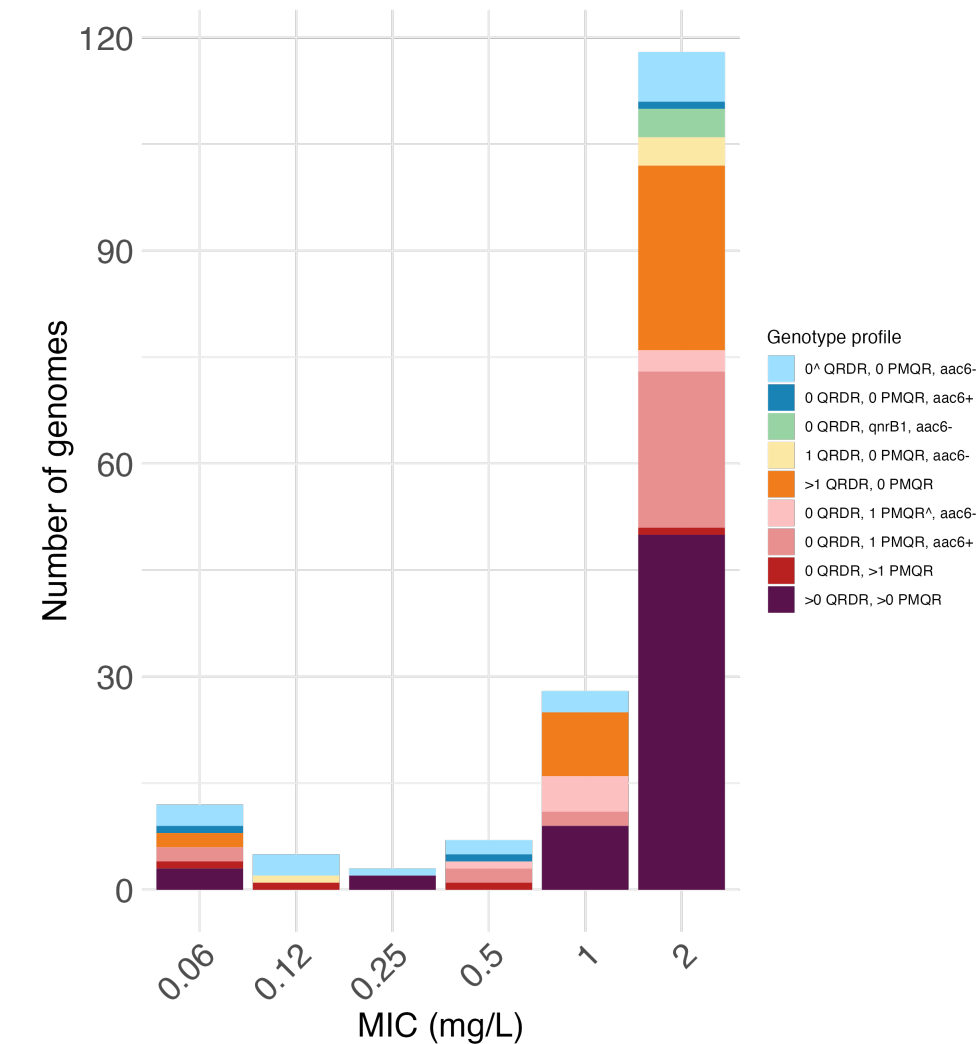
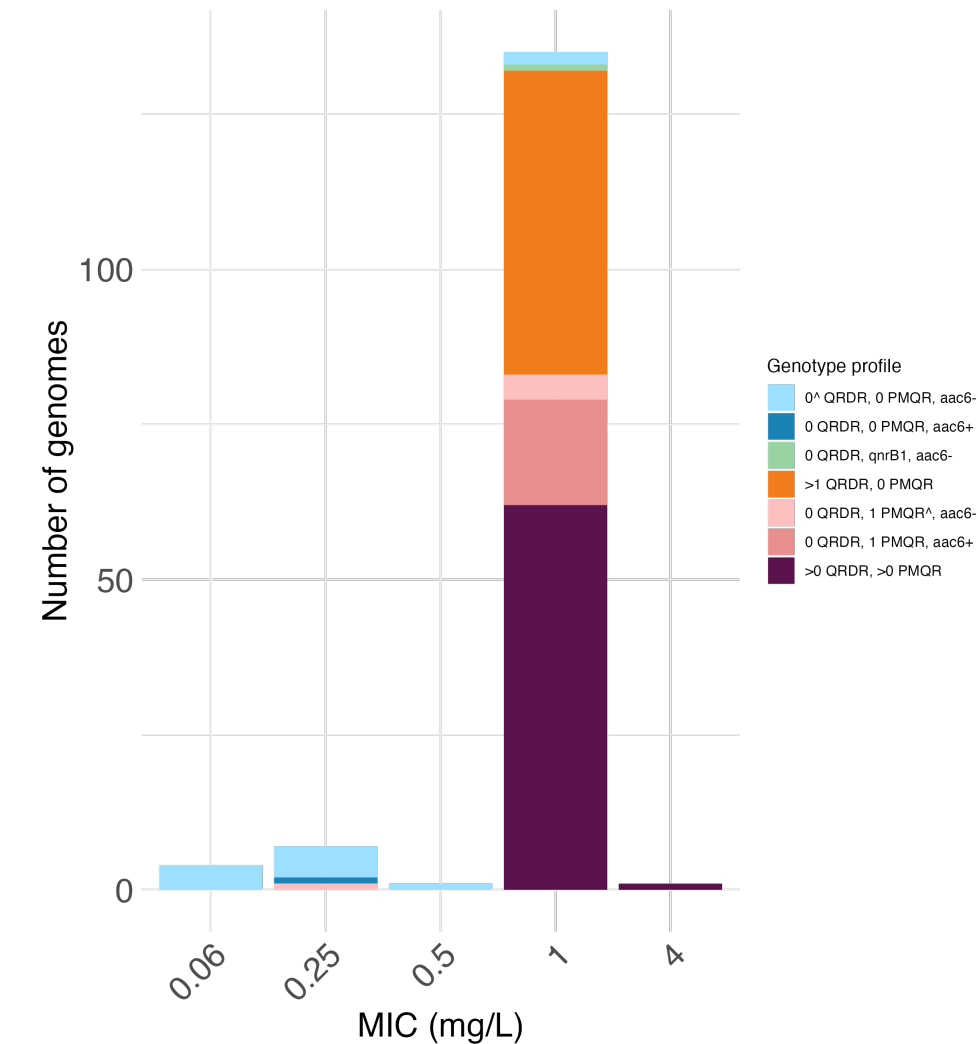


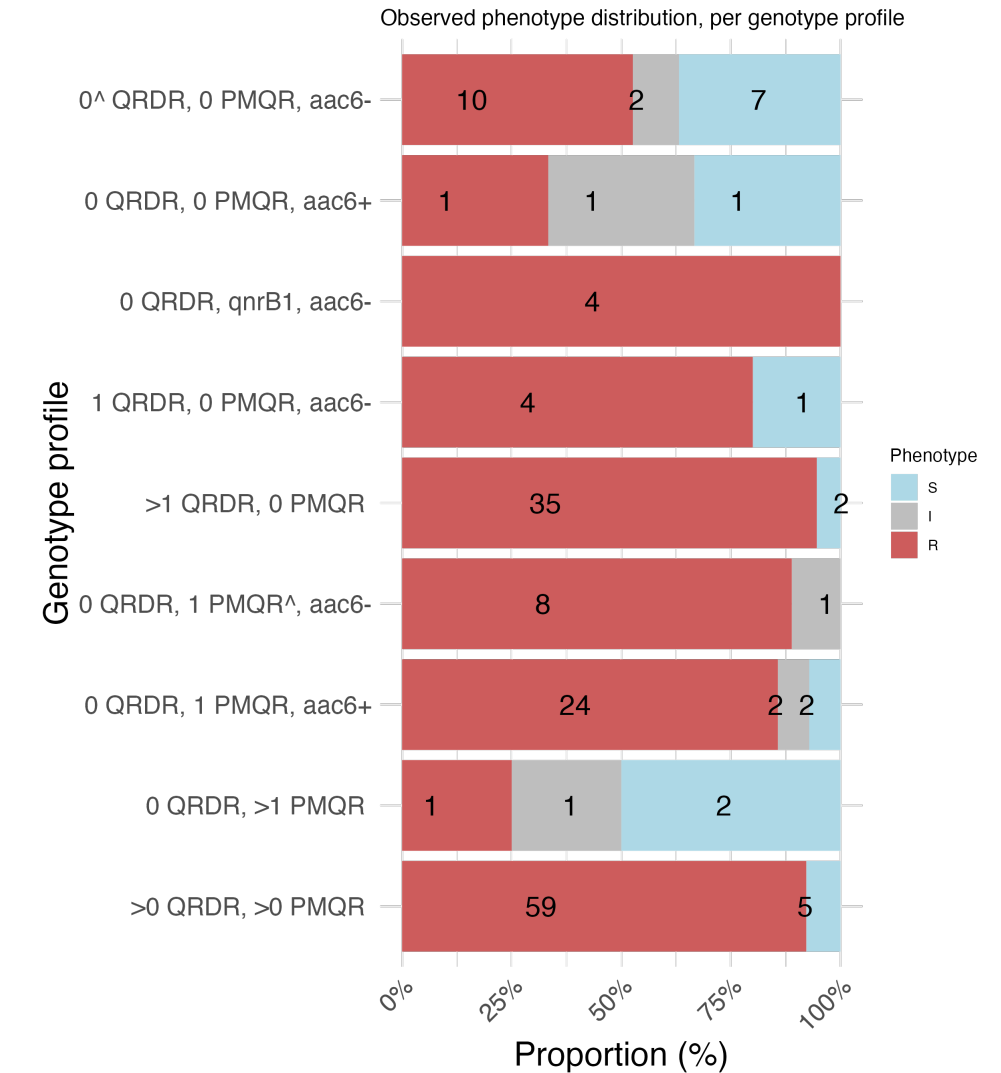
9) E (EUCAST, S/M) /
HURYC & CIBERINFEC (n=173) – MIC distribution vs.
genotype profile



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



10) E (EUCAST, S/M)
HURYC & CIBERINFEC (n=173)
– observed phenotype vs. genotype profile



12) F (EUCAST, MW/E)
HULP (n=148)
– observed phenotype vs. genotype profile

