Effects of Enhanced-Efficiency Nitrogen Fertilizers on Soil Microbial Biomass and Respiration in Tropical Soil Under Upland Rice Cultivation

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Abstract

While over-use of N fertilizers can suppress microbial biomass, application of urease inhibitors is known to be a potential way to rebuilt microbial diversity and improve soil functions. However, the hypothesis of this study is that the application of N fertilizers regardless of the source would increase soil microbial biomass and reduce soil respiration. A two-year field experiment was conducted to assess the effects of enhanced-efficiency N sources on soil microbial biomass, and soil respiration. The experiment was set up in a randomized block design in a $3 \times 4 + 1$ factorial scheme, with four replicates. Treatments comprised three sources (conventional uncoated urea, NBPT (N-(n-butyl) thiophosphoric triamide)-treated urea, and polymer-coated urea) and four rates (30, 60, 90 and 120 kg ha⁻¹) of N, in addition to a control treatment (no fertilizer application). Microbial biomass C (MBC) and microbial biomass N (MBN), and soil respiration (C-CO₂ and *q*CO₂) were determined in upland rice rhizosphere in each crop season. Responses of soil microbial properties to N fertilization were dependent on the N rates, but no significant effect of the N sources was observed. All measured parameters, except MBC in the first season and C-CO₂ in the second season, were increased with increasing N rates. However, the application of N higher than 60 kg ha⁻¹ suppressed soil microbial biomass, as well as soil respiration. Therefore, the lack of response by added urease inhibitors to the N sources indicate that optimizing N rates for upland rice production is a far more effective option for improving soil microbial community than using enhanced-efficiency N sources.

Keywords: soil microbial activity, fertilizer efficiency, N rate

1. Introduction

Nitrogen (N) is one of the most important elements in natural and agricultural ecosystems (Krivtsov et al., 2011). Increased N input has been a major contributor to higher crop yields over the last decades. In fact, the N availability directly influence the growth and abundance of microorganisms, which play a critical role in nutrient cycling and maintenance of soil structure for improved plant production. However, the N addition to agricultural fields can negatively affect microbial communities and restrict crop productivity depending on many factors, such as fertilizer type, N rate, soil pH, soil texture, and the crop (Lupwayi et al., 2012).

Previous studies reported that N fertilizers can have no effect or act either positively or negatively on soil microorganisms (Lupwayi et al., 2012; Yu et al., 2018). Li et al. (2013) found that microbial activity decreased in soils with increasing NH_4^+ -N concentration under application of ammonium sulfate and urea, suggesting that microbial functional diversity can occur with a N gradient. Zhalnina et al. (2015) reported that the application of ammonium-based fertilizers decreased soil pH, thus reduced microbial abundance because some species cannot tolerate acidic environment. In contrast, Liu et al. (2020) found that the content of microbial biomass C (MBC) and N (MBN) in the rhizosphere soil were increased following fertilizer applications.

Besides the contradictory influence of N fertilizers on the soil microbial biomass, the use of inorganic N fertilizers is essential in agricultural systems. To ensure that the addition of N fertilizer is beneficial to both crop production and soil microorganisms, effort has been put into finding ways to improve fertilizer N use efficiency and minimizing environmental impacts. Recommendations to enhance N use efficiency include the use of formulated forms of fertilizer containing urease and nitrification inhibitors to reduce NH_3 volatilization from urea hydrolysis (Engel et al., 2019). The application of urease inhibitors can reduce volatilization by temporarily preventing urease from breaking down urea, thereby leaving more NH_3/NH_4^+ available for plants to take up (Chien et al., 2008; Gioacchini et al., 2002).

The effect of urease inhibitors on urea hydrolysis and enhanced N use efficiency has been widely investigated. However, the impact of different N fertilizers containing urease inhibitors on soil microbial properties is still an open question. A two-yr field experiment was conducted to assess the effects of the widely used urease inhibitor, N-(n-butyl) thiophosphoric triamide (NBPT) and polymer-coated urea (PCU) on soil microbial biomass, and soil respiration. The hypothesis of this study is that the application of N fertilizers regardless of the source would increase soil microbial biomass and reduce soil respiration.

2. Materials and Methods

2.1 Experimental Site

The field experiment was carried out in two growing seasons (2013/14 and 2014/15) in an area (16°35'50"S; 49°16'40"W; 735 m a.s.l.) located in the Agronomy College at the Federal University of Goiás, State of Goiás, Brazil. The local climate is classified as Aw (seasonal tropical savanna), with a humid season from October to April and a dry one from May to September according to the Köppen classification. The average annual precipitation is 1500 mm, and the mean annual temperature is around 22.5 °C.

The soil was classified as typic dystrophic Red Latosol (LVd) in the Brazilian Soil Classification System (Santos et al., 2013), which corresponded to an Oxisol in the US Soil Taxonomy System [11]. Soil samples for characterization were collected from the 0- to 20-cm depth layer before the N fertilizer rates were applied using an auger. Prior to characterization, the samples were air dried and sieved through a 2-mm mesh and then analyzed following methodologies as proposed by Embrapa (1997, 2009). Some selected soil chemical properties and particle size distribution are given in Table 1.

Chemica	al analysis								
$\mathrm{pH}_{\mathrm{Ca}}$	Ca ²⁺	Mg ²⁺	K^+	Al ³⁺	H+A1	CEC	V	Р	OM
		cmol _c dm ⁻³					%	mg dm ⁻³	g dm ⁻³
5.58	2.15	1.05	0.312	0	.63	6.14	57.46	4.88	17.5
Textural	analysis								
Sand			S	ilt			Clay		
					g kg ⁻¹				
450			1	10			440		

Table 1. Selected chemical properties and particle size distribution of the soil at the experimental site

Note. pH_{Ca} : pH measured in 0.01 mol L⁻¹ CaCl₂(1:2.5); Ca²⁺, Mg²⁺, and Al³⁺ extracted by 1 mol L⁻¹ KCl; P and K extracted by 0.05 mol L⁻¹ HCl + 0125 mol L⁻¹ H₂SO₄ (Mehlich-1 extractor); H+Al extracted by 0.5 mol L⁻¹ calcium acetate buffered at pH 7; CEC: cation exchange capacity; V: base saturation; OM: organic matter, estimated from the organic carbon extracted by the Walkley-Black method. Textual Analysis conducted using the pipette method.

2.2 Experimental Design and Treatments

The experiment was set up in a randomized block design in a $3 \times 4 + 1$ factorial arrangement, with four replicates. Treatments comprised three sources (conventional uncoated urea, NBPT-treated urea, and PCU) and four rates (30, 60, 90 and 120 kg ha⁻¹) of N, in addition to a control treatment (no fertilizer application).

2.3 Field Experiment

Field was ploughed at 20 cm depth prior to seeding. Plots consisted of four 5 meters long rows, spaced 0.5 m apart, using the rice cultivar BRS Esmeralda, which has a moderate resistance to major diseases and a certain tolerance to water stress (Castro et al., 2014). Additionally, BRS Esmeralda is a relatively recent upland rice

cultivar developed by the breeding program coordinated by the Brazilian Corporation for Agricultural Research (Embrapa). The higher performance of BRS Esmeralda compared to other current cultivars is due to its high grain quality, good drought tolerance, high disease resistance and lodging resistance (Colombari et al., 2013). Further, this cultivar has a great stability and adaptability to a large range of soils, climates, and crop management on the Cerrado region, which may lead to a satisfactory yield performance in this study.

The net plot area was composed of the two central rows, considering the lateral rows as borders. The soil was prepared in both years by one plowing and one disk harrow leveling. The seeds of rice were sown manually 20 cm spaced apart in rows, with two or three seeds per hole. Nitrogen fertilizers were applied manually in two split doses: 50% at the seedling stage, and 50% at the tillering stage (~80 d after sowing). All treatments received 400 kg ha⁻¹ of the formula 00-20-20 as a basal fertilizer to supply phosphorus and potassium. Weed management consisted of hand weeding plots two times during the growing season. Rice was harvested in every growing season at the end of maturing stage (between 103 and 108 d after sowing).

2.3 Soil Sampling

At flowering stage, soil samples were collected from five points randomly at the 0-10 cm depth between the rows in each experimental plot at three different points to form a composite sample per replicate. The samples were homogenized using a 2 mm mesh sieve and prepared for microbial analysis. All samples were immediately stored in sealed plastic bags in a cooler and transported to laboratory and stored at 4 °C. All microbiological determinations were conducted within 3 d of sampling.

2.4 Microbial Biomass Measurements

Prior to microbial biomass measurements, the soil moisture was adjusted to about 80% water holding capacity. The chloroform fumigation extraction and measurements of soil microbial biomass C (MBC) and N (MBN) in fumigated and non-fumigated soil extracts were performed following Vance et al. (1987). Samples of 20 g of dry weight equivalent of soil was fumigated with ethanol-free chloroform (CHCl₃) at 25 °C for 24 h. Both fumigated and non-fumigated soils were separately extracted with 80 mL of 0.5 mol $L^{-1} K_2 SO_4$ by shaking for 40 min on a reciprocating shaker at 180 cycles per minute and then filtered using slow quantitative filter paper (Whatman No. 42).

An aliquot of the filtered extract (8 mL) was refluxed with 0.5 mol L^{-1} K₂Cr₂O₇ (2 mL), and 15 mL of a mixture containing H₂SO₄ and H₃PO₄ (2:1 v/v) in glass flasks. The mixture was burned at 100 °C for 30 min. Cold blank was not heated. Samples were then cooled and diluted to 50 mL with deionized water. The residual dichromate was measured by back titration with 0.4 mol L^{-1} ferrous ammonium sulphate solution using ferroin as an indicator. Soil organic C (extracted org-C) in the extracts was measured by the titration technique using the following equation:

Extracted org-C =
$$\frac{(H-S) \times M \times D \times E \times 1,000}{C \times A} \times \frac{K}{W \times dwt}$$
 (mg g⁻¹ dry soil) (1)

where, H = titration solution consumed by the hot (refluxed) blank (mL), S = titration solution consumed by the sample (mL), C = titration solution consumed by the cold (unrefluxed) blank (mL), M = normality of the K₂Cr₂O₇ solution, D = volume of K₂Cr₂O₇ solution added to the reaction mixture (mL), A = aliquot of the extract (8 mL), E = 3 (conversion of Cr^{VI+} to Cr^{III+}), K = volume of the extractant (mL) and dwt = dry weight of moist soil per gram. Microbial biomass C (MBC) was determined as follows:

where, CF is the extracted organic-C from fumigated sample, CNF is the extracted organic-C from the unfumigated sample, and KEc is the calibration factor (0.38) as proposed by Wardle (1994).

Microbial biomass N (MBN) was also examined by the fumigation-extraction technique. As such, fumigated and non-fumigated soil samples were extracted with K_2SO_4 and the filtered extract was measured for total N by using the Kjeldahl digestion procedure. For digestion, 1.0 g of soil was digested with a digestion mixture (K_2SO_4 :CuSO_4:Se) at the ratio of 1:0.1:0.01, and 3.0 mL of concentrated H_2SO_4 in each digestion tube at 80° C for 12 h. After digestion, the mixtures were carried out for distillation by pouring the samples into the steam distillation chamber of Kjeldahl with 10 mol L⁻¹ NaOH and 2% H₃BO₃. Then, samples from the distillation chamber were titrated against 0.0025 mol L⁻¹ H₂SO₄. Microbial biomass N (MBN) was determined as follows:

where, NF is the extracted organic-N from fumigated sample, NNF is the extracted organic-N from the unfumigated sample, and KEN is the calibration factor (0.54) as proposed by Wardle (1994).

2.5 Basal Soil Respiration

Soil respiration was determined by trapping the evolved CO₂ from N-treated samples in KOH. Briefly, soil samples of 20 g were incubated for 7 d at 25 °C in 500 mL vessels containing 10 mL of 0.3 mol L⁻¹ KOH. A vial containing 10 mL of 1 mol L⁻¹ KOH with the lid open to exclude CO₂ evolved from the soil served as control to account for the CO₂ trapped from the atmosphere. The concentration of CO₂ trapped in the KOH solution was measured by titration with 0.1 mol L⁻¹ HCl against phenolphthalein after carbonate precipitation with 3 mL of 0.5 mol L⁻¹ BaCl₂. Microbial metabolic quotient (*q*CO₂) was thereafter calculated as the ratio of basal respiration to MBC.

2.6 Statistical Analysis

Analysis of variance (ANOVA) was used to analyze the effects of N sources, N rates and their interaction on the microbial biomass pools, viz (MBC and MBN), soil basal respiration (C-CO₂), and metabolic quotient (qCO₂) from each growing season. Homogeneity of variances and normality tests were checked with the Bartlett's and Shapiro-Wilk tests. For qualitative factors (N sources), the means were compared by the Tukey's test at the p < 0.05 level when the F test proved significant, whereas the quantitative factors (N rates) were submitted to regression analysis. Sigmaplot 10.0 was used to create graphs.

3. Results and Discussion

3.1 Effects of N Fertilization on Biomass and Soil Respiration

The effects of N treatments during the two growing seasons (2013 and 2014) of upland rice crop on soil microbial abundance and activity are presented in Table 2. The enhanced-efficiency N fertilizers had no influence on the microbial soil properties at any growing season of the upland rice crop. On the other hand, the N rates significantly affected (p < 0.05) the MBN, C-CO₂ and qCO₂ in the first growing season (2013), whereas all microbial parameters except the C-CO₂ were affected by N rates in the second growing season (2014) (Table 2). No significant interaction effect of N sources and N rates occurred for the microbial parameters.

These results partially support the hypothesis of this study once soil microbial community was affected by N fertilization regardless of the N source. Ramirez et al. (2010) also found that the addition of different forms of N (NH_4NO_3 , (NH_2)₂CO (urea), KNO_3 , NH_4Cl , (NH_4)₂SO₄, $Ca(NO_3)_2$) yielded a similar microbial response to three distinct soils, and demonstrated that a range of fertilizer types have the same impacts on soil respiration.

Source of variation	DF	MS				
Source of variation		MBC	MBN	C-CO ₂	qCO_2	
2013						
N source (S)	2	13.17	9.90	0.60	0.00004	
N rate (R)	4	207.22	167.28**	4.53*	0.0003*	
$S \times R$	8	161.59	6.31	1.82	0.0002	
Residual	42	123.99	4.49	31.46	0.00006	
C.V. %		10	6.83	15.30	11.26	
2014						
N source (S)	2	4.27	0.50	1.77	0.0003	
N rate (R)	4	549.09*	95.28**	2.11	0.0003°	
$S \times R$	8	43.69	7.76	1.39	0.0002	
Residual	42	120.55	5.52	1.43	0.0002	
C.V. %		9.95	7.50	14.20	16.12	

Table 2. Analysis of variance for the effects of enhanced-efficiency nitrogen fertilizers applied at different N rates on microbial biomass carbon (MBC) and nitrogen (MBN), basal respiration (C-CO₂) and metabolic quotient (qCO₂) in two growing seasons (2013 and 2014) of upland rice (cultivar BRS Esmeralda)

Note. DF, degrees of freedom. MS, mean square. °, *, **, ^{ns} significant at the 10, 5 and 1% probability levels, respectively.

3.2 Soil Microbial Biomass

Soil MBC and MBN were highly variable across the two growing seasons and ranged from 101.95 to 120.65 mg C kg⁻¹ and 24.45 to 36.50 mg N kg⁻¹, respectively (Table 3). Nitrogen fertilizer treatments tended to increase soil microbial biomass with increasing N rate, as revealed by the highest values of MBC and MBN in the highest N

rates compared with the control treatment (Table 3). The positive effect of N addition on soil microbial biomass has been reported by other authors (Zhou et al., 2012; Geisseler & Scow, 2014; Geisseler et al., 2016).

In our study, the quadratic responses to N rates of MBN in the 2013 season (Figure 1a), as well as MBC and MBN in the 2014 season (Figures 2a and 2b), respectively, indicated from 75 to 214 kg N ha⁻¹ as the optimum rates for enhancing such properties. Hence, responses of soil microbial biomass were positive up to these N rates and negative thereafter, highlighting that microbial responses are more dependent on the rate of N application rather than the form of N.

Most studies on the effects of N fertilizers on soil microorganisms examine only one N application rate. A large number of studies based on long-term field experiments have proved that optimum fertilizer N application to crops neither resulted in loss of organic matter nor adversely affected microbial activity in the soil (Zhang et al., 2008; Lupwayi et al., 2010; Lupwayi et al., 2011; Singh, 2018). In a 2-yr study with N applied as urea to barley and corn at soil-test recommended rates, Lupwayi et al. (2012) reported either no effect or a positive effect of N on MBC and bacterial diversity, and they concluded that N fertilizer effects depend on the N application rate. Furthermore, it is largely reported in literature that addition of N above than the optimum rates not only adversely influence soil biological communities but also increase residual inorganic N, which can enhance C mineralization and lead to its loss (Singh, 2018). Therefore, the results reported here are in agreement with those reported in literature that fertilizer N applied at recommended rates probably does not cause negative effects on soil microorganisms.

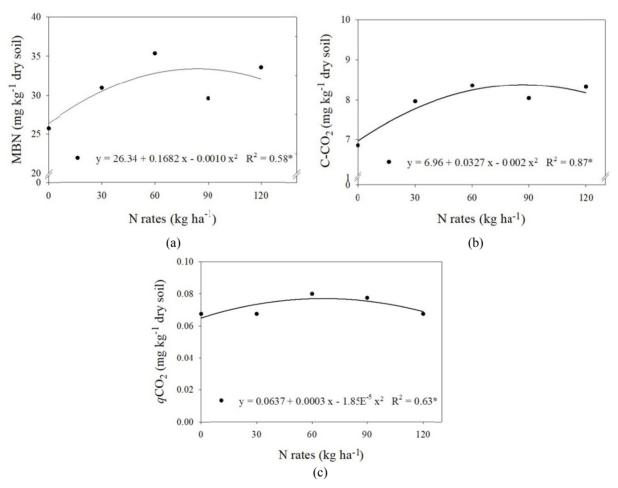


Figure 1. Effect of N application on (a) microbial biomass carbon (MBC), (b) microbial basal respiration (C-CO₂), and (c) microbial metabolic quotient (qCO₂) in soil cultivated with upland rice (cultivar BRS Esmeralda) in the 2013 season

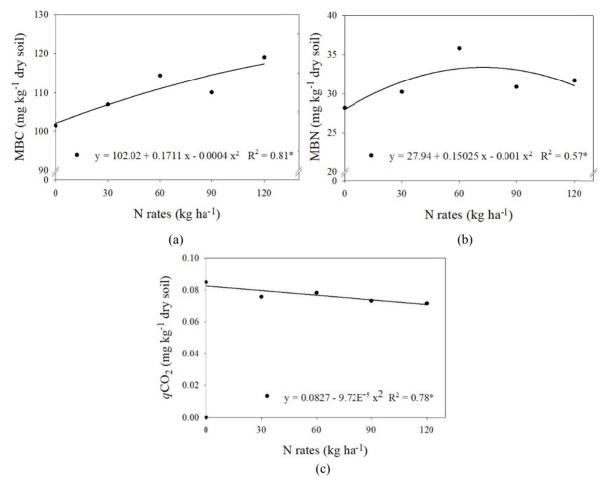


Figure 2. Effect of N application on (a) microbial biomass carbon (MBC), (b) microbial basal nitrogen (MBN), and (c) microbial metabolic quotient (qCO₂) in soil cultivated with upland rice (cultivar BRS Esmeralda) in the 2014 season

3.3 Soil Microbial Respiration

Basal respiration (C-CO₂) and microbial metabolic quotient (qCO₂) followed a quadratic response to N rates in the 2013 season (Figures 1b and 1c) and 2014 season in the case of qCO₂ (Figure 2c). The values of C-CO₂ and qCO₂ across the two growing seasons ranged from 6.58 to 9.70 mg C-CO₂ kg⁻¹ and 0.058 to 0.095 mg qCO₂ kg⁻¹, respectively (Table 3). Significantly lower values of C-CO₂ and qCO₂ were observed in the unfertilized control as compared to the highest N treatments, indicating positive effect of N addition on soil microbial respiration. In addition, a positive effect of N enrichment rates below 60 kg N ha⁻¹ on soil respiration was also observed, whereas N enrichment showed a negative effect at rates higher than 60 kg N ha⁻¹, indicating that a threshold N rate may occur for soil microbial respiration, which requires further study.

	N sources							
N rate (kg ha ⁻¹)		2013			2014			
	UU	PCU	NBPT	UU	PCU	NBPT		
$MBC (mg kg^{-1})$								
0	104.08	113.80	106.28	103.05	102.75	98.50		
30	109.55	109.73	101.95	105.23	102.03	103.45		
60	106.75	110.93	126.50	114.85	111.33	116.75		
90	112.28	108.73	109.98	111.73	106.75	111.55		
120	119.70	114.45	115.63	117.68	120.65	118.85		
$MBN (mg kg^{-1})$								
0	26.80	26.87	24.45	29.45	28.75	26.26		
30	30.89	31.83	30.03	29.60	31.51	29.59		
60	36.10	36.42	33.58	35.48	36.50	35.48		
90	27.91	30.36	30.42	31.35	30.38	30.90		
120	31.92	34.67	34.13	31.35	30.06	33.62		
$C-CO_2 (mg kg^{-l})$						· · · · · · · · · · · · · · · · · · ·		
0	7.21	6.78	6.58	8.00	8.15	8.82		
30	7.63	8.01	8.25	9.20	7.43	9.08		
60	8.57	7.30	9.23	7.88	8.35	8.14		
90	8.66	8.30	7.15	9.70	8.25	9.13		
120	8.00	8.15	8.82	8.80	8.58	7.95		
$qCO_2 (mg kg^{-1})$								
0	0.058	0.068	0.063	0.090	0.070	0.095		
30	0.068	0.073	0.080	0.078	0.078	0.078		
60	0.080	0.068	0.073	0085	0.073	0.078		
90	0.078	0.078	0.065	0.073	0.073	0.075		
120	0.068	0.073	0.078	0.075	0.073	0.068		

Table 3. Absolute amounts of microbial biomass carbon (MBC) and nitrogen (MBN), microbial basal
respiration (C-CO ₂), and microbial metabolic quotient (q CO ₂) in soils cultivated with upland rice (cultivar
BRS Esmeralda) in two growing seasons (2013 and 2014) under addition of enhanced-efficiency nitrogen
fertilizers applied at different N rates

Note UU: uncoated urea, PCU: polymer-coated urea, and NBPT: urease inhibitor NBPT-treated urea.

Applying N at rates above 60 kg ha⁻¹ exceeded the optimal N rate for microorganism activity, as indicated by the decrease in microbial respiration induced by the highest N rates (Figures 1b, 1c and 2c). Similar observations have been made before. In a 2-yr field experiment with a rain-fed rice cropping system, Oladele et al. (2019) reported that soil CO_2 efflux was increased with the addition of N rates up to 60 kg ha⁻¹, above which microbial respiration decreased because of the inhibitory effect of excess N. The results are also in line with Zhai et al. (2017) who found negative effects of N addition on microbial respiration in diverse land-use types after using solid granules of urea at varying N rates. An overall reduction in basal respiration with fertilization was also reported in five grassland ecosystems (Riggs et al., 2015; Strecker et al., 2015; Geisseler et al., 2016).

The decline in microbial CO_2 production over the 2-yr experiment under high N rates is an evidence of any indirect effect of increased N availability on soil respiration. Recently, Zhang et al. (2019) found a decrease in microbial respiration under the addition of 100 kg N ha⁻¹ in the form of urea in a grassland soil. These authors attributed their results to the increased C input into the soil through litter decomposition and root exudates promoted by the increases in available N, which in turn promotes plant growth and aboveground C accumulation. Li et al. (2015) have also observed that soil microbial respiration was adversely influenced by N addition due to the effect of N availability on soil C storage. It is well acknowledged that microbial activities in agricultural soils are mainly determined by C availability (Chantigny et al., 1999), revealing the importance of the regulatory effects of N fertilization on soil C, which enhance aboveground plant biomass thus suppressing soil microbial biomass.

4. Conclusions

Responses of soil microbial properties to N fertilization were dependent on the N rates, but no significant effect of the N sources was observed. All measured parameters, except MBC in the first season and C-CO₂ in the second season, were increased with increasing N rates. However, the application of N higher than 60 kg ha⁻¹ suppressed microbial biomass, as well as soil respiration. Therefore, the lack of response by added urease inhibitors to the N sources indicate that optimizing N rates for upland rice production is a far more effective option for improving soil microbial community than using enhanced-efficiency N sources.

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