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## Phosphomonoesterase activity under different microbiological soil properties

*In the present work, effects of microbiological soil properties and phosphomonoesterase activity are presented. The abundances of ammonifiers, fungi, actinomycetes, Azotobacter sp., cellulolytic microorganisms, microorganisms that solubilize organic and inorganic P compounds bound to Al, Fe and Ca, have been determined by the dilution method with application of selective nutrient mediums. The first five axes explains 86% of the total variance. The first axis explains 34.6% of variance and it appears to reflect the processes of microbiological mineralization and immobilization of phosphorus. The second axis explains 25.9% of variance and reflects microbial activity and organic matter accumulation, N biomasses, Azotobacter sp. abundance and acid and alkaline PME activities. Phosphomonoesterase activity in soil depend primarily on soil organic matter and biological factors and can be used as a parameter of plants' uptake of phosphorus, especially in conditions of its low concentration in soil.*

**Keywords:** microorganisms, soil, microbial biomass

### 1. INTRODUCTION

The transformations of organic and inorganic forms of P are closely interrelated and included in the soil P cycle. The availability of phosphorus is often controlled by activity of microorganisms, primarily their enzymes, which play the key role in biological activity of soil [1] and provide a biological assessment of soil [2]. Concept of soil quality includes different biological and microbiological parameters such as soil enzymes [3]. Enzymes synthesis in soil is continual; they have a great importance for agricultural production [4]. Different microbial populations might have developed with different turnover rates and P cycling rates through their biomasses [5]. Biological and biochemical control of P cycle in soil depend on organic phosphates mineralization, i.e. on acid phosphatase activity and on the content of inorganic phosphates. Enzymes play a key role in biochemical cycling of nutrients in soils [6]. Activity of soil enzymes is in correlation with properties of soil [7], vegetation [8], succession [9] microbial population [10].

One of most examined enzymes are phosphomonoesterases, who catalyze hydrolysis of orthophosphorous acid esters and are involved in processes of mineralization of organic P compounds [11]. These enzymes have an important

role in supplying of plants with available phosphorus. Soil phosphomonoesterases originate from plant roots and microorganisms [12]. The synthesis of these enzymes is in correlation with soil enzymatic activity [13]. The present work includes researches of abundance of various microorganism groups, microbiological biomass content in different soil types and phosphomonoesterase activities.

### 2. MATERIALS AND METHODS

The investigated soils belong to different types, subtypes, modes of utilization and sampling sites in Serbia (Table 1). Ten pedological profiles were opened per each type and subtype of soil, selecting representative ones, from which samples were collected along genetical horizons down to original substrate, except in the case of chernozem in Zemun Polje.

Acid and alkaline phosphomonoesterase (PME) activities were determined utilizing a modified universal buffers pH 6.5 and 11 according to Tabatabai [14]. Measurement of phosphomonoesterase activity was conducted in four replications and expressed in pmol p-nitrophenole g<sup>-1</sup> s<sup>-1</sup>.

Microbial biomass of C has been determined by fumigation-incubation method [15]. Microbial biomass was calculated as Fc/kc (15) where Fc is (CO<sub>2</sub>-C evolved from fumigated soil during 0-10 d incubation) - (CO<sub>2</sub> - C evolved from nonfumigated soil during 0-10 d incubation) and kc is 0.45 [16]. N biomass has been determined by incubation of fumigated and non-fumigated soil samples at 25 °C, at 60% WHC, for 14 days [17]. Extraction was performed with 2M KCl, and NO<sub>3</sub><sup>-</sup>-N and NH<sub>4</sub><sup>+</sup>-N in the extract were determined [18]. N biomass has been calculated on the basis of differences between net amount of min-N produced in fumigated,

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non-fumigated soils and before incubation, using risk factor  $k = 0.33$ . P biomass has been determined according to Brookes [19]. The initial incubation of the samples under moist conditions in the presence of soda lime for 10 days was followed

by fumigation [15]. Microbiological P was determined according to Murphy and Riley [20]. Correction for the adsorbed P during extraction was performed according to Brookes [19].

Table 1 - Types of the sampled soils

| No. prof | Soil type (soil subtype)                            | Depth (cm)                   | Site          | Vegetation type   | Texture class |
|----------|---|------------------------------|---------------|-------------------|---------------|
| 1        | Calcomelanosol<br>(Organo-mineral black soil)       | 0 - 20                       | Rajac         | Pasture           | Silty clay    |
| 2        | Calcomelanosol<br>(Brownized calcareous black soil) | 0 - 20<br>20 - 40<br>40 - 60 | Rajac         | Pasture           | Heavy clay    |
| 3        | Calcomelanosol<br>(Lessive calcareous black soil)   | 0 - 15<br>30 - 50<br>50 - 70 | Rajac         | Pasture           | Silty clay    |
| 4        | Solonetz  | 0 - 15<br>30 - 50<br>50 - 70 | Kumane        | Uncultivated land | Silty clay    |
| 5        | Chernozem   | 0 - 20<br>20 - 40            | Zemun Polje   | Maize             | Silty clay    |
| 6        | Chernozem   | 0 - 20<br>20 - 40            | Roman ditches | Soybean           | Light clay    |
| 7        | Humogley  | 0 - 30<br>30 - 70            | Becej         | Winter wheat      | Heavy clay    |
| 8        | Fluvisol  | 0 - 30<br>30 - 70            | Kaska forest  | Poplar plantation | Sand          |

Microbiological analyses included determination of ammonifiers, fungi, actinomycetes, *Azotobacter* sp. and cellulolytic microorganisms. The abundance of microorganisms solubilizing inorganic P compounds (PSM) was determined on glucose- asparagine substrate according to Muramtsov ( $10^{-4}$ ) with the following composition: asparagine 1g, glucose 10 g,  $K_2SO_4$  0.2 g,  $MgSO_4$  0.2 g, maize extract 0.02%, agar 20 g, and tap water 1000 ml. The substrate was supplied also with 1.5 g  $CaCO_3$  ( $PO_4$ )<sub>2</sub>, 0.264 g  $AlPO_4$  and 0.28 g

$FePO_4$ , for determination of PSM-Ca, Al and Fe. Abundance of microorganisms transforming organic phosphates (PSM-Po) was determined on Menkine substrate [21] with lecithine as a sole P source. As a mathematical model for the interrelation of the investigated parameters, correlation coefficient was calculated, expressing the strength of relationship between the investigated components. Starting from indicators defined in this way, we have examined the nature of factor spaces for microbiological properties of soils.

## 3. RESULTS

The abundance of ammonifiers, actinomycetes, *Azotobacter* sp., PSM-Po and PSM-Ca, Al and Fe is higher in chernozem and humogley compared to the remaining investigated soil types,

while the abundance of fungi is highest in calcomelanosol. Microorganisms solubilizing  $\text{FePO}_4$  are represented more poorly than those solubilizing  $\text{AlPO}_4$ , but more abundantly than PSM-Ca. (Table 2).

Table 2 - Microbiological soil properties

| Soil | PSM                      |       |       |      | Ammonifcators         | Fungi                   | Actino-mycetes          | <i>Azotobacter</i> sp.  | Cellulolytic microorg.  |
|------|--------------------------|-------|-------|------|-----------------------|-------------------------|-------------------------|-------------------------|-------------------------|
|      | Po                       | Fe    | Al    | Ca   |                       |                         |                         |                         |                         |
| Type | $(10^4 \text{ xg}^{-1})$ |       |       |      | $104 \text{ xg}^{-1}$ | $(104 \text{ xg}^{-1})$ | $(104 \text{ xg}^{-1})$ | $(102 \text{ xg}^{-1})$ | $(104 \text{ xg}^{-1})$ |
| 1    | 65.3                     | 60.7  | 80    | 39.6 | 57                    | 7.3                     | 0.7                     | 4.68                    | 0.7                     |
| 2    | 67                       | 65    | 86    | 41.3 | 92                    | 10.6                    | 2.7                     | 2.58                    | 0.7                     |
|      | 41.7                     | 55    | 73.7  | 32   | 58.7                  | 7.3                     | 1.3                     | 2.13                    | 0.5                     |
|      | 10.7                     | 8     | 20.7  | 3.3  | 20.7                  | 8.6                     | 0                       | 0.88                    | 2.4                     |
| 3    | 43.3                     | 33.3  | 64.7  | 26   | 47.3                  | 4                       | 0                       | 2.01                    | 0.8                     |
|      | 9.3                      | 11.3  | 14.7  | 4.6  | 9.3                   | 7.3                     | 0                       | 0.52                    | 0.5                     |
|      | 5                        | 3.3   | 3     | 2.6  | 4.7                   | 4.7                     | 0                       | 0.18                    | 1.7                     |
| 4    | 152.7                    | 141.3 | 229   | 60   | 165.3                 | 2.7                     | 30                      | 4.69                    | 7.3                     |
|      | 52                       | 34.7  | 62    | 31.5 | 98                    | 3.3                     | 10.6                    | 2.06                    | 1.5                     |
|      | 87                       | 56    | 53    | 22.6 | 75                    | 8.7                     | 2                       | 1.09                    | 1.3                     |
| 5    | 167                      | 102   | 110   | 26   | 190                   | 2                       | 28                      | 2.3                     | 12                      |
|      | 138                      | 90.6  | 87    | 20   | 147                   | 2.6                     | 12                      | 2.15                    | 13                      |
|      | 146.2                    | 105.6 | 150.5 | 67.9 | 186                   | 0.5                     | 36                      | 2.5                     | 23.2                    |
| 6    | 130                      | 86.2  | 135   | 59   | 157                   | 0.2                     | 18                      | 1.8                     | 14.5                    |
|      | 280                      | 180   | 320   | 53.1 | 242                   | 0.1                     | 36                      | 3.29                    | 11                      |
| 7    | 120                      | 65.3  | 170.9 | 30.9 | 47.8                  | 0.2                     | 12                      | 1.25                    | 10.4                    |
| 8    | 90                       | 42.6  | 155   | 47.1 | 77.1                  | 1.3                     | 4                       | 1.12                    | 0.7                     |
|      | 15                       | 7.9   | 84.3  | 53.3 | 11.6                  | 0.3                     | 1                       | 0.93                    | 0.1                     |

Activities of acid and alkaline PME are significantly correlated with C and N biomasses, but not with P biomass (Table 3).

Alkaline PME activity correlates positively with the abundances of all microorganism groups, while the acid PME is significantly correlated only with the abundance of *Azotobacter* sp. and fungi.

Principal components analysis was used to investigate relationships among biochemical and microbiological properties (Table 4). The first five axes explained 86.04% of the total variance. The

first axis explained 34.6% of variance and appears to reflect microbiological activity: abundance of cellulolytic microorganisms, PSM-Po and PSM-Fe and PSM-Al, aminoheterotrophs, actinomycetes, P biomass and available P content in soil. The second axis explains 25.9% of variance and reflects microbial activity and organic matter accumulation: positive correlation with humus and total N contents, biomasses of C and N, *Azotobacter* abundance and acid and alkaline PME activities; and a negative with pH.

Table 3 - Correlation coefficients between PME activity, microbiological biomass and microorganism's abundance

|                   | Acid   | Alkaline | Biomass C | Biomass | Biomass P |
|-------------------|--------|----------|-----------|---------|-----------|
|                   | PME    | PME      |           | N       |           |
| Acid PME          | 1      | 0.77a    | 0.62 a    | 0.71 a  | -0.1      |
| Alkaline PME      | 0.77 a | 1        | 0.79 a    | 0.72 a  | 0.28      |
| Biomass C         | 0.62 a | 0.79 a   | 1         | 0.63 a  | 0.44      |
| Biomass N         | 0.71 a | 0.72 a   | 0.63 a    | 1       | 0.16      |
| Biomass P         | -0.1   | 0.28     | 0.44      | -0.16   | 1         |
| PSM – Po          | -0.06  | 0.38     | 0.49b     | 0.01    | 0.86 a    |
| PSM – Fe          | 0.03   | 0.39     | 0.61 a    | 0.26    | 0.77 a    |
| PSM – Al          | 0.1    | 0.57 a   | 0.57 a    | 0.08    | 0.66 a    |
| PSM – Ca          | 0.16   | 0.45     | 0.45      | 0.19    | 0.39      |
| Ammonifiers       | -0.05  | 0.46     | 0.47 b    | 0.13    | 0.82 a    |
| Fungi             | 0.49 b | 0.18     | -0.14     | 0.37    | -0.58 b   |
| Actinomycetes     | -0.21  | 0.32     | 0.44      | 0.04    | 0.82 a    |
| Azotobacter sp.   | 0.53 a | 0.80 a   | 0.73 a    | 0.72 a  | 0.3       |
| Cellulolytic mic. | -0.3   | 0.17     | 0.34      | -0.21   | 0.82 a    |

a -  $P < 0.01$ b -  $P < 0.05$ 

Table 4 - Rotated factor matrix for soil microbiological properties

|                 | Factor 1 | Factor 2 | Factor 3 | Factor 4 | Factor 5 |
|-----------------|----------|----------|----------|----------|----------|
| PSM - Po        | 0.918a   | 0.059    | 0.033    | 0.166    | 0.202    |
| PSM – Fe        | 0.877 a  | 0.264    | 0.031    | 0.113    | 0.268    |
| PSM – Al        | 0.791 a  | 0.209    | 0.32     | 0.345    | -0.01    |
| PSM – Ca        | 0.545    | 0.283    | 0.676    | 0.037    | 0.017    |
| Ammonifiers     | 0.902 a  | 0.078    | 0.041    | -0.001   | 0.307    |
| Fungi           | -0.64    | 0.324    | -0.45    | 0.017    | 0.325    |
| Actinomycetes   | 0.932 a  | -0.01    | 0.074    | -0.05    | 0.066    |
| Azotobacter sp. | 0.446    | 0.700 a  | 0.163    | -0.07    | 0.342    |
| Cellul. mic.    | 0.854 a  | -0.18    | -0.06    | -0.32    | -0.14    |
| Biomass C       | 0.495    | 0.777 a  | 0.011    | -0.13    | -0.16    |
| Biomass N       | -0.05    | 0.834 a  | 0.011    | -0.24    | 0.265    |
| Biomass P       | 0.910 a  | -0.04    | -0.19    | 0.022    | -0.08    |
| Expl.Var        | 7.789    | 5.589    | 3.974    | 1.576    | 1.722    |
| Prp.Totl        | 0.325    | 0.233    | 0.166    | 0.066    | 0.072    |

a - Marked loadings are  $>0.700$

#### 4. DISCUSSION

Soil function mainly involves recycling of organic inputs and maintenance of physical structure. These processes are regulated by a suite of hierarchically organized factors [22] operated at nested scales of space and time in other: climate, soil (clay and nutrient status) properties, organic matter, macro- and microorganisms. PME activity, microorganism abundance and microbiological biomass have been determined in soils formed under the influence of various pedogenetic factors and which differ from one another by chemical and physical properties [13]. Alkaline PME is an enzyme of microbiological origin and depends on the number of microorganisms that synthesize it [23]. However, acid PME is produced by microorganisms and plant roots and thus it is not directly dependent on the abundance of microorganisms. Nakas [24] state that bacterial isolates have 10-100 times higher phosphatase activities than fungi isolates. The same authors state that the activity of alkaline phosphatase increases when amino acid is introduced in soil. High microbiological activity and PME exudation may decrease phosphatase activity. Few previous studies have addressed the decrease of PME activity [25] with soil depth. This conclusion is in correspondence with our results.

Dominant group of microorganism are ammonifiers, actinomycetes, *Azotobacter* sp. and PSM, especially in chernozem and humogley. Some reports [26, 27] confirmed also the highest abundance of same microbial groups in chernozem and humogley. The results of this investigation show that PME activity was not significantly correlated with biomass of phosphorus. However, Sparling [28] find significant correlation coefficients. Biological mechanisms play an important role in controlling the magnitude and forms of phosphorus release from the soil although they have been underrepresented in the past. Ammonifiers and PSM-Po mineralize organic nitrogen and phosphorus compounds bound to PSM-Al and Fe-phosphates. In the association of phosphorus components with organic matter there are Al-phosphates and Fe-phosphates (but not Ca-phosphates). Rojo [29] showed a significant correlation between the contents of organic phosphorus, Al-phosphates, Fe-phosphates and phosphatase activities. Through decomposition processes, P of biological origin supplies nearly 87% and weathering of primary inorganic P furnishes 13% of the total P taken up by plants [30]. Phosphorus held in the soil microbial biomass ( $P_{mic}$ ) is the most rapidly cycling organic P pool and may play an important role in plant nutrition, especially in natural eco- and low input agrosystems.

#### 5. CONCLUSION

Activities of acid and alkaline PME are significantly correlated with C and N biomasses, but not with P biomass. Alkaline PME activity correlates positively with the abundances of all microorganism groups, while the acid PME is significantly correlated only with the abundance of *Azotobacter* sp. and fungi.

This research may be helpful for studying of phosphatase activity in different soil types and can be used as a parameter of plants' uptake of phosphorus, especially in conditions of its low concentration in soil.

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## IZVOD

### AKTIVNOST FOSFOMONOESTERAZE U ZAVISNOSTI OD MIKROBIOLOŠKIH SVOJSTAVA ZEMLJIŠTA

U ovom radu ispitivan je uticaj mikrobioloskih svojstava zemljista i aktivnosti fosfomonoestereaze. Brojnost azotofiksatora, gljiva, aktinomiceta, *Azotobacter sp.*, celulolitičkih mikroorganizama, mikroorganizma koji razlazu organske i neorganske jedinjenja vezana za P, Al, Fe i Ca, utvrđene su postupkom razređivanja s primenom selektivnih hranjivih medija. Prvih pet osa objašnjava 86% ukupne varijanse. Prva osa objašnjava 34,6% varijanse, a odražava procese mikrobiološke mineralizacije i imobilizacije fosfora. Druga osa objašnjava 25,9% varijanse i odražava mikrobiološku aktivnost i nakupljanje organske materije, N biomase, brojnost *Azotobacter sp.* i aktivnosti kisele i alkalne fosfomonoestereaze.

Fosfomonoesterazna djelovanja u tlu primarno zavise od organske materije i bioloških faktora i može se koristiti kao parametar biljnog usvajanja fosfora, naročito u uslovima njegove niske koncentracije u tlu.

**Ključne reči:** mikroorganizmi, zemljište, mikrobiološka biomasa

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