

Significance

- Non-significant
- Adjusted p-value ≤ 0.05

log₁₀ (clade-level reads)

- **o** 1.0 O 1.5
- O 2.0
- O 2.5
- O 3.0 O 3.5

Ratio between unique minimisers found in sample and total clade–level minimisers in database 1.00

- 0.75
- 0.50
- 0.25
- 0.00