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MICROBIAL COMMUNITY PROFILES IN RESPONSE TO DIFFERENT SOIL MANAGERMENTS IN SANDY SOIL

SUMMARY

The growing human population has higher and higher food demand, which requires an increase in efficiency of agricultural production. Healthy and fertile soils are essential to satisfy this demand. The microbial community structure is an excellent indicator of the soil fertility and the diversity of bacteria and fungi. In our work we compared the effect of organic and conventional cultivation methods on the microbial community profiles of sandy soils in the Nyírség region, Hungary. These fields have topographical heterogeneity therefore the sampling sites were divided into top of hill and bottom of hill. Sampling was done in autumn 2013, from the 0-30 cm and the 30-60 cm soil depths. The phospholipid fatty acids (PLFA) were used for the monitoring of microbial community. PLFAs break down when the cell dies therefore these molecules show the community structure in a real time. In the 0-30 cm soil layer of organically managed field the PLFA structure was significantly different between top and bottom of hill, but the difference were low. In conventionally managed field, high differences were found between the PLFA groups measured in the top and bottom of hill. The PLFA values were higher in the top of hill in organic field, while in case of bottom of hills sites higher PLFA values were measured in conventional farming system. In the deeper soil layers the tendencies were found similar to the upper soil layer, but the measured values were lower.

Keywords: cultivation system, microbial community structure, PLFA, topography, soil depth

INTRODUCTION

The soil microbial community largely determines the soil fertility by influencing the dynamics of organic matter and nutrient cycles (Liu et al., 2006; Bowles et al., 2014). These microbes are responsible for breakdown of dead plant residues and turning those to the carbon cycle. Higher microbial activity and biodegradation were observed in the ecological or reduced tillage cultivation than in conventional management system (Ge et al., 2013; Mangalaserry et al., 2015).

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Therefore it is important to apply properly selected cultivation method, which results in favourable conditions for the soil microbes (Prasad *et al.*, 2016).

Before 1989 intensive farming was conducted in large fields with large volume of the used cheap pesticides and fertilizers. The harmful effects of agricultural chemicals to the biodiversity and human health have already known (Snedeker, 2001; Colborn *et al.*, 1993). Nowadays, in Hungary the use of sustainable agricultural methods is preferred (Birkás *et al.*, 2004).

The objective of this study was to compare the effects of ecological and conventional farming systems on microbiological properties of soil. We investigated the effects of studied management systems and the modification effects of relief of the studied area, and we observed the stronger effect of management compared to the relief.

MATERIALS AND METHODS

The sampling sites are located in the Nyírség region, in the north-eastern part of Hungary, where the main soil type is acidic sand (Arenosols). In this region the climate is moderately cold-dry. In the investigated year the main annual temperature was 12.5 °C and the annual precipitation was 486 mm in sampling areas. Because of the topographical heterogeneity of this region, samples were collected from top of hills and bottom hills.

Sampling sites were the followings:

1. Ecological, top of hill (ET) (214036°82'E, 475849°17'N, 156 m above sea level);
2. Ecological, bottom of hill (EB) (214031°48'E, 475848°81'N, 151 m above sea level);
3. Conventional, top of hill (CT) (214051°10'E, 475841°35'N, 158 m above sea level);
4. Conventional, bottom of hill (CB) (214054°64'E, 475842° 43'N, 153 m above sea level).

The studied areas located at the Research Institute of Nyíregyháza, Debrecen University, where the ecological crop production has been carrying since 1997. In the studied sampling period, rye was cultivated in both management systems. In both cultivation methods ploughing was applied up to 30 cm depth and deep loosening up to 60 cm depth in every 5th year. In conventional area the bottom of hill was liming in autumn 2012 because of the low pH (3.89±0.03) of soil.

Samples were collected in autumn 2013 (22 and 24 October), from two depths, 0-30 cm and the 30-60 cm, in four repetitions (one repetition was a composite sample of four point samples). Samples were stored frozen before analysis. The soil temperature was varied in the range between 10.9 – 17.5 °C in the sampling days.

Before the chemical analysis the larger plant roots were removed in the laboratory, and the air dried samples were sieved (Ø 2 mm). The pH_{KCl} was measured with a Hach-Lange, HQ411D type digital pH meter (Hach-Lange,

Loveland, Colorado, USA) where the soil : KCl ratio was 1 : 2.5. Total carbon (C) and nitrogen (N) content were determined by Dumas method (varioMax CNS, Elementar Analysensysteme GmbH, Hanau, Germany). The organic carbon content was calculated from the humus content, which was measured with a UNICAM UV2 spectrophotometer (Thermo Scientific, Waltham, Massachusetts, USA), after the digestion with potassium dichromate and cc. sulphuric acid.

The nitrite-nitrate-N content was measured with a FIA Star 5000 device (Foss, Hilleroed, Denmark), from potassium chloride extracts.

The PLFAs were prepared as described by White et al. (White et al., 1979) with some modifications, in four repetitions. Briefly, 10.00 g (frozen, then incubated and sieved) soil was extracted using a single-phase chloroform-methanol-phosphate buffer system, during 2h with a laboratory shaker.

Then we added chloroform and distilled water to this extract, and filtrated in celite. The filtrate was separated with shaker funnel for the bottom chloroform phase. After the sodium sulphate filtration and vacuum evaporation of the chloroform phase, the phospholipids were separated from neutral lipids and glycolipids using silica acid columns (Chromabond, Macherey Nagel, Germany), followed by methylation of the phospholipids. Samples were stored at -20°C until the analysis. Methyl nonadecanoate was used as internal standard after the methylation. The PLFAs were separated and identified using a gas chromatograph-mass spectrometer system (GC 6890N with MS 5975, Agilent, Santa Clara, CA, USA) with a 100 m Supelco SP-2560 column, in selected ion mode and scan mode as well (50-350 amu).

Gerstel MPS2 autosampler was used for the injection, the injection volume was 3 µl. The PLFAs 14:0, 15:0, 16:0 and 18:0 represented the general bacterial biomass. The branched, saturated PLFAs, as iC15:0, aC15:0, iC16:0, iC17:0, aC17:0 were used as Gram⁺ as well as the monoenoic and cyclopropane unsaturated C16:1n7c, C16:1n5c, C18:1n9c, cyC19:0 as Gram⁻ biomarkers. The 10MeC16:0 and 10MeC17:0 represented the Actinobacteria, and C18:2n6 was used as fungi marker. These PLFAs were used to calculate total PLFA concentration (nmol PLFA g⁻¹ dry soil). On the bases of measured PLFAs the ratios of these groups were calculated.

All measurements were reported as mean values with standard errors, where statistical analysis was done using internal repetitions respectively (n = 4). IBM SPSS Statistics 22.0 software package (IBM Inc., USA) were used for statistical analyses at P = 0.05 significance level.

One-way analysis of variance (ANOVA) was used for comparing the means of different sampling sites followed by Games-Howell and Tukey-b test. Correlations between investigated parameters were determined using Pearson's correlation.

RESULTS AND DISCUSSION

Generally, the measured values of pH_{KCl} fit in the pH range of most acidic sandy soils (4.02-7.01). Furthermore acidifying effect of fertilizer was also

observed (Pais et al., 1990) in conventional top of hill, but in conventional bottom of hill the liming was resulted in higher pH_{KCl} in autumn 2013. Compared to the two farming systems, favourable chemical values were measured in ecological field on the top of hill, except the inorganic N (Table 1), but at the bottom of hill the investigated parameters were higher in conventional field. The available nitrogen and carbon usually increased with increasing pH (Zhao et al., 2011). Generally, the most investigated parameters were higher in upper (0-30 cm) soil layer and also were higher in the bottom of hill, than in top of hill (except of nitrite-nitrate-N in ecological field). In ecological top of hill significantly higher total and organic carbon were observed and recycled plant residues increased the total nitrogen content and carbon: nitrogen ratio (Chen et al., 2000; Edmeades, 2003).

Table 1. The main chemical parameters of investigated soils

Sampling site	pH_{KCl}	total C *	organic C *	total N *	$\text{NO}_2^-\text{NO}_3^-$ -N*	C:N ratio
ET 0-30 cm	5.43±0.01c	0.497±0.02b	0.372±0.02b	0.055±0.00b	0.019±0.00b	9.026±0.06b
ET 30-60 cm	5.53±0.17C	0.332±0.06B	0.232±0.06B	0.040±0.00B	0.010±0.00C	8.241±0.17C
EB 0-30 cm	4.85±0.06b	0.818±0.01c	0.715±0.02c	0.093±0.00c	0.010±0.00ab	8.839±0.01ab
EB 30-60 cm	5.05±0.35B	0.721±0.01C	0.589±0.08C	0.083±0.00C	0.007±0.00B	8.658±0.35B
CT 0-30 cm	4.02±0.04a	0.346±0.00a	0.202±0.00a	0.040±0.00a	0.008±0.00a	8.751±0.06a
CT 30-60 cm	4.15±0.09A	0.222±0.06A	0.100±0.08A	0.029±0.00A	0.004±0.00A	7.488±0.09C
CB 0-30 cm	7.01±0.01d	0.943±0.02d	0.729±0.01c	0.096±0.00c	0.020±0.00b	9.778±0.09c
CB 30-60 cm	6.61±0.44D	0.735±0.08C	0.524±0.06C	0.081±0.00C	0.009±0.00BC	9.116±0.44D

^a Data are expressed as mean ± standard errors (n=12).

^b Within a column followed by different letters represent the differences by one-way ANOVA followed by Games-Howell test ($P < 0.05$).

The management practice affected not only the soil chemical parameters, but also the structure of microbial community. The nutrient supply method has strong effect on the community structure, through the differences of organic matter content, and the value of available substrates (Hartman et al., 2006), and the change of pH affect the structure of microbial community (Rousk et al., 2009). The General bacteria (2.12 ± 0.00 nmol PLFA g^{-1} dry soil), Gram⁺ (1.59 ± 0.00 nmol PLFA g^{-1} dry soil) and Gram⁻ (2.05 ± 0.00 nmol PLFA g^{-1} dry soil) bacteria, Actinobacteria (0.48 ± 0.00 nmol PLFA g^{-1} dry soil) and Fungi (0.32 ± 0.00 nmol PLFA g^{-1} dry soil) markers were highest in upper soil layer in conventional bottom of hill, while the lowest values were measured in conventional top of hill (Table 2). In case of these markers and the total PLFA significant differences were observed between two reliefs, but compared to the

two farming systems, the differences were lower in ecological, than the conventional farming system. Furthermore in ecological field the values of measured markers and total PLFA were higher in top of hill, but in conventional plots were higher in bottom of hill. The pH influences of microbial community (Bååth & Anderson, 2003) and liming could cause the increased PLFA values of CB site.

Table 2. Structure of microbial community in autumn 2013

PLFA markers	ET		EB		CT		CB	
	0-30 cm	30-60 cm	0-30 cm	30-60 cm	0-30 cm	30-60 cm	0-30 cm	30-60 cm
General bacteria ^a	1.30±0.00 c	1.07±0.00 C	1.2±0.00 b	0.69±0.00 A	0.75±0.00 a	0.72±0.00 B	2.12±0.00 d	1.08±0.00 D
Gram ⁺ bacteria ^a	1.05±0.00 c	0.98±0.00 D	1.04±0.00 b	0.43±0.00 A	0.53±0.00 a	0.55±0.00 B	1.59±0.00 d	0.73±0.00 C
Gram ⁻ bacteria ^a	0.97±0.00 b	0.74±0.00 C	1.08±0.00 c	0.46±0.00 B	0.50±0.00 a	0.45±0.00 A	2.05±0.00 d	0.84±0.00 D
Actinobacteria ^a	0.38±0.00 c	0.16±0.00 A	0.35±0.00 b	0.17±0.00 B	0.17±0.00 a	0.27±0.00 C	0.48±0.00 d	0.42±0.00 D
Fungi (C18:2n6) ^a	0.13±0.00 c	0.08±0.00 C	0.10±0.00 b	0.03±0.00 A	0.05±0.00 a	0.05±0.00 B	0.32±0.00 d	0.11±0.00 D
Gram ⁺ : Gram ⁻ ratio	0.92±0.00 a	0.76±0.00 A	1.05±0.00 c	1.05±0.00 C	0.95±0.00 b	0.82±0.00 B	1.29±0.00 d	1.16±0.00 D
Fungi : General bacteria ratio	0.10±0.00 c	0.07±0.00 C	0.08±0.00 b	0.04±0.00 A	0.07±0.00 a	0.07±0.00 B	0.15±0.00 d	0.11±0.00 D
Actinobacteria : General bacteria ratio	0.30±0.00 d	0.39±0.00 D	0.29±0.00 c	0.25±0.00 C	0.23±0.00 b	0.22±0.00 A	0.22±0.00 a	0.25±0.00 B
Total PLFA	3.83±0.00 c	3.29±0.00 D	3.77±0.00 b	1.78±0.00 A	2.00±0.00 a	1.92±0.00 B	6.56±0.00 d	3.02±0.00 C

^a Data are expressed as mean (nmol PLFA g⁻¹ dry soil) ± standard errors (n=4).

^b Within a raw different letters represent the differences of means according to one-way ANOVA followed by Tukey-b test ($P < 0.05$), the lowercases (a-d) are marked the results of 0-30 cm soil layer and capital letters (A-D) are marked the results of 30-60 cm soil layer.

Low proportion of easily available carbon source increases the value of Gram⁺ bacteria (Esperschütz et al., 2007), while values of Gram⁻ bacteria markers was increased when the quantity of easily available, unstable carbon forms were increased in the soil (Peacock et al., 2001). Therefore, the measured lower Gram⁺ : Gram⁻ bacteria ratio in ET and CT indicates higher value of easily available carbon forms in this sites.

The increasing organic matter input results in increase of fungi, thereby the increase of Fungi : Bacteria ratio (Frostegård et al., 1996). We measured higher Fungi values from upper soil layer in conventional bottom of hill and the ecological top of hill, but the organic carbon was higher in the two bottom of hill and the fungi was higher in the CB and ET, there was not significant relationship

between the two parameters. The value of fungi depends also on the pH. When the pH lower, the conditions are unfavourable for bacteria, and the fungi community increases (Rousk *et al.*, 2009). However, we not found negative correlation between the values of Fungi and pH (0.467, $P < 0.01$). The relationship between pH and general bacteria markers was significantly positive (0.455, $P < 0.01$)

Higher ratio of Actinobacteria: General bacteria in ecological field indicated the progressed decomposition process in sampling time, opposite in conventional field (Bastida *et al.*, 2013).

CONCLUSIONS

Ecological management had favourable effect on the chemical parameters and microbial community of sandy soils in Nyírség region. The lower differences between two reliefs in ecological field were resulted by favourable conditions and better buffer capacity of soil, providing protection against extreme environmental events. The positive effect of liming was also observed, like increasing pH and microbial biomass in conventional bottom of hill. However, additional studies are needed to understand changes of Fungi : bacteria ratio with increasing pH. Our results revealed that regular recycling of organic matter after harvesting without artificial fertilizer utilization could maintain the fertility of sandy soils.

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