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.

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