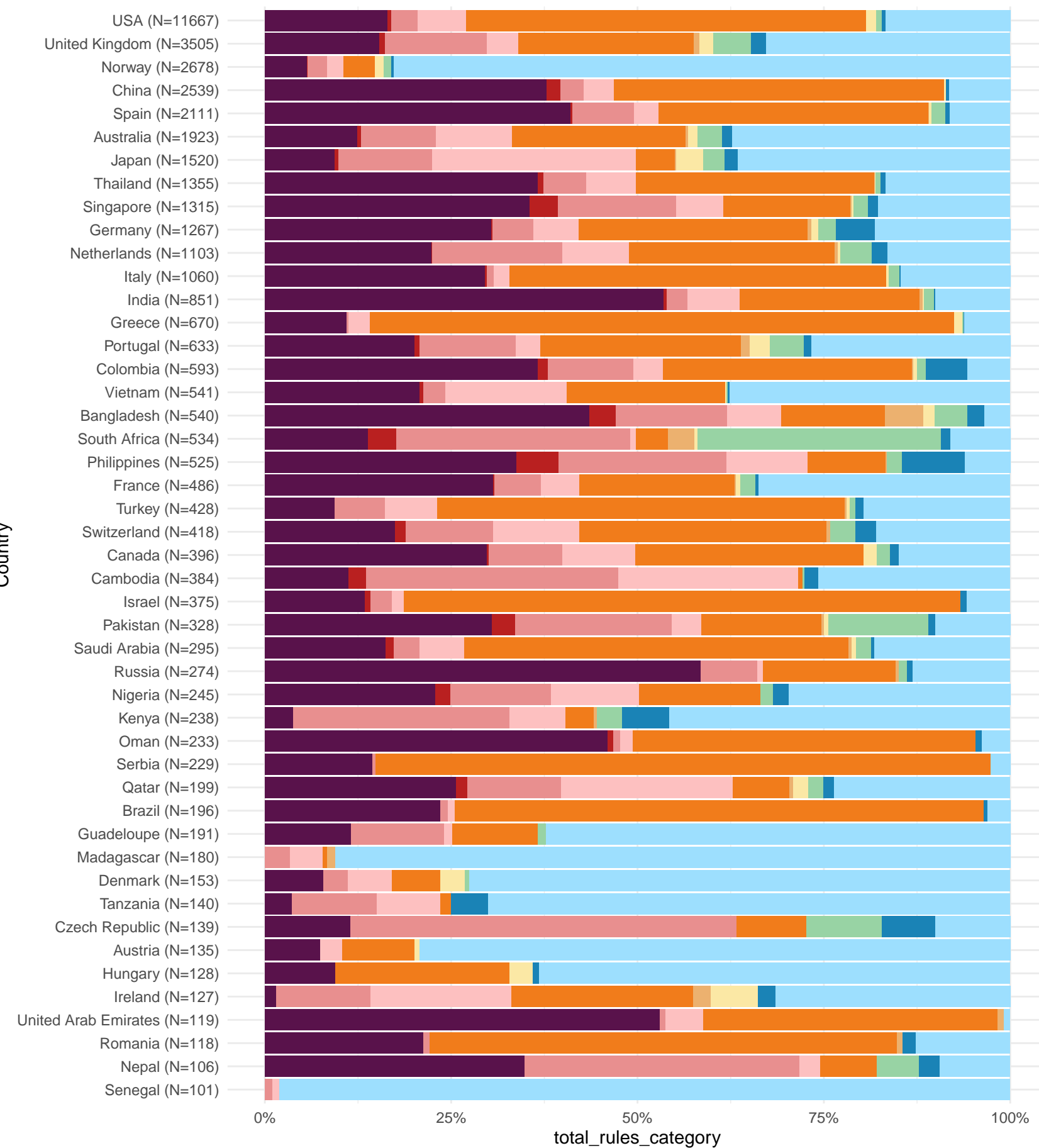
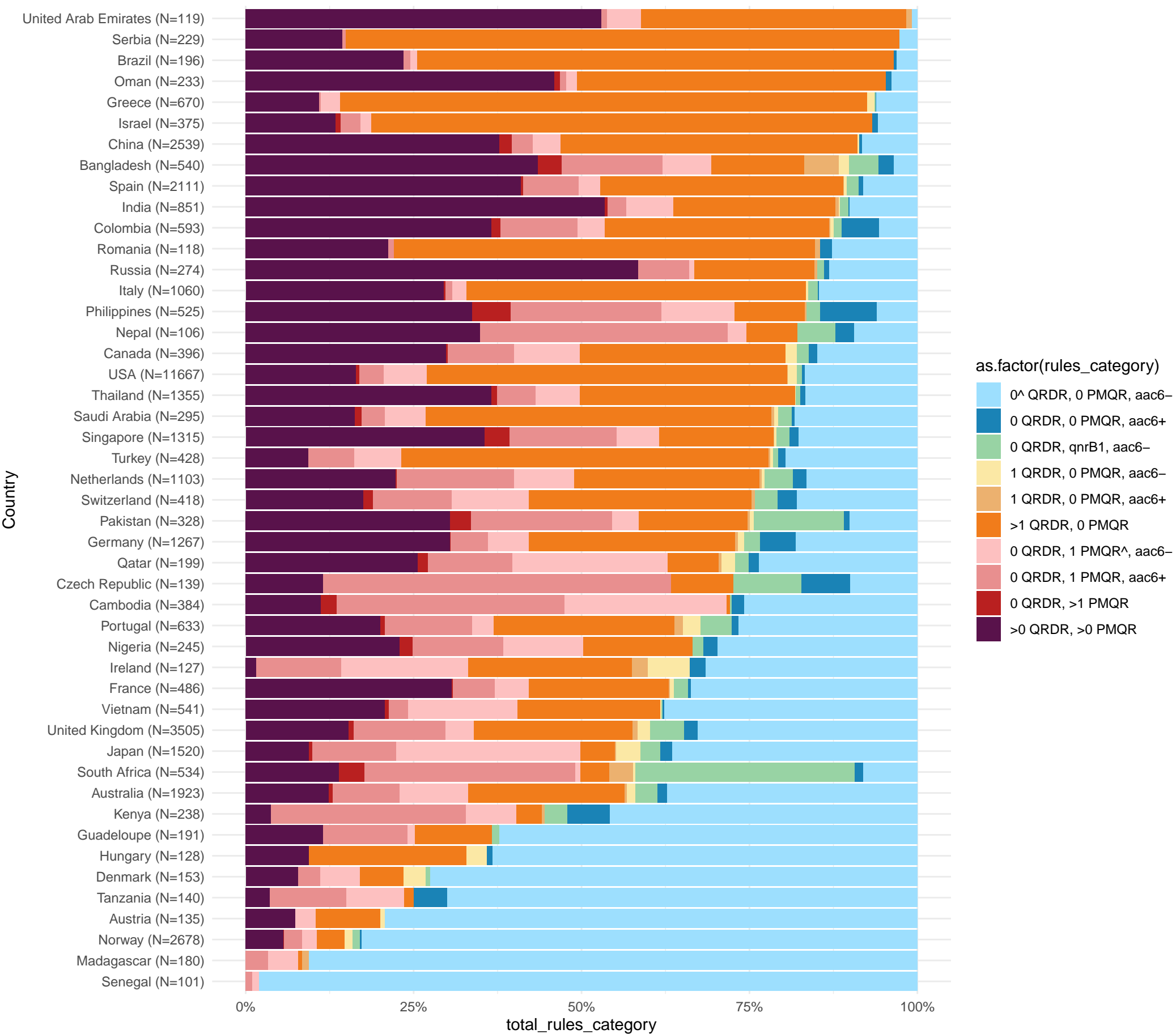


A) Country (>=100 genomes)



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



- as.factor(rules_category)
- 0[^] QRDR, 0 PMQR, aac6-
 - 0 QRDR, 0 PMQR, aac6+
 - 0 QRDR, qnrB1, aac6-
 - 1 QRDR, 0 PMQR, aac6-
 - 1 QRDR, 0 PMQR, aac6+
 - >1 QRDR, 0 PMQR
 - 0 QRDR, 1 PMQR[^], aac6-
 - 0 QRDR, 1 PMQR, aac6+
 - 0 QRDR, >1 PMQR
 - >0 QRDR, >0 PMQR