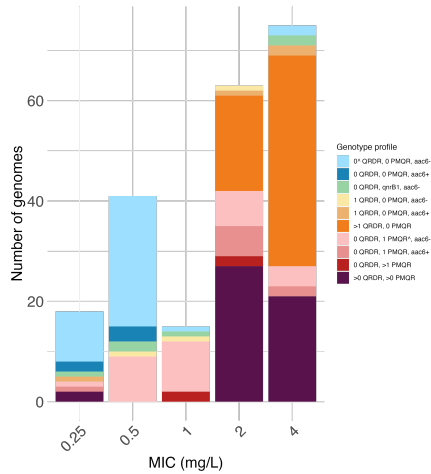
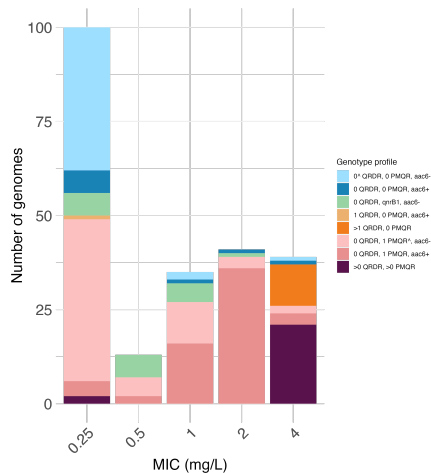


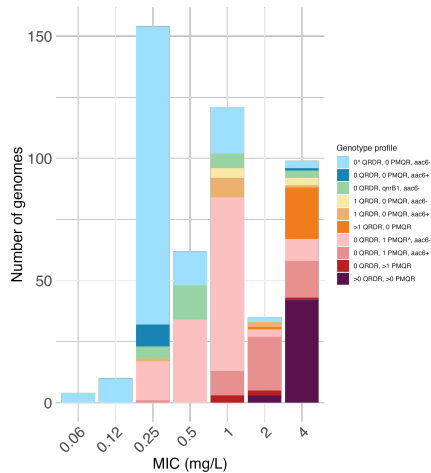
A) HowdenGorrie (N=212) – MIC distribution vs. genotype profile



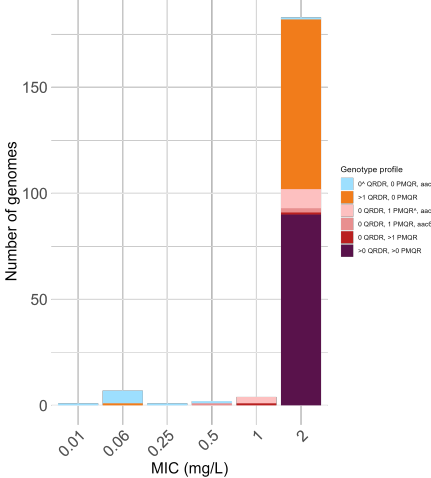
C) MBIRA (N=228) – MIC distribution vs. genotype profile



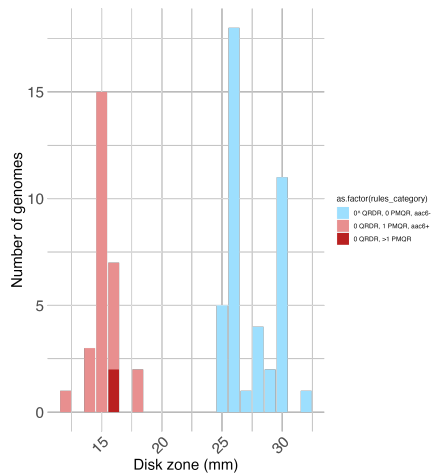
E) AGAR (N=198) – MIC distribution vs. genotype profile



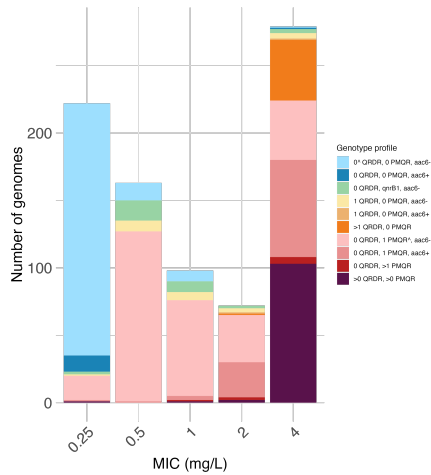
G) Milan SanRafaele (N=198) – MIC distribution vs. genotype profile



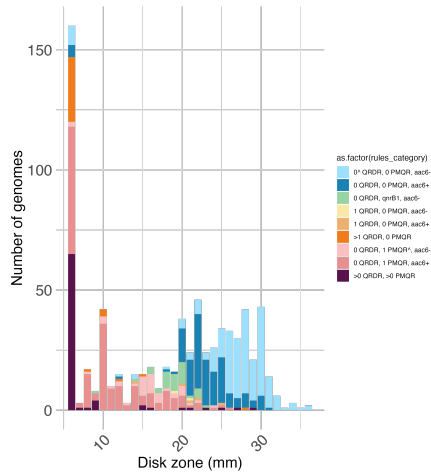
I) Hailu Ethiopia (N=70) – DD distribution vs. genotype profile



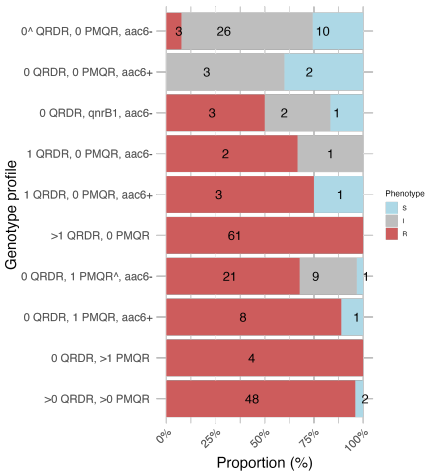
K) Kayama Japan (N=834) – MIC distribution vs. genotype profile



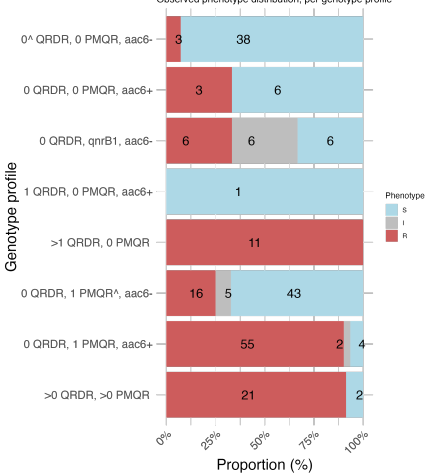
M) Feasey Malawi (N=737) – MIC distribution vs. genotype profile



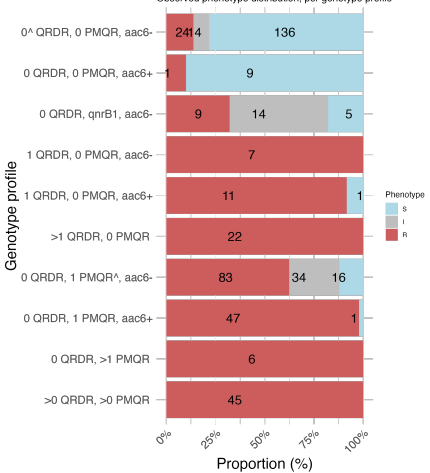
B) HowdenGorrie (N=212) – observed phenotype vs. genotype profile



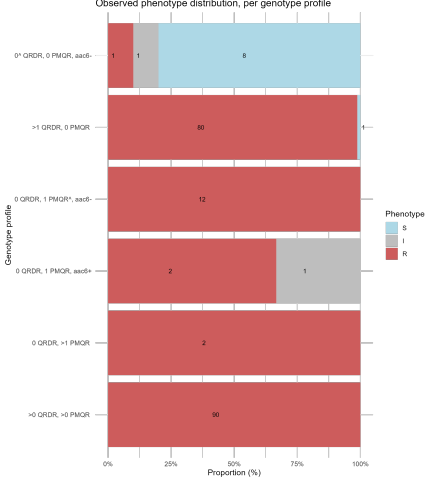
D) MBIRA (N=228) – observed phenotype vs. genotype profile



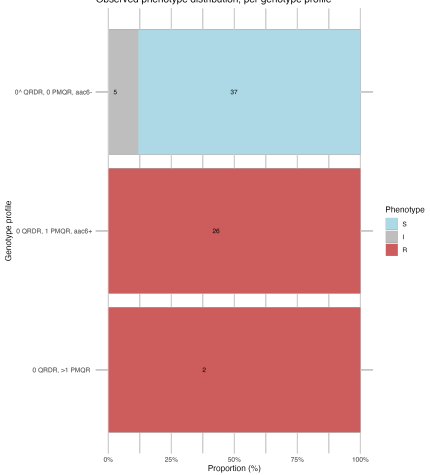
F) AGAR (N=485) – observed phenotype vs. genotype profile



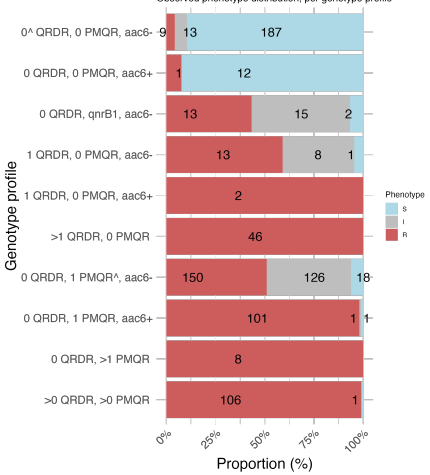
H) Milan SanRafaele (N=198) – observed phenotype vs. genotype profile



J) Hailu Ethiopia (N=70) – observed phenotype vs. genotype profile



L) Kayama Japan (N=834) – observed phenotype vs. genotype profile



N) Feasey Malawi (N=737) – observed phenotype vs. genotype profile

