

Biochemical fertility index and enzyme activity of mid-laterite soils in coconut-growing systems in Kerala (AEU 9)

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Research Article

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Abstract

Natural processes such as nutrient mineralization, organic matter production, soil fertility, and cycling are regulated by enzymes. The biological, chemical, and physical characteristics of the soil are closely related to them. The purpose of the experiment was to measure the levels of various enzymes found in soil in five different cropping systems involving coconut: Coconut+Fodder, Coconut+Banana, Coconut+Pepper, Coconut+Tuber, and Coconut+Vegetable. These systems were grown using either organic or conventional farming methods. The enzymes included dehydrogenase, acid phosphatase, glucosidase, protease, amylase, and catalase. The biologically productive agricultural system in Kerala's mid-laterite soils (AEU 9) is another important consideration. The levels of dehydrogenase, acid phosphatase, and catalase were higher in coconut + bananas grown under an organic farming system. The levels of glucosidase and amylase were higher in coconut + peppers grown under an organic farming system. The levels of protease were higher in coconut + tubers grown under an organic management system. Organic farming settings revealed that banana and coconut cropping were the most successful of all the cropping systems tested.

Keywords: coconut based cropping system, dehydrogenase, acid phosphatase, glucosidase, protease, catalase, amylase, enzyme activity number

1. Introduction

2. The primary sources of enzymes in soils include plants, soil, animals, and microbes that are covalently, cross-linked, copolymerized, adsorbed, and incorporated into the microcapsules of soil particles (Girish and Ajit, 2011) [12]. Because of their connection to soil biology, operational practicality, sensitivity, integrative nature, ease of measurement, and description as "biological fingerprints" of previous soil management, soil enzymes have been reported as helpful indicators of soil quality (Utobo and Tewari, 2014) [22]. The production, conversion, and breakdown of organic materials into forms that plants can digest, the breakdown of xenobiotics, the cycle of nitrogen and other elements, and the life cycle of soil microbes all depend on soil enzymes (Dick and Tabatabai, 1992) [9]. In order to assess soil quality, climate change, ecosystem damage, and toxification, soil enzyme activity have been used as indicators. Seasonal variations in soil enzyme activity have been linked to soil physiochemical characteristics, microbial community structure, vegetation, disturbance, and succession (Caldwell, 2005) [5]. One of the primary elements of soil enzymatic activities that participate in and guarantee the proper order of all the biochemical pathways in soil biogeochemical cycles are soil dehydrogenase enzymes. Because they are found intracellularly in all live microbial cells, dehydrogenases are among the most significant enzymes in the soil environment and are employed as a gauge of total soil microbial activity (Salazar et al., 2011) [17]. By moving protons and electrons from substrate to acceptors, the enzyme dehydrogenase oxidizes soil organic matter. Although this enzyme does not build up extracellularly in the soil, it is thought to be an essential component of intact cells (Das and Varma, 2011) [7]. Acid phosphatases play a significant role in the supply and metabolism of phosphate in plants by catalyzing the non-specific hydrolysis of inorganic phosphate from phosphate monoesters in pH ranges between 4 and 6 (Tabaldi et al., 2007) [19]. After hydrolysis and the release of free phosphate, organic phosphorus, which is plentiful in soil, may aid in the P feeding of microorganisms and plants. Phosphatase enzymes, which are released into soil, catalyze this process (Miller et al., 2001) [16]. Enzymes known as "phosphatases" that catalyze the hydrolysis of phosphoric acid esters and anhydrides are often responsible for the mineralization of P compounds in soil. Because it catalyzes the hydrolysis and biodegradation of different β -glucosidases found in plant waste decaying in the environment, Jn

B-glucosidase is a significant enzyme in soils (Martinez and Tabatabai 1997) [15]. An enzyme that breaks down cellulose is crucial to the soil organic carbon cycle (Esen, 1993) [11]. Protease activity plays a significant role in the ecology of microorganisms in the ecosystem and is a measure of the biological capacity of the soil for the enzymatic conversion of the substrate, regardless of the level of microbial activity (Burns, 1982) [4]. Soil protease, which is more active in forest soils and landfills with high humus and water contents, is important for nitrogen mineralization. Catalase breaks down peroxide, and its activity is influenced by the quantity of organic oxygen, microbial biomass, variations in CO₂, and the activity of dehydrogenase, amidase, glucosidase, and esterase in soils (Burns, 1982) [4]. In well-aerated conditions, soil catalase activity is increased (Brzezinska et al., 2005) [3]. Plants, animals, and microbes all produce α -amylases, although plants are the primary producers of β -amylase (Thoma et al., 1971) [21]. Widely found in plants and soils, amylase is an important component in the breakdown of starch, converting starch-like substrates to glucose and oligosaccharides, which in turn transforms starch to maltose (Thoma et al., 1971) [21].

3. Materials and Methods

4. In the mid-laterite soils of Kerala's Kollam district, soil samples were taken from five distinct coconut-based cropping systems under two farming techniques: organic and conventional (Agro Ecological Unit). AEU 9. Dehydrogenase, acid phosphatase, glucosidase, protease, catalase, and amylase activity were measured in each soil sample. The technique outlined by Casida et al., 1964 [6] was used to test the dehydrogenase activity. 3% 2, 3, 5-triphenyl tetrazolium chloride (TTC) was used to assess the dehydrogenase activity. Using triphenyl formazon (TPF) as a standard, a standard graph was created to determine the concentration of dehydrogenase in the sample. μg of TPF emitted g^{-1} soil 24 h^{-1} was used to express the enzyme activity. The technique described by Tabatabai and Bremner (1969) [20] was used to test the acid phosphatase activity of soils. 0.05 M p-nitrophenyl phosphate was used to evaluate the acid phosphatase activity. On a dry weight basis, the enzyme activity was measured in μg of p-nitrophenol emitted g^{-1} soil h^{-1} at 37.0°C and pH 6.5.

Using 0.5 M p-nitrophenyl β -glucopyranoside, the Eivazi and Tabatabai (1988) [10] technique was used to estimate the activity of glucosidase in soil samples. The glucosidase activity was measured in μg pnp D-glucosidase g^{-1} soil $\text{h}^{-1} \times 10^{-4}$. The Kunitz M. (1947) [14] technique was used to assess the soil protease activity, which is expressed in μM of amino nitrogen hydrolyzed g^{-1} soil h^{-1} . The Bach A.N. and Zubkova S.M. method was used to measure soil catalase activity, which was represented as a percentage of H₂O₂ hydrolyzed g^{-1} soil h^{-1} . The method for determining amylase activity in soil samples was described by Bernfeld (1955) [2].

Based on the activity of five distinct enzymes—dehydrogenase, catalase, acid phosphatase, protease, and amylase—the Biological Fertility Index for the various treatment combinations was calculated using the enzyme activity number established by Beck (1984). The formula was used to calculate the Enzyme Activity Number for each treatment. EAN is equal to $0.2\{\text{TPF} + \text{catalase} (\%) / 10 + \text{phenol} (\mu\text{g}) / 40 + \text{amino-N} (\mu\text{g}) / 40 + \text{amylase} (\%) / 20\}$.

Statistical Analysis

The data generated from the experiments were subjected to the Analysis of variance as per the design, Factorial CRD and their significance was tested using F test (Snedecor and Cochran, 1975) [18]. Critical difference (CD) was calculated at 0.05% probability levels using statistical tools.

5. Results and Discussion

6. According to the research, the coconut+banana cropping system under organic farming had the maximum dehydrogenase activity (462.52 μg of TPF released g^{-1} of soil 24 h^{-1}) (Table 1). Higher dehydrogenase activity may have resulted from an increase in the number of microorganisms, particularly bacteria, in the soil caused by an organic farming method. Wells et al. (2000) [24] found a similar rise in dehydrogenase activity with the addition of FYM/vermicompost, whereas the traditional farming system of Coconut + Tuber had the lowest dehydrogenase activity (101.25 μg of TPF released g^{-1} of soil 24 h^{-1}) (Fig. 1). The organic Coconut + Banana farming system has the greatest acid phosphatase activity (80.35 μg of p-nitrophenol emitted g^{-1} of soil 24 h^{-1}).

1) (Table 1 & Fig. 2). The low available P status may be the reason for the maximum acid phosphatase activity seen in these treatments. According to Versaw and Harrison (2002) [23], acid phosphatase secretion from plant roots increases if there is a signal signaling a P deficit in the soil. This enhances the solubilization and remobilization of PO_4 , hence altering the capacity to deal with P-stressed conditions. Coconut + Pepper ($4.1 \mu\text{g pnp D-glucosidase g}^{-1}\text{soil h}^{-1} \times 10^{-4}$) and Coconut + Tuber had the greatest β -glucosidase activity (Table 1). The organic farming approach was generally shown to be favorable for β -glucosidase activity. The findings that microbial development favors the release of β -glucosidase in this kind of environment (Dick, 1994) [8] and that the traditional farming system of Coconut+Vegetable showed the lowest β -glucosidase activity ($1.53 \mu\text{g pnp D-glucosidase g}^{-1}\text{soil h}^{-1} \times 10^{-4}$) further confirm this. (Fig. 3). Protease activity was greatest in the Coconut+Tuber cropping system ($188.84 \mu\text{M}$ of amino nitrogen hydrolyzed g^{-1} of soil h^{-1}) (Table 1 & Fig. 4). This might be explained by the greatest levels of water-holding capacity, organic carbon, mineralizable nitrogen, and accessible K. Soil fertility and the quantity of aerobic microorganisms have been linked to catalase activity, which is regarded as an indication of aerobic microbial activity. Catalase activity was lowest in Coconut + Vegetable (Fig. 6) and greatest in Coconut + Banana (2.85% of H_2O_2 hydrolyzed g^{-1} of soil h^{-1}) cropping system (Table 1), which was comparable to Coconut + Pepper (2.64% of H_2O_2 hydrolyzed g^{-1} of soil h^{-1}). According to the experiment, Coconut+Pepper had the maximum amylase activity ($13.82 \mu\text{M}$ of maltose g^{-1} of soil) under the organic management method (Table 1 & Fig. 5). Similar patterns were noted by Griffiths et al. (1999) [13] when examining the impact of different C substrate rates on amylase activity. Regarding the EAN as suggested by Beck (1984) [1], Table 2 shows that the primary impacts of Coconut + Banana (94.01) under an organic way of agriculture had the greatest EAN. The study clearly shows that the highest values of catalase (3.9% of H_2O_2 hydrolyzed g^{-1} of soil h^{-1}), acid phosphatase ($80.35 \mu\text{g}$ of p-nitrophenol released g^{-1} of soil 24 h^{-1}), and dehydrogenase ($462.52 \mu\text{g}$ of TPF released g^{-1} of soil 24 h^{-1}) observed in Coconut+Banana under organic system may have contributed to the highest enzyme activity.

Table 1: Enzyme activities of the soils of coconut based cropping systems AEU 9

Treatments	Dehydrogenase (μg of TPF released g^{-1} of soil 24 h^{-1})	Acid phosphatase (μg of p nitro phenol released g^{-1} of soil 24 h^{-1})	Glucosidase ($\mu\text{g pnp D-}$ glucosidase g^{-1} soil $\text{h}^{-1} \times 10^{-4}$)	Protease (μM of amino nitrogen hydrolysed g^{-1} of soil h^{-1})	Catalase ($\%$ of H_2O_2 hydrolysed g^{-1} of soil h^{-1})	Amylase (μM of maltose g^{-1} of soil)
Coconut + Fodder (C ₁)	221.46	52.44	3.06	84.94	2.02	8.98
Coconut + Banana (C ₂)	326.92	56.23	2.51	165.84	2.85	9.44
Coconut + Pepper (C ₃)	276.42	42.19	3.64	149.7	2.64	9.7
Coconut + Tuber (C ₄)	212.87	55.95	3.27	174.98	2.32	9.16
Coconut + Vegetable (C ₅)	184.3	45.76	2.12	172.17	1.96	9.01
Organic Farming (F ₁)	344.49	59.89	3.37	163.22	3.05	12.31
Conventional Farming (F ₂)	144.29	41.14	2.47	135.84	1.67	6.21
C ₁ F ₁	311.83	56	3.51	99.34	2.54	9.8
C ₂ F ₁	462.52	80.35	2.8	186.37	3.9	12.52
C ₃ F ₁	368.38	47.26	4.1	161.46	3.54	13.82
C ₄ F ₁	321.49	58.68	3.75	188.84	2.74	12.7
C ₅ F ₁	258.22	57.13	2.72	180.07	2.52	12.7
C ₁ F ₂	131.08	48.87	2.61	70.54	1.5	8.16
C ₂ F ₂	191.32	32.11	2.23	145.32	1.8	6.36
C ₃ F ₂	184.45	37.11	3.18	137.95	1.74	5.58
C ₄ F ₂	101.25	53.2	2.79	161.11	1.9	5.62
C ₅ F ₂	110.37	34.37	1.53	164.27	1.4	5.32
CD- C (0.05)	21.66	6.29	0.14	19.05	0.23	0.25
CD- F (0.05)	13.69	3.98	0.06	12.05	0.14	0.16
CD- C F(0.05)	30.63	8.91	0.14	NS	0.32	0.35

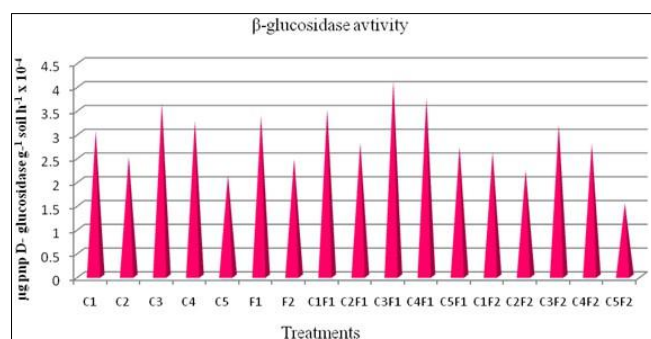


Fig 3: β -glucosidase activity of the soils under coconut based cropping system AEU 9

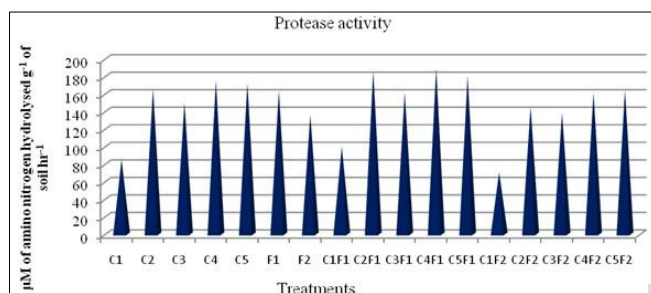


Fig 4: Protease activity of the soils under coconut based cropping system AEU 9

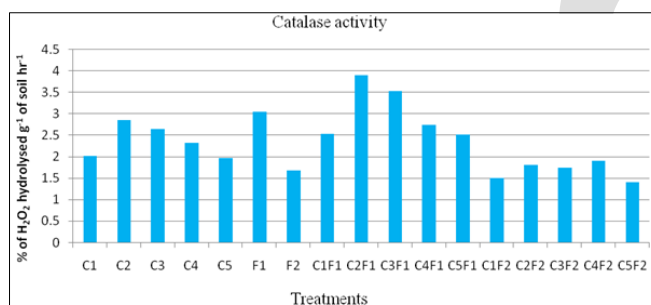


Fig 5: Catalase activity of the soils under coconut based cropping system AEU 9

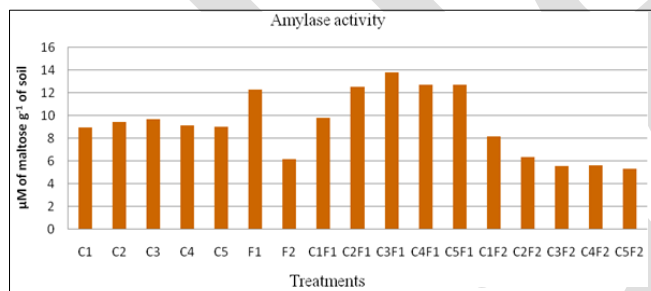


Fig 6: Amylase activity of the soils under coconut based cropping system AEU 9

7. Conclusion

8. The research found that out of all the cropping and farming systems tested, the organic method produced the most biologically fertile and long-term sustainable coconut and banana crops. This was due to the fact that the organic system had the highest enzyme activity numbers (EAN) and the highest levels of dehydrogenase, acid phosphatase, and catalase.

9. References

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