Spatially standardised generic richness (mean +/− 1 SD)
Classical rarefaction (sample size = 20)
100km grid cells, no covariate data required

Legend:
— Bivalves
— Brachiopods

Y-axis: Generic richness (0, 5, 10, 15, 20)

X-axis: Time (mya) — 500, 400, 300, 200, 100, 0

Timeline: Cm, O, S, D, C, P, Tr, J, K, Pg, Ng