



CHANGES IN GROWTH AND CHEMICAL DEFENCES UPON DEFOLIATION IN MAIZE

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Abstract—The effect of defoliation on growth and on levels and allocation patterns of hydroxamic acids (Hx) in maize seedlings was evaluated 6 days after treatment. No significant differences were found between defoliated and nondefoliated plants for the Hx concentration, relative Hx content and Hx-aglucone to Hx-glucoside ratio in shoots, roots and root exudates, with the exception of Hx concentration in shoots, which decreased upon defoliation. However, growth of defoliated seedlings was considerably higher than that of nondefoliated ones. These results indicate that maize responds to defoliation by allocating its resources mainly to growth rather than to defence. Since previous work described the opposite strategy in rye, responses to defoliation in both species are discussed in relation to current theories of plant defence. © 1998 Elsevier Science Ltd. All rights reserved

INTRODUCTION

Hydroxamic acids (Hx) present in wild and cultivated Poaceae [1, 2] play a role in plant resistance against herbivores and pathogens [3–5] and in allelopathy [6, 7]. Hx occur mainly as glucosides which are converted to the more active aglucones [8] by enzymatic hydrolysis upon tissue damage [9].

Research on plant responses induced by insect feeding or wounding has focused mainly on the increase in concentration of defensive compounds [10]. The induction of Hx after insect attack or localised artificial damage has been widely described [11–15], while their induction upon defoliation, a process analogous to grazing or harvesting practices, has been only recently reported [16, 17]. Collantes *et al.* [17] showed that rye (*Secale cereale* L.) seedlings responded to defoliation not by an increase in Hx concentration of aerial tissue but by: (i) an increase of exudation of Hx by the roots, (ii) increased relative allocation of Hx to roots and root exudates and (iii) transformation of Hx-glucosides into more toxic aglucones. The present work addresses the induction of Hx upon defoliation in maize (*Zea mays* L.) and compares it with the patterns found in rye [17].

RESULTS AND DISCUSSION

Quantification of DIMBOA in maize seedlings

The most abundant hydroxamic acid in maize is 2- β -O-D-glucopyranosyl-4-hydroxy-7-methoxy-1,4-benzoxazin-3-one (DIMBOA-Glc) [1]. Both DIMBOA-Glc and its aglucone were found in shoots and roots of maize seedlings, while in root exudates only the aglucone was found, in agreement with a previous report by Pérez and Ormeño [6]. The concentration of total hydroxamic acids in both nondefoliated and defoliated seedlings was considerably higher in shoots than in roots and in roots than in root exudates (Table 1). The same patterns were found for Hx in rye under the same experimental conditions [17].

Induction of HX upon defoliation

The induction of Hx upon defoliation was evaluated by assessing the following parameters: (i) Hx concentration (mmol/kg fresh-weight) in shoots, roots and root exudates (considered as the Hx content of root exudates divided by the fresh weight of roots), (ii) relative Hx content (% of Hx content in the whole plant) in shoots, roots and root exudates and (iii) Hx aglucone/glucoside ratio in shoots and roots. Results for defoliated and nondefoliated seedlings were compared by a one-way ANOVA.

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Table 1. Hydroxamic acids concentration (mmol/kg fresh-weight) in maize seedlings $\pm$ S.E.

	Nondefoliated	Defoliated	<i>P</i> -value*
Shoot	26.5 $\pm$ 1.2	12.1 $\pm$ 0.6	< 0.0001
Root	1.70 $\pm$ 0.21	1.28 $\pm$ 0.09	0.064
Exudates	0.03 $\pm$ 0.01	0.02 $\pm$ 0.01	0.39

*One-way ANOVA.

Nonsignificant differences were found between defoliated and nondefoliated seedlings for all assessed parameters except for the Hx concentration in shoots (Tables 1–3). Defoliated seedlings showed lower concentrations of Hx than nondefoliated ones. These results indicate that Hx in maize is not induced by defoliation, contrary to what occurs with Hx in rye under the same experimental conditions [16, 17].

Compensatory growth of defoliated seedlings

Maize seedlings were weighed before extraction. Shoot biomass (mg) of defoliated seedlings (747.78 $\pm$ 36.13) was significantly higher ($P < 0.0001$, one-way ANOVA) than that of nondefoliated ones (287.55 $\pm$ 20.08). Thus, while showing no Hx induction, maize seedlings subjected to defoliation experienced overcompensatory growth (*sensu* Paige and Whitham [18]). On the contrary, Hx induction was found in defoliated rye seedlings, which showed in addition subcompensatory growth, under the same experimental conditions [17].

The predominant investment of rye in chemical defence and of maize in compensatory growth as a consequence of defoliation may be attributed to the C₃ nature of rye and C₄ nature of maize [19]. Thus, on one hand C₄ plants are poorer food sources for herbivores than C₃ plants and hence should invest less resources in defence [20] and, on the other, according to the optimal defence theory, given a limited amount of resources, plants tend not to invest simultaneously in alternative functions (e.g. growth and defence) [21–23]. Moreover, since maize is a crop species which has been largely modified by breeding in contrast to rye [24], the overcompensatory growth and lack of induction of chemical defences of maize upon defoliation may be reflecting past selection aiming at increased crops yields. In addition, given that phytochemical induction is a widespread phenomenon in wild plants [10], our

Table 2. Hydroxamic acids relative content (% of whole plant content) in maize seedlings $\pm$ S.E.

	Nondefoliated	Defoliated	<i>P</i> -value*
Shoot	80.8 $\pm$ 2.3	82.9 $\pm$ 0.9	0.37
Root	18.7 $\pm$ 2.4	16.9 $\pm$ 0.9	0.47
Exudates	0.41 $\pm$ 0.10	0.23 $\pm$ 0.03	0.067

*One-way ANOVA.

Table 3. Hydroxamic acid aglucone to glucoside ratio in maize seedlings $\pm$ S.E.

	Nondefoliated	Defoliated	<i>P</i> -value*
Shoot	0.37 $\pm$ 0.08	0.49 $\pm$ 0.10	0.36
Root	0.22 $\pm$ 0.07	0.34 $\pm$ 0.01	0.089

*One-way ANOVA.

results may also reflect the erosion of genetic variability in maize through breeding [25].

EXPERIMENTAL

Plant material

Seeds of *Z. mays* cv. T55s from Tracy Seed, Chile, were germinated in individual plastic pots (100 ml) filled with sterilised sand (Anasac). Seedlings were kept in a room at 25 $\pm$ 3°C and 16:8 h light regime and irrigated with a nutritious solution (Anasac) containing N:P:K (6:4:3) and micronutrients. Half of the seedlings were defoliated 6 days after germination by cutting the plants just above the coleoptile. Nondefoliated and defoliated seedlings (seven seedlings per treatment) were evaluated 6 days after sowing and defoliation, respectively.

Extracts for analysis

Seedlings were carefully drawn from the sand to avoid root damage. Roots were washed with distilled water directly onto the pots to remove sand particles. Seedlings were weighed and then shoots and roots were separated and immediately macerated using mortar and pestle with *ca.* 300 mg sea sand in 1 ml 0.1 M glycine-HCl buffer pH 2. Plant parts were macerated immediately in order to avoid spurious conversion of Hx glucosides into aglucones due to the scission of the plant. Consequently, the weight of tissues was not directly determined but estimated from linear regressions of total plant weight vs shoot weight ($R^2 = 0.98$ for nondefoliated and $R^2 = 0.89$ for defoliated) and total plant weight vs root weight ($R^2 = 0.99$ for nondefoliated and $R^2 = 0.96$ for defoliated). The sand in the pots was washed with 100 ml distilled water to obtain root exudates. The washing solution was evaporated to dryness under vacuum at 45°C and the dry residue extracted with 1 ml *n*-BuOH. All extracts were centrifuged at 10,400g for 15 min and the supernatants stored in a freezer until analysis.

Chromatography

A 100 ml aliquot of each extract was directly injected into a HPLC fitted with an RP-100 LiChrospher-C18 column (5 mm i.d., Merck). Conditions were constant solvent flow (1.5 ml min⁻¹), the following linear gradients between solvents A (MeOH) and B (0.5 ml 85%

H₃PO₄ in 1 l H₂O): 0–9 min 30% A, 9–11 min 100% A, 11–15 min 30% A and detection at 263 nm. Both DIMBOA-glucoside and its aglucone were analysed. Only the aglucone was found in root exudates. Retention times for aq. extracts were 3.7 ± 0.1 and 4.8 ± 0.1 min for DIMBOA-glucoside and DIMBOA aglucone, respectively, and 1.9 ± 0.1 min for the DIMBOA aglucone in *n*-BuOH. Identity of the peaks was confirmed by coinjection of standards dissolved in water and *n*-BuOH, respectively.

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