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► To cite this version:

Nicolas Fanin, D Moorhead, Isabelle Bertrand. Eco-enzymatic stoichiometry and enzymatic vectors in decaying litter reveal differential C, N, P dynamics along a land-use gradient. 5. International EcoSummit: Ecological Sustainability: Engineering Change, Aug 2016, Montpellier, France. 2016. hal-02740564

HAL Id: hal-02740564

<https://hal.inrae.fr/hal-02740564>

Submitted on 2 Jun 2020

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Eco-enzymatic stoichiometry and enzymatic vectors in decaying litter reveal differential C,N,P dynamic along a land-use gradient

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Introduction

At a global scale, eco-enzymatic ratios were shown to provide a functional measure of the threshold at which control of community metabolism shifts from nutrient to carbon limitation¹. However at a finer scale, the controls of nutrient availability on microbial community structure and associated enzyme production are still poorly understood. Microbial decomposer communities cope with substrates that vary considerably in C:N:P stoichiometry compared to that of their own biomass. Changes in relative abundances of extracellular enzymatic activities (EEA) involved in C, N, and P cycling should reflect relative resource acquisition by these communities.

Objectives

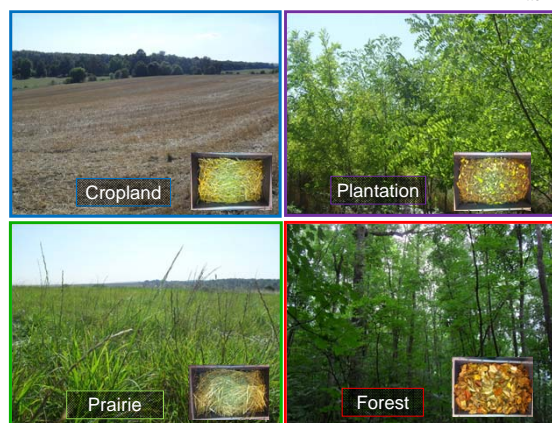
Evaluate the eco-enzymatic theory using a land-use gradient under controlled conditions.

Use length and angle of enzyme vectors² to explore relationships between EEA and mineralization rates, microbial metabolism and community structure.

Material & Methods

Full reciprocal transplant experiment

Litter and soil samples were collected in a one-hectare plot with a common soil substratum



4 litter species
(*Fagus sylvatica*, *Triticum aestivum*, *Robinia pseudoacacia*, *Festuca arundinacea*)

x

4 soils
(Forest, Culture, Plantation, Grassland)

x

3 sampling dates
(27, 97 and 202 days)

x

3 replicates

144 microcosms

(20°C, 60% WHC, 10 g C per g of soil)

Dynamic measure of:

- C Respiration
- Mineral N and Olsen P
- Microbial community structure (PLFAs)
- Enzymatic activities



Enzyme C (β -glu, α -glu, CBH, xylosidase)
Enzyme N and P (NAG, LAP, alkaline phosphatase)

Results

Vector length = $\sqrt{x^2 + y^2}$ x = proportion of C versus P acquiring enzyme activities; y = proportion of C versus N acquiring activities^{2,3}.
Angle = $\text{Atan2}(x, y)$

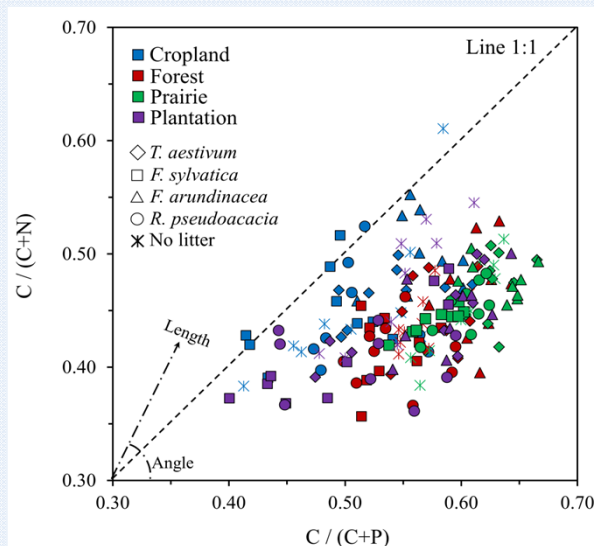


Figure 1: Eco-enzyme stoichiometry of the relative proportions of C to N acquisition versus C to P acquisition. C acquisition is represented by the sum of β -1,4-glucosidase (BG), β -D-cellobiosidase (CBH), β -xylosidase (XYL), and α -1,4-glucosidase (AG); N acquisition is represented by the sum of β -1,4-N-acetylglucosaminidase (NAG) and leucine aminopeptidase (LAP); and P acquisition is represented by alkaline phosphatase (AP) activity

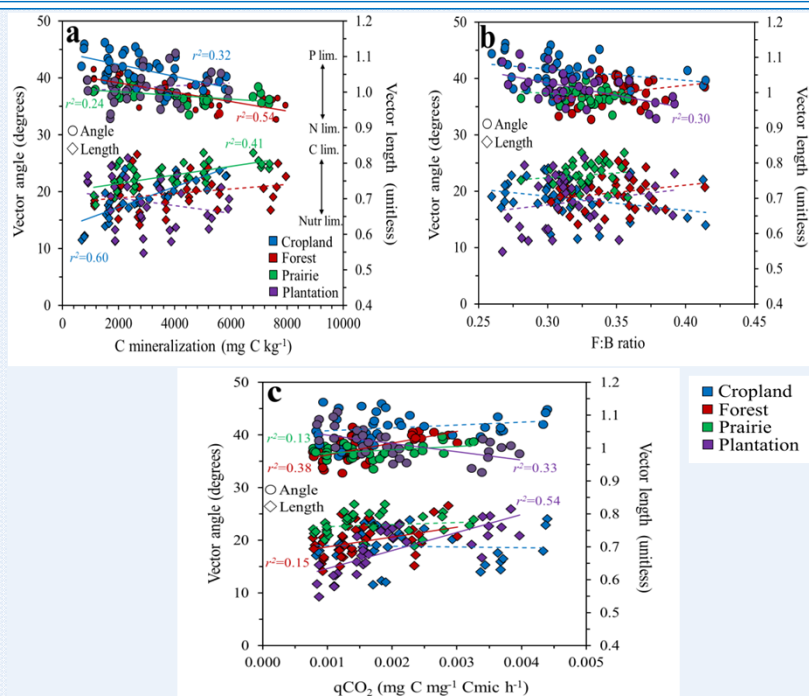


Figure 2: Relationships between vector lengths and angles with a) carbon mineralized, b) Fungi:Bacteria ratio, and c) metabolic quotient according to vector angle (circle) or length (diamond) (litter types combined). Full lines represent significant relationships and associated r^2 values.

Conclusions

- EEA showed that N requirements increased relative to P during litter decay but C requirements increased more rapidly than either N or P in most of these ecosystems.
- Shifts in EEA were related to changes in metabolic quotient (C respired per unit biomass) but not in fungi:bacteria ratios. Functional abilities of soil microbes may be more important than their identity for assessing their resource requirements.
- The use of EEA as a proxy of microbial resource demand improved our understanding of temporal shifts in resource requirements to microbial communities.

References

¹Sinsabaugh R.L., Hill B.H., Follstad J.J. (2009) Eco-enzymatic stoichiometry of microbial organic nutrient acquisition in soil and sediment. *Nature* 462: 795-798.

²Moorhead D.L., Sinsabaugh R.L., Hill B.H., Weintraub M.N. (2016) Vector analysis of ecoenzyme activities reveal constraints on coupled C, N and P dynamics. *Soil Biology & Biochemistry* 93: 1-7

This work is published in:

Fanin N., Moorhead D.L., Bertrand I. (2016) Eco-enzymatic stoichiometry and enzymatic vectors reveal differential C,N,P dynamics in decaying litter along a land-use gradient. *Biogeochemistry*, 129: 21-36