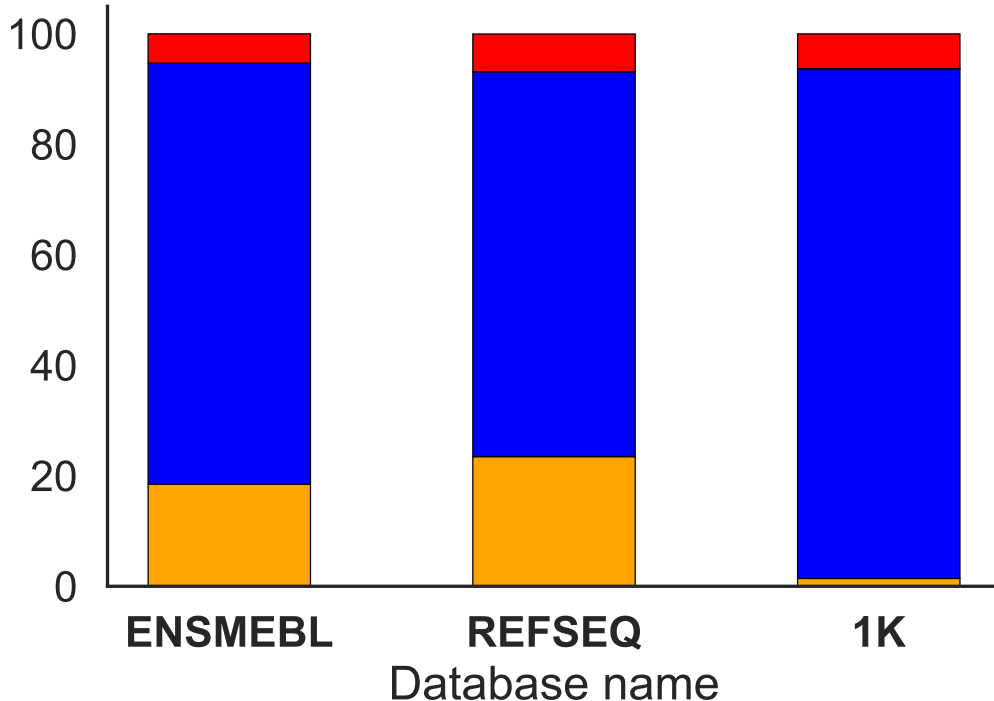


Number of species



- (1) Complete genome
- (2) Fragmented genome (Set of contigs)
- Mixture of (1) and (2)