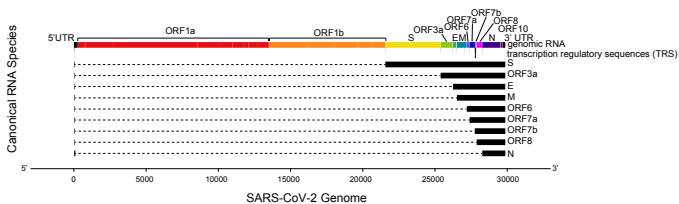
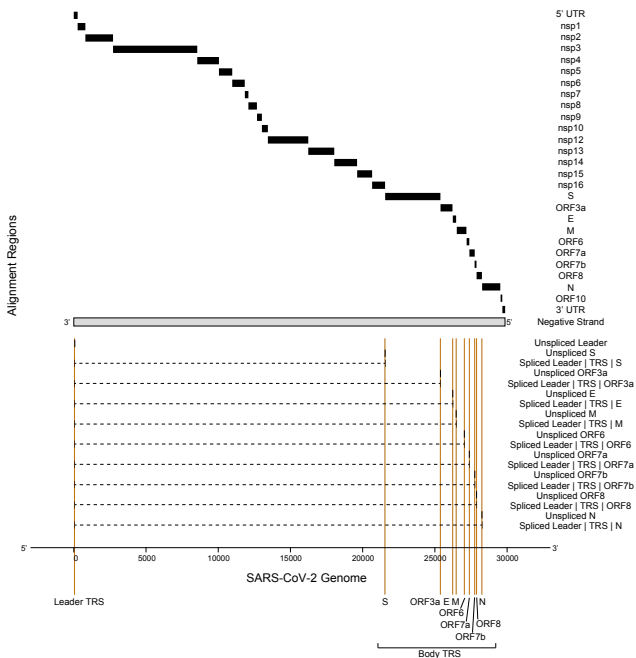


Canonical RNA Species in SARS-CoV-2 (from Kim et al. 2020)



GTF regions in custom genome



Distinguishing between reads that align across spliced vs. unspliced TRS sites

