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Importance of considering enzyme degradation for interpreting the response of soil enzyme activity to nutrient addition: insights from a field and laboratory study

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Abstract: Soil enzyme activity can be affected by both production and degradation, as enzymes can be degraded by proteases. However, the impact of nutrient addition on enzyme activity is often solely attributed to changes in enzyme production without fully considering degradation. Here, we demonstrated that the activities of β -1,4-glucosidase (BG), β -D-cellobiohydrolase (CBH), β -1,4-Xylosidase (BX), and β -1,4-N-acetyl-glucosaminidase (NAG) were comparable in nitrogen (N) and phosphorus (P) fertilized soils and the unfertilized control under field conditions, but the reduction in activity was substantially greater in the fertilized soils during short-term laboratory incubation. The results show that the interruption of the natural, continuous supply of organic matter or non-soil microbial-derived enzymes, which typically occurs under field conditions, leads to a more significant reduction in soil enzyme activities in fertilized soils compared to unfertilized control. This may be attributed to the higher abundance of protease in fertilized soils, resulting in faster enzyme degradation. Interestingly, P fertilization alone did not have a similar effect, indicating that N fertilization is likely the main cause of the larger decreases in enzyme activity during incubation in fertilized soils compared to unfertilized control soils, despite our study site being poor in P and rich in N. These findings highlight the importance of considering enzyme degradation when investigating material dynamics in forest ecosystems, including the impact of nutrient addition on enzyme activity, as enzyme production alone may not fully explain changes in soil enzyme activity.

Keywords: enzyme degradation; nitrogen; phosphorus; protease; forest soil; tropics

1. Introduction

In forest ecosystems, soil extracellular enzymes play essential roles in nutrient cycling and carbon (C) dynamics [1–4]. Litters deposited on the forest floor provide nutrients to the soil, but it is the enzymes that break down the organic matter, allowing organisms to reuse the nutrients. Soil extracellular enzymes are predominantly synthesized by soil microorganisms and plant roots, but leaf litter [5] or throughfall [6–8] can also contribute to the pool of soil extracellular enzymes.

In ecosystems where multiple enzymes coexist, the activity of soil extracellular enzymes (i.e., V_{max}) can serve as an indicator of enzyme abundance [9,10]. This enzyme

abundance reflects the net result of both enzyme production and degradation, as soil extracellular enzymes can be degraded by proteases [11,12]. However, the degradation of these enzymes is rarely measured, despite the fact that the degradation rate of enzymes is often used in ecosystem models. Allison [13] measured the degradation of enzymes by monitoring changes in enzyme activity using gamma-irradiated soils where enzyme production by microorganisms was halted. Schimel et al. [11], on the other hand, used chloroform fumigation to halt microbial enzyme production and observed enzyme decay ratios. Both studies reported a significant decrease in enzyme activity during the incubation period, typically within a few weeks, but the degree of decrease varied among different enzymes. Assuming dynamic equilibrium, these results suggest that a similar amount of enzymes is being continuously produced in the field as the amount that is degraded. In field conditions, plants continuously supply resources such as C and nutrients through litters and root exudates. Additionally, plants also provide enzymes to soils through litter layers [5] or throughfall [6,8]. However, the degree to which these resources or enzyme supplies contribute to the soil enzyme pool remains unknown. The first objective of this study is to investigate the extent to which enzyme activity decreases in an incubation conducted under laboratory conditions where these resource supplies are disrupted.

Changes in the availability of nutrients, such as phosphorus (P) and nitrogen (N), strongly influence soil enzyme activity, as well as other ecosystem processes such as primary production [14,15], soil respiration [16–18], and litter decomposition [19–21]. Therefore, numerous studies have investigated the effect of nutrient addition on soil enzyme activity, and multiple meta-analyses have been published demonstrating this impact [22–29]. For example, a meta-analysis has shown that N fertilization can stimulate enzymes related to N mineralization [26], whereas it has also been demonstrated that protein- and chitin-degrading enzymes may be suppressed by N fertilization due to a reduced need for N acquisition [30,31]. P addition has generally been found to suppress phosphatase activity [23,24]. On the other hand, it has been observed that phosphatase activity is often stimulated by N addition, which is attributed to the increased N supply available for synthesizing phosphatase [28,32,33]. Oxidative enzymes have been shown to be suppressed by N fertilization [23,25], but the precise mechanisms are still under discussion [34,35]. Thus, the impact of nutrient addition on enzyme activity has been extensively investigated to better understand material dynamics in forest ecosystems. However, it is important to note that the effects of nutrient addition on enzyme activity have often been attributed solely to changes in enzyme production without fully considering degradation [3], despite the fact that most of the studies have attributed the changes in enzyme activity affected by nutrient addition to changes in enzyme production. The assumption that changes in enzyme activity account for the impacts of nutrient addition on soil enzyme production can lead to a serious misinterpretation. Therefore, the second objective of our study is to demonstrate that the impact of nutrient addition on soil enzyme activity cannot be solely attributed to changes in enzyme production.

To achieve these objectives, we first tested how enzyme activity changes during incubation when there is a disruption in the continuous supply of organic matter or non-soil microbial-derived enzymes. Second, we compared the changes in enzyme activity during incubation in fertilized and unfertilized soils.

2. Materials and Methods

Study site

The study was conducted at the Heshan National Field Research Station of Forest Ecosystems in southern China [36]. Two 33-year-old plantation forests were chosen for the experiment, namely a leguminous acacia plantation (AA), dominated by *Acacia auriculiformis* A.Cunn. ex Beth. with the occasional occurrence of *Acacia mangium* Willd., and a non-leguminous eucalyptus plantation (EU), dominated by *Eucalyptus urophylla* S.T. Blake. Experimental plots were established in both forests. The soil type of both

plantations is acrisols, and the annual temperature and mean annual precipitation are 21.7 °C and 1580.4 mm, respectively [36]. This site has a high N deposition rate through rainfall, with an average of 43.1 ± 3.9 kg N ha⁻¹yr⁻¹ from July 2010 to June 2012 [37]. Table 1 shows the basic information of the study site.

Table 1. Basic information of the study site.

Properties	AA	EU
Soil total organic C (g kg ⁻¹) ^{a,c}	22.1 (2.3)	15.5 (0.9)
Soil total N (g kg ⁻¹) ^{a,c}	1.6 (0.1)	1.5 (0.1)
Soil NH ₄ ⁺ (mg kg ⁻¹) ^{a,c}	16.0 (1.0)	13.4 (2.0)
Soil NO ₃ ⁻ (mg kg ⁻¹) ^{a,c}	17.8 (1.4)	13.6 (2.6)
DOC (mg C kg ⁻¹) ^b	275.6 (3.5)	297.9 (3.6)
DN (mg N kg ⁻¹) ^b	38.5 (1.1)	41.8 (2.2)
Soil available P (mg kg ⁻¹) ^{a,c}	1.8 (0.2)	1.6 (0.3)
Soil pH ^{a,c}	3.83 (0.03)	3.91 (0.05)
Fine root biomass (g m ⁻²) ^{a,c}	92.9 (13.6)	74.4 (3.2)
Standing litter-layer mass (g m ⁻² yr ⁻¹) ^c	694.1 (25)	590.2 (14)
Basal area (m ² ha ⁻¹) ^d	12.5	16.7
Mean height of planted trees (m) ^e	12.2	11.5
DBH of planted trees (cm) ^e	15	11.1
Stem density (tree ha ⁻¹) ^e	2076	1962
Litterfall mass (g m ⁻² yr ⁻¹) ^e	841 (58)	870 (67)

AA, forest stands dominated by *Acacia auriculiformis* A.Cunn. ex Beth. with the occasional occurrence of *Acacia mangium* Willd. EU, forest stands dominated by *Eucalyptus urophylla* S.T. Blake. C, carbon. N, nitrogen. DOC, dissolved organic carbon. DN, dissolved nitrogen. DBH, Diameter at breast height.

^a Soil samples were obtained from surface soil at 0–10 cm depth both for chemical analysis and fine root biomass determination.

^b Data from Mori et al. 2021 [38]. Values are means with standard errors in parentheses (five replication of a composite soil).

^c Data from Zhang et al., 2012 [39]. Values are means with standard errors in parentheses (n=3).

^d Data from Zhang et al., 2012 [39]. The values were determined before the start of the fertilization experiment.

^e Data from Zhang et al., 2014 [37]. Litterfall mass data were taken from control plots.

Experimental design

In July 2010, a fertilization experiment was initiated with seven different treatments that involved the addition of N and/or P as amendments [37]: control with no fertilization (Con), medium-N fertilized (MN, 50 kg N ha⁻¹yr⁻¹), high-N fertilized (HN, 100 kg N ha⁻¹yr⁻¹), medium-P fertilized (MP, 50 kg P ha⁻¹yr⁻¹), high-P fertilized (HP, 100 kg P ha⁻¹yr⁻¹), medium-NP fertilized (MNP, 50 kg P ha⁻¹yr⁻¹ and 50 kg N ha⁻¹yr⁻¹), and high-NP fertilized (HNP, 100 kg P ha⁻¹yr⁻¹ and 100 kg N ha⁻¹yr⁻¹). Each plot consisted of three replicates and was 10 m × 10 m in size. P-fertilized plots were treated with sodium biphosphate (NaH₂PO₄) spray, and N-fertilized plots were treated with ammonium nitrate (NH₄NO₃). Both NaH₂PO₄ and NH₄NO₃ were dissolved in water and treated bi-monthly. Con plot was received commensurable water spray only. Nutrient addition began in August 2010 and is still ongoing. For this study, we focused on three treatments only: Con (hereafter Con-plots), HP (hereafter P-plots), and HNP (hereafter NP-plots). The basic chemical and biological properties in these three plots were shown in Table 2.

Soil sampling

In December 2016, six years after the commencement of the experiment, 0-5 cm soil samples were collected from ten randomly selected points in each plot (Con-plots, P-plots and NP-plots of AA and EU stands) using a soil corer (inner diameter 3.5 cm). The collected soil samples were then processed by passing them through a 2-mm sieve, with the removal of large organic matter. Soil pH was measured using a pH meter (METTLER TOLEDO FE20) according to a soil:water ratio of 1:2.5, and already reported by Wang et al. (2020) [36]. Soil pH values in Con-plots, P-plots, and NP-plots were 3.59 ± 0.035 , 3.68 ± 0.082 , and 3.59 ± 0.037 , respectively in AA stands, and 3.56 ± 0.044 , 3.69 ± 0.028 , and 3.53 ± 0.032 , respectively in EU stands [36]. Water contents of the soil samples in Con-plots, P-plots, and NP-plots were 0.23 ± 0.004 , 0.22 ± 0.004 , and 0.22 ± 0.007 , respectively in AA stands, and 0.23 ± 0.017 , 0.22 ± 0.007 , and 0.22 ± 0.007 , respectively in EU stands [36].

Table 2. Impacts of fertilization on soil chemical and biological properties.

Stand	Properties	Con-plot	P-plot	NP-plot
AA	Soil total organic C (g kg ⁻¹)	40.7(3)	45.3(4)	55.8(4)
	Soil total N (g kg ⁻¹)	2.2(0.1)	2.2(0.2)	2.0(0.2)
	C:N ratio	18.5 (1)	18.9 (3)	27.9 (3)
	Soil NH ₄ ⁺ (mg kg ⁻¹)	9.4(0.5)	11.9(0.7)	12.2(0.8)
	Soil NO ₃ ⁻ (mg kg ⁻¹)	7.7(0.9)	6.6(0.4)	11.3(1.0)
	MBC (µg C g ⁻¹)	330(31)	634(38)	446(34)
	MBN (µg N g ⁻¹)	67(12)	86(17)	52(14)
	Soil available P (mg kg ⁻¹)	2.9(0.3)	4.1(0.5)	4.0(0.1)
EU	Soil total organic C (g kg ⁻¹)	20.9(3)	33.9(2)	33.6(3)
	Soil total N (g kg ⁻¹)	1.6(0.1)	1.6(0.3)	1.7(0.1)
	C:N ratio	13.1 (2)	21.2 (2)	19.8 (1)
	Soil NH ₄ ⁺ (mg kg ⁻¹)	6.7(0.2)	5.2(0.8)	6.9(0.7)
	Soil NO ₃ ⁻ (mg kg ⁻¹)	5.6(0.5)	4.2(0.7)	6.0(0.6)
	MBC (µg C g ⁻¹)	378(33)	359(26)	350(20)
	MBN (µg N g ⁻¹)	78(8)	47(12)	80(10)
	Soil available P (mg kg ⁻¹)	2.6(0.1)	4.1(0.4)	4.0(0.5)

AA, forest stands dominated by *Acacia auriculiformis* A. Cunn. ex Beth. with the occasional occurrence of *Acacia mangium* Willd. EU, forest stands dominated by *Eucalyptus urophylla* S.T. Blake. C, carbon. N, nitrogen. P, phosphorus. MBC, microbial biomass carbon. MBN, microbial biomass nitrogen. Data were cited from Zhang et al. (2014) [37]. Soil samples were obtained from surface soil at 0–10 cm depth in July 2012. Values are means with standard errors in parentheses (n=3).

Enzyme assay of the sampled soils

Enzyme assays of β -1,4-glucosidase (BG), β -1,4-xylosidase (XYL), cellobiohydrolase (CBH), and β -1,4-N-acetyl-glucosaminidase (NAG) performed using the fluorescence enzyme assay described by Bell et al. [40] with minor modifications was already reported by Wang et al. [36]. Briefly, 1.5 g of fresh soil was added to 91 mL of acetate buffer (50 mM, pH 5.0) and blended for 1 min until well homogenized. Two-hundred microliters of the substrates labeled with 4-methylumbelliferone (MUB) (Table 3) were added to 800 µL of the homogenized sample solution in deep 96-well plates. The plates were incubated in the dark at 20 °C for 4 hours, followed by centrifugation at $2,090 \times g$. Then, 250 µL of the supernatant from the incubated mixture of sample and substrates were transferred into black 96-well plates. We measured the fluorescence of the incubated solution using a microplate spectrophotometer (Infinite M200 PRO) with a 365 nm excitation fil-

ter and a 450 nm emission filter. Enzyme activity was then calculated using standard lines prepared for each sample.

Table 3. Extracellular enzymes, abbreviation, and substrate used in the present study.

Enzyme	Abbreviation	Substrate
β-1,4-Glucosidase	BG	MUB-β-D-glucosid
β-1,4-Xylosidase	XYL	MUB-β-D-cellobioside
Cellobiohydrolase	CBH	MUB-β-D-xyloside
β-1,4-N-acetylglucosaminidase	NAG	MUB-N-acetyl-β-D-glucosaminide
MUB, 4-methylumbelliferone.		

Incubation and enzyme assay of the incubated soils

Soil samples from Con-plots, P-plots, and NP-plots (3.0 g each) were placed in plastic bottles. After the addition of 0.9 mL deionized water, the bottles were covered with polyethylene rap for preventing the evaporation of water [41] and put into the incubator. Soils were incubated for 7.5 days at 25 °C in the dark. After the end of the incubation, soils were frozen at -20 °C until the enzyme assay. Enzyme activity of the incubated soils was measured using the same methodology as enzyme assay of the sampled soils (i.e., measurement of soil enzyme activity at the field condition), but a different amount of soil (3.0 g).

Calculation and Statistics

The apparent enzyme decay rate was determined using first-order fitting, as shown by the equation:

$$E_{aft}= E_{bef} \times e^{-k}$$

where E_{aft} represents enzyme activity after incubation, E_{bef} represents enzyme activity before incubation, k is the apparent decay rate constant, and t is the incubation time (i.e., 7.5 days). It is important to note that the calculated k does not indicate the true decay rate constant of the enzyme, as enzymes are continually produced during incubation. Therefore, k should be smaller than the actual decay rate constant of the enzyme. We used a linear mixed-effects model to examine the effect of incubation time and fertilization on enzyme activity, with subplot as a random effect. We calculated the changes in enzyme activities before and after incubation and compared the values among Con-plots, P-plots, and NP-plots using one-way ANOVA, followed by Tukey’s HDS. All statistical analyses were performed using R version 4.2.2 [42].

3. Results

The enzyme activities in AA stands tended to decrease, particularly for BG, as suggested by a linear mixed-effects model analysis (Fig. 1). The apparent enzyme decay rate of BG, XYL, and NAG was 0.038, 0.00038, and 0.010, respectively, in Con-plots of AA stands. Meanwhile, the CBH activity in Con-plots of AA stands was elevated and could not be calculated (Fig. 1). In NP-plots of AA stands, all four enzymes had significantly lower activity than Con-plots or P-plots (Fig. 1), resulting in higher apparent decay rates (0.083, 0.083, 0.16, and 0.12 for BG, XYL, CBH, and NAG, respectively). In contrast, lower enzyme activities compared to Con-plots were not observed in P-plots of AA stands. In P-plots of AA stands, the apparent decay rates were 0.014, 0.0043, and 0.045 for BG, XYL, and NAG, respectively, and CBH activity was elevated and not calculable.

In EU stands, the decreases in enzyme activities tended to be larger compared to AA stands. In Con-plots of EU stands, the apparent enzyme decay rate of BG, XYL, CBH, and NAG was 0.047, 0.069, 0.10, and 0.053, respectively. However, due to the larger variability of data, these decreasing trends did not have statistical significance. Similar to AA

stands, enzyme activities tended to be higher in NP-plots, but the results were statistically insignificant due to large variances among the data (Fig. 2). The apparent decay rates in NP-plots of EU stands were 0.096, 0.10, 0.23, and 0.11 for BG, XYL, CBH, and NAG, respectively, and 0.03, 0.03, 0.08, and 0.04, respectively, in P-plots of EU stands.

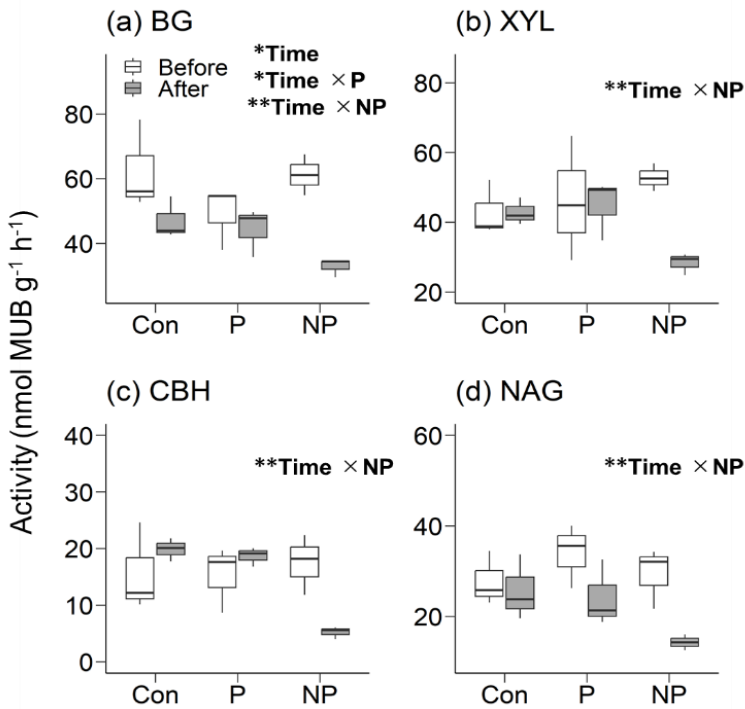


Fig. 1. The enzyme activity of β -1,4-glucosidase (BG), β -1,4-xylosidase (XYL), cellobiohydrolase (CBH), and β -1,4-N-acetyl-glucosaminidase (NAG) measured before and after a 7.5-day incubation in soils taken from Con-plots, P-plots, and NP-plots of AA stands. MUB, 4-methylumbelliferone.

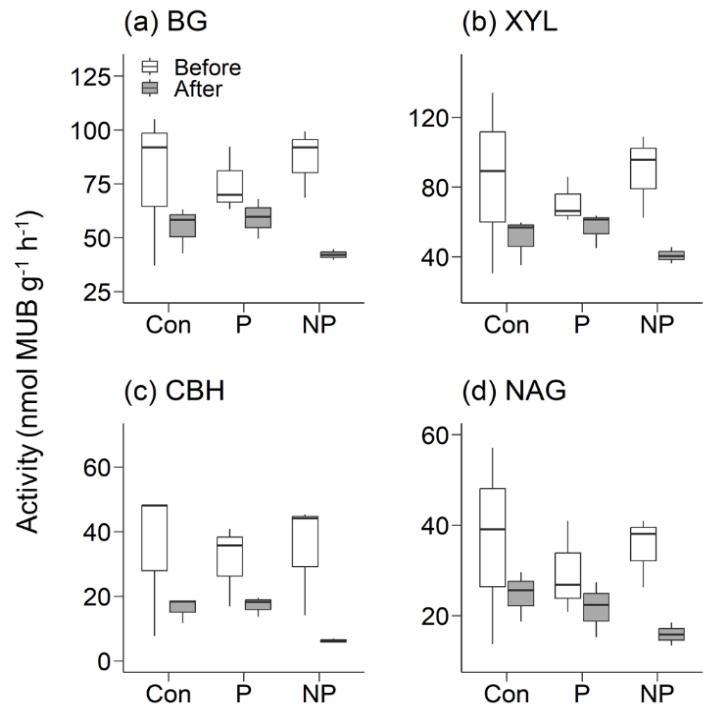


Fig. 2. The enzyme activity of β -1,4-glucosidase (BG), β -1,4-xylosidase (XYL), cellobiohydrolase (CBH), and β -1,4-N-acetyl-glucosaminidase (NAG) measured before and after a 7.5 day incubation in soils taken from Con-plots, P-plots, and NP-plots of EU stands. MUB, 4-methylumbelliferone.

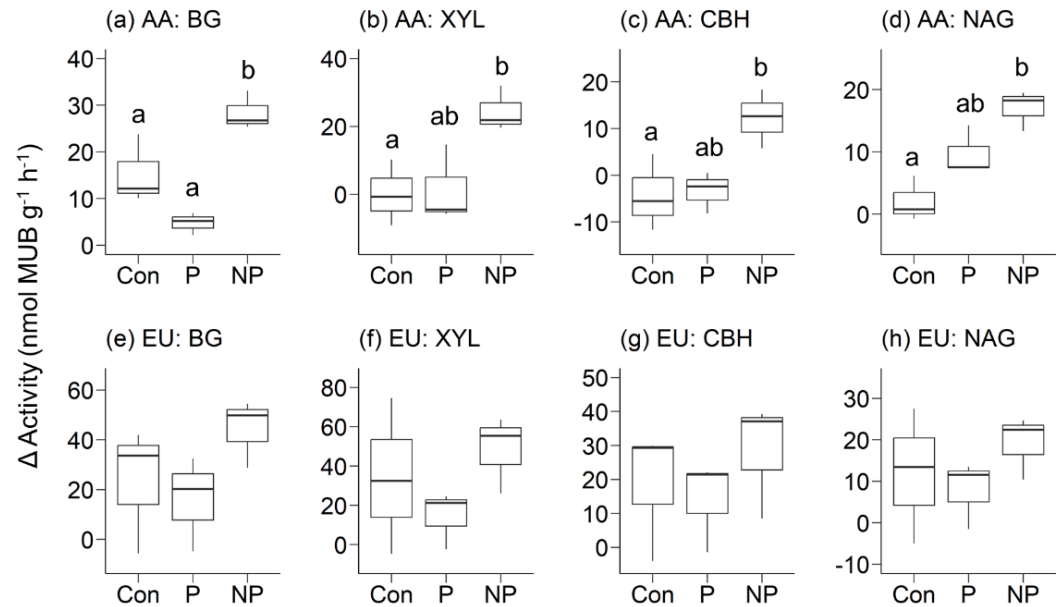


Fig. 3. The differences in enzyme activity values before and after the 7.5-day incubation (Δ Activity) of (a) β -1,4-glucosidase (BG), (b) β -1,4-xylosidase (XYL), (c) cellobiohydrolase (CBH), and (d) β -1,4-N-acetyl-glucosaminidase (NAG) in AA stands, and (e) BG, (f) XYL, (g) CBH, and (h) NAG in EU stands. MUB, 4-methylumbelliferone.

Differences in enzyme activity values before and after incubation were significantly larger in NP-plots in AA stands for all four enzymes: BG, XYL, CBH, and NAG (Fig. 3a-d). In EU stands, the averages of the differences in enzyme activity values before and after incubation were consistently highest in NP-plots, but the differences were not statistically significant (Fig. 3e-h).

4. Discussion

In our study, microorganisms continuously produced enzymes during incubation, because we did not inhibit soil microbial activity through gamma-irradiation [13] or chloroform fumigation [11]. However, we observed high decay rates of BG and NAG in Con-plots. The values were 0.038 and 0.047 in AA stands, and 0.010 and 0.053 in EU stands for BG and NAG, respectively, which were higher than those reported for most soils by Schimel et al. [11], where microbial activity had been excluded by chloroform fumigation. Our findings indicate that the activity of at least some enzymes could decrease significantly under laboratory conditions, despite the continuous production of enzymes by microorganisms during incubation. This decrease in enzyme activity in laboratory conditions could be attributed to the interruption of the natural, continuous supply of organic matter or enzymes provided by microorganisms in areas other than the soil, such as the litter layer[5] or phyllosphere [6], which occur in field conditions. This highlights the significance of a continuous supply of organic matter and non-soil microbial-derived enzymes in maintaining enzyme activity in forest soils.

In ecosystems where multiple enzymes coexist, the activity of soil extracellular enzymes (i.e., V_{max}) can serve as an indicator of enzyme abundance [9,10]. Therefore, various studies have utilized the ratios of different enzyme activities as indicators of microbial resource allocation for nutrient acquisition[33,43–47] (it should be noted that several

studies questioned this approach [48–53]). In these studies, accurately determining enzyme production by microorganisms, rather than enzyme activity, is essential because microbial resource allocation to obtain nutrients should be related to microbial resource allocation to produce enzymes. Our results demonstrate that enzyme activity (or enzyme pools) are maintained by a continuous supply of organic matter and non-soil microbial-derived enzymes in forest soils. Therefore, relying solely on soil enzyme activity measurements may lead to misleading conclusions regarding microbial resource allocation for nutrient acquisition.

Under field conditions in both AA and EU stands (i.e., before incubations as shown in Fig. 1 and 2), the activities of BG, XYL, CBH, and NAG were similar between NP-plots and Con-plots. However, a significant reduction in activity was observed during short-term laboratory incubation in AA stands, with a higher reduction in NP-plots than in Con-plots (Fig. 1, 2a-d). In EU stands, a similar trend was observed, although it was statistically insignificant due to high variability. These results show that the impact of interrupting the natural, continuous supply of organic matter or non-soil microbial-derived enzymes, which occurs under field conditions, on enzyme activities was higher in NP-plot soils compared to Con-plot soils. It is possible that the higher quantity of protease in NP-plot soils, as compared to the Con-plot soils, may have led to faster degradation of enzymes in the NP-plot soils. Because P fertilization alone did not have a similar effect observed in the NP-plot soils, it is likely that N fertilization is the main cause of the substantially larger decreases in enzyme activity during incubation in NP-plots compared to Con-plots. If this is the case, our findings challenge the assumption that the impacts of N fertilization are less crucial than those of P fertilization in tropical forest soils that are often considered N-rich and P-poor [16,54] (but see [55]). Future research should aim to determine the impact of nutrient addition on pure turnover rate of enzymes by excluding enzyme production by soil microorganisms, as previous studies have done [11,13].

5. Conclusion

In this study, we demonstrated that the activities of hydrolysable enzymes, specifically BG and NAG, decreased rapidly during incubation when the natural and continuous supply of organic matter or enzymes provided by microorganisms in areas other than the soil were disrupted. This highlights the importance of a continuous supply of organic matter and non-soil microbial-derived enzymes in maintaining enzyme activity in forest soils. Therefore, caution is necessary when assessing microbial resource allocation for nutrient acquisition (or energy) by measuring enzyme activity, as the measured values may not reflect enzyme production accurately.

We also demonstrated that the activities of four hydrolysable enzymes, i.e., BG, XYL, CBH, and NAG, were comparable in N and P fertilized soils and the unfertilized control under field conditions, but decreases in those activities were substantially greater in the fertilized soils during short-term laboratory incubation. Since a similar pattern was not observed in P-fertilized soils, our results suggest that the interruption of the continuous supply of organic matter or non-soil microbial-derived enzymes in the field leads to a more significant reduction in enzyme activities in N-fertilized soils compared to unfertilized control.

In conclusion, the present study highlights the importance of considering enzyme degradation when investigating material dynamics in forest ecosystems, including the impact of nutrient addition on enzyme activity, as soil enzyme activity may not always reflect enzyme production.

Author Contributions: Conceptualization, T.M.; software, T.M. and S.W.; validation, T.M. and S.W.; formal analysis, T.M., C.P. and S.W.; investigation, T.M., C.P., C.W. and S.W.; resources, W.Z. and J.M.; data curation, T.M., C.W., M.Z. and S.W.; writing—original draft preparation, T.M.; writing—review and editing, T.M., C.P., M.Z. and W.Z.; visualization, T.M.; supervision, W.Z. and

J.M.; project administration, W.Z.; funding acquisition, W.Z. All authors have read and agreed to the published version of the manuscript.

Funding: This study was financially supported by the National Natural Science Foundation of China (42077311, 42173077), Grant-in-Aid for JSPS Postdoctoral Fellowships for Research Abroad (28-601), and a grant from The Sumitomo Foundation (153082).

Data Availability Statement: Data are provided by corresponding author when requested.

Acknowledgments: We thank Shengxing Fu and Meifang Hu for their support for our field work.

Conflicts of Interest: The authors declare no conflict of interest. The funders had no role in the design of the study; in the collection, analyses, or interpretation of data; in the writing of the manuscript; or in the decision to publish the results.

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