

Microbial Diversity and Dynamics in the mother of Vinegar Revealed by Metagenomics

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The production of artisanal vinegars can be carried out using controlled methods, such as submerged cultures in bioreactors, or spontaneous methods, like Orleans system¹. In the latter, the fermentable substrates such as wine, juice, or a molasses solution are added, and upon contact with the liquid in the bioreactor, the fermentation process begins. During this process, Acetic Acid Bacteria (AAB) are responsible for the biotransformation of ethanol into acetic acid², forming a biofilm known as the "mother of vinegar," which acts as a starter culture and a microenvironment for other AAB, Lactic Acid Bacteria, and yeasts.

METHODOLOGY. To observe the microbial diversity and shifts in the population throughout the fermentation process, population dynamics were evaluated by counting viable cultivable microorganisms on days 0 and 7 of fermentation, using MRS and standard agar media with the pour plate technique. In addition, DNA was extracted from the mother of vinegar on the same fermentation days, and amplification of the 16s and 18s ribosomal genes was carried on for metagenomic analysis.

CONCLUSION. Given that microorganisms involved in fermentation often require specific nutritional conditions and produce a variety of metabolites, traditional media are of limited use. In this context, metagenomics proves to be an effective tool for studying microbial diversity and population dynamics in fermented products.

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