

Number of genomes

20

15

10

5

0

10

20

30

40

Disk zone (mm)

as.factor(rules\_category)

- 0<sup>^</sup> QRDR, 0 PMQR, aac6-
- 0 QRDR, 0 PMQR, aac6+
- 0 QRDR, qnrB1, aac6-
- >1 QRDR, 0 PMQR
- 0 QRDR, 1 PMQR<sup>^</sup>, aac6-
- 0 QRDR, 1 PMQR, aac6+
- >0 QRDR, >0 PMQR

