

Significance

Non-significant

• Adjusted p-value ≤ 0.05

log₁₀ (clade-level reads)

O 1.0 O 1.5

O 2.0 O 2.5

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

0.75

0.50

0.25

0.00