

Model

$\log(\sigma) = \log(\text{fragment size}) + (\log(\text{fragment size}) | \rho | \text{Study})$

$\log(\sigma) = 1 + (1 | \rho | \text{Study})$

$\log(\sigma) = \log(\text{fragment size}) + (\log(\text{fragment size}) | \text{Study})$

$\log(\sigma) = 1 + (1 | \text{Study})$

homoscedastic σ

0.000

0.025

0.050

0.075

Standardized species richness ~ fragment size slope (average)

