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Cover figure: Dice coefficient-based clustering of the TGGE fingerprints of the full-scale membrane bioreactor (MBR) during the spring, summer and autumn of 2009. A: aerated bioreactor; X: anoxic bioreactor; M: internal markers (A2). The dotted lines mark the clustering similarity level of internal markers of each clustering analyses, indicating the limit to consider significant differences between samples. The scale bar indicates the percentage of similarity and the numbers in nodes represent the cophenetic correlation coefficient values. See the article "Exploring the links between population dynamics of total and active bacteria and the variables influencing a full-scale membrane bioreactor (MBR)” by C. Gómez-Silván et al.