



Microbial Environmental Genomics (MEG) (Methods in Molecular Biology)

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This volume seeks to understand how organisms and gene functions are influenced by environmental cues while accounting for variation that takes place within and among environmental populations and communities. *Microbial Environmental Genomics (MEG)* guides readers through methods to analyse the diversity of different organism types (archaea, bacteria, fungi, protists and microfauna), interactions between fungi and trees, and methods to identify and characterize functions and functional diversity of both pro- and eukaryotes. Written for the *Methods in Molecular Biology* series, chapters include introductions to their respective topics, lists of the necessary materials and reagents, step-by-step, readily reproducible laboratory protocols, and tips on troubleshooting and avoiding known pitfalls.

Authoritative and practical, *Microbial Environmental Genomics (MEG)* will serve as a primary research reference for researchers and research managers in environmental microbiology working in the expanding field of molecular ecology and environmental genomics.

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