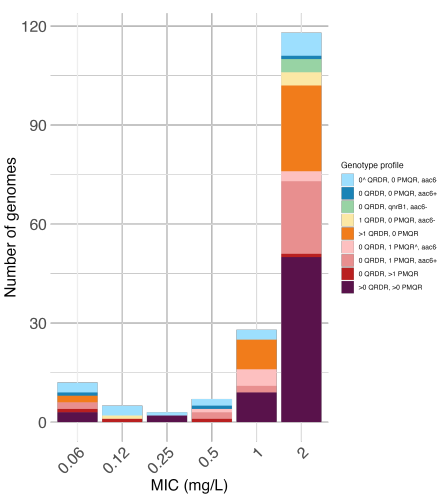
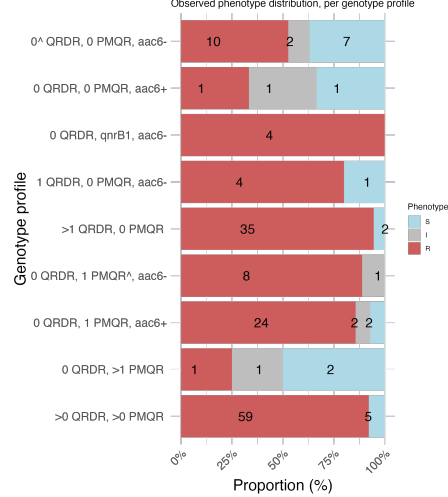


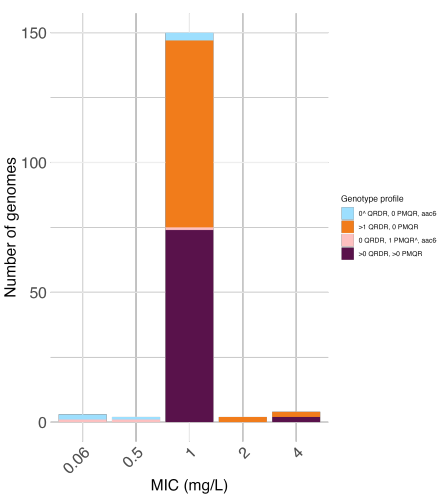
Q) HURYC & CIBERINFEC (n=173) – MIC distribution vs. genotype profile



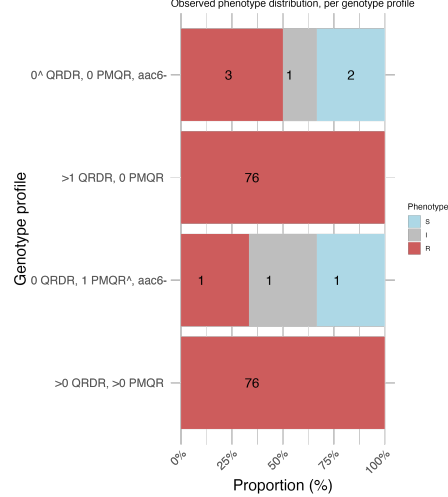
R) HURYC & CIBERINFEC (n=173) – observed phenotype vs. genotype profile



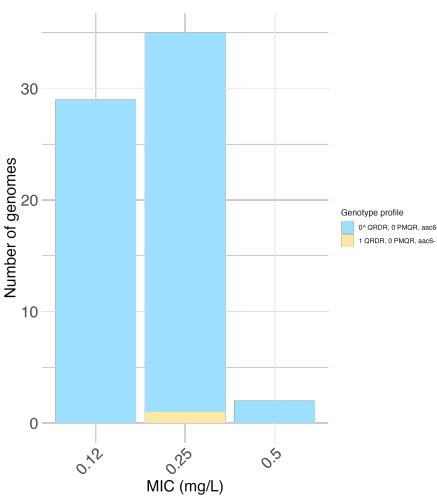
S) IZSAM (n=161) – MIC distribution vs. genotype profile



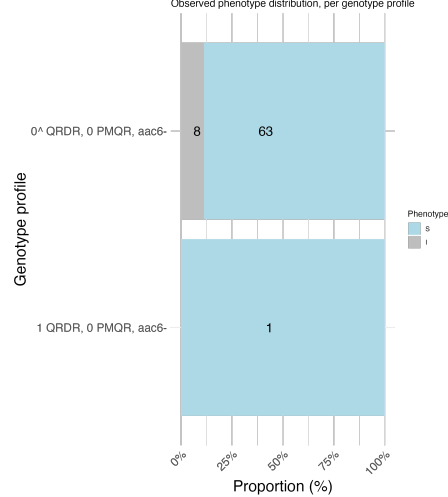
T) IZSAM (n=161) – observed phenotype vs. genotype profile



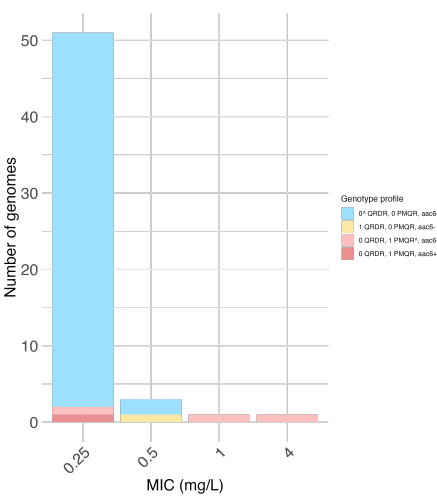
U) Oxfordshire* (n=72) – MIC distribution vs. genotype profile



V) Oxfordshire* (n=72) – observed phenotype vs. genotype profile



W) SCHARKI (n=56) – MIC distribution vs. genotype profile



X) SCHARKI (n=56) – observed phenotype vs. genotype profile

