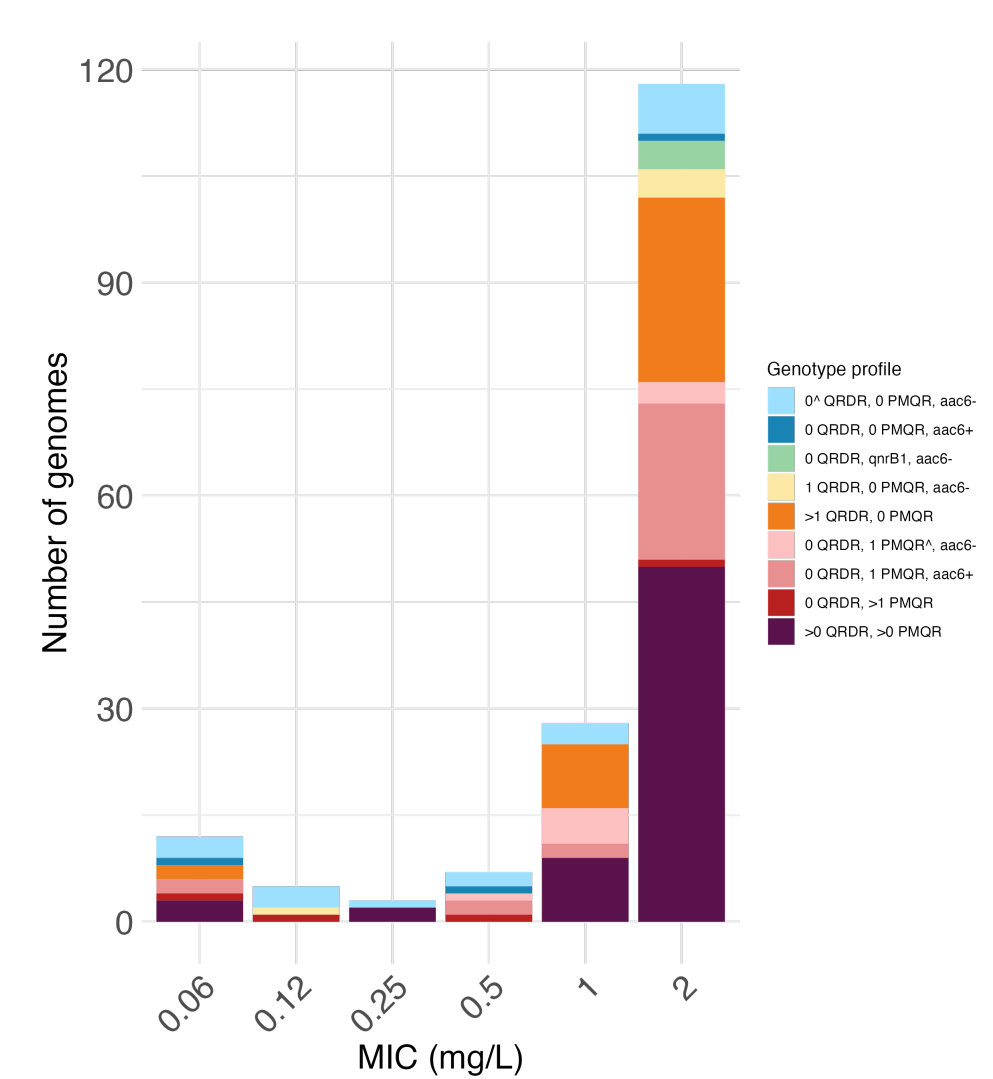
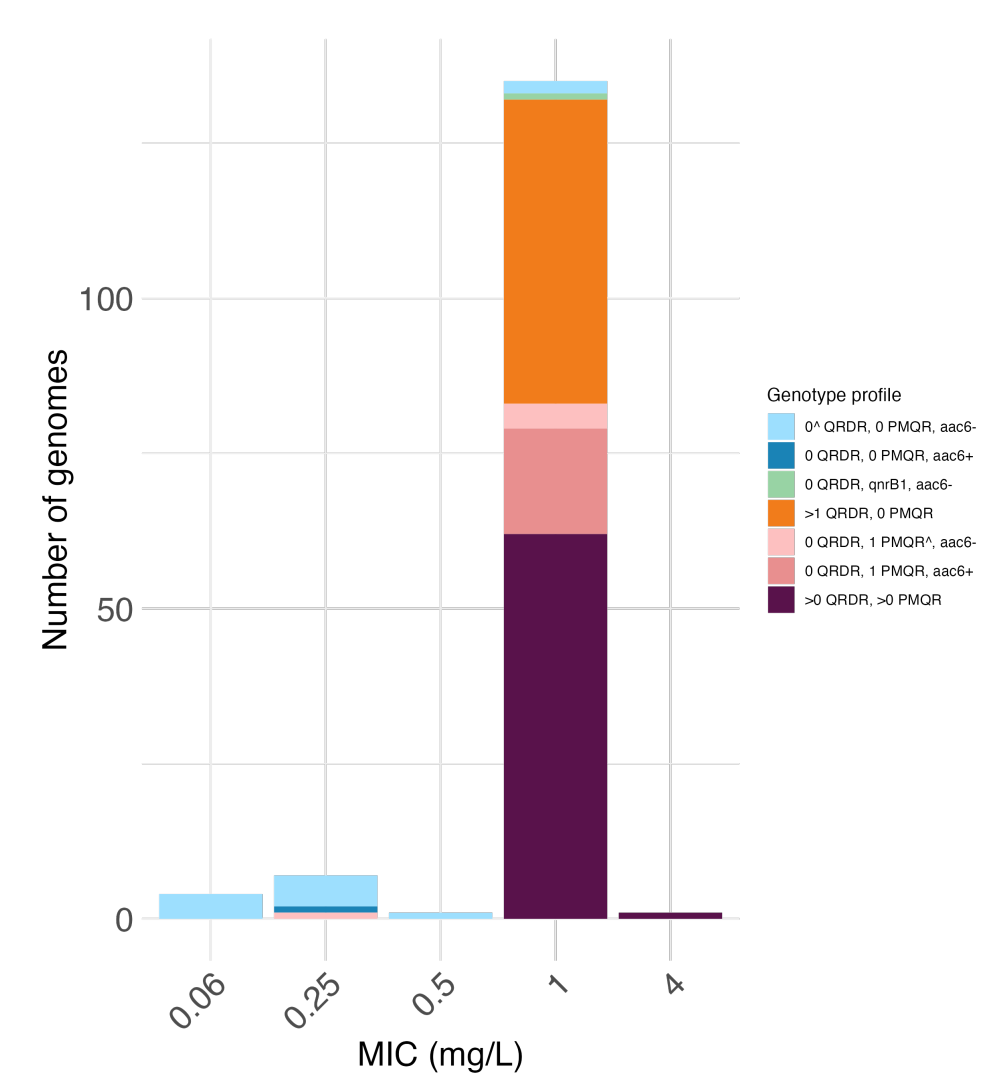


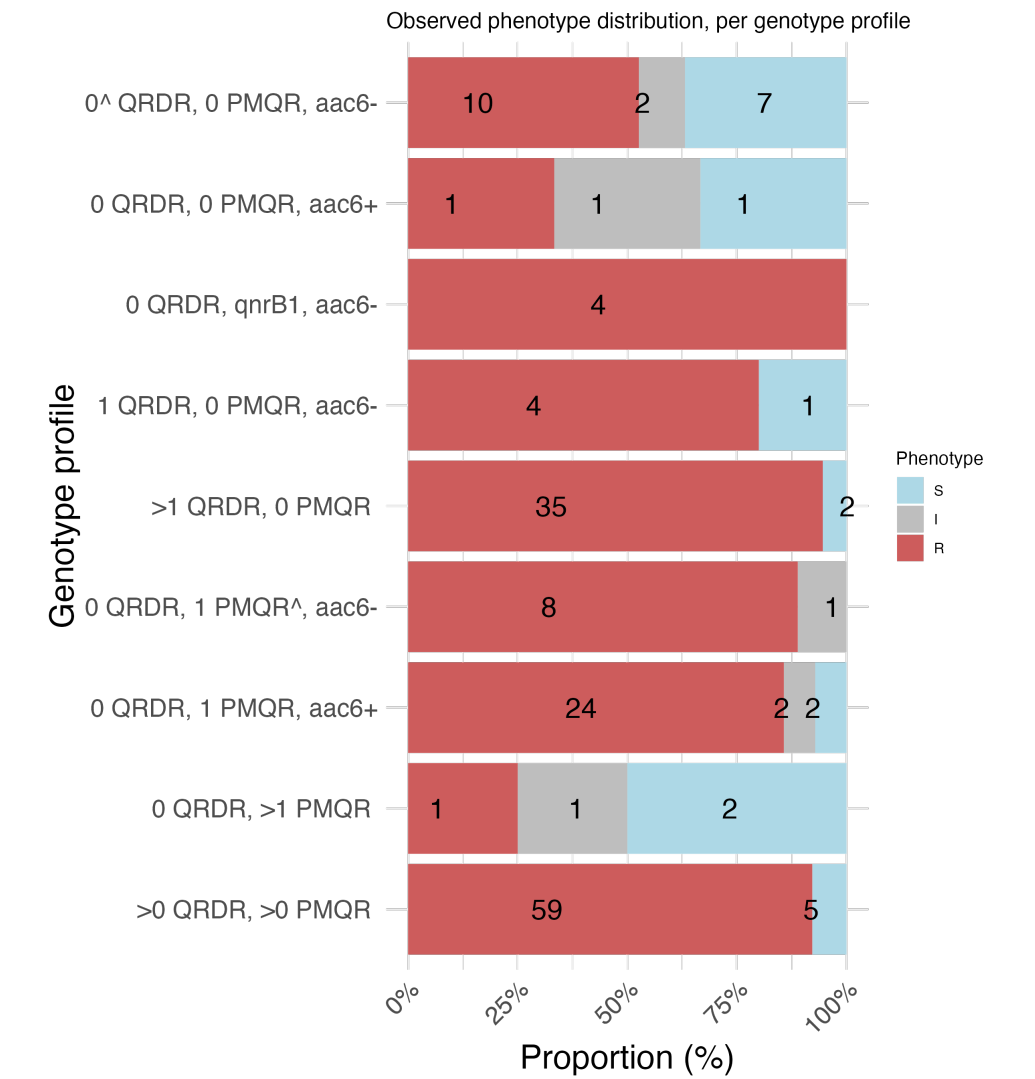
ix) E (EUCAST) /
HURYC & CIBERINFEC (n=173) – MIC distribution vs.
genotype profile



xi) F (EUCAST) /
HULP (n=148) – MIC distribution vs.
genotype profile



x) E (EUCAST)
HURYC & CIBERINFEC (n=173)
– observed phenotype vs. genotype profile



xii) F (EUCAST)
HULP (n=148)
– observed phenotype vs. genotype profile

