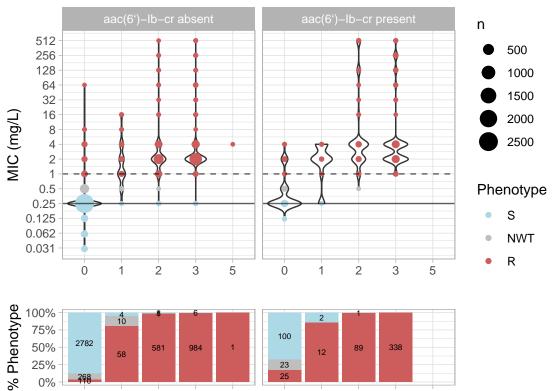
a) No. QRDR mutations vs phenotype (in absence of PMQR genes)



b) No. PMQR genes vs phenotype

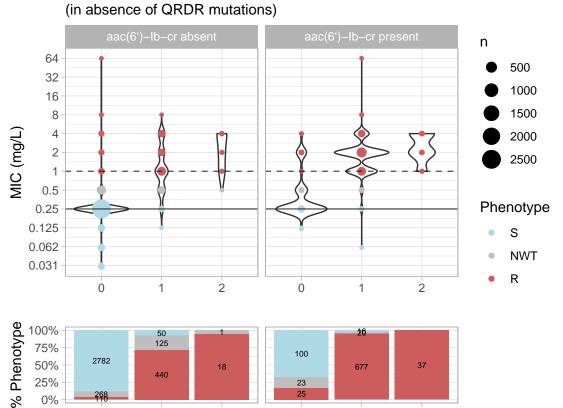
i

50%

25%

0%

0%



Number of PMQR genes

Number of QRDR mutations