

Special Issue

Microbial Diversity under Forage Silage Fermentation

Message from the Guest Editors

Ensiling is a common way to preserve the moist forage crop. During ensiling process, various kinds of microorganisms are involved in silage fermentation, and their structure and diversity greatly influenced the end-product quality. The interactions and selection between microorganisms produce specific metabolic patterns during ensiling. Therefore, study on microbial community structure, function and succession process is vital to understanding silage fermentation. Nowadays, the rapid development of metagenomics, microbial culturomics and transcriptomics techniques has allowed us to delve into microbial diversity, contributing to give a completeness view of culturable and nonculturable silage-associated bacteria. Therefore, a deeper insight into the microbial diversity during ensiling process remains a challenging subject for improving silage quality and develop new silage additives. This Special Issue is seeking contributions including comprehensive reviews and experimental research about the microbial diversity of the fermentation process and interactions amongst microbial species/groups or microbial ecology of forage silages.

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Diversity (ISSN 1424-2818) is a scholarly journal that covers all areas of diversity research. Our distinguished editorial board and refereeing process ensures the highest degree of scientific rigor for publishing. Original research articles and timely reviews are released online, with unlimited free access.

We invite papers and reviews on multidisciplinary topics of diversity that bridge organismic diversity (systematics, biodiversity, phylogeny, population genetics, and evolution) and molecular diversity (phytochemistry and biophysics).

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