

# Microbial enzyme-catalyzed processes in soils and their analysis

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## ABSTRACT

Currently, measuring enzyme activities in soils or other lignocellulose-based materials is technically feasible; this measurement is particularly suitable for evaluating soil processes of biopolymer (cellulose, hemicelluloses, lignin, chitin and others) degradation by microbes and for assessing cycling and mobilization of principal nutrients including nitrogen, phosphorus and sulfur. With some considerations, assay methods can provide reliable information on the concentration of enzymes in soil or the rates of enzyme-catalyzed processes. Enzyme analyses in recent studies demonstrated a high level of spatial variability of soil enzyme activity both in depth and in space. The vertical gradients of enzyme activities are most developed in forest soils. Furthermore, enzyme activity in soils is regulated by seasonally-dependent variables such as temperature, moisture and the input of fresh litter. While several enzymes are widely produced by different groups of soil microorganisms, some of them can be used as indicators of the presence or activity of specific microbial taxa.

**Keywords:** assay methods; ecology; extracellular enzymes; fungi; lignocellulose; litter; soil microorganisms

Soil is an important component of all terrestrial ecosystems, as well as a main source of production in agriculture and forestry. Its function is essential for maintenance of the global biogeochemical cycles for all important nutrients, and thus, the processes in soils affect many other components of ecosystems, both biotic and abiotic. To understand the functioning of soils and to prevent soil damage due to anthropogenic and climatic factors, it is important to have suitable instruments for the assessment and quantification of soil processes performed by soil microorganisms and other members of the soil biota.

## Microbial processes in soils catalyzed by enzymes

Enzymes are major components of biological soil processes, such as the degradation of organic

compounds, their mineralization and the liberation or recycling of nutrients including nitrogen, phosphorus, sulphur and other essential metals. The activity of hydrolytic enzymes and ligninolytic oxidases and peroxidases directly affects the rates of transformation of soil biopolymers into compounds that are accessible for microorganisms and plants. The study of enzymatic activities in environmental samples (soil, litter, lignocellulose or other matrices) is a useful tool for assessing the functional diversity of soil microbial communities or soil organic mass turnover (Kandeler et al. 1999). Measuring enzyme activities in soil has a long tradition in connection with evaluating soil fertility and quantifying processes in natural and seminatural ecosystems with a high turnover of organic compounds, such as in forest and grassland soils. An overview of enzymes targeted in soils is summarized in Table 1. The list includes enzymes involved in the mobilization of nutrients, N, P or S,

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Supported by the DAAD-PPP scheme between the Czech Republic and Germany, Project No. D9-CZ17/08-09, by the Ministry of Education, Youth and Sports of the Czech Republic, Projects No. LC06066, OC155 and OC08050, by the Ministry of Agriculture of the Czech Republic, Project No. QH72216, and by the Institutional Research Concept of the Institute of Microbiology of the ASCR, v.v.i., Project No. AV0Z50200510.

from complex organic substrates and an intricate array of enzymes that participate in transforming biopolymers, including plant cell wall polymers such as cellulose and hemicelluloses along with other polysaccharides available in soils and litter. While the first group of enzymes is produced by a wide variety of soil microorganisms and some of them also secreted by plant roots, the production of several polymer-degrading enzymes is often ascribed to fungi (Miller et al. 1998, de Boer et al. 2005, Hättenschwiler et al. 2005, Steffen et al.

2007, Baldrian 2008a, Šnajdr et al. 2008a). Some ligninolytic enzymes, namely Mn-peroxidase and lignin peroxidase, are produced exclusively by saprotrophic species of fungi from *Basidiomycota*. Enzyme activity measurements can thus be an indirect assessment of the activity of a specific group of microorganisms in the soil (Hofrichter 2002, Baldrian 2008a).

Due to the effects of external disturbance on their activity, enzymes can serve as sensitive indicators of soil quality (Dick 1994, Dick et al. 1996,

Table 1. Overview of enzymes studied in soil and litter. Abbreviations: ABTS - 2,2'-azinobis-3-ethylbenzothiazoline-6-sulfonic acid; AMC – amidomethylcoumarin; DMAB - 3,3-dimethylaminobenzoic acid; MUF – methylumbelliferone; pNP – *p*-nitrophenol; VaOH – veratryl alcohol

Process	Enzyme <sup>1</sup>	EC	Assay	Method <sup>2</sup>	Reference
Cellulose degradation	endoglucanase	3.2.1.4	depolymerization	E	Lynd et al. (2002), Baldrian and Valášková (2008)
	cellobiohydrolase	3.2.1.91	pNP, MUF	D	Lynd et al. (2002), Baldrian and Valášková (2008)
	β-glucosidase	3.2.1.21	pNP, MUF	D	Lynd et al. (2002), Baldrian and Valášková (2008)
Degradation of hemicelluloses	endoxylanase	3.2.1.8	depolymerization	E	Collins et al. (2005), Biely and Puchart (2006)
	endomannanase	3.2.1.78	depolymerization	E	Collins et al. (2005), Biely and Puchart (2006)
	β-glycosidases	3.2.1.XX	pNP, MUF	D	Collins et al. (2005); Biely and Puchart (2006)
	esterases	3.2.1.XX	pNP, MUF	D	Biely and Puchart (2006)
Polysaccharide degradation	endochitinase	3.2.1.14	depolymerization	E	Seidl (2008)
	N-acetylglucosaminidase	3.2.1.52	pNP, MUF	D	Seidl (2008)
	α-glucosidase	3.2.1.3	pNP, MUF	D	Seidl (2008)
Lignin transformation	Mn-peroxidase	1.11.1.13	Mn <sup>2+</sup> , DMAB, ABTS	E	Hofrichter (2002)
	lignin peroxidase	1.11.1.14	VaOH	E	Martínez et al. (2005)
	laccase (phenoloxidase)	1.10.3.2	ABTS, guaiacol	E	Baldrian (2006)
	H <sub>2</sub> O <sub>2</sub> -producing enzymes	1.1.3.XX	various compounds	E	Martínez et al. (2005)
N acquisition	proteases	3.4.21.XX	depolymerization	E	Rao et al. (1998)
	aminopeptidases	3.4.11.XX	pNP, AMC	D	Kilcawley et al. (2002)
	urease	3.5.1.5	urea	D	Klose and Tabatabai (1999a)
P acquisition	phosphomonoesterase	3.1.3.1	pNP, MUF	D	Hayes et al. (2000)
	phosphodiesterase	3.1.4.1	pNP, MUF	D	Hayano (1977)
S acquisition	arylsulfatase	3.1.6.1	pNP, MUF	D	Klose and Tabatabai (1999b)

<sup>1</sup>For more details about the properties of enzymes, refer to [www.brenda-enzymes.info/](http://www.brenda-enzymes.info/) or (Schomburg and Schomburg 2007)

<sup>2</sup>D – direct; E – in extract

Marx et al. 2001). This relationship is emphasized by the fact that changes in enzymatic activity are the fastest response to external disturbances (Vepsäläinen et al. 2001). This observation was confirmed by many studies on the response of enzymes to environmental pollution, including the effects of heavy metals or organic xenobiotics (Kandeler et al. 1999, Baldrian et al. 2000, Burns and Dick 2002, Effron et al. 2004, Roy et al. 2004, Baldrian 2008b). It was proposed that a simultaneous estimation of multiple enzyme activities can be a suitable indicator of soil microbial activity (Bolton et al. 1985).

### **Methodological considerations for measuring enzyme activities in soils**

A variety of methods for measuring enzyme activities in soils have been proposed. The techniques often differ in the mode of detection (spectrophotometry, fluorescence, radiolabelling), the reaction conditions (temperature, use of buffers, time of reaction), and/or in the use of a variety of reaction substrates for measuring the enzyme activity, even for a single enzyme (Tabatabai 1994, Alef and Nannipieri 1995, Gianfreda and Bollag 1996, Schinner et al. 1996, Burns and Dick 2002). Unfortunately, generally accepted standard procedures still do not exist.

The most widely used substrates for the quantification of a relatively large group of exo-cleaving hydrolases are the molecules based on *p*-nitrophenol, 4-methylumbelliferone and amidomethylcoumarin. *p*-nitrophenol (pNP) is used for spectrophotometric techniques; the substrate exhibits a relatively low sensitivity, and the color reaction is inapplicable with dark or turbid samples. 4-methylumbelliferone (MUF) and amidomethylcoumarin (AMC) are substrates that are utilized with fluorescence detection. They are highly sensitive, although phenolic compounds (e.g. humic substances) can cause fluorescence quenching. Moreover, the fluorescence of MUF is strongly pH-dependent. The weaknesses of the methods using MUF or AMC can be overcome by choosing a suitable set of standards. Quantification of endo-cleaving polysaccharide hydrolases is usually assayed through the production of reducing sugars in the reaction or by analyzing the liberation of a soluble dye from a dyed polymer. With oxidative enzymes, the appearance of a colored product is typically measured using spectrophotometry.

Enzyme assays are either direct, when the reaction substrate is added to the studied system,

or indirect, when enzymes are recovered from soil and assayed afterwards. A direct assay overcomes any problems with enzyme extractability. A significant fraction of the enzyme is often not extractable from soils due to binding to soil components like microbial biomass (Schlosser et al. 1997, Valášková and Baldrian 2006) or abiotic soil material (Claus and Filip 1990, Quiquampoix et al. 1993). In soil with high clay content, the extractable fraction of total activity may account for less than a few percent of the total (Vepsäläinen 2001). Higher extraction efficiencies were reported from organic-rich horizons of forest soils and in plant litter (Šnajdr et al. 2008b). Measurements in extracts are better defined biochemically, but they only target the extractable fractions of enzymes. Unfortunately, unlike most hydrolases, enzymes performing endocleavage of polysaccharides (e.g. endoglucanases, endoxylanases or endochitinases) and oxidative enzymes can only be quantified in extracts. After enzyme recovery using extraction, desalting and dialysis should be performed to remove inhibitory small molecular mass compounds that may inhibit enzyme assay. These types of compounds include heavy metals (Baldrian and Gabriel 2002) or humic substances that competitively inhibit assays targeting lignolytic enzymes (Zavarzina et al. 2004).

A range of extraction buffers were proposed for use with soils (Lang et al. 1997, Criquet et al. 2000, Criquet et al. 2002, Baldrian and Gabriel 2003). Although some of them seem to work well with a wider variety of soils, when working in a single soil type, it is still advisable to test the extraction solutions with that specific soil. At minimum, the solutions to be tested should include phosphate and acetate buffers of pH 7, 0.1mM CaCl and distilled water.

Some authors use 'ambient conditions' in their enzyme assays, which means that the enzymes are tested at temperatures measured on site and in unbuffered solutions of substrates with the aim to reflect the *in situ* conditions and to measure the actual rates of enzymatic reactions (Burns and Dick 2002, Toberman et al. 2008). However, none of these techniques is able to completely reconstruct the local environment in terms of soil moisture content or osmolality, and its usefulness is thus questionable. It is recommended to use well-defined conditions in terms of pH (buffer composition) and temperature that result in biochemically-defined reaction kinetics. This type of experiment allows for a comparison of results across studies and in soils with different properties.

To approximate *in situ* conditions, it is possible to research the pH and temperature profiles of enzyme activities for the soil of interest and to use the results for estimating reaction rates.

Enzyme activity is extremely variable in space, the spatial variability usually being composed of vertical and horizontal components. The vertical gradient of soil properties includes soil organic matter content and quality, microbial biomass and physical condition (moisture content and pH). This results in a spatial gradient of enzyme activities, which are most often decreasing with increasing soil depth (Trasar-Cepeda et al. 2000, Prietzel 2001, Andersson et al. 2004, Wittmann et al. 2004, Baldrian et al. 2008, Šnajdr et al. 2008b). Vertical stratification of enzyme activities is particularly pronounced in highly stratified soils (forest soils) and less pronounced in soils under tillage (agricultural soils) or in soils with high activity of mixing soil biota, like earthworms. Sharp gradients are often found even within individual soil horizons at a centimeter scale (Šnajdr et al. 2008b). To overcome the effects of spatial variation, it is important to collect well-defined samples in terms of soil material (horizon) or depth (in centimeters) and to use composite samples. Previous studies showed that a set of five or six total samples, composed of three subsamples (soil core segments) can give the representative mean activity in forest or grassland soils (Baldrian et al. 2008, Šnajdr et al. 2008b).

In previous studies, seasonal climatic factors, such as temperature or soil moisture content, were also identified as the main factors responsible for

observed seasonal differences (Dilly and Munch 1996, Criquet et al. 2000, 2002, 2004, Wittmann et al. 2004, Niemi et al. 2005, Baldrian et al. 2008). This factor must be taken into account during the interpretation of enzyme activity data, and several sampling times during a season are usually needed to cover the annual variability of soil enzymatic processes. Moreover, to exclude random climatic effects, seasonality studies should be repeated during the following years.

### Recommended procedure

The following procedures are recommended for the routine measurement of enzyme activities in soils, litter or similar lignocellulose materials, such as naturally decomposing agroresidues. These methods were tested over a relatively wide range of soils with varying properties, vegetation and management practices, and they yielded satisfactory results (Vepsäläinen et al. 2001, Niemi and Vepsäläinen 2005, Baldrian et al. 2008, Šnajdr et al. 2008b).

**Direct Assay.** For the enzyme assays, 1 g fresh weight of soil is homogenized in 50 ml of 50mM sodium acetate buffer, pH 5.0, using an UltraTurrax (IKA Labortechnik, Germany) for 3 min at 8000 rev/min in an ice bath. For more solid litter particles, litter must be cut into pieces before homogenization. For the alkaline phosphomonoesterase assay, the reaction buffer is 0.5M tris-acetate at pH 8.0. When adding the reagents, the soil slurry is used at final dilution of 1:100.

Table 2. The enzymes determined by direct incubation of soil samples

Enzyme	Substrate
$\beta$ -Glucosidase	4-methylumbellyferyl- $\beta$ -D-glucopyranoside
Cellobiohydrolase	4-methylumbellyferyl-N-cellobiopyranoside
$\beta$ -Xylosidase	4-methylumbellyferyl- $\beta$ -D-xylopyranoside
N-Acetylglucosaminidase	4-methylumbellyferyl-N-acetylglucosaminide
$\alpha$ -Glucosidase	4-methylumbellyferyl- $\alpha$ -D-glucopyranoside
Esterase	4-MUF-acetate
Phosphomonoesterase	4-methylumbellyferyl-phosphate
Phosphodiesterase	bis-4-methylumbellyferyl-phosphate
Lipase	4-MUF-heptanoate
Arylsulfatase	4-methylumbellyferyl sulfate potassium salt
Alanine aminopeptidase	L-alanine-7-amido-4-methylcoumarin
Leucine aminopeptidase	L-leucine-7-amido-4-methylcoumarin

The activities of soil enzymes are assessed using specific artificial substrates (glycosynth, UK; Table 2) by measuring the fluorescence. Substrates (100  $\mu$ l, in DMSO), at a final concentration of 500  $\mu$ M, are added to 100  $\mu$ l of diluted soil slurry samples in a 96-well plate in three to four replicates. For the background fluorescence measurement, 100  $\mu$ l of the soil slurry is combined with 100  $\mu$ l of 4-methylumbelliferone (MUF) or 7-amino-4-methylcoumarin (AMC) standards to correct the results for fluorescence quenching. The standards are diluted to give final concentrations of 0, 0.1, 0.5, 1.0, 5.0, 10, 25 and 50  $\mu$ M in final volumes of 200  $\mu$ l (Vepsäläinen et al. 2001).

The multiwell plate is incubated at 40°C, and fluorescence is read after 5 min and 125 min. The fluorescence can be conveniently measured by any fluorescence reader, like the Infinite (TECAN, Austria), using an excitation wavelength of 355 nm and an emission wavelength of 460 nm (Niemi and Vepsäläinen 2005). The quantitative enzyme activities are calculated on the basis of fluorescence measurements. The fluorescence values, after subtraction of a blank, are compared with standard curves of MUF and AMC. If necessary, a substrate blank can also be used. One unit of enzyme activity is defined as the amount of enzyme forming 1  $\mu$ M of reaction product per min.

**Enzyme extraction and measurement in extracts.** The highest recovery of soil enzymes is usually achieved with 100mM phosphate buffer, pH 7.0 for soil and plant litter (Šnajdr et al. 2008a) or with 50mM sodium acetate buffer for specific lignocellulose types like wheat straw and other agroresidues (Lang et al. 1997). Homogenized samples of soil or litter material are extracted at 4°C for 2 h on an orbital shaker (100 rpm) with the appropriate buffer (16:1 w/v). They are then filtered through Whatman # 5 filter paper and desalted using PD-10 desalting columns (Pharmacia, Sweden), according to the manufacturer's protocol, to remove inhibitory small molecular mass compounds. The desalted samples can be stored at 4°C for two to three days or at -18°C for two to three weeks until enzyme activity analysis.

Laccase activity is measured by monitoring the oxidation of ABTS (2, 2'-azinobis-3-ethylbenzothiazoline-6-sulfonic acid) in citrate-phosphate (100mM citrate, 200mM phosphate) buffer (pH 5.0) at 420 nm (Bourbonnais and Paice 1990). Manganese peroxidase (MnP) is assayed using succinate-lactate buffer (100mM, pH 4.5). MBTH (3-methyl-2-benzothiazolinone hydrazone) and DMAB (3,3-dimethylaminobenzoic acid) are oxidatively coupled by

the enzyme, and the resulting purple indamine dye is detected spectrophotometrically at 595 nm. The results can be corrected by the activities of samples without manganese (for MnP), where the manganese sulfate is substituted with an equimolar amount of ethylenediaminetetraacetate (Ngo and Lenhoff 1980). One unit of enzyme activity is defined as the amount of enzyme forming 1  $\mu$ M of reaction product per min. Lignin peroxidase is usually not assayed in soils, which is due to the fact that the production of this enzymes is limited to basidiomycete fungi growing on wood. Their presence in soil is not anticipated (Hildén et al. 2008).

Endoglucanase, endoxylanase, endomannanase, endochitinase and proteases can be easily measured with azo-dyed carbohydrate substrates, including carboxymethyl cellulose, birchwood xylan, galactomannan (Megazyme, Ireland), azochitin or azocasein (Sigma, USA) following the protocols of the substrate suppliers. The reaction mixture contains 0.2 ml of 2% dyed substrate in 200mM sodium acetate buffer (pH 5.0) and 0.2 ml of the sample. The reaction mixture is incubated at 40°C for 20–120 min and the reaction is stopped by adding 1 ml of ethanol followed by 10 s of vortexing and 10 min of centrifugation (10 000  $\times$  g) (Valášková et al. 2007). The amount of released dye is measured at 595 nm, and the enzyme activity is calculated according to standard curves correlating the dye release with the release of reducing sugars.

Hydrolytic enzymes in extracts can be essentially measured using the same method as soil homogenate samples (see above). Alternatively, cellobiohydrolase,  $\beta$ -glucosidase,  $\beta$ -xylosidase and *N*-acetylglucosaminidase can be assayed using *p*-nitrophenyl- $\beta$ -D-cellobioside *p*-nitrophenyl- $\beta$ -D-glucoside, *p*-nitrophenyl- $\beta$ -D-xyloside and *p*-nitrophenyl-*N*-acetyl- $\beta$ -D-glucosaminide, respectively. The reaction mixture contains 0.16 ml of 1.2mM PNP-substrate in 50mM sodium acetate buffer (pH 5.0) and 0.04 ml of the sample. Reaction mixtures are incubated at 40°C for 20–120 min. The reaction is stopped by adding 0.1 ml of 0.5M sodium carbonate, and the absorbance is read at 400 nm (Valášková et al. 2007).

Acid phosphatase and arylsulfatase are assayed using 2 g/l *p*-nitrophenylphosphate or 50mM nitrophenylsulfate in 50 mM sodium acetate buffer (pH 5.0) (Tabatabai and Bremner 1970, Šnajdr et al. 2008a). One unit of enzyme activity is usually defined as the amount of enzyme releasing 1  $\mu$ M of *p*-nitrophenol per min. The *p*-nitrophenol-based assays can also be adopted for direct incubations under similar conditions (Baldrian et al. 2008).

However, the soils with a high content of organic matter interfere with the assay due to the release of soluble colored substances, and the sensitivity of this method is lower than that based on 4-methylumbelliferone by at least one order of magnitude.

### Other methods targeting microbial processes in soils

In addition to a direct analysis of enzyme activities, enzymes are also involved in processes assayed using substrate-induced respiration. In this method, a substrate is added to soil, and production of CO<sub>2</sub> or consumption of O<sub>2</sub> is measured. Mineralization of polymeric substrates can only be achieved by the action of extracellular enzymes (Alef and Nannipieri 1995). Recently, a convenient multiwell plate system was developed for the simultaneous measurement of substrate-induced respiration of numerous substrates at once. The assay can be used with both standard and radiolabeled compounds (Chapman et al. 2007). Enzymes are also targets of physiological profiling methods of soil communities based on substrate oxidation, such as the Biolog™ system (Garland and Mills 1991). However, this process is based on selection and cultivation of a subpopulation of utilizers of individual test compounds, and results relate more to the potential than to the actual rate of an *in situ* process. Moreover, it seems to be better suited for the study of individual microbial isolates compared to complex communities (Garland and Mills 1991, Preston-Mafham et al. 2002). Another physiological profiling device, the ApiZym™ system, is directly oriented to enzyme activities, but its applicability is limited solely to individual isolates of microorganisms. It has not yet come into routine use for microbial ecology and is more often used for typing of clinical isolates (Gruner et al. 1992).

Molecular methods may qualitatively target the abundance of enzyme-encoding genes (DNA analysis) or transcribed sequences (mRNA analysis). These approaches have been combined, for example, in the study of laccase sequences in hardwood forest soils (Luis et al. 2005a, Luis et al. 2005b). Cloning of PCR transcripts or microarrays containing target sequences are typically used in combination with DNA or RNA extraction. Microarrays targeting sequences for functional genes, the GeoChip, have come into use during the last few years, and they tend to target several thousands of functional genes that also cover a

significant portion of enzyme-catalyzed processes (He et al. 2007). Quantitative PCR of gene sequences or mRNA transcripts was recently used for quantification of some enzyme-encoding genes, including phenoloxidase-laccase (Hassett et al. 2009, Lauber et al. 2009). However, the connection between transcript or gene copy level and enzyme activity is indirect and is biased by genes encoding inactive molecules. Proteomic methods would definitely represent a suitable tool for a direct study of enzymes and their properties. However, they are still limited to qualitative analyses and are mostly reduced to *in vitro* cultures since soil seems to be much too complex for routine use. In the near future, molecular tools should enable us to link the diversity and quantity of the soil nucleic acid pool that encodes for enzymes with the identity of their producers, and these data will hopefully increase our understanding of the participation of individual microbes in the transformation of nutrients in soils.

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Received on July 1, 2009

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