Immune Stimulation *via* Wounding Alters Chemical Profiles of Adult *Tribolium castaneum*

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Abstract - Group-living individuals experience immerse risk of disease transmission and parasite infection. Especially in social, but also in some non-social insects, disease control with immunomodulation takes place not only *via* individual immune defenses, but also *via* infochemicals such as contact cues and (defensive) volatiles to mount a group level immunity. However, little is known about whether an activation of the immune system leads to changes in chemical phenotypes, which may mediate these responses. We here asked whether individual immune experience resulting from wounding or injection of heat-killed *Bacillus thuringiensis* leads to changes in the chemical profiles of female and male adult red flour beetles (*Tribolium castaneum*), which are non-social but gregarious. We analyzed insect extracts using GC-FID to study the chemical composition of (1) cuticular hydrocarbons (CHCs) as candidates for the transfer of immunity-related information between individuals *via* contact, and (2) stink gland secretions, with target analysis of benzoquinones as main active compounds regulating 'external immunity'. Despite a pronounced sexual dimorphism in CHC profiles, wounding stimulation led to similar profile changes in males and females with increases in the proportion of methyl-branched alkanes compared to naïve beetles. While changes in the overall secretion profiles were less pronounced, absolute amounts of benzoquinones were transiently elevated in wounded compared to naïve females. We suggest that the changes in different infochemicals may mediate immune status signaling in the context of both internal and external immune responses in groups of this non-social insect, thus showing parallels to social immunity.

Key Words - Group-living insect, Immune defense, Benzoquinones, Cuticular hydrocarbons (CHC), Immune status signaling

INTRODUCTION

In group-living animals increased contacts of individuals facilitate the spread of diseases (Kappeler et al. 2015). Thus, the individual ability to respond to immune stimuli and limit infection, but also the recognition of the health-status of group members are vital (Kappeler et al. 2015). Especially eusocial insect species show remarkable physiological and/or behavioral group-level responses to disease, such as distancing (Stockmaier et al. 2021) or hygiene behaviors (Rosengaus et al. 1998; Pull et al. 2018) leading to 'social immunity' (Cremer et al. 2018). Insight into individual and group level immunity across insect groups with different types of social structure is highly valuable. However, there have been only a few studies on the phenomenon of social transfer of immunity in non-social insects (Gallagher et al. 2018), and its underlying mechanisms. Horizontal transfer of immunity among conspecifics is often mediated *via* chemicals (Milutinović and Schmitt 2022). Nevertheless, little is known about the direct effects of immune stimuli like wounding and/or pathogen exposure on chemical phenotypes of individuals, which can potentially signal the immune status into the group and/or serve direct defensive functions.

Chemical cues can be crucial for healthy animals to identify diseased conspecifics across taxa (Csata et al. 2017; Pull et al. 2018; Behringer et al. 2020). As one group of such cues acting as contact infochemicals, changes in profiles of cuticular hydrocarbons (CHCs) may play a major role. CHCs cover the surface of insects and consist of complex mixtures of straight-chain, methyl-branched, and unsaturated hydrocarbons (Blomquist and Ginzel 2021). They serve as pheromones or allelochemicals for recognition of species, sex, mutualistic partners, or hosts, and for nestmate recognition and regulation of social interactions in eusocial insects (van Zweden and d'Ettorre 2010; Leonhardt et al. 2016). Many behavioral reactions to immunological and disease cues of nestmates in social insects, and even in a non-social insect species, could be directly related to changes in CHC profiles (Richard et al. 2008; Nielsen and Holman 2012; Baracchi et al. 2012; Hernández López et al. 2017; Alciatore et al. 2021). However, in an ant species immune experiences resulted in changes in the CHCs, but their role in the observed nestmate responses were unclear, suggesting volatile chemicals or behavioral cues as mediators (Alciatore et al. 2021). These findings underline that different infochemicals can be involved in information transfer among conspecifics (Müller et al. 2020).

In addition to contact cues, volatile compounds are known to largely mediate interactions

between organisms, both as pheromones detectable over large distances and/or with direct defensive

effects against pathogens or predators (Yew and Chung 2015). For example, odors emitted by diseased

brood induce hygienic behavior in bees (Swanson et al. 2009), and ant nest volatiles can have

antifungal effects (Wang et al. 2015). The release of defensive secretions into the environment can be

regarded as 'external immunity' (Cotter and Kilner 2010; Otti et al. 2014), i.e., putatively, a process of

niche construction (Odling-Smee et al. 2013; Müller et al. 2020). External immunity *via* secretions has

not only been found in several social species but also in various group-living (gregarious) insects, e.g.

beetles from the Tenebrionidae family (Cotter and Kilner 2010). Their stink gland secretions contain

mostly benzoquinones, which, along with further substances, have inhibitory effects on microbes and

thus serve a hygienic function (Yezerski et al. 2007; Li et al. 2013). This phenomenon is interpreted as

parental care towards larvae (Trivers 1972; Clutton-Brock 1991), but also other adults from the group

could benefit from secretions of conspecifics (Gokhale et al. 2017), resulting in herd-protection (Cotter

and Kilner 2010). Acting as pheromones, benzoquinones can trigger aggregation in low

concentrations, but also mediate dispersal, being repellent and autotoxic in high concentrations

(Chapman 1926; Roth and Howland 1941; Sokoloff 1974). However, the cascade from internal

immune experience to external immunological responses is rarely studied.

The red flour beetle, *Tribolium castaneum* (Coleoptera: Tenebrionidae), is a non-social, but

group-living insect that lives in and consumes flour, and serves as a model species (Campbell et al.

2022), also for studying resistance and immune reactions to various types of stimuli (e.g., Altincicek et

al. 2008; Wegner et al. 2008). Exposure to inactivated pathogen or a sublethal dose of living pathogen

has previously been shown to induce immune 'priming', which is a form of innate immune memory in

invertebrates (Little and Kraaijeveld 2004; Netea et al. 2016; Milutinović and Kurtz 2016), leading to

enhanced protection (Roth et al. 2009; Tate and Graham 2017; Khan et al. 2017), associated with

strong and specific transcriptional responses in *T. castaneum* (Greenwood et al. 2017; Ferro et al.

2019). Beyond that, even after experience of wounding alone this species shows social transfer of

immunity, as the interaction of naïve beetles with wounded conspecifics induces an enhanced activity

of the important immune effector phenoloxidase (Peuß et al. 2015). However, it is unclear if CHCs or

other infochemicals could act as cue for conspecifics to detect immune-challenged beetles. Moreover,

T. castaneum engages in external immune defenses, releasing benzoquinone-rich stink gland

secretions into the flour shared with conspecifics (Prendeville and Stevens 2002; Yezerski et al. 2007;

Joop et al. 2014). The role of the main compounds, methyl- and ethyl-1,4-benzoquinone (MBQ and

EBQ, respectively) and their carrier 1-pentadecene (Loconti and Roth 1953; Villaverde et al. 2007),

for inter- and intraspecific interactions in *Tribolium* beetles is well described (Blum 1981; Yezerski et

al. 2007; Verheggen et al. 2007; Duehl et al. 2011). For intraspecific communication, a sex-specific

function was suggested for benzoquinones in *T. castaneum*, being specifically involved in female

density-dependent competition (Khan et al. 2018). However, to our knowledge it is yet unknown how

immune experiences shape overall secretion profiles, whether these responses differ between females

and males, and whether potential changes are persistent over time.

We here examined changes in infochemicals of red flour beetles resulting from different immune

experiences, i.e., wounding alone or in combination with the injection of heat-killed *Bacillus*

thuringiensis bv. *tenebrionis* (bacterial beetle pathogen, 'priming' treatment) in comparison to naïve

beetles. We analyzed insect extracts using gas chromatography flame ionization detection and

expected changes in (1) the CHC profiles, which may act as a cue for the immune status of

conspecifics and (2) the gland secretion profiles, in particular quantities of EBQ, MBQ and 1-

pentadecene, which are involved in external immunity and population regulation.

Model Organisms. Our study was done on the outbred strain Croatia 1 (CRO1) of *T. castaneum* which

was kept at 30 ºC, 70% humidity and a 12 h:12 h light:dark cycle in plastic boxes containing heat-

sterilized (75 ºC for at least 24 hours) organic wheat flour (Bio Weizenmehl Type 550, dm-drogerie

markt) with 5% brewer's yeast powder (hereafter the flour-yeast mixture is referred to as "flour")

(Milutinović et al. 2013). For this experiment, around 2,000 one-month old adult beetles were allowed

to mate and oviposit for 24 h in approximately 350 g flour.

A culture of *Bacillus thuringiensis* bv. *tenebrionis* was obtained from Bacillus Genetic Stock

Center (Ohio State University, USA). The species is a bacterial pathogen of beetles, and immune

priming *via* injection of this bacterium has been demonstrated in previous work (e.g., Ferro et al.

2019). Heat-killed vegetative cells at 1×10^9 cells mL^{-1} used for the immune stimulation treatment

were prepared as previously described (Roth et al. 2009; Ferro et al. 2017) (see Methods S1 for

details).

Immune Treatment. Ten days after oviposition, all experimental beetles were individually placed into

sterile 96-well plates (Type F, Sarstedt) containing 0.08 g flour per well to minimize any injuries due

to conspecific cannibalism or mating (Park et al. 1965; Via 1999; Alabi et al. 2008). One week post

eclosion, virgin beetles of both sexes were divided into three treatment groups: Untreated as the

handling control ('naïve'), injected with sterile PBS ('wounded') and heat-killed *B. thuringiensis* bv.

tenebrionis injected ('primed'). For the latter two treatments, 18.4 nL (~ 20,000 cells per injection in

the 'primed' treatment) of solution was injected into the body cavity of adults with a nanoinjector

(Drummond Nanoject II) between the head and pronotum laterally towards the abdomen.

Subsequently, beetles were transferred into clean 96-well plates without flour, to reduce the

contamination of samples by faecal material and flour.

For the profiling of CHCs at 18 h post treatment, we prepared n = 6 samples per sex and

treatment with four beetles pooled into each sample. The sampling time was chosen based on a

previous study that reported transfer of immunity within 18 h post treatment (Peuß et al. 2015). For the

stink gland secretion collection at 24 h and 72 h post treatment, we obtained n = 8 samples of

individual beetles per sex, treatment and time point. These two time-points were chosen to test the

time course of the stink gland responses to the treatment.

Extraction of Beetle Chemicals. For extraction of the CHCs, four beetles per sample were pooled and

freeze-killed at -20 ℃. Samples were shaken in 120 µL of *n*-hexane (GC-MS grade, Merck)

containing 5 µL of eicosane (99.5%, Sigma-Aldrich) solution (0.2 mg mL^{-1} in *n*-hexane) as internal

standard at 4 ℃ for 10 min. After centrifugation, 100 µL of each extract was dried under reduced

pressure and resolved in 80 µL of *n*-hexane. The volume of *n*-hexane was scaled down accordingly in

three samples, where less than four beetles were available. For each set of samples (secretions and

CHCs), four blank samples were prepared alongside with the beetle samples but did not contain

beetles. For extraction of stink gland secretions, individual beetles were transferred to tubes that were

immersed in ice water at 3 ℃ for 3 min to stimulate the release of secretions (Joop et al. 2014).

Beetles were then freeze-killed at -20 ℃ and shaken in 65 µL ice-cold acetone (LC-MS grade, Sigma-

Aldrich) containing 0.05 mg mL^{-1} octadecane (98.5%, Sigma-Aldrich) as internal standard at 4 ℃ for

5 min. After centrifugation, 50 µL of each extract were kept at -20 ℃ until analysis.

Chemical Analysis, Data Pre-processing and Feature Identification. All samples were analyzed using

a gas-chromatograph coupled with a flame ionization detector (GC-FID-2010 Plus, Shimadzu) and

equipped with a VF-5ms column (30 m × 0.25 mm × 0.25 µm, with 10 m EZ-guard column, J&W

Agilent Technologies). For the analysis of CHCs, 1 µL from each sample was injected with a split

ratio of 2 at a constant nitrogen flow of 1.02 mL min^{-1}. The column oven temperature started at 100

℃, first increased to 230 ℃ with a rate of 20 ℃ min^{-1}, then increased to 260 ℃ with a rate of 5 ℃

min^{-1}, further increased to 277 ℃ with a rate of 1 ℃ min^{-1}, and finally increased to 320 ℃ with a rate

of 5 ℃ min^{-1}, which was held for 9.9 min. For the analysis of stink gland secretions, a sample volume

of 1 µL was injected with a split ratio of 3 at a constant stream of hydrogen flow of 1.13 mL min^{-1}.

The column oven temperature was set at 50 ℃, held for 2 minutes and increased to 300 ℃ with a rate

of 10 ℃ min^{-1}, followed by a holding time of 10 min. Additionally, two alkane mix standards, C8-20

and C8-C40 (both Sigma-Aldrich), were analyzed using both chromatographic methods. Standards of

MBQ (98% purity), methyl-1,4-hydroquinone (99%, both Sigma-Aldrich) and ethyl-1,4-hydroquinone

(97%, Abcr) were analyzed using the method for gland secretions. The hydroquinones are the form to

which benzoquinones convert to (Yezerski et al. 2000) and could thus appear in the datasets.

Data pre-processing was done using the GCsolution Postrun software (Version 2.30.00,

Shimadzu), and retention time alignment of peaks in R (version 4.0.3, R Core Team 2020) using the

package *GCalignR* (Ottensmann et al. 2018); for feature selection criteria and details on settings see

Methods S2. Data were normalized to the internal standard and relative feature values calculated

(proportion of total amount per sample). For identification, the retention times of peaks from the

alkane mix standards were used to calculate the retention index (RI) of detected features according to

(Kováts 1958). Putative identification of CHCs was done by comparing the RIs with those published

in previous studies (Lockey 1978; Alnajim et al. 2019; Awater-Salendo et al. 2020). Benzoquinones

(BQs) as well as other major secretion compounds previously described in *T. castaneum* glandular

secretions were identified by comparing their RIs to previous studies (Li et al. 2013; Lehmann 2015)

and by comparisons to the analyzed commercial standards.

Statistical Analyses. All statistical analyses were done using R (version 4.0.3, R Core Team 2020). To

visualize dissimilarities between groups of beetles, non-metric multidimensional scaling (NMDS)

based on pairwise Kulczynski distances (package *vegan*, version 2.5-7, Oksanen et al. 2020), was

applied to the relative datasets (Wisconsin-square-root-double-standardized) of CHCs and secretions,

respectively. After testing assumptions, permutational multivariate analysis of variance

(PERMANOVA) was performed to test for the influence of treatment on the dissimilarities, partly

combined with post hoc comparisons (Martinez Arbizu 2020) including corrections (Benjamini and

Hochberg 1995) (see Methods S3). Analyses of treatment effects were done separately for sexes and in

secretion datasets also separately for time points. Additionally, sex differences in the CHC profiles of

naïve females *versus* naïve males were addressed using PERMANOVA.

Responses of CHC profiles of females and males to treatments were visualized in heatmaps using

the package *pheatmap* (Kolde 2019). Therefore, log_2 fold change values were calculated for all CHCs

and replicates in comparison to the median value of the naïve female or male beetles and data plotted

as heatmaps. Moreover, heatmaps comparing CHC profiles of naïve females and naïve males were

generated based on z-scores for each CHC. All heatmap plots included Euclidean cluster analysis.

For the secretions, we additionally identified features most important for profile differences

between treatment groups within females and males 24 h after treatment using a conditional random

forest classification (for details see Methods S3) (Strobl et al. 2009a, b; De Moraes et al. 2014). The

relative amounts of these features, as well as the absolute amounts of MBQ, EBQ and 1-pentadecene

(1-C15-ene), were analyzed for an influence of treatment within females and males at both time points.

We tested the model assumptions (see Methods S3), and either fitted linear models with appropriate

transformations revealed using the package *bestNormalize* (Peterson 2021), or performed non-

parametric Kruskal-Wallis tests. When significant treatment effects were founds, Benjamini-

Hochberg-corrected post hoc comparisons were performed between treatment groups.

RESULTS

Cuticular Hydrocarbons. In total, 20 CHCs were putatively identified in the dataset. These consisted

of seven *n*-alkanes (C25-C31), six 3- and 4-methyl-branched alkanes and seven internally methyl-

branched alkanes of which one was a dimethyl alkane (Table S1). Significant immune treatment

effects on the CHC profile were found for both sexes (Table 1). In both females and males, the CHC

profiles of naïve beetles differed from those of immune treated beetles, but separated stronger from

profiles of wounded than of primed beetles (Fig. 1a,c). Accordingly, in post hoc tests on datasets of

both females and males, CHC profiles of naïve beetles differed significantly from those of wounded,

but not from those of primed beetles, while CHC profiles of wounded and primed beetles were quite

similar (Table 1). Interestingly, despite general sex differences (Fig. S1), the same CHCs were overall

involved in separation of naïve from wounded and primed beetles in females and males (Fig. 1). As

inferred from the CHC scores in the NMDS plot and heatmaps showing treatment-related differences

relative to naïve beetles, higher proportions of shorter methyl alkanes (C26-C29) characterized CHC

profiles of wounded and primed beetles in comparison to those of naïve beetles (Fig. 1). The CHC

profiles of naïve beetles had higher proportions of comparatively longer *n*-alkanes (C29-C31) and also

of 3-MeC31 (Fig. 1).

NMDS plots of CHC profiles of naïve females *versus* males resulted in a clear separation of the

sexes (Fig. S1a), which was mirrored in a significant difference (PERMANOVA; $F = 13.5$, $R^2 = 0.58$,

$P = 0.002$). Most obviously, the CHC profiles of female beetles were characterized by higher

proportions of rather short-chain methyl alkanes, especially of the cluster comprising 10-/11-/12-

MeC26, 3-MeC27 and 13-MeC27 (Fig. S1a,b). In contrast, male CHC profiles harbored higher

proportions of rather long-chain *n*- and methyl alkanes, especially of the cluster comprising 3-MeC29,

n-C30 and *n*-C31 (Fig. S1a,b).

Stink Gland Secretions. For analyses of the dataset acquired 24 h after treatment, 26 chemical features

were included, of which 24 were also considered in the dataset acquired 72 h after treatment (Table

S2). The benzoquinones MBQ and EBQ could clearly be identified, but the corresponding

hydroquinones were not detected in the dataset. The other five identified features were alkenes with

one or two double bonds (Table S2). The amounts of benzoquinones, together with 1-C15-ene made

up the major part of the overall secretion composition (Table S2).

For secretion profiles of female beetles 24 h after treatment, the treatment effect was marginally

non-significant (PERMANOVA, $F = 2.0$, $R^2 = 0.17$, $P = 0.057$). As visible in the NMDS plot, this

tendency resulted mainly from separation of naïve and wounded females (Figs 2a, S2). Analyzing

male beetles sampled 24 h after treatment, the composition of the overall secretions seemed visually

quite similar in all three treatment groups (Figs 2c, S3), and there was no significant treatment effect

($F = 1.4$, $R^2 = 0.12$, $P = 0.180$). To explore if individual features from the profiles respond to immune

challenge in females *versus* males, random forest analysis was applied for both sexes separately.

Random forest analysis of female beetles revealed one unidentified feature, RI 1015, as important for

separation of treatment groups (Fig. S4a). The relative amount of this feature was significantly

influenced by treatment in females (*Kruskal-Wallis test*; 2 df, $\chi^2 = 9.84$, $P = 0.034$; males: 2 df, $\chi^2 =$

1.72, $P = 0.490$) and lower in both primed and wounded beetles compared to naïve ones (Fig. S4b).

The unidentified feature RI 1473 was suggested as important in males (Fig. S5a). The relative amount

of this feature was significantly influenced by treatment in males (*Kruskal-Wallis test*; 2 df, $\chi^2 = 9.27$,

$P = 0.034$; females: 2 df, $\chi^2 = 3.43$, $P = 0.370$), being higher in wounded than in naïve beetles (Fig.

S5b). In the dataset acquired 72 h after treatment, neither females (PERMANOVA, $F = 0.6$, $R^2 =$

0.105, $P = 0.802$) nor males ($F = 0.6$, $R^2 = 0.06$, $P = 0.854$) revealed statistical or visual effects of

treatments on the secretion profiles (Fig. 2b,d) and thus further random forest analysis was not

performed.

Furthermore, we compared the absolute amounts of the major compounds MBQ, EBQ and their

carrier 1-C15-ene. In the secretions acquired after 24 h, an effect of treatment was only significant for

females, but not for males in these three compounds (Table 2, Fig. 3). In post hoc tests, the absolute

amounts of all three compounds differed significantly between naïve and wounded females (Table 2),

being higher in wounded individuals (Fig. 3). The amounts of MBQ and EBQ were by trend also

higher in wounded than in primed females and this difference was significant for 1-C15-ene (Table 2,

Fig. 3). In the dataset acquired 72 h after treatment, no treatment effects were found in females or

males (Table 2). Interestingly, in secretions collected after 72 h the benzoquinones in primed, not in

wounded females reached the highest average amounts (Fig. 3).

DISCUSSION

As group-living animals that share the same food resources and live in proximity, flour beetles are prone to increased disease transmission. We investigated the influence of different immune stimuli on profiles of different infochemicals in *T. castaneum*. Wounding led to similar changes in CHC patterns of females and males, potentially involved in transferring information on the immune status within the group. Wounding-induced changes in gland secretion compounds that could directly mediate external immunity and behavioral responses were mainly restricted to females. The response to priming was lower than to wounding across both types of infochemicals, emphasizing the complexity of physiological responses, especially regarding internal (individual) *versus* external (group) immunity.

The CHC profiles of the beetles were hypothesized to be altered upon immune stimulation *via* wounding and/or priming. Indeed, we found significant differences in the CHC profiles of naïve *versus* wounded females and males. These changed profiles may act as a potential cue for the detection of immune experience in conspecifics, leading to the previously described social transfer of immunity (enhanced phenoloxidase activity) from wounded to naïve *T. castaneum* during cohabitation (Peuß et al. 2015). Moreover, in that study cohabitation with wounded beetles resulted in reduction in expression of two heat shock protein genes, *Hsp83* and *Hsp90*, in naïve conspecifics (Peuß et al. 2015) within the same time span, in which profile changes occurred in the present study. As the molecular chaperone HSP90 serves as an evolutionary capacitor by mediating the storage and release of cryptic genetic variation (Rutherford and Lindquist 1998), such down-regulation has the potential to lead to increased adaptability. However, a causal relationship between changes in CHCs and transfer of information on wounding, including the consequences for social immunity and evolutionary capacitance, needs to be proven in additional bioassays.

The CHC profiles of immune-stimulated beetles in the present study, especially the wounded ones, were characterized by higher proportions of shorter methyl-branched alkanes. This is in line with studies on other insect species, in which specifically methyl alkane CHCs were altered by immune challenge (Richard et al. 2008; Baracchi et al. 2012; Alciatore et al. 2021), and could therefore contribute to social transfer of immunity. Despite higher metabolic costs for these CHCs (Nelson

1993), a generally more important role of methyl alkanes in comparison to linear *n*-alkanes as

interspecific recognition cues is well documented in social insects (van Zweden and d'Ettorre 2010).

In larvae of *T. castaneum*, wounding caused an upregulation of multiple candidate fatty acid synthase

genes (Behrens et al. 2014), which are involved in the synthesis of methyl alkanes (Blomquist and

Ginzel 2021; Holze et al. 2021). Assuming that CHC synthesis is regulated in comparable ways in *T.*

castaneum larvae and adults, larvae may carry equivalent information on the immune status in their

CHCs profiles. Moreover, the wounding-induced alterations in methyl alkanes in adult CHCs observed

in the present study may then result from similar, active processes as suggested for the larvae. Methyl

alkanes have furthermore been shown to directly mediate interspecific interactions in larvae of *T.*

castaneum, acting as host detection cues for parasitoids (Awater-Salendo et al. 2020). Taken together,

methyl-branched CHCs are very likely an important factor also in intraspecific transfer of information

in both larvae and adults of *T. castaneum.*

Furthermore, we could demonstrate a sexual dimorphism in the CHC profiles of naïve *T.*

castaneum adults, which is also well documented in various other invertebrate species (Jallon and

David 1987; Thomas and Simmons 2008; Müller and Müller 2016). However, CHC profiles seemed to

change in similar directions in *T. castaneum*. In contrast, in the non-social species *T. molitor* changes

of the CHC profiles after immune challenge were sex-specific, and the resulting (sexual) signals

towards conspecifics resulted in advantages for the wounded males (Nielsen and Holman 2012),

interpreted as terminal investment (Sadd et al. 2006; Nielsen and Holman 2012; Gallagher et al. 2018).

The similarity in the CHC profile changes across sexes in the present study, potentially equivalently

displaying wounding experience, may be beneficial for an effective transfer of information into the

group, and could thus be regarded as 'social trait'. To investigate the potential of CHCs to signal the

immune status to conspecifics and their species-specific ecological functions, studies are needed in

both sexes of various eusocial *versus* non-social insects.

Flour beetles perform niche construction by secreting quinones-rich stink gland secretions into

the flour, which influences microbial growth (Van Wyk et al. 1959; Sokoloff 1974) as a form of

external immunity (Joop et al. 2014; Otti et al. 2014). Considering this as parental care (Trivers 1972;

Clutton-Brock 1991), we expected that adult females and/or males may modulate their secretions upon

detection of pathogen risks to protect their offspring which lack the glands. We found an increase in

the absolute amount of the benzoquinones (MBQ, EBQ) and their carrier 1-C15-ene in the secretions

of wounded females 24 h after the treatment. The finding that particularly females may engage in

niche construction fits theory, proposing more resource investment in immunity and longevity

(Bateman 1948; Rolff 2002) or into brood protection (Trivers 1972) in females than in males.

However, these effects lasted for less than 72 h after immune treatments. Potential metabolic costs

associated with quinone production and individual survival costs caused by autotoxicity of quinones in

high concentrations (Yezerski et al. 2004; Joop et al. 2014; Gokhale et al. 2017), may explain the

transience of this effect. Moreover, on the group level enhanced quinone production may also increase

herd-protection (Cotter and Kilner 2010), because under certain circumstances external immunity can

be interpreted as public good (Yezerski et al. 2004; Joop et al. 2014; Gokhale et al. 2017). However,

increased quinone levels in the environment may provide benefits especially to non-secreting *T.*

castaneum conspecifics, which represents a social dilemma (Yezerski et al. 2004; Joop et al. 2014;

Gokhale et al. 2017). Thus, there is likely a complex trade-off between benefits and various individual

cost of quinone production, determining the time course of responses to immune experience. This

study underlines the importance to distinguish between sexes and time points when exploring

responses to immune challenge, potentially influencing niche construction.

Apart from serving as antimicrobials, benzoquinones and alkenes in stink gland secretions also

lead to dose-dependent behavioral responses in conspecifics of *Tribolium* beetles (Suzuki 1985;

Verheggen et al. 2007). For example, together with an aggregation pheromone, 4,8-dimethyldecanal,

benzoquinones and alkenes mediate dispersal (Ogden 1969; Verheggen et al. 2007; Duehl et al. 2011).

If wounding occurs at high population densities, an enhanced secretion of benzoquinones can signal

overcrowding (Faustini and Burkholder 1987; Verheggen et al. 2007). In accordance with the results

of our study, females appear particularly sensitive to perceive certain other cues, for example the

secretions of other females in the flour, responded by upregulating their own quinone production

(Sonleitner and Gutherie 1991; Khan et al. 2018). High amounts of quinones can even suppress the

fecundity of conspecifics, which has huge consequences for population density regulation (Sonleitner

and Gutherie 1991; Khan et al. 2018). We determined two important, yet unidentified compounds

from secretion profiles using random forest analysis, which revealed significant sex-specific treatment

effects on their relative amounts. The modulation of these compounds after immune challenge,

potentially together with other cues, may serve yet unknown sex-specific functions, as described for

pheromones in *T. molitor* (Rantala et al. 2002; Sadd et al. 2006; Nielsen and Holman 2012). Further

quantification of quinone amounts actually present in the flour environment, together with

identification of other potentially relevant compounds could help elucidating how the beetles regulate

their environment and/or population density *via* their stink gland secretions.

　　Comparing different types of immune stimulation, in a previous study wounding alone (i.e.,

pricking control) resulted in weaker immune gene expression than treatment with alive *B.*

thuringiensis bv. *tenebrionis* (Behrens et al. 2014). Here, however, the response of *T. castaneum* to

priming was weaker than the response to wounding across both types of infochemicals, CHCs and

quinones. Firstly, wounding may quickly establish a transient physiological state of heightened

'alertness', leading to responses with quinones as broadly-acting antimicrobials (Joop et al. 2014;

Pedrini et al. 2015; Sawada et al. 2020) and potentially displayed to conspecifics in changed CHC

profiles. Meanwhile, immune priming *via* injecting heat-killed bacteria may carry more information

and induce additional physiological and immunological responses that mediate more tailored defenses

against certain types of pathogens (Roth et al. 2009; Ferro et al. 2019). Secondly, potential competition

for tyrosine precursors between both the quinone production and the phenoloxidase cascade activated

upon bacterial challenge (Behrens et al. 2014; Joop et al. 2014; Otti et al. 2014), may result in a trade-

off between external and internal immunity. Initially prioritizing individual internal immunity over

external immunity, which confers protection also on adjacent conspecifics, may be beneficial for this

non-social species. The aforementioned possible reasons together with the finding that 72 h after

treatment primed beetles had the highest average amounts of quinones, hint to a delay or shift in

immunological processes rather than absence of responses in primed beetles. This may indicate a

possible more durable enhanced protection representing a 'primed' physiological state. Furthermore,

the common patterns of a more pronounced response to wounding than to priming in both CHCs and

quinone secretions may suggest that the modulation of both types of chemicals shares a common

physiological cascade. For example, cytochrome P450 genes can be involved in CHC modification in

insects (Chen et al. 2016) and have been shown to be crucial for quinone and alkene production in *T.*

castaneum (Li et al. 2013; Lehmann 2015). Further analyses are needed to explore immune responses

in CHCs and secretion compounds and their associated metabolic processes upon wounding *versus*

priming over time.

In conclusion, our study highlights the potential role of chemical profiles in mediating the

intertwined processes of immunity and niche construction in the red flour beetle. Immune stimulation

triggered changes in potentially important infochemicals which can have fitness consequences for both

the individual and the group. Furthermore, our results emphasize the high specificity of responses

towards different types of immune stimuli and the importance to focus on sex-specific functions of

chemicals. More generally, regarding the special lifestyle of a group-living but non-social insect, these

results bear fundamental relevance for questions of common and specific mechanisms of disease

control in groups and societies across taxa.

Data availability

Raw data and R codes are provided in the supplementary information. The GC-FID raw data will be released upon publication of the paper in the MetaboLights repository (Haug et al. 2013, 2020), being available with the accession number MTBLS2277 at www.ebi.ac.uk/metabolights/MTBLS2277.

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Methods S2 *Details on Data Processing*

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Fig. S1 NMDS plot and heatmaps of cuticular hydrocarbon profiles from naïve *T. castaneum*

Fig. S2 NMDS plot of glandular secretion profiles from immune treated female *T. castaneum*

Fig. S3 NMDS plot of glandular secretion profiles from immune treated male *T. castaneum*

Fig. S4 Mean decrease in accuracy plot from random forest analysis of glandular secretion profiles of immune treated female *T. castaneum* and relative amounts of relevant feature RI 1015

Fig. S5 Mean decrease in accuracy plot from random forest analysis of glandular secretion profiles of immune treated male *T. castaneum* and relative amounts of relevant feature RI 1473

Table S1 Cuticular hydrocarbons and their relative composition in immune treated *T. castaneum*

Table S2 Features putatively identified from glandular secretions of *T. castaneum* beetles

Table S3 Candidate fatty acid synthase genes upregulated post wounding in *T. castaneum* larvae

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Statements and Declarations

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Competing interests

No competing interests declared.

Author Contributions

JK and CM conceived the ideas. JK, CM, LJT, LKL and RR planned the experimental procedure. LKL and RR carried out the experiments and analyzed the raw data together with LJT. BM, LJT and LKL carried out the statistical analyses. All authors contributed to writing of the article and approved the submitted version.

 Figure captions

 Fig. 1 NMDS plots (a,c) and heatmaps (b, d) of cuticular hydrocarbon (CHC) profiles from female

 (**a,b**) and male (**c,d**) beetles of *Tribolium castaneum*. Beetles (n = 6 per sex and treatment) were

 untreated ('Naïve'), or treated by injection of phosphate buffered saline (PBS; 'Wounded') or of PBS

 with heat-killed *Bacillus thuringiensis* bv. *tenebrionis* ('Primed') 18 h before extraction. All plots are

 based on relative amounts of 20 individual CHCs. For NMDS plots, datasets were transformed with

 Wisconsin square root double standardization and dissimilarities are based on pairwise Kulczynki

 distances. Data points represent profiles of individual beetles, larger symbols show centroids for

 treatment groups. Arrows range from zero to positions of expanded weighted average scores of CHCs.

 Heatmaps show increase and decrease in the proportion of individual CHCs in wounded and primed

 females (**b**) and males (**d**) in relation to the median value of female or male naïve beetles as $\log_2$ fold

 change. Clustering of CHCs is based on Euclidean distances

 Fig. 2 NMDS plots of glandular secretion profiles from female (**a**, **b**) and male (**c**, **d**) beetles of

 Tribolium castaneum of different immune treatment groups. Beetles (n = 7-8 per sex and treatment)

 were untreated ('Naïve'), or treated by injection of phosphate buffered saline (PBS; 'Wounded') or of

 PBS with heat-killed *Bacillus thuringiensis* bv. *tenebrionis* ('Primed') 24 h (**a**, **c**) or 72 h (**b**, **d**) before

 extraction. Plots are based on relative amounts of 26 individual chemical features released from

 glands. Datasets were transformed with Wisconsin square root double standardization. Dissimilarities

 are based on pairwise Kulczynki distances. Data points represent profiles of individual beetles, larger

 symbols show centroids for treatment groups

 Fig. 3 Amount of methyl-1,4-benzoquinone (MBQ), ethyl-1,4-benzoquinone (EBQ) and 1-

 pentadecene (1-C15-ene) from glandular secretions of female and male beetles of *Tribolium*

 castaneum at two time points (24 h, 72 h) after treatment. Beetles (n = 7-8 per sex, treatment and time

 point) were untreated (naïve, 'N'), or treated by injection of phosphate buffered saline (PBS; wounded,

689 ‘W’) or of PBS with heat-killed *Bacillus thuringiensis* bv. *tenebrionis* (primed, ‘P’) 24 h or 72 h

690 before extraction. Boxes show 25th and 75th percentiles, lines are medians, crosses are means,

691 whiskers mark minimum and maximum within 1.5-fold interquartile ranges. Dots represent values of

692 individual beetles

Figures

Fig. 1

698   **Fig. 2**

Fig. 3

704   **Tables**

705   **Table 1** Effects of treatment on the composition of CHC profiles in female and male *Tribolium*

706   *castaneum* beetles in a permutational analysis of variance (overall tests and pairwise comparisons)

	F	R^2	P
treatment (females)	2.5	0.25	**0.023**
naïve vs wounded	4.8	0.33	**0.006**
naïve vs primed	2.0	0.17	0.153
wounded vs primed	0.6	0.06	0.630
treatment (males)	2.4	0.24	**0.009**
naïve vs wounded	3.8	0.28	**0.009**
naïve vs primed	2.1	0.18	0.110
wounded vs primed	1.2	0.10	0.317

Beetles (n = 6 per sex and treatment) were untreated ('naïve'), or treated by injection of phosphate
buffered saline (PBS; 'wounded') or of PBS with heat-killed *Bacillus thuringiensis* bv. *tenebrionis*
('primed') 18 h before extraction. The datasets included relative values of 20 CHCs and were
transformed with Wisconsin square root double standardization. Dissimilarities are based on pairwise
Kulczynki distances. Significant *P*-values shown in bold.

Table 2 Effects of treatment on absolute amounts of methyl-1,4-benzoquinone (MBQ), ethyl-1,4-

benzoquinone (EBQ), and 1-pentadecene (1-C15-ene) from glandular secretions of female and male

beetles of *Tribolium castaneum* in linear models (overall tests) and Tukey HSD post hoc tests

(pairwise comparisons)

	MBQ			EBQ			1-C15-ene		
24 h after treatment	df	*F/t*	*P*	df	*F/t*	*P*	df	*F/t*	*P*
treatment (females)	2	5.36	**0.014**	2	5.12	**0.016**	2	9.02	**0.002**
naïve vs wounded	1	3.16	**0.015**	1	3.07	**0.018**	1	4.23	**0.001**
naïve vs primed	1	0.79	0.439	1	0.70	0.492	1	1.69	0.107
wounded vs primed	1	2.27	***0.052***	1	2.27	***0.052***	1	2.40	***0.040***
treatment (males)	2	0.37	0.694	2	0.60	0.559	2	1.31	0.292
72 h after treatment									
treatment (females)	2	1.13	0.343	2	1.23	0.313	2	0.12	0.887
treatment (males)	2	0.03	0.972	2	0.05	0.956	2	0.09	0.912

716 Beetles (n = 7-8 per sex, treatment and time point) were untreated ('naïve'), or treated by injection of
717 phosphate buffered saline (PBS; 'wounded') or of PBS with heat-killed *Bacillus thuringiensis* bv.
718 *tenebrionis* ('primed') 24 h or 72 h before extraction. Significant *P*-values shown in bold, marginally
719 significant *P* values in bold and italics. Column *F/t* shows *F* value in linear models and *t* value in
720 Tukey HSD post hoc tests.