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Soil microbial biomass, activity, and community structure as affected by mineral P fertilization in two grassland soils Yichao SHI^{*1}, Noura ZIADI¹, Roger LALANDE¹, Chantal HAMEL², Julie LAJEUNESSE³, Jean LAFOND³, Gilles BÉLANGER¹

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Introduction

- Soil microbiota is a major driver of soil formation, nutrient cycling, and organic matter turnover.
- Reports on how phosphorus (P) fertilization affect



Results and discussion

microbial properties in arable soils are contradictory. Zhong and Cai (2007) report that mineral P fertilizer increased microbial biomass and diversity, while others found no significant effect on the composition of soil microbial communities (Hamel et al. 2006; Shi et al. 2012, 2013).

Soil microorganisms response to mineral P fertilization in grasslands is poorly understood.

Objective

To assess the effect of mineral P fertilization on soil microbial biomass, activity, and community structure in timothy-based grasslands of eastern Canada.

Materials and Methods

Site description

Two sites in timothy (*Phleum pratense* L.) were seeded in

Figure 2. Effect of P fertilization on soil dehydrogenase and alkaline phosphamonoesterase activity.

2009.

---- a Kamouraska clay at Lévis; ---- a Labarre clay loam at Normandin.

Experimental design

✤Three P rates (0, 20, and 40 kg P ha⁻¹; P0, P20, P40, respectively) were applied in the spring of each year starting in 2010 with three replications.

Soil sampling and measurements

✤In 2013, soils were sampled to a depth of 10 cm in June, Aug.–Sept., and Oct. at both sites.

Soil microbial biomass C (SMB-C), N (SMB-N) and P (SMB-P), dehydrogenase and alkaline phosphamonoesterase (Alk-PO₄), and phospholipids fatty acids (PLFA) were analyzed as in Shi et al. (2012).

Soil chemical properties (pH, total C and N, Mehlich-3 P) were determined (*Table 1*).

Statistical analysis

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Data were analyzed with the MIXED procedure of SAS.
The relative abundance of PLFA was analyzed using principal component analysis (PCA) and MANOVA.

Figure 1. Effect of P fertilization on soil microbial biomass C (SMB-C), N (SMB-N), and P (SMB-P).



Figure 3. Ordination plots of the microbial community structure (based on phospholipids fatty acids) as determined by PCA.

P fertilization had no significant effect on SMB-C and SMB-N (*Fig. 1*), and on dehydrogenase and Alk-PO₄ at both sites (*Fig. 2*).

SMB-P was greater with P40 than with P0 and P20 at Normandin, but was not affected by P fertilization at Lévis (*Fig. 1*). The same trend was also observed in soil Mehlich-3 P (*Table 1*).

The soil microbial community structure was significantly influenced by sampling dates, but not affected by P fertilization (*Fig. 3*), as previously reported by Shi et al. (2013) in a long-term corn-soybean rotation.

Table 1. Effect of P fertilization on soil pH, total C and N, and Mehlich-3 P.

	Lévis				Normandin			
Treatments	рН	Total C	Total N	Mehlich-3 P	рН	Total C	Total N	Mehlich-3 P
	-	g	kg ⁻¹	mg kg ⁻¹	-	g k	g^{-1}	mg kg ⁻¹
P0	6.0a	44.1a	3.5a	11.2a	5.1a	23.3a	1.9a	16.8b
D2 0	$6 0 \circ$	11 50	260	10.2	5 20	$\gamma \gamma \gamma_{\alpha}$	1 0 0	20.01





One year of data suggests that four years of P fertilization has

limited effects on the composition and function of the soil microbial community of timothy-based grasslands in eastern Canada.

 P20
 6.0a
 44.5a
 3.6a
 19.2a
 5.2a
 23.2a
 1.8a
 28.8b

 P40
 6.0a
 44.0a
 3.4a
 28.8a
 5.2a
 23.2a
 1.8a
 28.8b

*Means followed by different letters in each column are significant different ($\alpha < 0.05$).



Hamel et al. 2006. Soil Biol. Biochem. 38:2104–2116. Shi et al.
2012. Appl. Soil Ecol. 62:14–23. Shi et al. 2013. Biol. Fertil. Soils.
49: 803–818. Zhong and Cai. 2007. Appl. Soil Ecol. 36: 84–91.

