CHANGES IN THE STRUCTURE OF ACTINOMYCETE POPULATIONS IN THE RHIZOSPHERE OF VICIA SATIVA SPECIES

MARINEL PAȘCA¹, LUMINIȚA COJOCARIU¹, HORABLAGA MARINEL¹, DESPINA-MARIA BORDEAN¹, DRAGOS NICA¹, FILIMON MARIOARA NICOLETA², GERGEN IOSIF¹, AURICA BREICA BOROZAN¹

¹ Banat's University of Agricultural Sciences and Veterinary Medicine 300645 Timisoara, 119 Aradului Way
²West University of Timisoara, 300223 Timisoara, 4 Blvd. V. Parvan, Romania; <u>borozan_a@yahoo.com</u>

ABSTRACT

It is a known fact that species of legumes improve the soil they are grown on, but at the same time, they produce the so-called rhizosphere effect or rhizodeposit that has a selective effect on the microorganisms which are considered "fertility effectors" for soil. From the three studied area the highest number of actinomycetes was found in edaphosphere and the lowest number in the area influenced by roots. Among the few factors under research for the purpose of this paper, humus and potassium were observed to have the strongest impact on this group. Humidity is a factor that could change the competition between soil microorganisms and plants in the soil for N and it could affect the stability of aggregates.

Keywords: actinomycetes, rhizodeposition, rhizosphere, edaphosphere, Gause medium

INTRODUCTION

Rhizosphere microbial communities are important for plant nutrition and plant health (MARSCHNER ET AL., 2004). The increased use of cereal/legume crop rotation has been advocated as a strategy to increase cereal yields of subsistence farmers in West Africa, and is believed to promote changes in the rhizosphere that enhance early plant growth (ALVEY ET AL., 2003). Rhizosphere is influenced by the region, soil and the roots of plants with high microbial activity (HILTNER, 1904). Plant roots secrete a large variety of compounds that they release into the rhizosphere, which leads to unique micromedia for microorganisms. Rhizodeposits differ in relation with plant species and plant developmental stage (WHIPPS, 2001; RENGEL, 2002). Interactions and biochemical exchanges that take place between plants and microorganisms in the soil have already been described and analysed (PINTON ET AL., 2007). Competitiveness is fierce among the microorganisms in this area under the influence of plant roots; this makes it possible for intimate associations to be realized between these organisms and plants (HARTMANN ET AL., 2009).

The exudates produced by plant roots select and influence the development of bacterial and fungal populations in their vicinity (GRAYSTONE ET AL., 1996; YANG and CROWLEY, 2000; WHIPPS, 2001). The stimulation of actinomycetes in the rhizospere has never been studied in detail. There exists a general observation that actinomycetes are less stimulated by the rhizosphere effect than the bacteria, but when the number of antagonistic actinomycetes increases in this area, bacteria are inhibited (LECHEVALIER, M., 1989B).

Of actinomycetes, genera *Nocardia* and *Streptomyces* play an important part in phosphorus solubilisation .Edaphosphere is the area that is not influenced by plant roots. Sporogenic bacteria and actinomycetes are larger in numbers than any other type.

The microbial population in the rhizosphere is influenced by the interaction between the type of soil, plant species and its stage of development (MARSCHNER ET AL., 2001, 2004).

The same authors state that the bacteria in the rhizosphere are also affected by the complex interaction among the type of soil, plant species and the location in relation to the root. In some situations, the effect of the soil type on the microbiota in the rhizosphere can be stronger as compared with that of some plant species (SINGH ET AL., 2007), but there are also cases when plant species have a greater influence on the structure of microbial populations (WIELAND ET AL., 2001, GRAYSTON J. SUSAN AND CAMPBELL D. COLIN, 1996).

Humidity is the factor that could change the competition between soil microorganisms and plants in the soil for N (LIPSON and MONSON, 1998). It could affect the stability of aggregates (LAVEE ET AL., 1996), the intensity of humectation-rehumectation cycles, and in their turn, they could affect root secretions (GORRISEN ET AL., 2004).

The interactions among plants, soils and microorganisms are well known. Nevertheless, few studies have been made in order to understand the microbial diversity, the way soil functions and the influence of the cultivated plant.

MATERIAL AND METHOD

The soil under study is moderately gleyic eutric cambisol found in Banat area and cultivated with a vetch species (*Vicia sativa*). The depth for sampling soil was between 0 and 20 cm. The samples were taken from the rhizosphere of the cultivated plants, from the edaphosphere and a control variant. The samples were processed in laboratory conditions. We isolated the actinomycetes using the method of decimal dilutions and sowing the suspension of culture medium Gause 1. The incubation of the samples was at a temperature of 28 °C for five days (STEFANIC, 2006). The data were statistically analyzed using PAST 2.14 (HAMMER ET AL, 2001).

RESULTS AND DISCUSSIONS

The experimental data obtained after the incubation period were interpreted statistically and they are represented graphically below (*Figures 1-6*).



Figure 1. Evolution of studied parameters

Of the three areas under study, edaphosphere (68.96 CFU/g soil) presents the largest number of actinomycetes, followed by the control variant. The smallest number of actinobacteria is to be found in the rhizosphere (20.22 CFU/g soil).



Figure 2. Diversity profiles

The edaphosphere is the area that is not influenced by plant roots. This area is dominated by sporogenic bacteria and by actinomycetes. A large variety of media were used in order to isolate and count the actinomycetes in the rhizosphere, but also to compare cultures and numbers of actinomycetes, eubacteria and fungi found in the soil (BASIL ET AL., 2004).



Figure 3. PCA graphical representation of studied parameters

Figure 2 is presenting linear variations for pH, humidity coefficient, humus and potassium content and it is visible that humus and potassium content is influenced by the pH and humidity of soil. UFC is presenting a different profile, but it's visibly influenced by humus and potassium content

PCA analysis is presented in figure 3. Humidity and pH describe the batch, UFC is characteristic to edaphosphere while rhizosphere is characterized more by humus and potassium content (*Figure* 3). Soil humidity is the key factor that influences the microbial activity in the soil and the processes of decomposition of organic matter (BRADY and WEIL 2002). The variance of the values is 76.344% for the first PC and 23.656 % for the second PC. The PCA loadings for the first axis are presented in *Figure* 4.



Figure 4. PCA correlations of the first component loadings

The pH, humidity, humus and K contents show positive correlations while UFC presents negative correlation (*Figure* 4).

The PCA loadings for the second component (*Figure* 5) present positive correlation for UFC, pH and humidity content while humus and potassium content present negative correlations.



Figure 5. PCA correlations of the second component loadings

This module is used for plotting taxon abundances in descending rank order on a logarithmic (Whittaker plot) scale. This will give a straight descending line in the Whittaker plot. Fitting is by simple linear regression of the logarithm abundances.



Figure 6. Log Abundance Model of the UFC (k=0.4585; chi^2= 0.0365; p(same) =0.849)

CONCLUSIONS

The experimental data obtained are confirmed by the bibliography we studied for the purpose of this paper. The data show that actinomycetes are dominant in the area that is not influenced by legume roots and that their numbers decrease in numbers in the rhizosphere of *Vicia sativa*. The study shows that, of all the factors of influence considered, humus and potassium influence CFU/g soil in a positive way.

As for edaphosphere, the statistical data show a correlation with the pH and the humidity coefficient.

REFERENCES

ALVEY S., YANG C. H., BUERKERT, A., CROWLEY, D. E. (2003): Cereal/legume rotation effects on rhizosphere bacterial community structure in west african soils. Biol Fertil Soils (2003) 37:73–82.

BASIL A. J., STRAP J. L., KNOTEK-SMITH H. M. AND CRAWFORD D. L. (2004): Studies on the microbial populations of the rhizosphere of big sagebrush (*Artemisia tridentata*). Journal of Industrial Microbiology and Biotechnology 31, 278–288.

BRADY N., WEIL R. R. (2002): The Nature and Properties of Soils, 13th ed. Prentice-Hall, Upper Saddle River, NJ, 960 p.

GRAYSTONE S. J., VAUGHAN D., JONES D. (1996): Rhizosphere carbon flow in trees in comparison with annual plants: The importance of root exudation and its impact on microbial activity and nutrient availability. Applied Soil Ecology 5, 29-56.

GRAYSTON J. SUSAN AND CAMPBELL D. COLIN (1996): Functional biodiversity of microbial communities in the rhizospheres of hybrid larch (*Larix eurolepis*) and Sitka spruce (*Picea sitchensis*) Tree Physiology 16, 1031—1038.

GORISSEN A., TIETEMA A., JOOSTEN N.N., ESTIARTE M., PENUELAS J., SOWERBY A., EMMETT B.A., AND BEIER C. (2004): Climate change affects carbon allocation to the soil in shrublands. Ecosystems 7, 650–661.

HAMMER Q., HARPER D.A.T., AND P. D. RYAN (2001): PAST: Paleontological Statistics Software Package for Education and Data Analysis. Palaeontologia Electronica 4(1): 9pp.

HARTMAN A., SCHMID M., VAN TUINEN D. AND BERG G. (2009): Plant-driven selection of microbes. Plant. soil, 321, 235-257.

HILTNER L. (1904): Uber neuere erfarungen und problem auf dem gebiet der bodenbakteriologie und unter besonderer berucksichtigung der grundung und brache. Arbeiter der Deutsche Landwirtschafts-Gesellschaft; vol. 98, pp. 59–78.

LAVEE H., SARAH P., IMESON A.C. (1996): Aggregate stability dynamics as affected by soil temperature and moisture regimes. Geografiska Annaler Series A—Physical Geography 78A, 73–82.

LECHEVALIER M. (1989B): Actinomycetes in agriculture and forestry. In: Goodfellow, M., Williams, S.T. and Williams, M.M. (eds), Actinomycetes in Biotechnology. Academic Press, London, UK, pp. 327–358.

LIPSON D.A., MONSON R.K. (1998): Plant–microbe competition for soil amino acids in the alpine tundra: effects of freeze–thaw and dry–rewet events. Oecologia 113, 406–414.

MARSCHNER P., YANG C.H., LIEBEREI R., CROWLEY D.E. (2001): Soil and plant specific effects on bacterial community composition in the rhizosphere. Soil Biology & Biochemistry 33, 1437–1445.

MARSCHNER PETRAR, CROWLEY DAVID AND CHING HONG YANG (2004): Development of specific rhizosphere bacterial communities in relation to plant species, nutrition and soil type. Plant and Soil 261: 199–208.

PINTON R., VERANINI Z., NANNIPIERI P. (2007): The rhizosphere. Biochemistry and organic substances at the soil-plant interface. New York, USA. Taylor Francis Group, LLC.

RENGEL Z. (2002): Genetic control of root exudation. Plant and Soil 245, 59–70.

SINGH, B.K., MUNRO, S., POTTS, J.M., MILLARD, P., (2007): Influence of grass species and soil type on rhizosphere microbial community structure in grassland soils; Applied Soil Ecology 36, 147–155.

STEFANIC GH. (2006): Metode de analiza a solului (biologica, enzimatica si chimica). Probleme de agrofitotehnie teoretica si aplicata. Institutul de cercetare-dezvoltare agricola Fundulea, XXVIII, 5-25.

YANG C.H. AND CROWLEY D.E. (2000): Rhizosphere microbial community structure in relation to root location and plant iron nutritional status. Applied and Environmental Microbiology 66, 345–351.

WHIPPS J.M. (2001): Microbial interactions and biocontrol in the rhizosphere. Journal of Experimental Botany 52, 487–511.

WIELAND G., NEUMANN R., BACKHAUS H. (2001): Variation of microbial communities in soil, rhizosphere and rhizoplane in response to crop species, soil type and crop development. Applied and Environmental Microbiology 67, 5849–5854.