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1	Body distribution of toxic peptides in larvae of a pergid and an argid sawfly		
2	species		
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28 Abstract

29 Larvae of most Pergidae and Argidae (Symphyta: Tenthredinoidea) species contain toxic peptides such as pergidin and lophyrotomin. Here, larval hemolymph and organs of the 30 pergid Lophyrotoma zonalis and the argid Arge pagana were analysed by liquid 31 chromatography-tandem mass spectrometry. The major identified peptides were pergidin 32 and 4-valinepergidin in L. zonalis, whereas pergidin and lophyrotomin in A. pagana. The 33 storage period prior to chemical analysis was longer for the samples of the pergid than the 34 argid species, which influenced peptide concentrations. In both species, however, the 35 36 peptides occurred in decreasing order of concentration, first in the hemolymph, then in the integument, while minor amounts of the peptides were detected in other organs such as gut 37 and fat body. By separating the cuticle of the pergid from the remaining integument, the 38 peptides were found in equivalent amounts in each of these two body structures. The results 39 40 suggest that the peptides play an important role in the defence of these sawfly larvae against predators. 41 42 43 Keywords Lophyrotoma zonalis; Arge pagana; hemolymph; integument; LC-MS/MS; pergidin; lophyrotomin 44

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47 Introduction

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49 Insects that use chemical defence against predators store the harmful compounds in specific areas of the body (Dettner 2015). Specialized epidermal exocrine glands are often the site 50 51 both of production and storage of chemicals produced de novo (Noirot and Quennedey 1974). If an insect sequesters chemicals from its food, the chemicals are in the gut before 52 53 passing through the gut epithelium, to be stored in the hemolymph or other compartments (Opitz and Müller 2009). Endosymbionts can also produce compounds used by their host in 54 anti-predator defence. The evidence of such systems remains restricted to the genus 55 Paederus Fabricius, 1775 (Coleoptera: Staphylinidae) harbouring Pseudomonas Migula, 1894 56 57 bacterial symbionts that produce the toxin pederin and that are acquired by cannibalism (Kellner 2003; Maleki-Ravasan et al. 2019). 58

59 Argidae and Pergidae (Symphyta: Tenthredinoidea) sawfly larvae have been studied by 60 analytical chemistry since the 1970s (Leonard 1972; Oelrichs et al. 1977). Most argid and pergid species are known to contain some or all of the peptides: pergidin (Perg, molecular 61 weight (MW, in g/mol) = 864), 4-valinepergidin (VPerg, MW = 850), dephosphorylated 62 pergidin (dpPerg, MW = 784), and two closely related peptides both called lophyrotomin 63 (LGIn, MW = 1039, and LGIu, MW = 1040) (Boevé et al. 2014). The biosynthetic origin of the 64 sawfly peptides remains unknown, but they are most probably produced by endosymbionts 65 (Oelrichs et al. 1999). In the pergid *Lophyrotoma interrupta* (Klug, 1814), the oral discharge 66 of an oily fluid contains about ten times the concentration of lophyrotomin compared to 67 whole larvae (Oelrichs et al. 1992). The hemolymph, integument and gut were shown to be a 68 paralytic feeding deterrent by testing their extracts from the argids Arge pagana (Panzer, 69 1797) and Arge pullata (Zaddach, 1859) against ant workers of Myrmica rubra (Linnaeus, 70 71 1758), which indirectly indicates the presence of peptides (Petre et al. 2007). Nearly no peptides are detected in the excrements of Arge berberidis Schrank, 1802 (Boevé and 72 73 Rozenberg 2019). Aside from these fragmentary data, to date no analysis has been done to 74 determine the precise location of peptides within the body. 75 Here, we localized the peptides within the larval body of *L. zonalis* (Rohwer, 1910) and *A.* 76 pagana by collecting hemolymph and dissecting organs. The samples were analysed by

liquid chromatography-tandem mass spectrometry (LC-MS/MS). The results are discussed in
an ecological context. Some practical questions about storing samples over a long period are
also discussed.

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82 Materials and Methods

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Sawfly larvae from two populations were collected in the field: one of *L. zonalis* (30 March
2008; Brisbane, Australia; by JLB), the other of *A. pagana* (16 September 2016; Aalter,
Belgium; by Filip De Block). Voucher specimens are kept at the Royal Belgian Institute of
Natural Sciences (collection reference codes: P2865 and P4207, respectively). The larvae
were weighted on an analytical balance at 0.1-mg precision (Kern ABS 120–4, Kern & Sohn
GmbH, Balingen, Germany). Hemolymph samples were collected from live larvae and stored
in 100% ethanol. The hemolymph of *L. zonalis* was sticky and collected with bits of filter

91 paper. That of A. pagana was collected with glass capillaries and weighed. Body part samples 92 were obtained from larvae that were frozen, thawed, and dissected in distilled water to 93 isolate the following organs: the digestive tract, or gut, the integument (plus the attached longitudinal muscles), and additional organs such as salivary glands and fat body. Peptide 94 quantities and concentrations per L. zonalis larva were unknown in 2008, because 95 procedures of single-larva extraction and LC-MS/MS analysis of the peptides had not yet 96 been developed. Thus, we opted on the side of caution for this species and collected 97 hemolymph and organs from three larvae per sample. In contrast, each sample of A. pagana 98 consisted of a single larva. The gut of L. zonalis larvae was large enough to be separated in its 99 three parts: foregut, midgut, and hindgut. The cuticle of this species could generally be 100 101 detached from the rest of the integument (indicating that the larvae were in a pre-moulting phase), to which longitudinal muscles were associated (Table 1). All samples from L. zonalis 102 103 were stored at -30 °C, and those from *A. pagana* at -80 °C, until their extraction.

The extraction procedure was developed by Boevé et al. (2014). In the present study, two extractions were made from each sample of *A. pagana*, and three from each one of *L. zonalis*, independent of the larval fresh weight (FW). The gathered pooled extracts, each making up 3 ml and 4 ml, respectively, were stored at -80 °C. Those from *A. pagana* were diluted 20 times, those from *L. zonalis* 30 times, and these aliquots were then again stored at -80 °C until chemical analysis.

110 To summarize, there are three events preceding the LC-MS/MS analyses: 1) collection of hemolymph plus dissection of organs, 2) successive extractions of the samples resulting in a 111 pooled extract, and 3) triplicated dilution leading to aliquots. The time elapse between 112 113 points 1 and 2 was especially long, over nine years for the samples of *L. zonalis* (Table 1). Equipment and methodology corresponded exactly to those described by Boevé et al. 114 (2018). In short, three aliquots of each pooled extract were analysed by high-performance 115 LC-MS/MS. The LC system (Thermo Fisher Scientific, San Jose, California, USA) was equipped 116 with an Accela 1250 pump and an Accela autosampler. Separation of peptides was 117 performed on a C18-HL Alltima column (150 x 2.0 mm i.d., 3 µm; Grace, Deerfield, Illinois, 118 119 USA) using a linear gradient from 90% H₂O (with 1% CH₃CN and 0.1% HCOOH)/10% CH₃CN to 120 10% H₂O in 31 min; the flow rate was 0.2 ml min⁻¹. 121 Standards of the peptides Perg, VPerg, dpPerg and LGIn were purchased from Biosyntan

122 GmbH (Berlin, Germany) at > 95% purity. They were analysed by full MS/MS on the $[M+H]^+$

123 ions by a Q-Exactive mass spectrometer (Thermo Fisher Scientific). Calibration curves were 124 constructed over four concentrations in the range of 1–1000 ng/ml, and proved to be linear, with r^2 values > 0.9990. Peptide concentrations in the samples were determined by 125 comparing the ratio of a peptide peak area in the sample solution with the ratios of its peak 126 areas in the standard solutions. These concentrations (in ng/ml) were converted to 127 micrograms per sample and averaged over the three replicates, the final concentration was 128 expressed in % of larval FW. Thus, this concentration was calculated by using for A. pagana 129 the FW of the larva, and for *L. zonalis* the summed FWs of the three larvae from which the 130 hemolymph and organs were pooled to make a sample. 131

The single sample of integument including cuticle from *L. zonalis* (Table 1) was discarded when peptide concentrations were summed over organs. Wilcoxon signed-rank tests were performed using VassarStats (Lowry 2019).

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137 **Results**

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The average concentration of the four peptides (Perg, VPerg, dpPerg, and LGIn) together was
0.333% FW in *L. zonalis* and 0.116% FW in *A. pagana*. The major compounds were 0.204%
FW VPerg and 0.117% FW Perg in the former, and 0.080% FW LGIn and 0.033% FW Perg in
the latter species.

143 For *L. zonalis*, the concentration of the four peptides was 0.238% FW in the hemolymph, 144 0.078% FW in the integument by considering the samples of integument without cuticle plus 145 the samples of cuticle, and 0.016% FW in the gut (Fig. 1). For A. pagana, it was 0.095% FW in 146 the hemolymph, 0.018% FW in the integument (i.e. including cuticle), and 0.002% FW in the gut. For this species, the concentration of each LGIn and Perg was higher in the hemolymph 147 (0.066% FW LGIn and 0.027% FW Perg) than in the integument (0.013% FW LGIn and 0.005% 148 FW Perg), and higher in the integument than in the gut (0.001% FW LGIn and 0.001% FW 149 Perg) (W = 21, P = 0.05 four times, Wilcoxon signed-rank tests, two-tailed; n = 6 larvae). For 150 both species, the ratio between Perg and LGIn concentrations was similar across hemolymph 151 152 and organs (Fig. 1). Lophyrotoma zonalis larvae contained a similar peptide concentration in the cuticle 153

154 (0.025% FW VPerg and 0.013% FW Perg) compared with the rest of integument plus the

layer of longitudinal muscles (0.024% FW VPerg and 0.013% FW Perg; Fig. 1). For both

species, nearly no peptides were detected in the fat body and salivary glands (Fig. 1). The

157 weight was only quantifiable for the hemolymph of *A. pagana*, and it reached a mean ±

standard deviation of 14.5 ± 2.8 mg FW, thus representing 21% of the larval FW (Table 1).

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161 **Discussion**

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The chemical profiles of the hemolymph and integument were dominated by VPerg and Perg 163 in L. zonalis, and LGIn and Perg in A. pagana. These profiles are the same as those of whole 164 larvae of the corresponding species (Boevé et al. 2014). Thus, the distribution of the 165 peptides within the body qualitatively reflects that of a whole larva. In theory, considering 166 167 either all body compartments together or a whole larva should lead to equivalent peptide concentrations. This, however, was not the case, since the concentration of the summed 168 169 hemolymph and organs for each species L. zonalis and A. pagana was lower (0.333 and 170 0.116% FW, respectively) than from a whole larva (0.719 and 0.162% FW, respectively; Boevé et al. 2014). For each of the species, this difference may be due to variation between 171 populations (Boevé et al. 2014, 2018), to a loss of compounds during collection of the 172 hemolymph and dissection of larvae, and/or to a degradation of compounds during the 173 174 subsequent storage of samples and pooled extracts. The period between gathering the 175 samples and their extraction was much longer for *L. zonalis* (112 months) than *A. pagana* (10 176 months), and the loss in peptide concentration was also higher in *L. zonalis* than *A. pagana*. 177 Thus, a long storage period of biological samples in ethanol may cause peptide degradation. 178 In natural conditions, however, these compounds are known to be stable to enzymatic degradation (Oelrichs et al. 2001). 179

For both species, the highest peptide yields were from the hemolymph, followed by the integument. In *A. pagana*, the FW of the hemolymph represented ca. 20% of the larval FW. By measuring the FW of an *A. pagana* larva just after moulting and of its exuvia, it was found that the exuvia represented ca. 2% of the larval FW (J-L Boevé, unpublished result). These results combined with those gathered for *L. zonalis* lead to the conclusion that the cuticle, although of a low FW, contains high amounts of toxic peptides. This is in agreement with defensive function of the peptides (Oelrichs et al. 1999, Petre et al. 2007). Predators,

187	especially small ones, first come into contact with the insect's integument, and if biting, then
188	with its hemolymph. Toxic peptides in <i>L. zonalis</i> and <i>A. pagana</i> were detected in low
189	quantities in the gut and some other organs, and nearly none were detected in the
190	excrements (Boevé and Rozenberg 2019), which supports the idea that they play a defensive
191	role against attacking predators. Other peptides with an antimicrobial activity can be found
192	on the cuticle of other hymenopterans (Otti et al. 2014). More research about the toxic
193	peptides is needed to understand their function in relation to their body distribution, as well
194	as whether and, if so, how the compounds are transported within the larval body.
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196	Compliance with ethical standards
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198	Conflict of interest The authors declare that they have no conflict of interest.
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- **Table 1** Data about the samples of *Lophyrotoma zonalis* and *Arge pagana* sawfly larvae,
- 248 analysed by LC-MS/MS
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	L. zonalis	A. pagana
Larval fresh weight (mg)	152.1 ± 32.8	68.6 ± 14.4
Date of dissection	March–April 2008	September 2016
Gut	-	6 [1]
Foregut	5 [3]	-
Midgut	5 [3]	-
Hindgut	5 [3]	_
Hemolymph	5 [3]	6 [1]
Integument including cuticle	1 [3]	6 [1]
Integument without cuticle	3 [3]	-
Cuticle	3 [3]	_
Salivary glands & fat body	_	6 [1]
Fat body	4 [3]	_
Date of extraction	August 2017	July 2017
Date of dilution	August 2017	July 2017
Date of LC-MS/MS analysis	August 2017	July 2017

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251 The larval fresh weight is given as mean ± standard deviation. The date of dissection

corresponds to the period of collecting hemolymph and dissecting organs. For samples of

253 hemolymph, organs or organ parts, the number of samples analysed by LC-MS/MS is given

without square brackets, and each sample comprised a number of larvae that is given

255 between square brackets. Note that samples of integument include associated longitudinal

256 muscles. (–) No sample analysed. Dates are given for the extractions leading to pooled

257 extracts, the dilutions leading to aliquots, and the LC-MS/MS analyses

259 Figure caption

- 261 **Fig. 1** Peptide concentrations in the larval organs and hemolymph of *Lophyrotoma zonalis*
- and Arge pagana. The peptides are pergidin (Perg), 4-valinepergidin (VPerg),
- 263 dephosphorylated pergidin (dpPerg), and lophyrotomin (LGIn). The data are given as mean
- and standard deviation of percentage of larval fresh weight (FW). Details about the organs
- and hemolymph are given in Table 1. Note that in *L. zonalis* a single sample corresponded to
- the integument including cuticle

