

Figure S1A. Genotyping of the EBV+ sequencing reads aligned to human genome. The pie charts display the distribution of mapped reads from EBV+ patient samples.

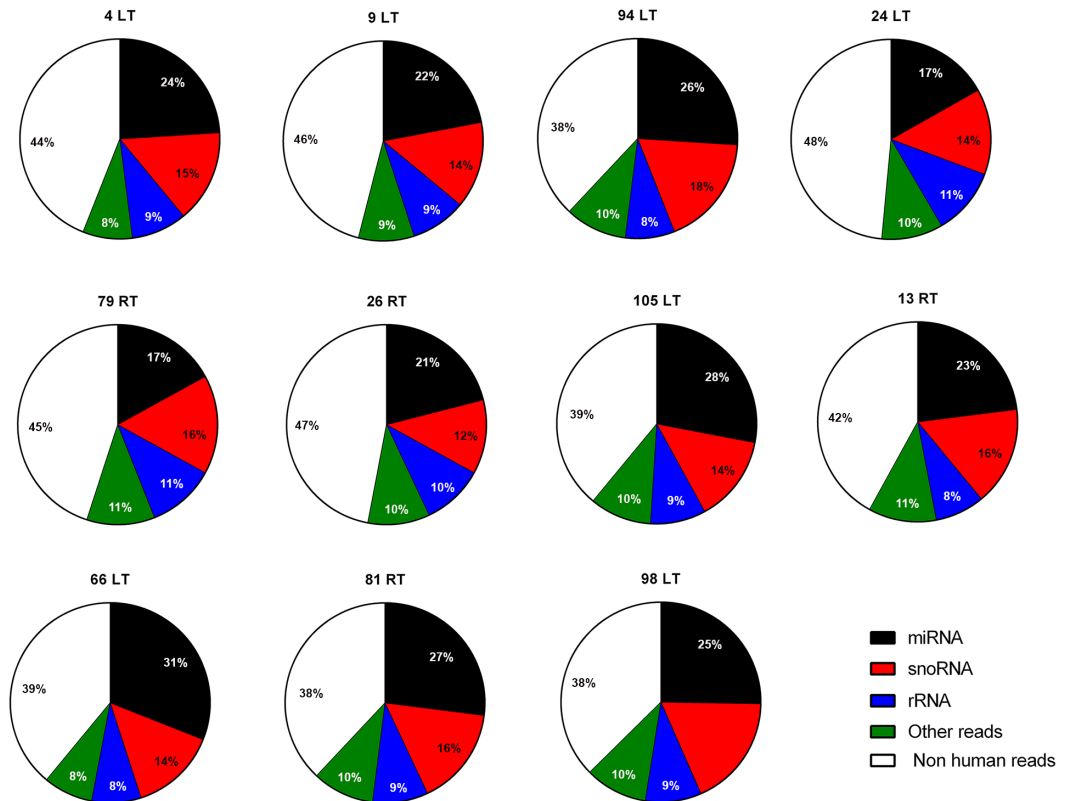


Figure S1B. Genotyping of the HAdV sequencing reads aligned to human genome. The pie charts display the distribution of the mapped reads from HAdV+ patient samples.