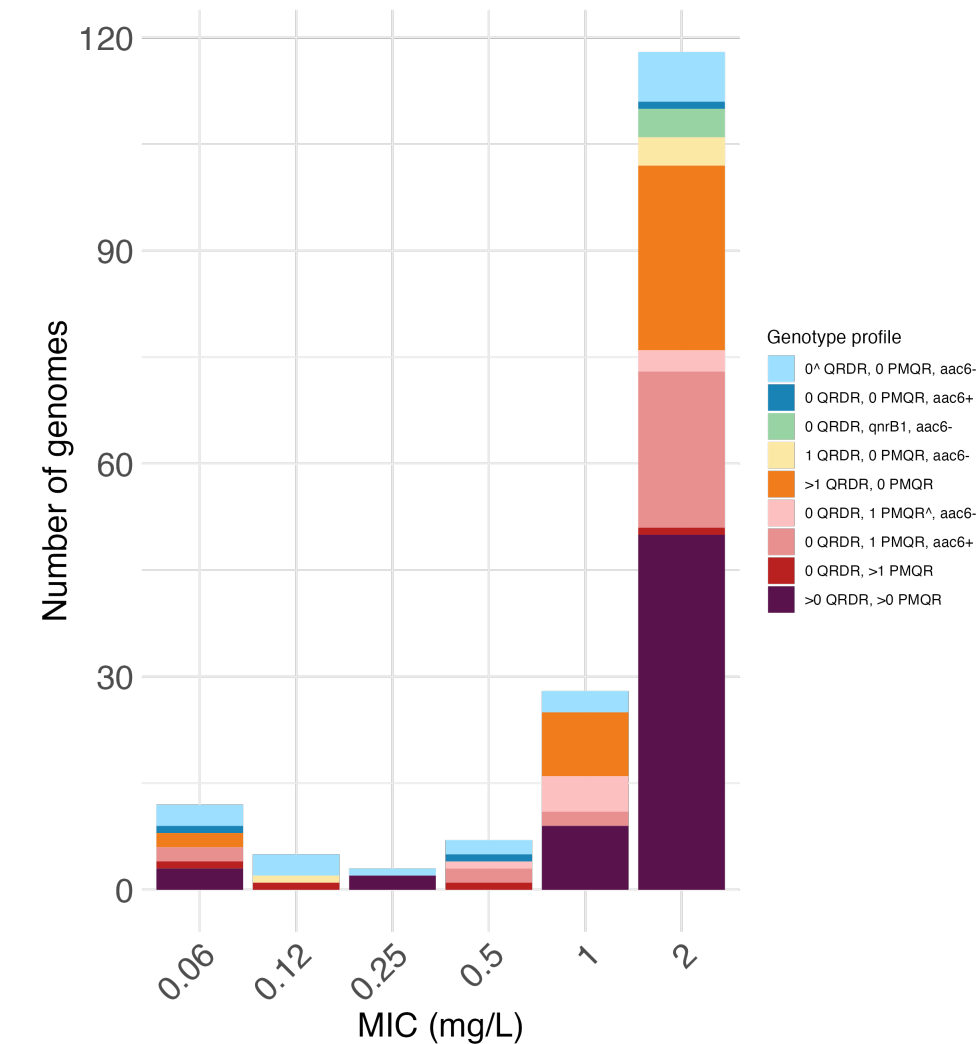
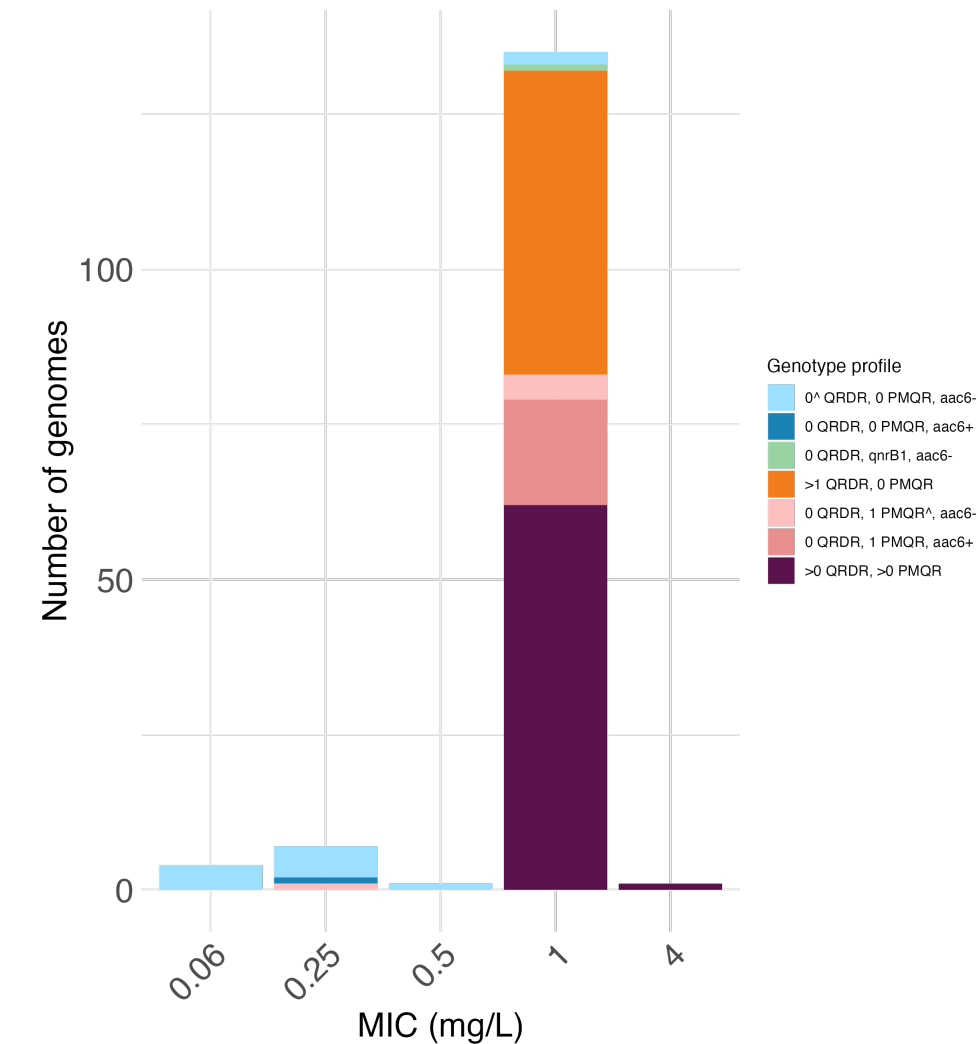


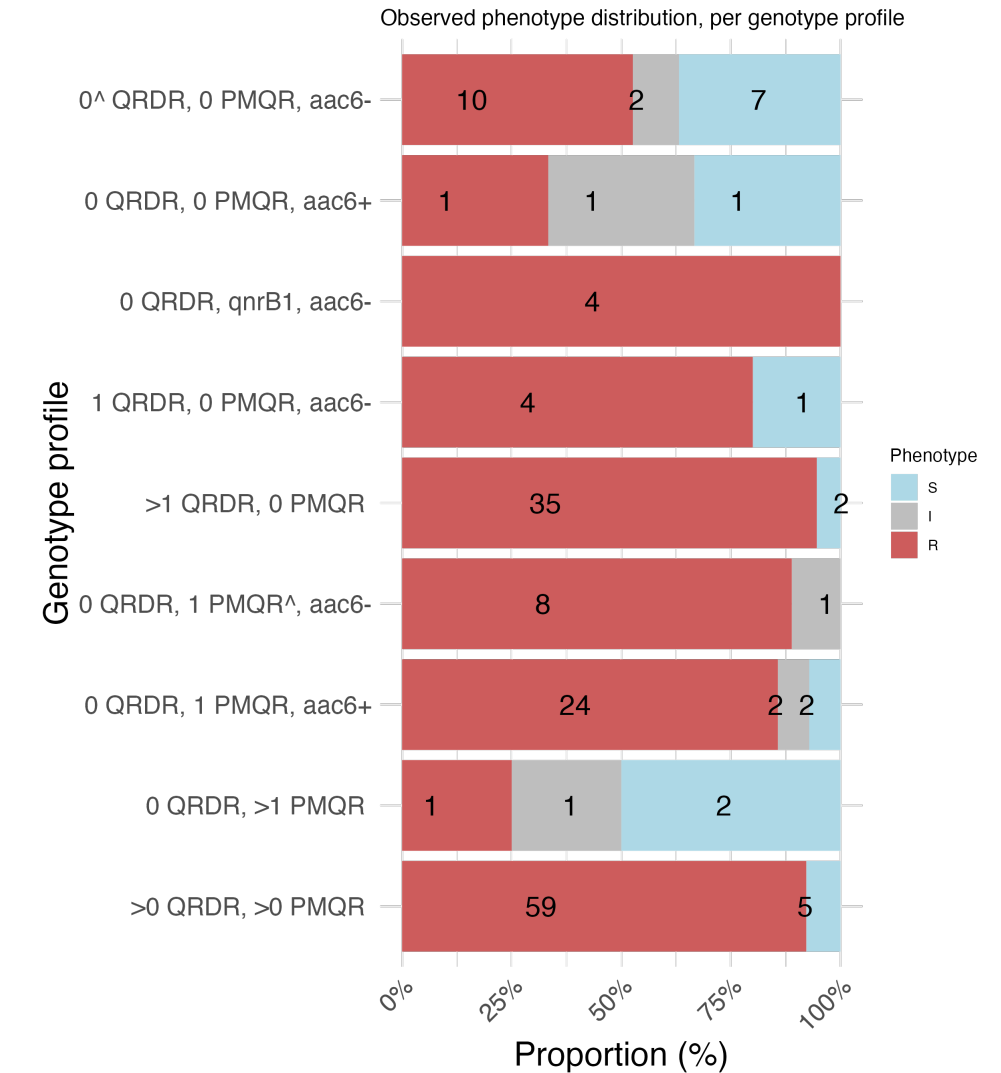
ix) E (EUCAST) /  
HURYC & CIBERINFEC (n=173) – MIC distribution vs.  
genotype profile



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

