

Standard resemblance choices

A detailed discussion of the competing properties of different resemblance matrices is outside this manual's scope (see [L&L](#), CiMC Chapters 2 & 16, or Clarke KR, Somerfield PJ, Chapman MG 2006, *J Exp Mar Biol Ecol* 330: 55-80). Novice users are recommended to take one of the main options (the defaults): Bray-Curtis similarity for biological assemblage data; Euclidean distance (having first normalised) for physico-chemical, biomarker or morphometric data etc., in which variables are not on comparable ranges or the same measurement scale at all; and (non-normalised) Euclidean distance for body- and particle-size histograms (first standardised), growth curves etc.

Revision #3

Created 30 May 2024 00:17:13 by Arden

Updated 15 January 2025 00:59:24 by Abby Miller