



Supplementary Figure 6. Shifts in the transcript abundance patterns of eukaryotic and giant viruses. Each plot shows a principal components (PC) analysis ordination of normalized transcript patterns across A) Eukaryotic *RPB1* candidates colored by date sampled B) *Nucleocytoviricota* total genome recruitments colored by date sampled C) Eukaryotic *RPB1* candidates colored by depth sampled D) *Nucleocytoviricota* total genome recruitments colored by depth sampled.