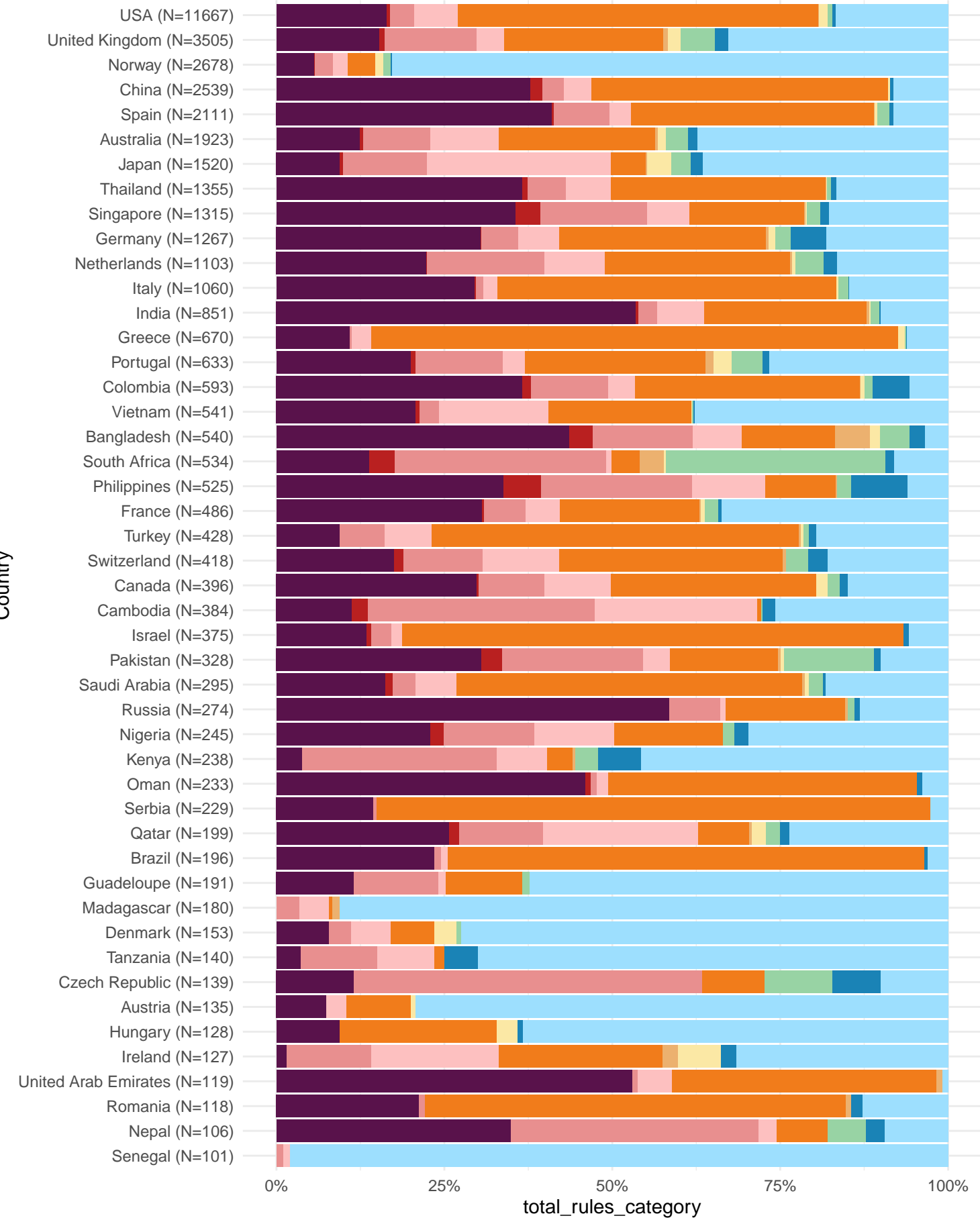


A) Country (>=100 genomes)



B) Country (>100 genomes) sorted by predicted resistance rate

