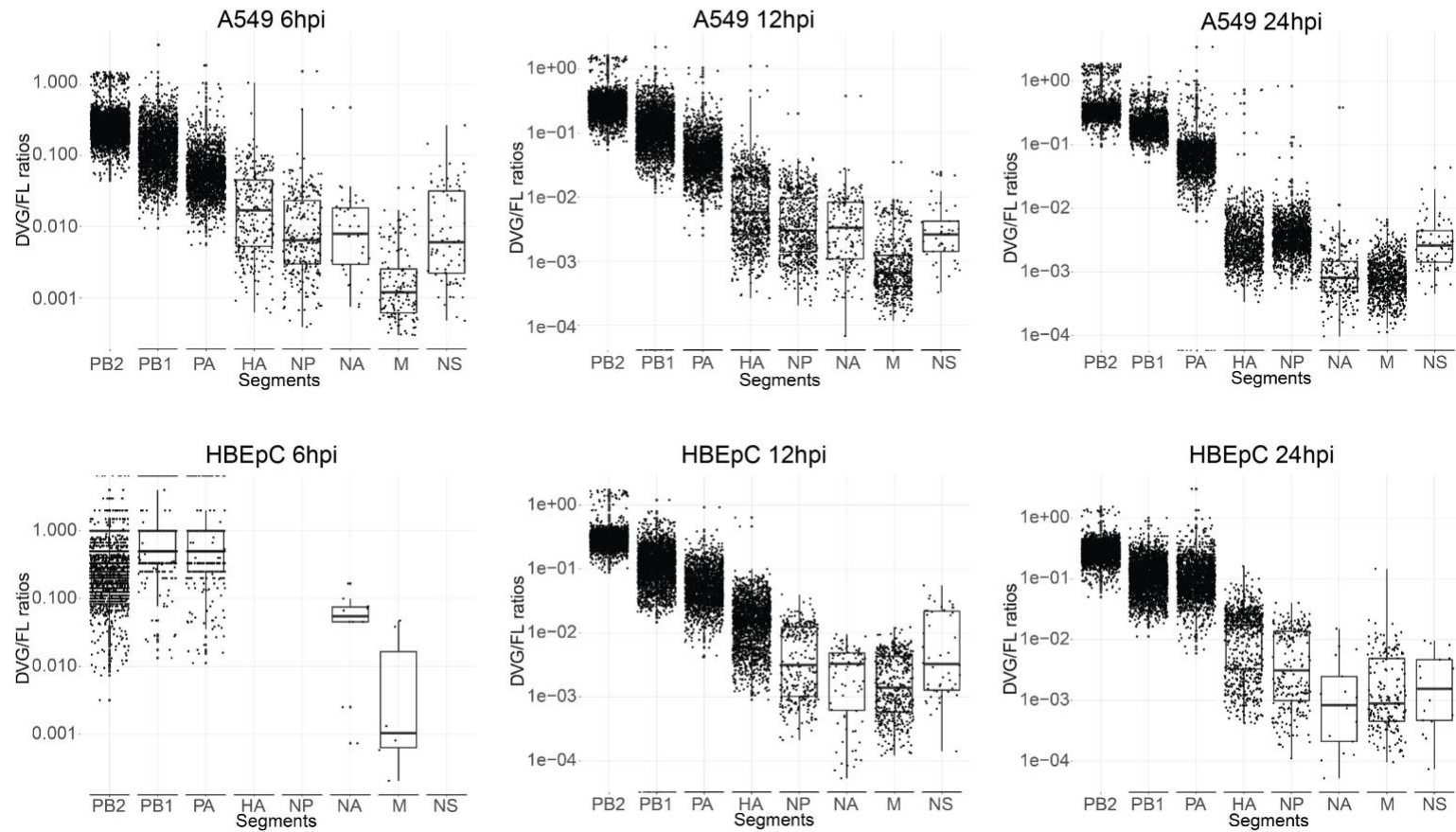


Supplementary File 1.



Supplementary File 1. Distribution of DVG/FL ratios for each viral segment in two cell types over the course of the infection. DVG transcripts represented by gap-spanning reads were grouped by their junction coordinates and filtered by their frequency in each sample (i.e., ≥ 10 times)