Resemblance calculation

Resemblance is the general term in PRIMER used to cover (dis)similarity or distance coefficients. The next stage in both the Fal nematode and environment runs of the **Basic multivariate analysis** wizard was to create an appropriate triangular resemblance matrix between all pairs of samples. This is a run of **Analyse>Resemblance** on the pre-treated (transformed or normalised) worksheet. Relevant defaults will be suggested, given the Data type, i.e. (Measure•Bray-Curtis) for biota and (Measure•Euclidean distance) for environmental variables, and (Analyse between•Samples) in both cases. There are, however, nearly 50 other possible choices on this dialog, see Section 5.

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