

Number of genomes

15

10

5

0

15

20

25

30

Disk zone (mm)

as.factor(rules_category)

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

