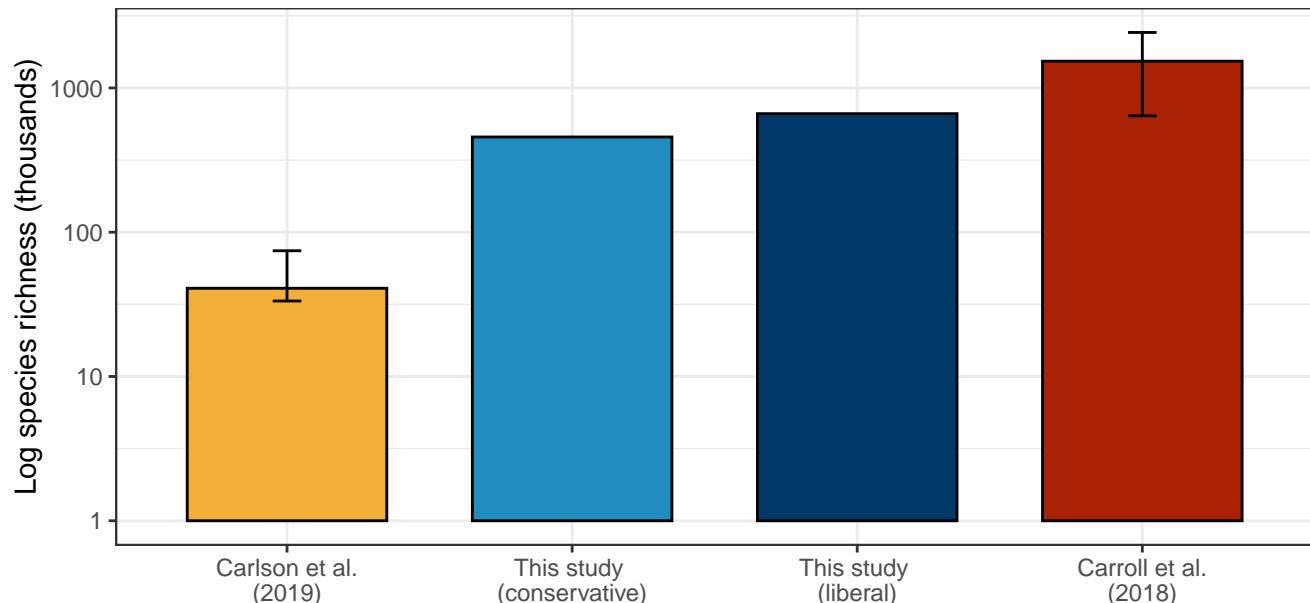
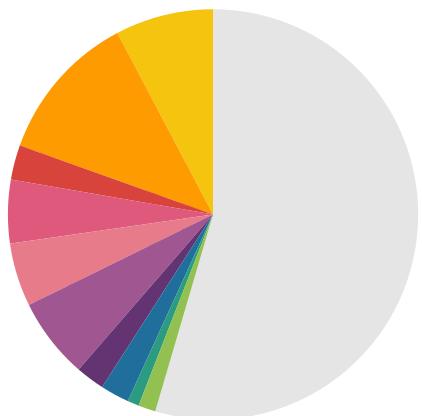


# Estimated global diversity of mammalian viruses



Carroll et al. (2018)

Raw metagenomic samples  
uncorrected for host range



This study (2025)

Global estimates broken down  
for vertebrate–infective groups

