

Number of genomes

150

100

50

0

10

20

30

Disk zone (mm)

as.factor(rules\_category)

- 0^ QRDR, 0 PMQR, aac6-
- 0 QRDR, 0 PMQR, aac6+
- 0 QRDR, qnrB1, aac6-
- 1 QRDR, 0 PMQR, aac6-
- 1 QRDR, 0 PMQR, aac6+
- >1 QRDR, 0 PMQR
- 0 QRDR, 1 PMQR^, aac6-
- 0 QRDR, 1 PMQR, aac6+
- >0 QRDR, >0 PMQR