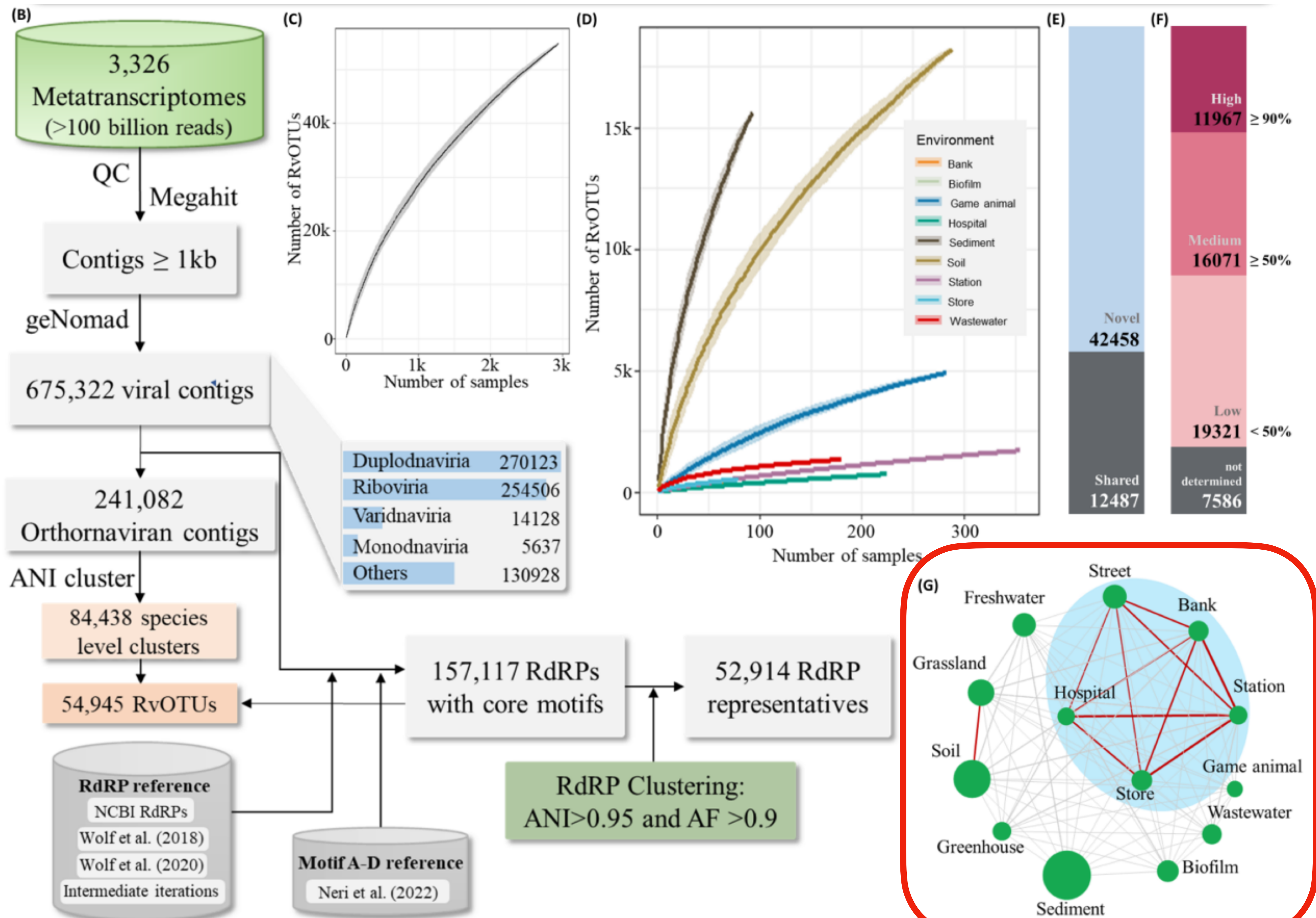


MetaCov Workflow

Environment
sample variability



Phylogenetic reconstruction of the RNA virome using RdRP

