R code of the neutral community model (NCM) is available at <https://github.com/Weidong-Chen-Microbial-Ecology/Stochastic-assembly-of-rivermicroeukaryotes> (Chen et al., 2019).

R code of PER-SIMPER and Dispersal Niche Continuum Index (DNCI) is available at <https://github.com/Corentin-Gibert-Paleontology/DNCImper> (Vilmi et al., 2021).