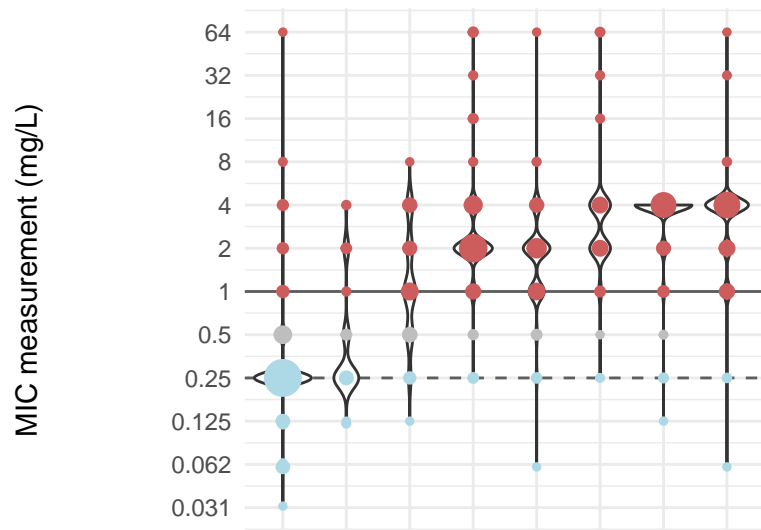
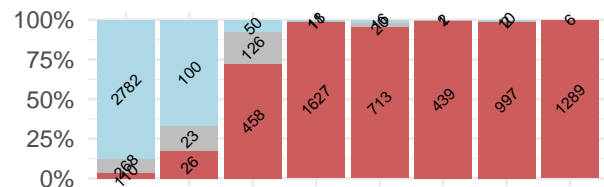


a) MIC data



Percent of genomes

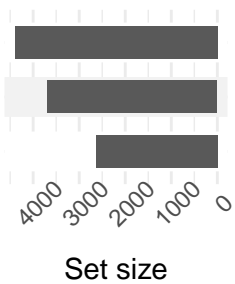


QRDR mutations

PMQR genes

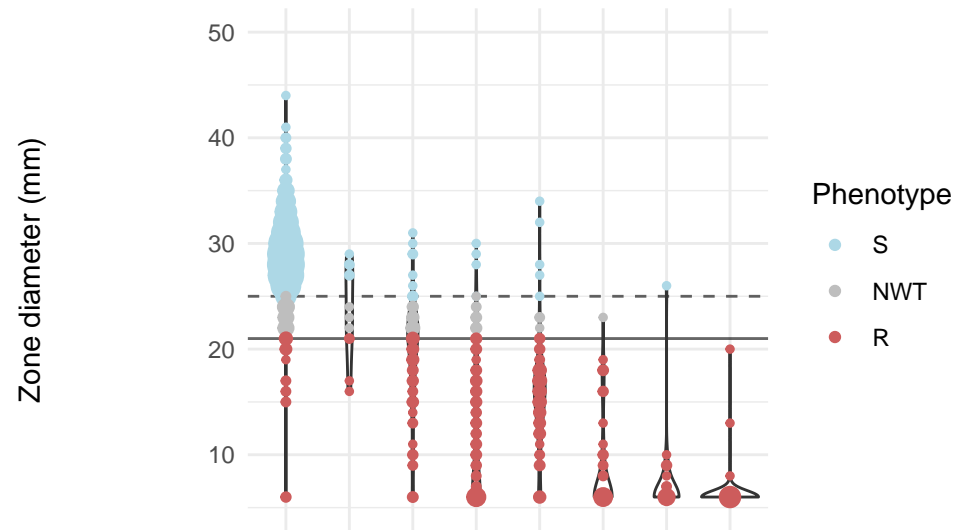
aac(6')-Ib-cr

group

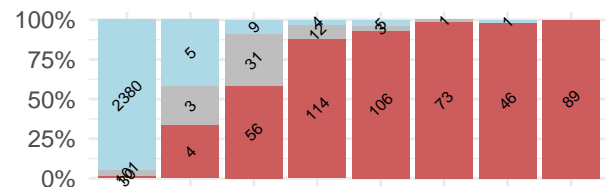


Set size

b) Disk diffusion data



Percent of genomes

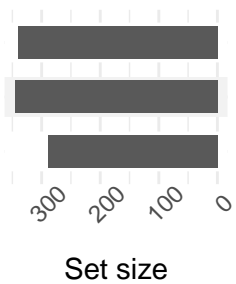


QRDR mutations

PMQR genes

aac(6')-Ib-cr

group



Set size

Phenotype

S

NWT

R