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Functional evidence for physiological mechanisms to circumvent neurotoxicity of cardenolides in an adapted and a non-adapted hawk-moth species

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Because cardenolides specifically inhibit the Na⁺K⁺-ATPase, insects feeding on cardenolide-containing plants need to circumvent this toxic effect. Some insects such as the monarch butterfly rely on target site insensitivity, yet other cardenolide-adapted lepidopterans such as the oleander hawk-moth, *Daphnis nerii*, possess highly sensitive Na⁺K⁺-ATPases. Nevertheless, larvae of this species and the related *Manduca sexta* are insensitive to injected cardenolides. By radioactive-binding assays with nerve cords of both species, we demonstrate that the perineurium surrounding the nervous tissue functions as a diffusion barrier for a polar cardenolide (ouabain). By contrast, for non-polar cardenolides such as digoxin an active efflux carrier limits the access to the nerve cord. This barrier can be abolished by metabolic inhibitors and by verapamil, a specific inhibitor of P-glycoproteins (PGPs). This supports that a PGP-like transporter is involved in the active cardenolide-barrier of the perineurium. Tissue specific RT-PCR demonstrated expression of three PGP-like genes in hornworm nerve cords, and immunohistochemistry further corroborated PGP expression in the perineurium. Our results thus suggest that the lepidopteran perineurium serves as a diffusion barrier for polar cardenolides and provides an active barrier for non-polar cardenolides. This may explain the high *in vivo* resistance to cardenolides observed in some lepidopteran larvae, despite their highly sensitive Na⁺K⁺-ATPases.

1. Introduction

Over the course of evolution, plants have evolved a vast diversity of secondary plant compounds many of which act as chemical weapons against herbivores. In return, herbivores have developed strategies to overcome plant defences. Mechanisms of resistance in insects are numerous and include detoxification of toxins by enzymes, excretion, exclusion (gut barriers) and target site insensitivity [1,2].

In this study, we focus on insect resistance to plant-produced cardenolides (aka cardiac glycosides), a specific class of plant toxins [3,4]. Cardenolides are specific inhibitors of the Na⁺K⁺-ATPase, a ubiquitous animal enzyme that is essential for many physiological processes [5,6].

Several herbivorous insects, including the monarch butterfly (*Danaus plexippus*) not only feed on cardenolide-containing plants, but also sequester the toxins and thus derive protection against predators [7]. The Na⁺K⁺-ATPase of *D. plexippus* is altered by specific amino acid substitutions, which significantly reduce its cardenolide susceptibility (target site insensitivity; [8,9]). In earlier studies, however, we found that lepidopterans that are adapted to cardenolides sometimes possess cardenolide sensitive Na⁺K⁺-ATPases [10,11]. Moreover, among cardenolide-adapted Lepidoptera, the monarch butterfly actually seems to be an exceptional case [9,12,13].

In Lepidoptera, Na⁺K⁺-ATPase is predominantly expressed in the nervous tissue. The concomitant occurrence of dietary cardenolides in the caterpillars' haemolymph, therefore, renders the interface between insect blood and nervous

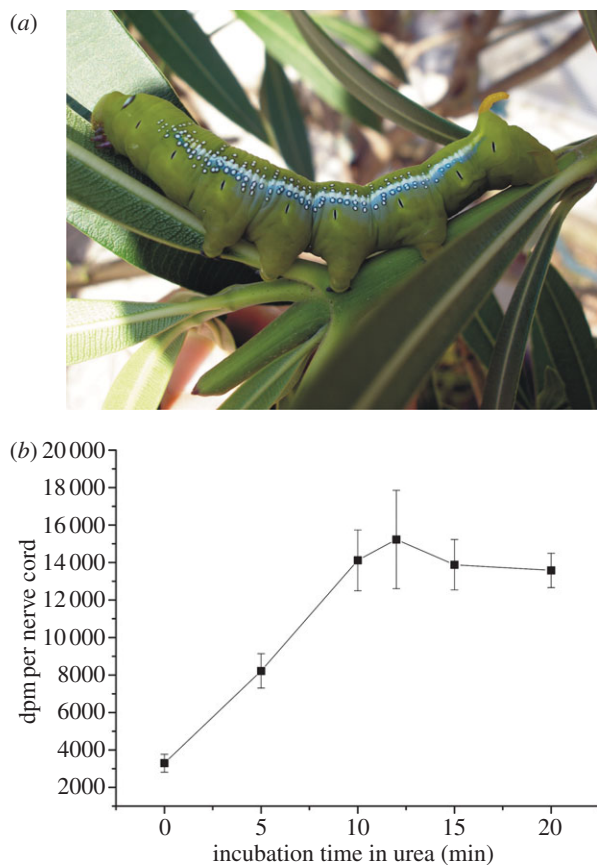


Figure 1. Binding of ^3H -ouabain to the isolated nerve cord of *D. nerii* caterpillars: (a) caterpillar of the oleander hawk-moth (*D. nerii*). The major host plant of this species is *N. oleander*, whose marked toxicity is based on cardenolides. (b) Disruption of the perineurial barrier with urea in isolated nerve cords of *D. nerii* (each datapoint represents the mean (\pm s.d.) of three independent incubations, i.e. 18 caterpillars were used in total). (Online version in colour.)

tissue especially important [11]. The ventral nerve cord of insects is, like in other organisms, surrounded by the perineurium, a tissue maintaining ionic conditions required for the excitability of neurons, which may be different from the composition of the haemolymph (low K^+ and high Na^+ required in the extracellular space of the nerve cord versus an approximate 1:1 ratio in the haemolymph; [14]). Additionally, this tissue is believed to function as a blood–brain barrier for toxic plant compounds present in the herbivores’ haemolymph [15]. However, thus far there is only limited functional evidence for such a protective function of the perineurium.

In this study, we test whether the perineurium can function as a barrier to dietary cardenolides absorbed into the haemolymph and thus potentially contribute to resistance to these toxins. We used the oleander hawk-moth (*Daphnis nerii*; figure 1a), a cardenolide specialist that feeds primarily on oleander (*Nerium oleander*), a plant rich in these toxins [16]. For comparison, we included *Manduca sexta*, a related species which is not adapted to cardenolides.

Daphnis nerii caterpillars have relatively low levels of oleander cardenolides in their body (approx. 150–200 μg at maximum; [16]). Nonetheless, owing to their highly sensitive Na^+K^+ -ATPase even minute amounts of cardenolides in the haemolymph can be fatal [10]. Because oleander and other cardenolide-containing plants have cardenolides with a wide range of polarity, we speculated that different mechanisms may be important in insect resistance to diverse cardenolides. For example, the perineurium forms a diffusion

barrier for polar cardenolides as was shown for *M. sexta* [17]. Nonetheless, non-polar cardenolides that are able to use the transcellular pathway [18] might require an active barrier mechanism (i.e. efflux transporters). Both mechanisms are tested here by physiological experiments.

In the mammalian brain, P-glycoprotein (PGP) is one of the most important efflux transporters [19] with an amazingly wide substrate spectrum including the cardenolide digoxin [20]. This 170-kDa membrane bound protein, a member of the ABC (ATP-binding cassette)-transporter superfamily, extrudes xenobiotic compounds from cells driven by ATP hydrolysis. In *M. sexta*, PGP was already suggested to be involved in nicotine resistance [15]. In other insects, it is believed to mediate resistance to insecticides or xenobiotics [21,22]. We therefore tested whether a PGP-like transporter may be involved in the physiological blood–brain barrier of the hawk-moth nerve cord using the well-known PGP inhibitors quinidine and verapamil. Immunohistochemical assays with monoclonal antibodies were further used to visualize PGP as well as Na^+K^+ -ATPase occurrence in the nerve cord. Moreover, an analysis of *M. sexta* expressed sequence tags (EST) data followed up by tissue-specific RT-PCR confirmed the occurrence of PGP-like transporters in the perineurium.

In summary, our investigations address the relative importance of passive and active mechanisms in protecting the hawk-moth nervous system from potent plant toxins.

2. Material and methods

(a) Radiochemicals and inhibitors

^3H -ouabain (12 Ci mmol^{-1} , dissolved in 9:1 ethanol:toluene, or 30 Ci mmol^{-1} , dissolved in ethanol) was purchased from GE Healthcare (Freiburg, Germany) and Perkin Elmer (Rodgau, Germany). ^3H -digoxin was purchased from Perkin Elmer (40 Ci mmol^{-1} , dissolved in ethanol). Both ouabain and digoxin most likely do not occur in larval host plants of *D. nerii*, but were used owing to their commercial availability and strongly differing polarity. 2,4-dinitrophenol (2,4-DNP; Fluka, Taufkirchen, Germany), carbonyl cyanide 3-chlorophenylhydrazone (CCCP; Sigma, Taufkirchen, Germany), verapamil hydrochloride (Sigma) and quinidine (Sigma) were used as 0.05 M stock solutions in ethanol. In our binding experiments, we used ^3H -cardenolide concentrations of 0.35 and 0.7 μM , respectively. We decided to use such low amounts because Rubin *et al.* [17] observed non-specific binding of ^3H -ouabain to native nerve cords of *Manduca* at concentrations above 10 μM . We decided not to refer our disintegrations per minute (dpm) values to protein content throughout the experiments because protein determination proved to be dependent on storage time (at -20°C) post-experiment. Referring to nerve cords as experimental units, on the other hand, proved to be highly reliable because the (simultaneously determined) protein content of 18 *D. nerii* nerve cords (eight ganglia each, see below) averaged 63.01 μg with a standard deviation of 9.74. The small standard deviations of our treatment groups throughout the experiments give further evidence that this approach provides reliable data that are not biased by size differences. Therefore, the radioactivity measured in our experiments is expressed as dpm per nerve cord. All data used in the inhibitor experiments are provided in the electronic supplementary material.

(b) Diffusion barrier

To test for a diffusion barrier to polar cardenolides, we followed the experimental design described by Rubin *et al.* [17], who disrupted the perineurium of *M. sexta* by treatment with urea.

Caterpillars of *D. nerii* (European origin) were raised on greater periwinkle (*Vinca major*), which is devoid of cardenolides, at 23°C (16 L : 8 D cycle). Prior to dissection, last instar caterpillars were chilled on ice and decapitated. Ventral nerve cords were removed, placed in cold incubation buffer (125 mM NaCl, 5 mM MgCl₂, 0.5% bovine serum albumin (BSA) and 12.5 mM imidazole, pH 7.3), cleaned from adherent tissue and trimmed to a chain of eight ganglia plus intervening connectives (abdominal ganglia plus metathoracic ganglion; [23]). For each of three replicates a series of six caterpillars was used. One nerve cord of a series was used as a control and was kept in incubation buffer at room temperature for the duration of the urea treatment. The additional five nerve cords were immersed in 3 M urea in incubation buffer for 5, 10, 12, 15 or 20 min, respectively. The cords were then washed twice with incubation buffer for at least 5 min each. Following urea treatment, cords (including the control cord) were individually incubated in 100 µl incubation buffer with 0.7 µM ³H-ouabain for 1 h at 37°C. After incubation, cords were washed in an excess volume of 10 mM imidazole (pH 7.3) for 30 min on ice. Each cord was then transferred to 200 µl 0.2 M NaOH/1 per cent SDS and digested overnight. To 150 µl of this extract 3 ml liquid scintillation cocktail (Ultima Gold, Perkin Elmer) were added and radioactivity determined in a liquid scintillation counter (Wallac 1409, easy count mode). The remainder of each sample was stored at -20°C for later protein determination with the bicinchoninic acid (BCA) assay (Thermo Scientific) using BSA as a standard.

(c) Active barrier

(i) *Manduca sexta*

Eggs of *M. sexta* were kindly supplied by Dr. Markus Huß (University of Osnabrück). Caterpillars were reared on gypsy moth diet (MP Biomedicals) supplemented with streptomycin, chloramphenicol, methyl benzoate and formalin (26°C; 16 L : 8 D cycle). Only last instar caterpillars before reaching the wandering stage were used.

(ii) *Daphnis nerii*

Caterpillars of *D. nerii* (origin Thailand) were raised from eggs at 27°C at 13 L : 11 D cycle. Hatched caterpillars were initially fed with *V. major* later transferred (second instar) to *N. oleander* and raised to the last instar.

Nerve cords of both species were dissected as described above and maintained until incubation on ice in *Manduca* saline: 5.0 mM K₂HPO₄, 10.0 mM MgCl₂, 1.0 mM CaCl₂, 10.0 mM NaCl, 10.0 mM KOH, 7.4 mM L-proline, 7.7 mM tri-potassium citrate, 2.8 mM disodium succinate, 2.0 mM glucose, 175.0 mM sucrose, 5.6 mM malic acid, 10.0 mM HEPES, pH 6.7 [24]. Again, the posterior eight ganglia were used. To test the hypothesis that the nerve cords of *M. sexta* and *D. nerii* possess an energy-driven barrier that prevents cardenolides from reaching the Na⁺K⁺-ATPase, the metabolic inhibitors 2,4-DNP and CCCP were applied. To test whether a PGP-like transporter is involved in this barrier, we used verapamil and quinidine that are well-known competitive PGP inhibitors [25]. All inhibitors were dissolved in ethanol and applied at a final concentration of 1 mM. Controls were incubated with an equivalent amount of ethanol. The concentration of ³H-digoxin in the assay was 0.35 µM (ethanol concentration 3.36%). Each nerve cord was incubated in a volume of 100 µl *Manduca* saline at 37°C. After 30 min, tubes were placed on ice, the radioactive solution was removed, 1 ml of cold 10 mM imidazole (pH 7.3) added and mixed by vortex stirring. After replacing the washing buffer once, tubes were inverted and kept on ice for 30 min. The short washing step was performed to remove adherent radioactive solution, whereas the long washing step was performed to remove unbound ³H-digoxin [17]. In an additional experiment (data not shown) we found that nearly all adhering radioactivity is removed

from the tissue after the 30 min washing step. After washing, the samples were lysed and radioactivity counted as described above.

(d) Statistical analysis

If necessary, data were squared or log-transformed to achieve homogeneity of variances (Levene's test) and approximately normal distributions (Shapiro-Wilk). Data were analysed by ANOVA using a randomized block design with the experiment as blocking factor. Post hoc comparisons are based on Tukey's honestly significant difference (HSD) test. All statistical tests were performed with SPSS (Statistical Package for the Social Sciences, IBM).

(e) Comparison of digoxin versus ouabain permeability

This experiment was performed to demonstrate the different permeability of the perineurium of *D. nerii* caterpillars for ouabain and digoxin. As incubation buffer, physiological saline without energy sources (PBS: 137 mM NaCl, 2.7 mM KCl, 4.3 mM Na₂HPO₄, 1.4 mM KH₂PO₄, pH 7.4; [26]) was used, otherwise the assay followed the procedures described above. Control nerve cords were incubated in buffer with 0.7 µM ³H-digoxin or ³H-ouabain only. In parallel, nerve cords were incubated with the labelled compounds plus CCCP (1 mM). CCCP was added to disable active transport processes and get an estimate of the amount of cardenolides infiltrating the nerve cord by diffusion.

(f) Immunohistochemistry

(i) P-glycoprotein-like transporter

Nerve cords of chilled *D. nerii* caterpillars (last instar) were dissected and immersed in PBS. Tissues were fixed for 1 h at room temperature in Lana's fixative (15% picric acid, 4% paraformaldehyde (PFA) in 0.5 M sodium phosphate buffer, pH 7; [15]). After fixation, tissues were washed three times for 10 min each in PBS and successively cryoprotected in 5, 10 and 15 per cent sucrose in PBS for 1 h each. Following cryoprotection tissues were embedded in optimal cutting temperature (OCT) compound (Sakura, Alphen aan den Rijn, The Netherlands) frozen in isopentane in liquid nitrogen and stored at -80°C until sectioning. Sections of 16 µm were cut on a Leica CM 1950 cryostat and allowed to dry at room temperature. Slides were stored at -80°C until use. The anti-PGP antibody C-219 (Abcam, Cambridge, UK; dissolved in PBS) was applied at a concentration of 10 µg ml⁻¹. In the control sections, the primary antibody was omitted. The primary antibody was detected with the NOVADetect DAB (3,3'-diaminobenzidine)-Substrate Kit (Dianova, Hamburg, Germany). Stained sections were shortly washed with deionized water, transferred into 80 per cent ethanol (via 60% ethanol) and mounted in Euparal. Sections of ganglia were inspected under a Zeiss Axioskop 2 and photographed with a Zeiss AxioCam colour camera.

(ii) Na⁺K⁺-ATPase

To visualize the target site of cardenolides in the hawk-moth ganglia (*M. sexta*) we followed the methods described in [27] and [11]. For the specific detection of Na⁺/K⁺-ATPase in paraffin sections, we used the monoclonal antibody α5 (developed by D.M. Fambrough, maintained and distributed by the Developmental Studies Hybridoma Bank, University of Iowa, USA).

(g) Molecular phylogenetic analyses

The coding sequences of three PGP-like transporters of *Trichoplusia ni* [28] were used to identify homologous *M. sexta* sequences in a collection of preassembled *M. sexta* ESTs (H. Vogel 2011, unpublished data). The corresponding amino acid sequences are given in the electronic supplementary material, figure S1.

Available PGP-like transporters were downloaded from GenBank (accession date: May 2012), including the three PGP-like transporters of *Drosophila melanogaster* (MDR49, MDR50 and MDR65; [29,30]), as well as several PGP-like transporters of other insects and crustaceans. A complete sequence list including accession numbers is given in electronic supplementary material, table S1. The amino acid sequences were aligned with MAFFT using the G-INS-i routine [31] and the alignment was processed with Gblocks v. 0.91b [32]. Gblocks settings and the final alignment are given in electronic supplementary material, figure S2. Phylogenetic analyses were performed with MRBAYES v. 3.1 [33] using the WAG model [34]. Metropolis-coupled Markov chain Monte Carlo sampling was performed with one cold and three heated chains. Two independent runs were performed for 1 million generations. Trees were sampled every 100th generation and posterior probabilities were estimated on the final 7500 trees (burnin = 2500). Mammal PGPs, which are known to be homologous to the PGP-like transporters of *D. melanogaster* [29,30] were used to root the phylogram for visualization purpose.

(h) RT-PCR

Total RNA was extracted from nerve cords (eight hindmost ganglia, tissues from 2–3 individuals pooled) and midguts of *M. sexta* caterpillars (last instar) with the RNeasy plus kit (Qiagen, Hilden, Germany). In both cases, three independent RNA extractions were performed (biological replicates). Amounts of RNA were assessed by reading the absorption at 260 nm and subsequently confirmed by denaturing gel electrophoresis. Equivalent amounts were transcribed into cDNA with Superscript III (Invitrogen, Darmstadt, Germany) using a combination of dT-17 and random hexamer primers. Amplification was performed using a standard protocol (Invitrogen Taq Polymerase; PCR: 95°C for 45 s, 52°C for 60 s, and 72°C for 60 s; 40 cycles). Gene specific oligonucleotide primers were: 5'-TTGACGGCAGTGT GACGATAG-3' and 5'-CCTTCAGGAGACGTTTGCATC-3' for *M. sexta* PGP-like transporter I, 5'-TCAAGATGTAGAGCCCG TGGT-3' and 5'-CCAGCGGTAGTGAAGGTTGAG-3' for *M. sexta* PGP-like transporter II, and 5'-TTCGGTGGCGCAGTTTATAGT-3' and 5'-TCTTGTGCCCATCTTCTTTC-3' for *M. sexta* PGP-like transporter III. Primer specificity was confirmed by sequencing the corresponding PCR products (GATC, Konstanz, Germany).

3. Results

(a) Diffusion barrier

In our experiment with *D. nerii* caterpillars, we found a corresponding result to that described for *M. sexta* [17]: ouabain binding to the isolated *D. nerii* nerve cord linearly increased with time over the first 10 min of incubation in 3 M urea (figure 1b). After 10 min, the curve reached a plateau either indicating that a limit of permeabilization was achieved, or complete permeabilization of the diffusion barrier and saturation of the ouabain binding sites.

(b) Active barrier

In contrast to polar ouabain, the more lipophilic digoxin is known to permeate cell membranes [18]. Therefore, we assume non-polar cardenolides to necessitate active barrier mechanisms that prevent them from entering the nerve cord.

Application of the metabolic inhibitors (ionophores) 2,4-DNP and CCCP on the isolated nerve cord of *M. sexta* significantly enhanced tissue binding of ³H-digoxin (figure 2a) to 1.7-fold and 2.3-fold, respectively, of the untreated control. For *D. nerii*, we tested only the more effective inhibitor CCCP

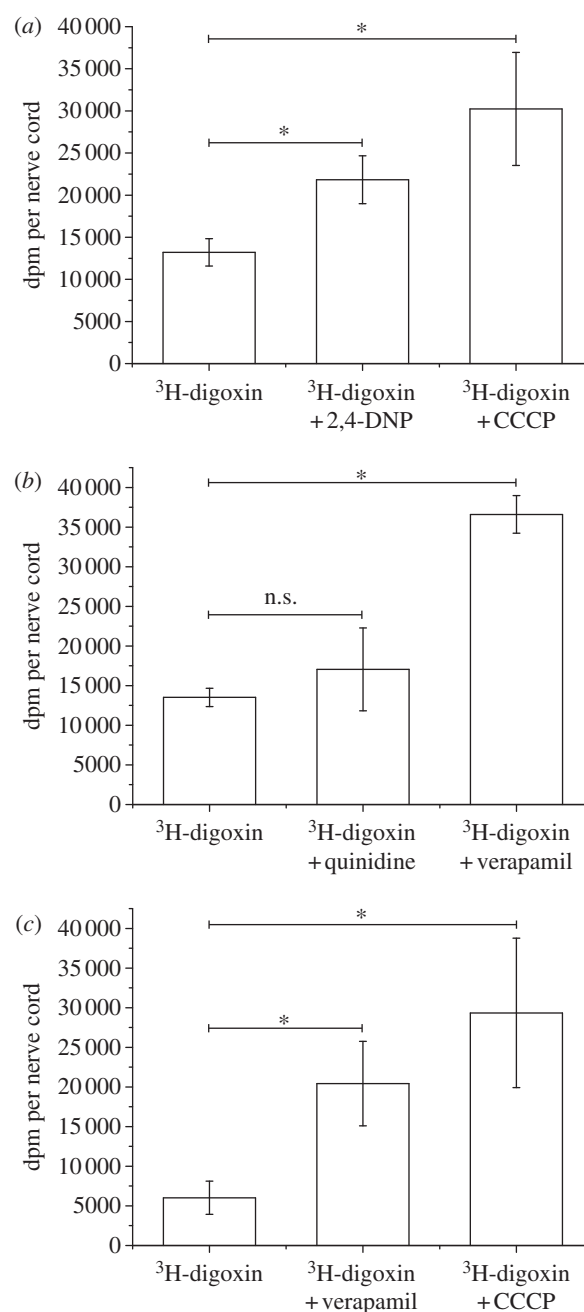


Figure 2. Influence of inhibitors on ³H-digoxin binding in isolated nerve cords: (a) effect of the metabolic inhibitors 2,4-DNP and CCCP on nerve cords of *M. sexta* ($n = 5$; mean \pm s.d.). (b) Effect of the PGP inhibitors quinidine and verapamil on nerve cords of *M. sexta* ($n = 5$; mean \pm s.d.). (c) Influence of the metabolic inhibitor CCCP and the PGP inhibitor verapamil on ³H-digoxin binding in isolated nerve cords of *D. nerii* ($n = 6$; mean \pm s.d.). Asterisks above horizontal lines indicate significant differences ($p < 0.05$, randomized block ANOVA with experiment as blocking factor, followed by Tukey HSD tests). Each data point represents the mean of 5 nerve cords per treatment (b and c) or 6 nerve cords per treatment (c) \pm s.d.

(figure 2c). Here, the increase of ³H-digoxin binding was even stronger than that in *M. sexta* (4.5-fold compared with 2.3-fold).

To assess the involvement of a PGP-like transporter in the active barrier, the widely used PGP inhibitors verapamil and quinidine were tested for their effect on digoxin binding to the nerve cord. For *M. sexta*, verapamil enhanced ³H-digoxin binding to the nerve cord 2.7-fold. Quinidine produced a similar trend (figure 2b) which was, however, not significant. For *D. nerii*, only verapamil was applied which again significantly increased ³H-digoxin binding. Interestingly, as with CCCP, this

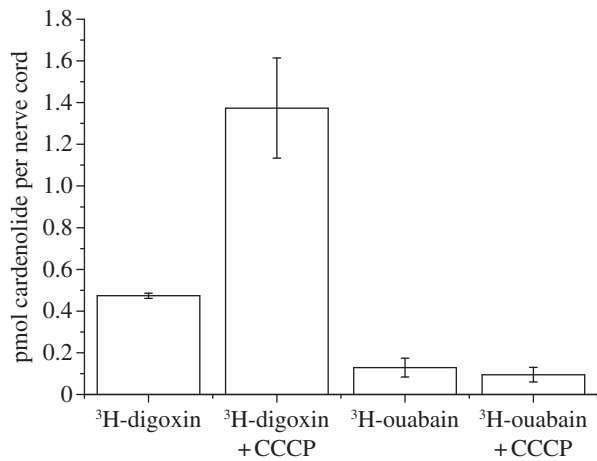


Figure 3. Diffusion of digoxin and ouabain into the nerve cord of *D. nerii*. As the two cardenolides have different specific activities, dpm values are not directly comparable. Data are, therefore, presented as pmol cardenolide/nerve cord (instead of dpm per nerve cord) to allow for quantitative comparisons. Nerve cords were incubated in equal concentrations of ouabain and digoxin, respectively (0.7 μ M). Each data point is the mean of three nerve cords per treatment (i.e. 12 caterpillars were used in total).

effect was also stronger than in *M. sexta* (3.4-fold compared with 2.7-fold).

To exclude the possibility that verapamil simply permeabilizes the perineurium, *M. sexta* nerve cords were incubated with digoxin and digoxin plus verapamil or with ouabain and ouabain plus verapamil, respectively. If verapamil simply disrupted the passive diffusion barrier, the polar ouabain was also expected to gain access and bind to the nervous tissue to a larger extent upon verapamil treatment. This was, however, not the case (see the electronic supplementary material, figure S3).

(c) Comparison of digoxin versus ouabain permeability

When nerve cords of *D. nerii* were incubated with either cardenolide alone or in combination with CCCP, the latter significantly increased digoxin binding but not ouabain binding (figure 3). This supports our conclusion that digoxin can infiltrate the nerve cord by diffusion when active efflux carriers are disabled. The polar ouabain, however, is excluded by a diffusion barrier and its binding is not increased when the metabolic poison CCCP is applied. Interestingly, the active barrier was still functional in this experiment (and excluding most digoxin when not blocked by CCCP), although we here used PBS instead of *Manduca* saline. In contrast to the latter, the former lacks energy sources thus indicating that intrinsic energy levels are sufficient to maintain functionality of the efflux carriers for the duration of the experiment.

(d) Immunohistochemical detection of a P-glycoprotein-like transporter

The widely used anti-PGP antibody C219 binds to a conserved epitope of the protein [35]. We applied this antibody to cryosections of ganglia from *D. nerii* caterpillars and found specific staining (brown precipitate) only in the periphery of the respective ganglion (figure 5a).

(e) Identification of P-glycoprotein-like transporters expressed in the nerve cords

Three proteins with at least 75 per cent identity to either one of the PGP-like transporters of *T. ni* were identified in the ESTs of *M. sexta*. We here arbitrarily refer to these corresponding proteins as PGP-like transporters I, II, and III. The phylogenetic analysis (figure 4a) shows that the lepidopteran PGP-like transporters I form a monophyletic clade with *D. melanogaster* MDR50 and several other insect proteins (posterior probability 1.0). Lepidopteran PGP-like transporters III form a monophyletic clade with *D. melanogaster* MDR65 (posterior probability 0.97), which are in a sister group position with a monophyletic clade that among other insect proteins includes *D. melanogaster* MDR49 (posterior probability 1.0). Lepidopteran PGP-like transporters II are in a sister group position to a monophyletic clade, which comprises both *D. melanogaster* MDR49 and MDR65 (posterior probability 0.98). RT-PCR analyses show that all of the lepidopteran PGP-like transporters are expressed in the nerve cord of *M. sexta*, whereas in the gut a comparable strong expression appears to be restricted to PGP-like transporter I (figure 4b).

(f) Immunohistochemical detection of Na⁺K⁺-ATPase

Application of the monoclonal anti-Na⁺K⁺-ATPase antibody α 5 revealed a strong signal in larval nerve cords of *M. sexta* ganglia (figure 5b). The occurrence of Na⁺K⁺-ATPase is apparently restricted to the neurons within the ganglion and no specific signal could be observed in the perineurium.

4. Discussion

In our study we focused on *D. nerii* and *M. sexta*, two closely related species that differ in their host plants and the secondary compounds they are typically exposed to. Whereas *M. sexta* is naturally not exposed to dietary cardenolides, *D. nerii* is an oleander specialist and encounters high concentrations of cardenolides of a wide polarity range in its natural diet, oleander. This species does not sequester cardenolides as do other specialists such as the monarch butterfly. The presence of only low amounts of cardenolides in the body [16] could be achieved by a relative impermeable gut membrane as has been observed in generalist insects such as *Schistocerca* and *Periplaneta* [1]. Such impermeability is not surprising in the case of polar cardenolides, which are unable to passively cross the gut membrane, yet in these species the guts are even impermeable to the markedly non-polar cardenolide digitoxin. As the Na⁺K⁺-ATPase of *D. nerii* is highly susceptible to cardenolides [10], additional mechanisms are needed to avoid even low amounts penetrating into the haemolymph.

An earlier study on cardenolide-adapted caterpillars revealed that Na⁺K⁺-ATPase is largely restricted to the caterpillars' nervous tissue [11]. Accordingly, we found that *D. nerii* caterpillars can tolerate high levels of ouabain if injected into the larvae's body cavity [10]. We, therefore, postulated a mechanism that prevents cardenolides from reaching the Na⁺K⁺-ATPase within the nervous system and thus, focused here on the interface between cardenolide-containing haemolymph and the ventral nerve cord.

Our data revealed that ouabain gains access to the Na⁺K⁺-ATPase only when the nerve cord of *D. nerii* has been treated with urea. As urea is believed to disrupt the

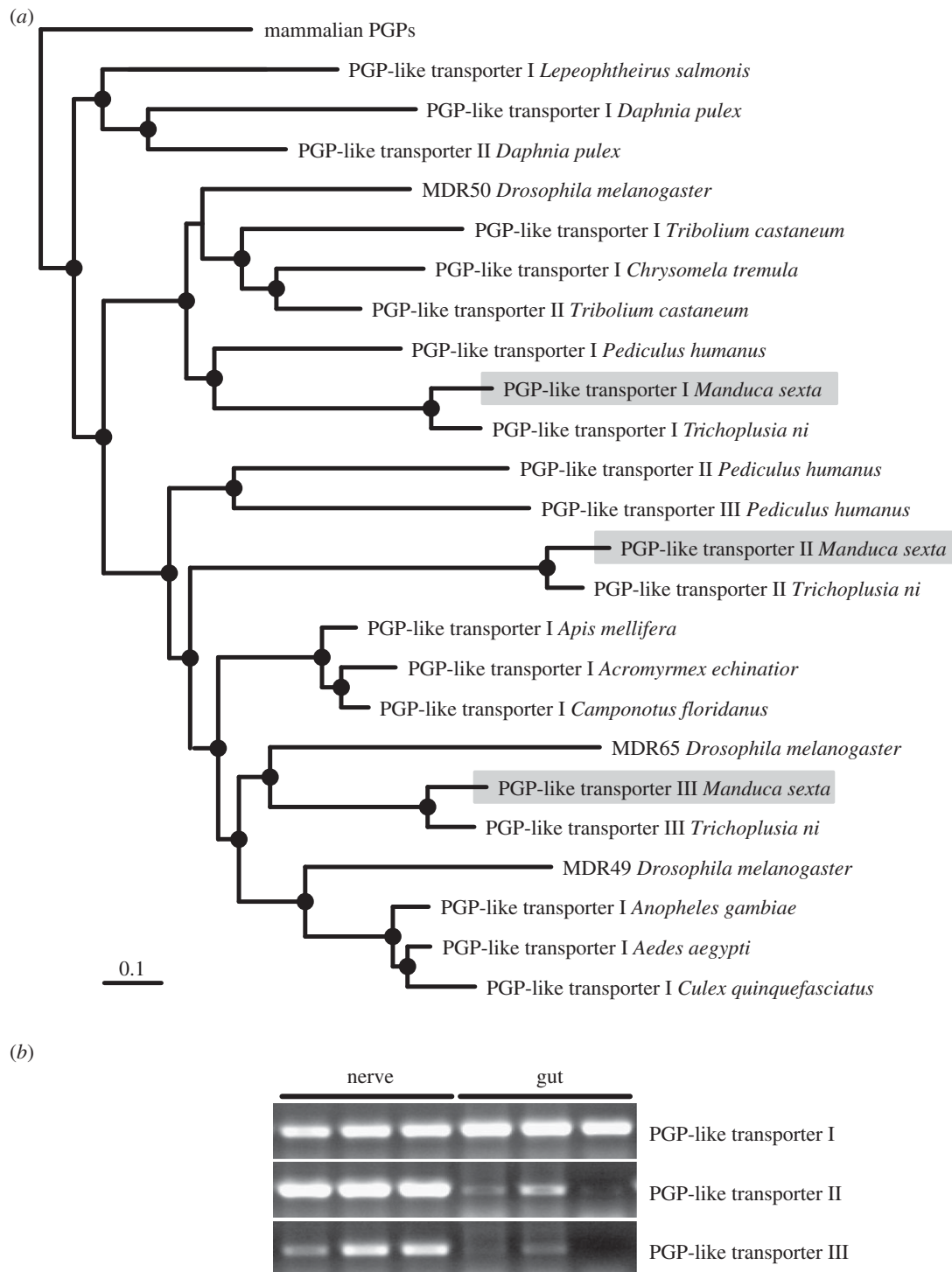


Figure 4. Molecular characterization of *M. sexta* PGP-like transporters: (a) phylogenetic relationships among arthropod PGP-like transporters. Except for *M. sexta*, available sequences were retrieved from GenBank and one was arbitrarily chosen if several sequences per genus existed. In accordance with previous studies, the PGP-like transporters of *D. melanogaster* were named multi drug resistance (MDR) protein and the mammalian PGPs were used to root the phylogram [29,30]. Nodes with Bayesian posterior probabilities higher than 0.95 are indicated by filled black circles, and the scale bar equals 0.1 expected substitutions per site. (b) Detection of PGP-like transporter encoding mRNAs in nervous and midgut tissue of *M. sexta* (biological triplicates under identical PCR-conditions).

lepidopteran perineurium [17], our results provide evidence that the native, intact perineurium is not permeable for ouabain and we can assume that this applies also to other relatively polar cardenolides. Therefore, the target site of cardenolides, the $\text{Na}^+\text{K}^+\text{-ATPase}$, is shielded from polar cardenolides present in the haemolymph. We here observed a similar time-dependent increase of ^3H -ouabain binding to the urea-treated nerve cord as described by Rubin *et al.* [17] for *M. sexta* to nerve cords of *D. nerii*, a species that actually has to cope with dietary cardenolides.

The diffusion barrier is probably constituted by the cells of the perineurium that form tight junctions [14] impeding the paracellular pathway for diffusing compounds. This barrier is most likely not selective for cardenolides, but represents a diffusion barrier for any polar compound. The ionic composition of the haemolymph would not be suitable for nervous function and thus, the perineurium is assumed to be responsible for the maintenance of the necessary ion concentrations in the nerve cord's extracellular space [14]. Therefore, the impermeability to ouabain of the perineurium



Figure 5. Immunohistochemical detection of a PGP-like transporter and Na^+K^+ -ATPase on sections of larval lepidopteran ganglia: (a) Frozen section of a *D. nerii* ganglion treated with the anti-PGP antibody C219. Specific labelling (brown precipitation, black arrow) can be seen in the periphery of the ganglion. Above, treatment; below, control (primary antibody omitted). (b) Paraffin section of a ganglion of a *M. sexta* caterpillar. Orange (white asterisk): specific label of Na^+K^+ -ATPase by the antibody $\alpha 5$. Blue (white arrow): nuclei stained with 4',6-Diamidino-2-phenylindole dihydrochlorid (DAPI). (Online version in colour.)

of *D. nerii* is likely not a specialization. Nonetheless, the LD_{50} of injected ouabain for *Schistocerca* and *Periplaneta* is as low as 4.4 and 0.6 μg per individual, respectively [36]. This might mean that the perineuria of *Schistocerca* and *Periplaneta* are not as tight to ouabain as the one of the hawk-moths. However, the Na^+K^+ -ATPase in these insects may not be restricted to the nervous tissue, but may also occur in other tissues, e.g. in the gut and the Malpighian tubules [37–39].

However, coping with dietary cardenolides also implies that non-polar forms have to be excluded from the nerve

cord. To test for the existence of an active barrier for non-polar cardenolides in the hawk-moth perineurium, we used the relatively non-polar cardenolide digoxin, which is known to passively permeate cells. By the application of the ionophores CCCP and 2,4-DNP, we aimed at blocking the respiratory chain in our test tissue, thus interrupting the supply of ATP. Consistent with the notion of an active barrier that protects the Na^+K^+ -ATPase, we found a higher binding of digoxin to the nerve cords when the metabolic inhibitors CCCP or 2,4-DNP were applied. This fits with the hypothesis of an active efflux mechanism: when the energy supply is depleted, digoxin can no longer be actively removed from the cells and reaches its target site.

Carriers of the PGP family are strong candidates to mediate the observed effect: in the mammalian brain they constitute the most important part of the blood–brain barrier by extruding infiltrating compounds [19]. Furthermore, Mayer *et al.* [40] demonstrated that PGP is responsible for excluding digoxin from the brain of wild-type mice. PGPs are members of the *mdr* gene family of which at least three genes are present in the *Drosophila* genome ([41] and references therein). In *M. sexta* (this study) and other lepidopteran species [28] three PGP-like transporters were identified which are homologous to those of *D. melanogaster*. All of these are expressed in the *Manduca* nerve cord, and are thus potential candidates for the efflux carriers evidenced here.

To test whether PGP-like transporters are involved in the energy-driven digoxin barrier, we incubated nerve cords of *M. sexta* and *D. nerii* with two of the most widely used PGP inhibitors, quinidine and verapamil. These compounds are known to elevate the plasma level of digoxin in humans when co-administered with this drug and this phenomenon is primarily attributed to the inhibition of PGP ([40], and references therein). Both in *D. nerii* and in *M. sexta*, the application of verapamil increased the amount of digoxin bound to the nervous tissue. These observations suggest that the efflux barrier for digoxin is mediated by a PGP-like transporter. When comparing the data of the two hawk-moth species, it is conspicuous that digoxin binding under control conditions is about twice as high in *M. sexta* as in *D. nerii*. At this point, however, it is difficult to judge whether this difference is due to quantitative or qualitative differences in the perineurial barrier of both species and there may also be size differences between both species.

The presence of a PGP-like transporter is furthermore demonstrated by our immunohistochemical data that revealed specific binding of the anti-PGP antibody C219 in the periphery of *D. nerii* ganglia. The presence of this protein in the nervous system of *M. sexta* was already demonstrated [15] though on the whole, the data on the occurrence of PGP-like transporters in insects is limited. Our knowledge about the involvement of PGPs in the exclusion of plant compounds in herbivorous insects is still insufficient, yet the excretion of nicotine in Malpighian tubules of *Manduca* caterpillars [42] was already suggested to be based on a PGP homologue [43]. Especially, the wide substrate range suggests a potential key role for these transporters in the resistance of herbivorous insects to toxic secondary plant compounds. PGP-like transporters might, in addition, not only be part of the blood brain barrier, but also be responsible for rendering insect guts impermeable to some plant toxins. They could in theory enable generalist species to cope with a wide array of diverse toxic secondary plant compounds.

The mechanisms of cardenolide exclusion described here, however, may not be the only mode of resistance to

cardenolides. It is known that cardenolides are metabolically modified within the insect body [16,44]. If regions of the molecule are affected, which mediate biochemical interactions, metabolism also results in detoxification. In addition, excretion by the Malpighian tubules can be expected to reduce haemolymph levels of cardenolides. The situation is further complicated by the fact that digoxin is not only substrate to PGP, but also to organic anion transporting polypeptides (Oatps) and potentially even additional carriers [45].

Thus, our picture of how specialists cope with cardenolides appears to be a diversity of strategies, with several often used

together, but potentially with distinct combinations among different herbivores.

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