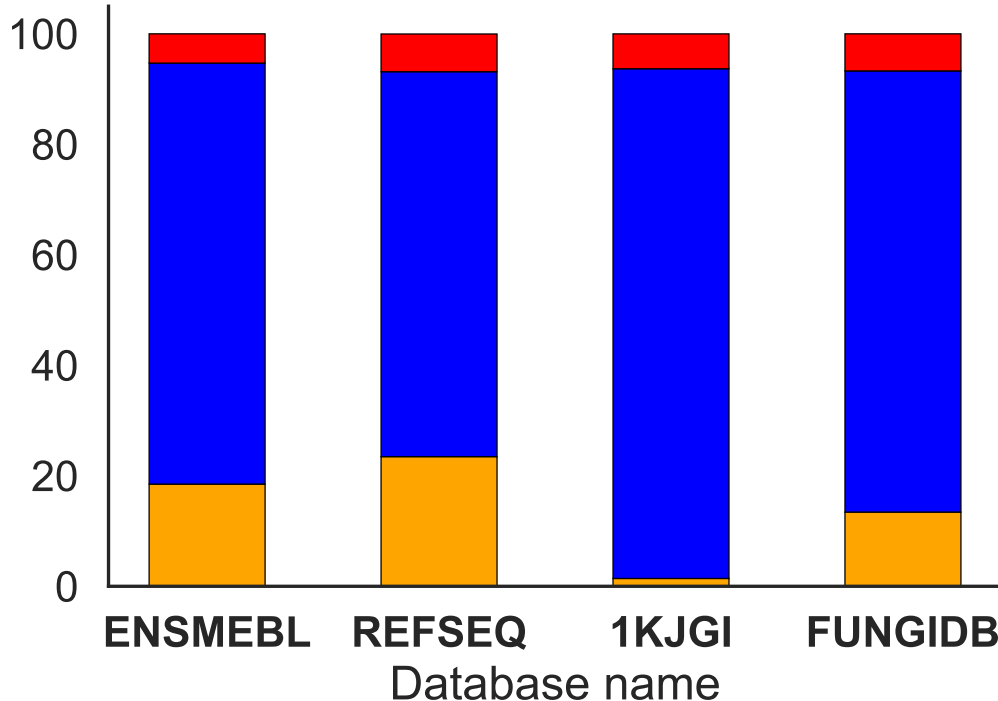


Number of species



(1) Complete genome



(2) Fragmented genome (Set of contigs)



Mixture of (1) and (2)