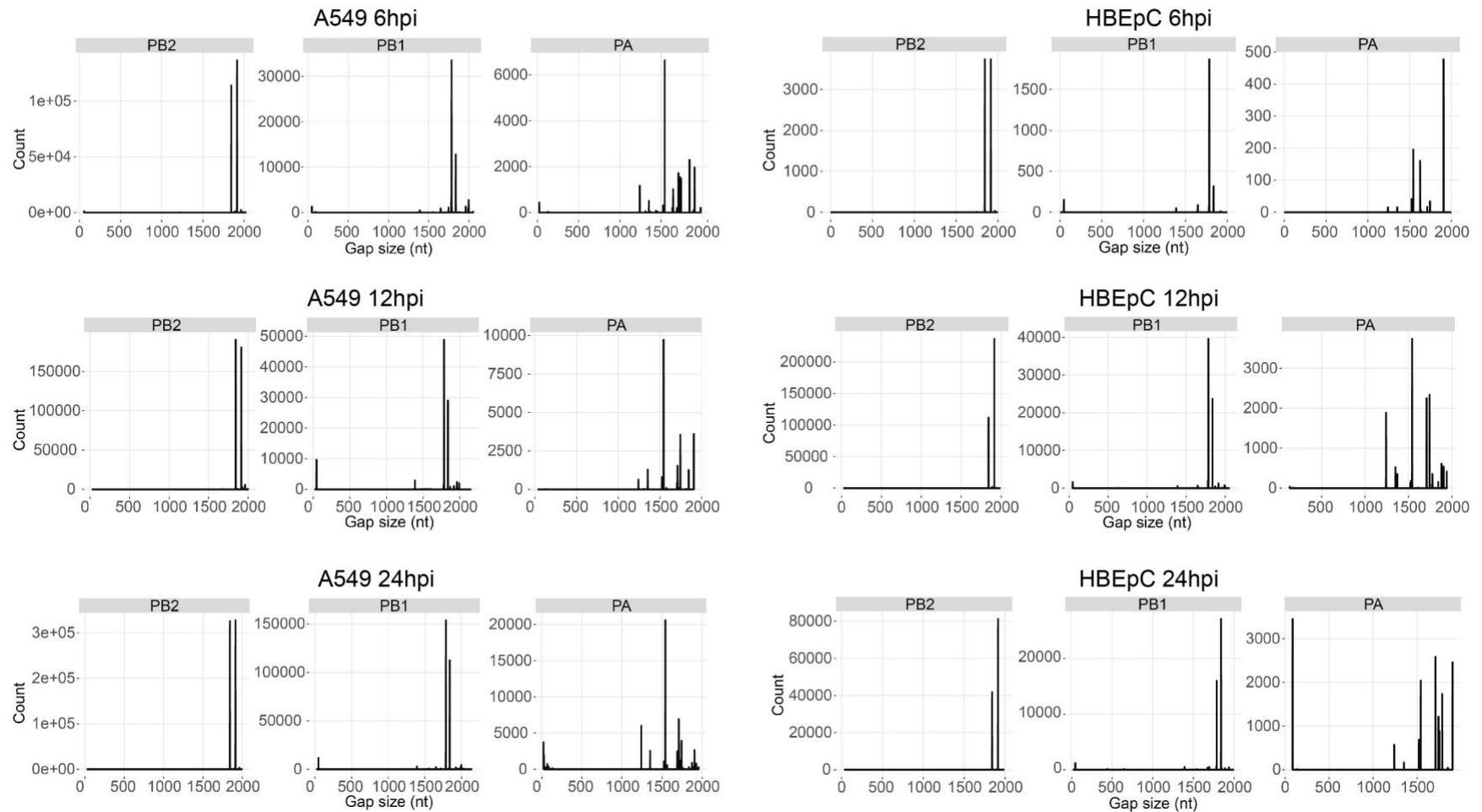


Supplementary File 9.



Supplementary File 9. Distribution of the size of the deletions in the gap-spanning reads aligned to the polymerase segments in two cell types over the course of the infection. The frequency of each size detected in a sample was represented by the height of the bar shown on the y-axis.