



# *Root herbivore performance suppressed when feeding on a jasmonate-induced pasture grass*

Article

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# 1 Short Communication

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## 3 Root herbivore performance suppressed when feeding on a 4 jasmonate induced pasture grass

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19 **Running title:** Jasmonate suppression of root herbivores

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21

### 22 Abstract

- 23 1. Plants defend themselves from insect herbivore attack using a range of physical and chemical defences which  
24 are in many cases regulated by phytohormones such as jasmonates. While much more is known about how  
25 jasmonates regulate defence against aboveground herbivores (e.g. herbivores of leaves), there is increasing  
26 interest in how they influence belowground defences.
- 27 2. For the Poaceae, most belowground studies focus on highly domesticated cereals. Here we demonstrate how  
28 exogenous application of methyl jasmonate (MeJA) to the leaf blades of a non-domesticated pasture grass  
29 (*Microlaena stipoides*) caused a more than two-fold decrease in relative growth rates (RGR) of a root-feeding  
30 chafer (*Dermolepida albohirtum*). MeJA treatment did not affect root consumption rates, but substantially  
31 reduced the efficiency of conversion of ingested food to body mass.
- 32 3. Non-targeted metabolomics identified significant changes in the metabolome of MeJA-induced plants, with  
33 three compounds (a galactolipid, a trihydroxy fatty acid and a lysophospholipid) found to be correlated with  
34 herbivore RGR, although their roles in herbivore defence remain uncertain.

35 4. This study suggests that an important Australian pasture grass can become better defended against root  
36 herbivores via enhanced jasmonate activity.

37 **Key Words** – belowground herbivore; grass; jasmonic acid; metabolomics; *Microlaena stipoides*; root herbivory.

38

## 39 **Introduction**

40 In the last decade, we have seen significant advances in our understanding of how plant hormones regulate plant  
41 defences against insect herbivores (Wu & Baldwin, 2010). Jasmonates are recognised as key regulators of chemical  
42 pathways that activate genes associated with herbivore defence aboveground (Jander & Howe, 2008). Compared to  
43 herbivores feeding aboveground, we know comparatively less about how jasmonate-regulated defences operate  
44 belowground against root herbivores (Erb *et al.*, 2012a; 2012b). While this situation is improving, the majority of  
45 studies addressing phytohormone induction and root herbivores are mainly concerned with domesticated plants,  
46 particularly cereals (Erb *et al.*, 2012a; Erb *et al.*, 2013; Lu *et al.*, 2015). These studies suggest that jasmonates, while  
47 less inducible than in the shoots, regulate root resistance to herbivores. Lower induction of jasmonates indicate that  
48 defensive machinery of the roots may be more sensitive of jasmonate activity or synergistic signals are involved.  
49 Salicylic acid, for example, which can dampen jasmonate-based defence, does not appear to be induced by root  
50 herbivores so its deficiency may indirectly boost jasmonate activity (Johnson *et al.*, 2016). In addition to being crop  
51 pests, root herbivores can be important within natural ecosystems because of their sheer abundance. The collective mass  
52 of pasture scarab beetles, for example, can outweigh that even of domesticated mammals grazing aboveground when  
53 considered per unit area (Frew *et al.*, 2016a).

54

55 A first step towards understanding jasmonate-regulated defences against root herbivores in pasture grasses would be to  
56 establish whether exposure of plant tissues to jasmonsates affects herbivore performance. Exogenous application of  
57 methyl jasmonic acid (MeJA) is a common approach for achieving this (Erb *et al.*, 2013; Lu *et al.*, 2015). We  
58 investigated whether MeJA application to an important pasture grass native to Australia, *Microlaena stipoides* (syn.  
59 *Ehrharta stipoides*), affects the performance and feeding behaviour of a root-feeding chafer insect. We undertook non-  
60 targeted metabolomic screening of MeJA-treated and untreated root tissue to explore whether changes in the root  
61 metabolome were related to root herbivore performance.

62

## 63 **Methods and Materials**

64 *Plants*

65 Forty weeping meadow grass (*Microlaena stipoides*) plants were grown from seed (Native Seeds, VIC, Australia) in all-  
66 purpose potting mix (Richgro, WA, Australia) in pots (70 mm diameter, 135 mm deep) for c. 32 weeks. The study was  
67 conducted in a naturally-lit glasshouse chamber (3 × 5 × 3 m; width × length × height) with UV transparent plexiglass  
68 walls and roof. Air temperature was regulated at 30 °C (±4 °C) and fell to 22 °C (±4 °C) at night. Humidity was  
69 controlled at 60 % (±6 %). Plants were irrigated with c. 70mL of water three times a week and provided with a single  
70 dose (2g) of Osmocote Controlled Release fertilizer. Twenty plants were selected at random and were treated with  
71 MeJA (95%, Sigma Aldrich) by applying 1mL of a solution 100µg/mL MeJA (Sigma Aldrich) in Tween 20 ± 0.1% to  
72 the leaf blades, just above the soil surface, with a pipette. Control plants received 1mL of Tween 20 ± 0.1% only. This  
73 was repeated 24 and 48 hours later. Plants were then separated from the soil and were washed free of soil for feeding  
74 assays.

75

#### 76 *Feeding Assays*

77 We used the greyback chafer (*Dermolepida albohirtum*) as a model grass-feeding root herbivore. This species is native  
78 to Australia and feeds in pastures but has also become a significant pest of sugarcane (Frew *et al.*, 2016a). Feeding  
79 assays were conducted as described by Frew *et al.* (2016b). In summary, individual third instar larvae were starved for  
80 24 h, weighed and placed in a Petri dish (14 cm diam) with c. 4 g of fresh root material taken from one of 10 plants  
81 treated with MeJA or 10 control plants, each selected at random. Larvae fed for 24 h, then starved for 12 h to ensure all  
82 frass was expelled, before being reweighed. Evaporative water loss from roots during the assay was accounted for  
83 (Frew *et al.*, 2016b).

84

85 Relative growth rate (RGR) of larvae was calculated as: mass gained (g)/ initial mass (g)/time (days). Relative  
86 consumption (RC) is an estimate of the mass of root material ingested over the 24 h period relative to initial body mass.  
87 It was calculated as: food ingested (mg change in dry mass) / mean body mass over experimental period (mg fresh  
88 mass). Efficiency of conversion of ingested food (ECI) was calculated as: mass gained (mg change in fresh body  
89 mass)/food ingested (mg change in dry mass) × 100.

90

#### 91 *Metabolomics sample preparation*

92 Root material not used for the feeding assays was snap frozen before being freeze dried and ball-milled to a fine  
93 powder. Five glass beads were added to 20 mg of each ground root sample, which was then shaken in a tissue lyser for  
94 4 min in 500 µL methanol:milliq water:formic acid (75:24.5:0.5 v/v) at 30 Hz. This was centrifuged and the supernatant  
95 was extracted for analysis.

96 Untargeted profiling was carried out by ultra-high performance liquid chromatography-quadrupole time-of-flight mass  
97 spectrometry according to a protocol adapted from Gaillard *et al.* (2018). The separation was performed at a flow rate of  
98 0.5 mL/min using a Waters Acquity UPLC HSS T3 column (2.1 × 100mm, 1.7 µm particle size) maintained at 40°C.  
99 The following gradient of mobile phase A (water+0.05% formic acid) and B (acetonitrile+0.05% formic acid) was  
100 applied: 0-60% B in 6.0 min, 60-100% B in 2.5 min, holding at 100% for 2.0 min and reequilibration at 0% B for 3.0  
101 min. The injection volume was of 2.5 µL. Mass spectrometric detection was performed in negative electrospray over a  
102 mass range of 85-1200 Da using independent data acquisition (MS<sup>E</sup>). A pool of all samples was injected as quality  
103 control using the following sequence: two injections at the beginning of the batch, then one injection every 20 samples  
104 and at the end of the batch. Peak selection was performed as described by Gaillard *et al.* (2018). Compounds that were  
105 significantly up/down-regulated by JA were further selected for identification. Putative identification was achieved on  
106 the basis of (i) determination of molecular formulae from accurate mass measurements and (ii) mass spectral  
107 fragmentation characteristics and comparison with existing databases.

108

#### 109 *Statistical analysis*

110 RGR was log transformed and analysed with a one-way ANOVA with MeJA-induction as the fixed factor. RC and ECI  
111 were analysed with Kruskal-Wallis tests. Compounds of potential interest relating to MeJA-induction were initially  
112 identified using partial least squares discriminant analysis (PLS-DA). Compounds with a PLS-DA PC1 score > 0.1 (i.e.  
113 the compounds that contribute to the majority of the dataset variance associated with MeJA-induction) were selected,  
114 before being analysed with one-way ANOVAs. Potential associations between concentrations of these compounds and  
115 RGR were examined with Pearson's correlation tests. All analysis was conducted with the R statistical package.

116

## 117 **Results and Discussion**

118 The RGR of larvae feeding on MeJA-induced grass roots was significantly lower than for those feeding on untreated  
119 plants (Fig. 1a;  $F_{1,18} = 12.24$ ,  $P = 0.003$ ), with most larvae losing mass when feeding on MeJA-induced plants. There  
120 was no significant difference in RC with herbivores eating similar amounts of root tissue from MeJA-induced and non-  
121 treated plants (Fig. 1b;  $H_1 = 1.463$ ,  $P = 0.226$ ). As a result of this, ECI was much lower ( $-27.15 \pm 12.12$ ) when feeding  
122 on MeJA-induced plants compared to non-treated plants ( $1.18 \pm 15.68$ ) (median values  $\pm$  inter-quartiles;  $H_1 = 10.57$   $P <$   
123  $0.001$ ). In short, herbivores were feeding at similar rates on MeJA-induced and non-induced *M. stipoides*, but were  
124 unable to obtain adequate nutrition when feeding on MeJA-induced plants. This indicated that herbivores were not  
125 deterred from feeding on MeJA-induced plants (at least in the short-term) but their performance was being adversely  
126 affected by consuming such root tissue, which is at least compatible with enhanced production of defensive compounds

127 in the roots. Assays that aim to quantify feeding metrics of insect herbivores are often standardised to 24 hr (e.g.  
128 Slansky, 1985; Frew *et al.*, 2016b), which was sufficient to detect suppression of herbivore performance in this study.  
129 We should, however, be cautious about extrapolating our findings over longer periods since larval *D. albobirtum* can  
130 live for over a year (Frew *et al.*, 2016a) and it is unknown how long the effects of MeJA-induction persist for in *M.*  
131 *stipoides*. Nonetheless, exogenous MeJA application to rice resulted in comparable reductions in the growth of root  
132 herbivores (cucumber beetle, -50% and rice water weevil, -100%) when feeding for 7 and 20 d, respectively (Lu *et al.*,  
133 2015).

134

135 In total, 14 primary and secondary metabolites in the roots of *M. stipoides* either increased or decreased in response to  
136 MeJA-induction, three of which were significantly, though not strongly, correlated with root herbivore RGR (Table 1).  
137 MeJA-induction caused concentrations of a galactolipid (a digalactosylmonoacylglyceride, 18:2-0:0-DGMG;  
138  $C_{33}H_{58}O_{14}$ ) to decrease, and this compound was positively correlated with RGR (Table 1). Galactolipids, including their  
139 aldehydes, can be major sources of essential fatty acids (Ohlsson, 2000) but can sometimes act as defences against  
140 insect and echinoderm herbivores (Deal *et al.*, 2003). MeJA-induction also caused significant increases in  $C_{19}H_{36}O_5$ , (a  
141 monounsaturated trihydroxy fatty acid) and  $C_{27}H_{52}NO_9P$  (a linoleoyl-lyso phosphocholine), both of which were  
142 negatively correlated with insect RGR (Table 1). Unsaturated fatty acids can be toxic to insects in bioassays (Harada *et*  
143 *al.*, 2000), including root herbivores (Bernklau *et al.*, 2016). However, a saturated trihydroxy fatty acid, phloionolic  
144 acid ( $C_{18}H_{36}O_5$ ) is a known constituent of suberin in cork (Pereira, 2007), and the monounsaturated  $C_{19}H_{36}O_5$  trihydroxy  
145 fatty acid observed here may therefore reflect an increase in suberinisation of roots. Mono- and  
146 digalactosyldiacylglycerols also play roles in the regulation of systemic acquired resistance in plants (Gao *et al.*, 2014).  
147 Perhaps the best candidate to explain the decreased herbivore performance is the linoleoyl lysophosphocholine, as some  
148 compounds in this class have been shown to be highly cytotoxic to human cancer cell lines (Niezgoda *et al.*, 2015).

149

150 To our knowledge, this is the first study to show that exogenous application of MeJA to a non-domesticated pasture  
151 grass has negative impacts on a root herbivore. Using metabolomics, we established that the concentrations of three  
152 compounds both changed in response to MeJA-induction and were correlated (either positively or negatively) with root  
153 herbivore performance. The exact structures of these compounds could not be resolved, and regardless, are relatively  
154 unstudied so it is unproven whether these are linked to the observed declines in herbivore performance and feeding  
155 efficiency. Further work and different analytical approaches such as detailed lipid profiling may shed light on this.

156 Nonetheless, we were able to establish that MeJA-induction impaired root herbivore growth and thus resulted in roots  
157 that were potentially better defended; this was previously unknown for non-domesticated grasses. In reporting these

158 findings, we aim to stimulate further interest in characterising belowground plant defences in non-domesticated as well  
159 as domesticated grass species.

160

161 **Author contributions.** Project design: SNJ, IH, BDM. Data collection and statistical analysis: JMWR. Metabolomics  
162 analysis: GG. All authors contributed manuscript preparation.

163

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