

Abstract

Microbial-explicit models of soil carbon (C) and nitrogen (N) cycling have improved upon simulations of C and N stocks and flows at site-to-global scales relative to traditional first-order linear models. However, before we can draw conclusions from microbialexplicit models about the future behavior of soils in a changing world, we need to thoroughly investigate model behavior with existing data and understand the impact of model development decisions on predictive outcomes. We used the MIcrobial-MIneral Carbon Stabilization Model with coupled N cycling (MIMICS-CN) to explore several ways of interrogating a model with data. We simulated C and N losses from litterbags in the Long-term Inter-site Decomposition Experiment (LIDET) while simultaneously comparing simulated values of soil pools and fluxes against ranges from a continent-wide data synthesis. We also discuss the impact of the way soil experiments are interpreted in the context of models, the importance of evaluating both equilibrium and transient model behavior, and the impact of assigning temperature sensitivity to 3 different aspects of microbial physiology.

Model formulation

MIMICS-CN represents C and N flow through metabolic and structural litter, oligotrophic and copiotrophic microbes, and physically protected, chemically protected, and available SOM pools. C dynamics are driven by reverse Michaelis-Menten kinetics, while N dynamics are driven by input and microbial C:N. N leaves the model as leaked inorganic N and C leaves the model as respired CO_2 .



Figure 1. MIMICS-CN model structure and stoichiometric parameters unique to the coupled C-N model. CNs = C:N of structural litter, CNm = C:N of metabolic litter, CNr = C:N of copiotrophs, CNk = C:N of oligotrophs, and NUE = nitrogen use efficiency of both microbial groups.



Figure 2. LIDET sites included in model simulations. Map borrowed from: Harmon, M. E. et al. (2009). Long-term patterns of mass loss during the decomposition of leaf and fine root litter: an intersite comparison. Global Change Biology, 15: 1320–1338.

Different mechanisms of soil microbial response to global change result in different outcomes in the MIMICS-CN model

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model was spun up to equilibrium at Harvard Forest and run for 15

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