Eating in a losing cause: anorexia and macronutrient manipulation by a

cockroach fails to boost immunity

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Summary

1. Host-pathogen interactions can lead to dramatic changes in host feeding behaviour. One aspect of this includes self-medication, where infected individuals consume substances such as toxins, minerals or secondary compounds or alter their macronutrient consumption to enhance immune competence. Another aspect includes illness-induced anorexia, which is a general mechanism adopted by animals following infection. Anorexia is thought to assist host immunity directly or by limiting the nutritional resources available to pathogens.

2. Here, we recorded macronutrient preferences of the global pest cockroach, *Blatta orientalis* to investigate how shifts in host macronutrient dietary preference and quantity interact with immunity following sublethal bacterial infection.

3. We compared the survival of uninfected and infected individuals on diets enriched with carbohydrate (C) or protein (P) and conducted food-choice experiments after challenging hosts with the common entomopathogenic soil bacterium *Pseudomonas entomophila*. We then carried out a quantitative proteomic analysis and an antimicrobial activity assay of hemolymph from infected individuals that had been restricted to diets with defined macronutrient compositions.

4. We find that diets enriched for P reduce cockroach survival. However, following immune challenge by *P. entomophila*, cockroaches significantly reduced their overall nutrient intake, particularly of carbohydrates, and increased the ratio of protein (P:C) consumed. Surprisingly, these behavioural shifts did not significantly improve cockroach immunity or

survival, with negligible differences in immune protein abundance and antimicrobial

activity between infected individuals placed on P- or C- rich diets.

5. We show that cockroach feeding behaviour can be dynamically modulated by a pathogen,

resulting in an illness-induced anorexia-like feeding response and preference for P-

enriched food. Interestingly however, our results also demonstrate that these responses

hold no special advantage for the host's immune response. The lack of a benefit following

the host's behavioural shift highlights a possible decoupling of dietary regulation from

immunity in these invasive animals.

Introduction

Microbe symbioses form a fluctuating but universal backdrop to animal life. However, the

evolutionary processes that drive animal hosts and their symbionts, including pathogens,

operate at different scales and often in opposing directions (Dawkins & Krebs 1979), with

the animal immune system acting as a key interface between host and symbiont ecology

(Schmid-Hempel 2003). In addition to the core immune system, behavioural mechanisms

have attracted increasing attention for their ability to co-ordinate host responses to

infection (Simpson *et al.* 2015; Wong *et al.* 2015). While behaviour is the primary means by

which animals interact with the biotic environment, its importance for a wide range of

immune-related functions has only relatively recently come to the fore.

Hosts can respond behaviourally before infection has even taken place. This can include

avoidance of pathogen transmission areas (e.g. defecation sites) and deterrence of disease

vectors (Hart 2011; Moore 2013). Other prominent examples include activities falling within

the category of "social immunity", which among insects can include pathogen detection

alarm behaviours (Rosengaus *et al.* 1999); grooming of conspecific group members

(Rosengaus *et al.* 1998; Reber *et al.* 2011); removal (Armitage *et al.* 2016) or even

destruction of infected individuals (Yanagawa *et al.* 2011). Such mechanisms are well

documented in many social insect lineages, where they contribute significantly to a number

of prophylactic mechanisms operating within societies (Schmid-Hempel 1998; Cremer *et al.*

2007). Other prophylactic social behaviours include the collection of secondary antimicrobial

compounds to prevent microbial growth in the nest environment (Castella *et al.* 2008;

Simone *et al.* 2009), in addition to the direct use – typically via feeding – of antimicrobials in

both individual and transgenerational prophylaxis (Lefevre *et al.* 2010; Lefevre *et al.* 2012;

Milan *et al.* 2012; de Roode *et al.* 2013; Kacsoh *et al.* 2013).

Once infection has occurred, the first and principal line of defence is the immune system.

Here, behavioural defensive adaptations can also play an important role in regulating or

augmenting the response to infection. As with prophylaxis, the role of feeding behaviour has

increasingly been viewed as a key mechanism by which animals can respond to infection

(Abbott 2014). Here, the selection of novel antimicrobial compounds, or the enrichment of

specific dietary elements can be employed as therapeutic treatment against pathogens (de

Roode *et al.* 2013). Fruit flies use ethanol therapeutically as well as prophylactically to

combat parasitoid wasp infection (Milan *et al.* 2012) whereas parasitoid fly-infected

Grammia caterpillars mix pyrrolizidine alkaloid-producing toxic plants into the normal diet to

assist parasitoid clearance, which comes at the expense of body growth (Singer *et al.* 2004;

Smilanich *et al.* 2011).

Infection-induced adaptive changes to feeding behaviour can also involve modifications to

the quantity and composition of macronutrients in the diet. Anorexia is a well-documented

response to infection in both vertebrates (Johnson *et al.* 1993; Konsman *et al.* 2002) and

invertebrates (Adamo *et al.* 2007; Ayres & Schneider 2009) and is thought to assist hosts in

limiting nutritional resources available to pathogens (Kluger & Rothenburg 1979). Anorexia

may also help by activating components of the immune system that are enhanced under

conditions of nutritional stress, such as autophagy (van Niekerk *et al.* 2016a; van Niekerk *et*

al. 2016b). In recent years, the balance of macronutrients itself has been examined as a way

for animals to regulate the response to infection. In particular, the proportion of protein has

been shown to be an important criterion in animal choice of diet following infection. In

Spodoptera moths, larvae select a diet enriched in protein following infection with a

generalist gram-positive bacterium and a host-specific DNA virus (Lee *et al.* 2006; Povey *et*

al. 2009; Povey *et al.* 2013), leading to enhanced antimicrobial activity in both cases. By

contrast, diets enriched in carbohydrate were selected when *Tenebrio* beetles and *Grammia*

caterpillars were infected with a rat tapeworm (Ponton *et al.* 2011) and a parasitoid fly

(Mason *et al.* 2014), respectively. In the latter study, this behaviour was also associated with

an enhanced melanization response.

The use of macronutrients by hosts to regulate immunity could in principle apply to any

animal that is not an obligate food specialist. However, virtually nothing is known about the

relationship between macronutrient diet choice and immunity outside of a handful of

holometabolous, mostly herbivorous, insect larvae. Holometabolous insects undergo

complete metamorphosis consisting of distinct larval, a pupal and an adult winged phase,

which are typically correlated with vastly different ecologies and corresponding

physiological, morphological and immunological conditions (McMahon & Hayward 2016). By

contrast, hemimetabolous insects undergo progressive molts where each larval instar closely

resembles the adult (Sehnal *et al.* 1996). Our understanding of the interface between diet,

behaviour and immunity would greatly benefit from studying animals that vary widely in

their taxonomy, ecology and development.

Our study addresses this by examining the interaction between macronutrient feeding

behaviour and immunity in the omnivorous cockroach, *Blatta orientalis*. We investigate the

macronutrient preferences of individuals in response to a range of sublethal immune

challenges, before examining the impact of macronutrients on host survival, immune

competence and finally, the expression of the host's proteome, which captures a critical

component of the host's response to a pathogen. Overall, our findings suggest that diet is

decoupled from immune regulation in cockroaches.

Materials and Methods

INSECTS AND BACTERIA

A breeding culture of sequential *Blatta orientalis* cohorts was established at the Federal

Institute for Materials Research and Testing (BAM) in June 2015, initially obtained from the

collection at the Federal Environment Agency, Berlin, which consists of a mixed population

of four independent genetic backgrounds maintained for 50 generations. Each generation

consists of a minimum of 150 breeding pairs of cockroaches to minimize the effects of

inbreeding. Each experimental cohort was maintained for approximately 190 days in the

dark at 26 °C and 50 % humidity. Prior to being placed on experimental (artificial) diets,

animals were reared on a mixture of 77.0 % dog biscuit powder, 19.2 % oat flakes and 3.8 %

brewer's yeast and supplied with water *ad libitum* and weekly with apple and carrot slices.

All experiments were conducted with males to minimise changes in physiology associated

with oogenesis. Each individual was used only once in each experiment. For the food choice

experiment and the survival on enforced diets, individuals from 3 different cohorts were

used. The generalist gram-negative bacterial pathogen *Pseudomonas entomophila* (strain

L48; DSM No. 28517) which is able to infect a variety of insect orders (Vallet-Gely *et al.* 2010)

was obtained from the Leibniz Institute DSMZ-German Collection of Microorganisms and Cell

Cultures. Bacteria were stored at -70 °C until use in experiments.

ARTIFICIAL DIETS

The artificial diets used in this study are based on isocaloric diets, as described elsewhere

(Lee *et al.* 2006; Povey *et al.* 2013), which was slightly modified to suit cockroach needs. We

employed two diets containing 35 % carbohydrate (C) and 7 % protein (P) or *vice versa*. The

carbohydrate portion consisted of sucrose while the protein portion consisted of casein,

peptone and albumin from eggs in a 3:1:1 ratio. Remaining ingredients are listed in

Supplementary Tab. 1. Diet blocks of approximately 0.125 cm³ in size were dried at 50 °C for

2 days before being weighed and given to experimental cockroaches.

BACTERIAL INOCULATION

About 200 µl of an overnight culture of *P. entomophila* was mixed in 10 ml fresh liquid

medium (according to DSMZ instructions) and incubated at 28 °C and 140 rpm to an OD_{600nm}

of 0.55, representing 1.5×10^8 CFUs (determined by plating). The desired concentrations of

bacteria were subsequently obtained by diluting bacteria in insect Ringer's solution (0.024 g

calcium chloride, 0.021 g potassium chloride, 0.01 g sodium hydrogen carbonate, 0.45 g

sodium chloride, 200 ml dest. water). Cockroaches were anaesthetised with CO_2, abdomens

swabbed with 70 % ethanol, then injected with 2µl of bacterial solution directly into the

hemocoel using a glass capillary needle inserted between the 3rd and 4th abdominal

segment. Sublethal infections (high: 5.8 x 10^5, low: 5.8 x 10^3) and lethal (4.5 x 10^6) doses

were determined in pre-experiment injection assays.

DIET CHOICE FOLLOWING SUBLETHAL INFECTION

From each of 3 cohorts, 40 *B. orientalis* males (120 in total) were given free choice of

macronutrients by placing them together with 1 block of known weight of each artificial diet.

Individuals were kept for three days to accustom them to artificial diets, and to obtain a

baseline P:C ratio preference. Thereafter food blocks were collected, placed at 50°C until

completely dry, and then their weight loss was determined, equating to the amount eaten

by the cockroach. Experimental cockroaches were assigned randomly to one of the following

sublethal treatments (40 per treatment): 1) High infection (injected 5.8 x 10^5 *P. entomophila*

CFUs); 2) Low infection (injected 5.8 x 10^3 *P. entomophila* CFUs); 3) Wounding control

(injected Ringer's solution); 4) Unmanipulated control. Cockroaches were then placed on

new food blocks of both diets of known weight. The blocks were replaced daily for four days

and their loss of weight was again determined after drying at 50°C.

SURVIVAL ON ENFORCED DIET

From each of 3 cohorts, 20 *B. orientalis* males were placed on protein-rich diet (35 % P; 7 %

C) and another 10 were placed on carbohydrate-rich diet (7 % P; 35 % C). All individuals were

supplied with water *ad libitum*. Survival was checked twice weekly; food blocks and water

were changed once a week over 150 days.

SURVIVAL ON ENFORCED DIET FOLLOWING LETHAL INFECTION

One hundred and fifty *B. orientalis* males were equally assigned to one of the following

lethal treatments: 1) Infection (injected 4.5 x 10^6 *P. entomophila* CFUs); 2) Wounding control

(injected Ringer's solution); 3) Unmanipulated control. Half of the individuals from each

treatment were randomly assigned to either a protein- (35 % P; 7 % C) or carbohydrate-

enriched (7% P; 35% C) artificial diet and supplied with water *ad libitum*. Survival of each

individual was recorded for six days.

IMMUNE ASSAYS

Immune challenge and hemolymph collection

One hundred and twenty *B. orientalis* males were immune-challenged by injecting 2 µl

Ringer's solution containing 5.8 x 10^5 *P. entomophila* CFUs. Half were assigned to the

protein- (35 % P; 7 % C) and the other half to the carbohydrate-enriched (7 % P; 35 % C)

artificial diet and supplied with water *ad libitum*. After 24 hrs, the hemolymph from each

individual was collected by cutting the first 2 leg pairs and placing the cockroach head-first

into a spin-column (Sigma-Aldrich) in a 1.5 ml tube containing propylthiouracil (to inhibit

phenol-oxidase activity). They were then centrifuged at 500 *g* for up to 5 min or until 10 µl of

hemolymph was collected. The hemolymph of 10 individuals of the same treatment was

pooled resulting in 6 pools per treatment and stored at -70 °C until needed.

Bacteria growth inhibition assay

The bacterial growth inhibition of the cockroach hemolymph was measured using a plate

reader assay. First, 10 µl Mueller-Hinton broth were added to each well of a 384-well

polypropylene plate. Then 10 µl hemolymph was loaded in the second column of the plate.

One well contained the hemolymph of one pool of animals (in total 12 wells loaded with

hemolymph). Wells in the first column which did not contain hemolymph served as the

negative control. A seven-step serial dilution of the hemolymph was performed (with the

last 10 µl being discarded) and 10 µl *P. entomophila* in Mueller-Hinton broth with an OD_{600} of

0.005 was added to each well containing hemolymph, as well as to a further column not

containing hemolymph, which served as a positive control for unsuppressed bacterial

growth. OD_{600} was measured in a plate reader (BioTek) every 10 min for 16 hrs.

PROTEOMIC ANALYSIS BY MASS SPECTOMETRY

A detailed description of protein sample preparation and liquid chromatography-mass

spectrometry and data processing is described in Supplementary file 1. Initially, we carried

out *de novo* transcriptome sequencing to generate a peptide database for *B. orientalis*.

Briefly, RNA was extracted from *B. orientalis* by homogenizing individuals in pre-cooled Trizol

(Thermo Fisher Scientific) and recovering RNA using chloroform extraction and isopropanol

precipitation, and treatment with TurboDNase (Ambion) according to manufacturer's

instructions. mRNA libraries were enriched and prepared using a NEXTflexTM Rapid

Directional mRNA-seq Kit protocol (Bioo Scientific) before being sequenced on an Illumina

NextSeq500/550 platform at the Berlin Center for Genomics in Biodiversity Research

(BeGenDiv). Raw data were processed and annotated as described elsewhere (He *et al.*

2018) (Supplementary file 1). For proteomic analysis, protein identification and label-free

quantification was performed using MaxQuant (v1.6.0.1) with Andromeda search engine

(Cox & Mann 2008; Cox *et al.* 2011; Tyanova *et al.* 2015). Raw data were matched against an

in-house protein database of *B. orientalis* created by *de novo* transcriptome sequencing (see

above). Trypsin was selected as enzyme allowing a maximum of two missed cleavages. The

minimum peptide length was set to 7 amino acids and the false discovery rate (FDR) for

peptide and protein identification was set to 0.01.

STATISTICAL ANALYSIS

P:C ratios, the amounts of P and C eaten as well as total consumption differences between

treatments were analysed on the first day following infection using Bonferroni-corrected

Wilcox rank sum tests.

The food-choice data were analyzed using a Generalized linear mixed model (GLMM) with an

underlying beta family distribution. Analyses were run in the glmmADMB package v0.8.3.3

(Fournier *et al.* 2012; Skaug *et al.* 2014) in R 3.4.3 (R Core Team 2018) in conjunction with

the R2admb package v0.7.16 (Bolker *et al.* 2017). GLMMs examined whether a response

variable consisting of proportion of P consumed (amount of P eaten divided by the amount

of diet available) or proportion of C (amount of C eaten divided by the amount of diet

available) was influenced by treatment (high infection; low infection; wounded; and

unmanipulated) and day post infection, as well as an interaction between treatment and

day. Minimal adequate models were derived by stepwise-model simplification and

comparison via ANOVA. Individual and cohort were treated as random effects to account for

multiple measurements and origin. Comparisons among treatment levels were carried out

with post-hoc Tukey tests using a Bonferroni correction, using package multcomp v1.4-8

(Hothorn *et al.* 2008). Individuals 16, 48 82, 84 and 91 were removed prior to analysis due to

the presence of fungal growth on the artificial diet blocks.

The effect of treatment (infection and control) and diet (P- and C-enriched) on survival were

analysed using Cox proportional hazard models in R 3.3.1 with the survival package v2.39-5

(Therneau 2015). Bacterial growth inhibition data were analyzed using R package nparLD

v2.1 (Noguchi *et al.* 2012) using default parameters.

Results

DIET CHOICE FOLLOWING SUBLETHAL INFECTION

Before wounding or infection, cockroaches of all treatments preferred an average P:C ratio

of approximately 0.28 (Fig. 1A). The unmanipulated animals preferred this ratio over the

course of the experiment. By contrast, highly infected individuals selected a P:C ratio of

approximately 1.3 whereas low infected and wounded cockroaches preferred an

intermediate ratio of approximately 1.0 P:C on the first day post-infection (p.i.) (Wilcox rank

sum test: high vs. low: W= 595.0, p = 0.098; low vs. wounded: W = 412.5, p > 0.1; wounded

vs. unmanipulated: W = 713.5, p < 0.001). All manipulated groups returned to baseline P:C

ratios by day 4 p.i..

Final minimal GLMMs consisted of the fixed terms treatment and day without an interaction

since the model with a treatment*day interaction did not significantly improve the model

(ANOVA for model comparison, p > 0.1). Cockroaches that were wounded differed

significantly from unmanipulated cockroaches in their preferred P proportion on the first day

following treatment (P proportion chosen: 0.14 vs 0.06, wounded vs. unmanipulated,

respectively: z = -3.348 p = 0.005), as did cockroaches infected with a high (P proportion

chosen: 0.17 vs 0.06, high vs. unmanipulated respectively: z = -7.416, p < 0.001) or low

bacterial dose (P proportion chosen: 0.14 vs 0.06, low vs. unmanipulated respectively: z = -

2.809, p = 0.029). Cockroaches infected with a high bacterial dose also selected a higher

proportion of P compared to individuals exposed to both a low bacterial dose (P proportion

chosen: 0.17 vs 0.14, high vs. low respectively: z = -3.718, p = 0.001) or to wounding (P

proportion chosen: 0.17 vs 0.14, high vs. wounded respectively: z = -2.808, p = 0.029).

However, individuals that were wounded or were infected with a low bacterial dose did not

select a significantly different proportion of P (P proportion chosen: 0.14 vs 0.14, low vs.

wounded respectively: z = 0.625, p = 1.000). Concerning the proportion of C, the pattern is

slightly different. Only highly infected cockroaches on the first day p. i. differed significantly

from unmanipulated (C proportion chosen: 0.16 vs 0.26, high vs. unmanipulated

respectively: z = 5.270, p < 0.001). All the other treatments were not significantly different

from each other (Supplementary Tab. 2).

P, C as well as total food consumption varied significantly over the course of the experiment

(Fig. 1B-D). Individual cockroaches ate on average 0.6 mg P and 2.2 mg C under

unmanipulated conditions. Conversely, in all manipulated groups the amount of P eaten

declined immediately after infection (Wilcox rank sum test: wounded vs. unmanipulated: W

= 77.0, p < 0.001; low vs. unmanipulated: W = 66.5, p < 0.001; high vs. unmanipulated: W =

76.0, p < 0.001). The same accounts for the consumption of C (Wilcox rank sum test:

wounded vs. unmanipulated: W = 11.0, p < 0.001; low vs. unmanipulated: W = 7.0, p < 0.001;

high vs. unmanipulated: W = 6.0, p < 0.001). As a result, the total amount eaten was reduced

in the same way immediately following treatment (Wilcox rank sum test: wounded vs.

unmanipulated: W = 0.0, p < 0.001; low vs. unmanipulated: W = 0.0, p < 0.001; high vs.

unmanipulated: W = 0.0, p < 0.001). By the 2nd day, consumption across all groups began to

recover, reaching pre-treatment levels by the 4th day p.i..

SURVIVAL ON ENFORCED DIET WITH AND WITHOUT LETHAL INFECTION

The median (50 %) survival time for *B. orientalis* males placed on a P-rich diet was 82 days,

whereas the mortality of males placed on a C-rich diet did not exceed 30 % over the course

of the experiment (150 days) (Fig. 2A). By the end of the experiment males restricted to P-

rich diet showed a significantly higher mortality (86.44 %) compared to those on C-rich diet

(27.59 %; Cox proportional hazard regression P vs. C: z = 5.974, *p* < 0.001). When injected

with a lethal dose of *P. entomophila*, cockroaches started to die on day 1 p.i. (Fig. 2B), and

diet did not have a significant effect on survival following infection (Cox proportional hazard

regression P vs. C: p > 0.10). In both cases around 70 % of the individuals had died by day 6

p.i..

BACTERIA GROWTH INHIBITION ASSAY

The inhibitory effect of the hemolymph (N = 6 per dilution per treatment) of immune-

challenged *B. orientalis* males on bacterial growth was not diet-dependent (ANOVA: C 1:4 vs.

P 1:4: F_1 = 0.13, *p* > 0.10; C 1:8 vs. P 1:8: F_1 = 0.02, *p* > 0.10; Fig. 3) resulting in similar

suppression of bacterial growth regardless of whether cockroaches ate protein- or

carbohydrate-rich diet, with only dilution level influencing bacterial growth: hemolymph

diluted 1:4 and 1:8 suppressed *P. entomophila* growth for the first 600 min and 420 min,

respectively, after which bacteria started to slowly replicate. By the end of quantification,

bacteria exposed to the 1:8 hemolymph dilutions (from both P and C-enforced diets) had

reached a significantly higher OD than in controls (ANOVA: C 1:8 vs. Control: F_1 = 6.51, p =

0.011; P 1:8 vs. Control: F_1 = 4.59 p = 0.032) whereas both 1:4 dilutions did not.

PROTEOMIC ANALYSIS BY MASS SPECTOMETRY

A total number of 3514 peptide hits were identified and assembled into 750 proteins by

MaxQuant. After filtering, 387 different proteins were identified and quantified in the

hemolymph of infected *B. orientalis* males fed on a P-rich vs. a C-rich diet (N = 6 per

treatment) (Supplementary Tab. 3), of which 65 are putative immune-related genes. Overall,

Apolipophorin was the most abundant protein making up approximately 70 % of the whole

hemolymph protein content. Other highly abundant proteins were Transferrin, Gelsolin,

Heterochromatin-associated protein MENT and an Insulin-like growth factor-binding protein

complex. We identified 17 proteins that showed significant changes in abundance following

diet treatment (Fig. 4 and Supplementary Tab. 3). Infected individuals on a C-rich diet were

significantly enriched for Hexokinase type II, which is involved in carbohydrate metabolism

(glycolysis) (Yanagawa 1978), in addition to Carbonyl reductase I-like, which is involved in

NADPH-dependent reduction of active substrates including endogenous and xenobiotic

carbonyl compounds (Hoffmann & Maser 2007). Additionally, Tropomyosin which is a

calcium-dependent regulator of muscle contraction (Pomés *et al.* 2007), and Acyl-CoA-

binding protein, which carries out lipid-binding transport and suppresses glucose-induced

insulin secretion (Færgeman *et al.* 2007) were more abundant. Furthermore, a L-galactose

dehydrogenase-like protein was enriched but its function is not known in insects. Conversely

infected individuals on a P-rich diet were significantly enriched for Alpha-amylase, which is

involved in carbohydrate metabolism (Terra & Ferreira 1994) and Proteasome subunit alpha

type-3, which is involved in protein degradation (Rivett 1993). Additionally, Hemolymph

lipopolysaccharide-binding protein-like (2 isoforms), which binds carbohydrates (foreign

particles) (Jomori & Natori 1991) and Extracellular superoxide dismutase, which carries out

superoxide metabolic processing (Felton & Summers 1995) were detected. Glutamine

synthetase is involved in glutamate and glutamine catabolism and biosynthesis (Smartt *et al.*

1998) while Adenylate kinase isoenzyme 1 and Hexamerin are associated with ATP

metabolism (Fujisawa *et al.* 2009) and amino-acid and energy storage, respectively

(Burmester 1999). There was also an enrichment of Ankyrin-1, although its function in

insects remains unclear.

Discussion

Under normal conditions, extensive protein consumption shortens the lifespan of many

insects including ants, honeybees and flies (Lee *et al.* 2008; Dussutour & Simpson 2009;

Fanson *et al.* 2009; Grandison *et al.* 2009; Cook *et al.* 2010; Pirk *et al.* 2010), a finding that is

corroborated in our and another study of cockroaches (Hamilton *et al.* 1990). Here, we find

that male *B. orientalis* cockroaches showed 45 % higher mortality (Fig. 2A) when restricted

to a P- vs a C-rich diet. One explanation for this consistent observation across study

organisms is that elevated levels of proteins increase TOR signaling. TOR serves as a nutrient

sensor linked to macronutrient intake and metabolism, causing a broad anabolic response

that is life-shortening over the long term (reviewed in (Simpson & Raubenheimer 2009).

Other explanations could relate to the toxic effects of breaking down nitrogenous products,

and the enhanced production of mitochondrial radical oxygen species, DNA and protein

oxidative modifications, membrane fatty acid composition and mitochondrial metabolism

(reviewed in (Simpson & Raubenheimer 2009). The higher abundance of extracellular

superoxide dismutase in cockroach males fed on a P-rich diet (Fig. 4; Supplementary Tab. 3)

supports this explanation.

Unsurprisingly, male cockroaches consumed low amounts of protein under normal

conditions. But this preference shifted dramatically following infection. As with caterpillars

(Povey *et al.* 2013), highly infected male cockroaches switched to a P-rich diet. Furthermore,

cockroaches adapted their feeding behaviour to the severity of the infection. Lowly infected

and wounded (Ringer-injected) individuals chose an intermediate (balanced) P:C ratio,

whereas uninfected and highly infected individuals chose a significantly enriched diet for C

and P, respectively. This demonstrates that cockroaches quantitatively regulate their

behavioural response to infection. Additionally, our data suggest host-driven adaptation as

opposed to pathogen manipulation because wounded individuals also reduced their

carbohydrate intake. Wounding elicits a localized immune response in insects (Haine *et al.*

2007), suggesting a form of prophylactic behavior since it is likely that microbes can enter

the hemolymph via damaged cuticle (Siva-Jothy *et al.* 2005).

In contrast to caterpillars and other organisms which can modulate their immune response

with diet, changes in cockroach dietary preference did not influence any of the immune

parameters measured. In caterpillars, a shift from a carbohydrate- to a protein-biased diet

following *Bacillus subtilis* (gram-positive) or baculovirus infection led to an increase of

antibacterial and phenoloxidase activity and hemocyte density, and resulted in higher

survival (Povey *et al.* 2009; Povey *et al.* 2013). By contrast, a switch to a protein-biased diet

did not influence cockroach hemolymph antimicrobial activity (Fig. 3) or survival (Fig. 2B) nor

have a major impact on the production of immune related proteins (Fig. 4; Supplementary

Tab. 3). Only two hemolymph lipopolysaccharide-binding protein isoforms, which may play a

role in pathogen recognition by binding foreign particles (Jomori & Natori 1991) were more

abundant in the hemolymph of P-rich fed infected cockroaches, but overall our findings

indicate that the behavioural changes adopted by cockroaches are limited in their ability to

alter infection outcome.

The overrepresentation of proteins participating in C- and P metabolism in C- vs P-rich diets,

respectively (Fig. 4), demonstrate that the diets altered cockroach physiology in the

expected direction. For example, the higher abundance of alpha-amylase in the hemolymph

of *B. orientalis* males feeding on P-rich diet shows these individuals were metabolizing lower

quantities of C. Alpha-amylase is thought to be involved in the breakdown of glycogen,

which is the major glucose storage compound in animals. It is employed if not enough

carbohydrates are present in the diet (Mohamed 2004).

Overall, our results suggest that cockroaches may not be able to self-medicate using

macronutrients, but that they do engage in a typical anorexia response, as has been shown

in macronutrient self-medication in caterpillars (Adamo *et al.* 2007; Povey *et al.* 2009; Povey

et al. 2013). Illness-induced anorexia offsets physiological trade-offs between launching

immune responses and food digestion. A previous study demonstrated that crickets reduce

their food intake, especially for lipids, following infection with the bacterium *Serratia*

marcescens (Adamo *et al.* 2010). High hemolymph lipid levels are associated with decreased

concentrations of monomeric apolipophorin III, a lipid transporter, and higher susceptibility

to *S. marcescens* infection (Adamo *et al.* 2008). In other insects, anorexia can have a direct

impact on immunity. For example, in *Drosophila*, starvation can modify antimicrobial peptide

production and lead to reduced melanisation (Ayres & Schneider 2009).

The apparent lack of a link between macronutrient dietary selection and male cockroach

immunity is unexpected. One possible explanation is that future food availability and quality

may be far less predictable in omnivorous pest organisms like cockroaches (Raubenheimer &

Jones 2006). A recent genomic study reports major expansions of cockroach gene families

linked to chemoreception, detoxification and innate immunity (Li *et al.* 2018), indicating that

adaptations in these pathways may have been essential for enabling cockroaches to thrive in

unpredictable, antigen-rich environments. Indeed, while cockroach survival was reduced on

an enforced P-rich diet, a negative effect could only be observed well over 40 days after

exposure. In such organisms, there could be a clear advantage in reducing regulatory

interactions between host diet and immunity. An additional point to consider is that in

contrast to several previous studies, we performed experiments on adult individuals and not

larvae, which have different resource allocation strategies and consumption rates in general

(Boggs 2009). In holometabolous insects, most of the resources in larvae are allocated to

growth, maintenance and storage whereas in adults, they are allocated to reproduction and

maintenance. Consequently, there has been great emphasis on a trade-off between growth

and immunity in the larval stage of herbivorous insects (reviewed in (Singer *et al.* 2014)).

Here, the need for fast growth competes with the requirement to provide protection from

parasite and pathogen-induced mortality. Given that proteins and amino acids are a crucial

limiting factor in herbivorous diets, we argue that a trade-off between two essential life-

history parameters that depend strongly on P – growth and survival – could be particularly

pronounced in herbivorous insects (Schoonhoven *et al.* 2005; Simpson & Raubenheimer

2012).

In conclusion, we find that *B. orientalis* males modulate their macronutrient feeding

behaviour following infection by dramatically reducing food intake and favouring a diet

containing a higher P-ratio. We also show that a P-rich diet eventually leads to significantly

reduced host lifespan, and that male cockroaches avoid such a diet under normal conditions.

To our surprise, the observed behavioural response to immune challenge did not positively

influence the antimicrobial activity or proteomic profile of host immunity. Overall, our

findings support the concept of a generalized host-directed response to microbial challenge

in cockroaches, based on anorexia and the limitation of carbohydrate intake. Such a

response is likely to be beneficial to the host, but primarily as a means of avoiding

contaminated food and reducing pathogen access to resources, rather than facilitating cross-

talk with the immune system. Our findings indicate that dietary composition and immunity

may be decoupled in cockroaches resulting from adaptations to detoxification and innate

immunity linked to survival in antigen-rich and nutritionally diverse environments. Our study

highlights the importance of understanding variation in natural diet, development, and

ecology when exploring the link between nutrition and animal immunity.

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Authors' contribution

625 T.S. and D.P.M. conceived and coordinated the study; T.S. designed and conducted the
626 experiments; T.S. and D.P.M. wrote the manuscript; S. H. and P. R. J. conducted the
627 proteomic database preparation by de novo transcriptome sequencing; A.R.-R. guided the
628 sample preparation for the proteomic analysis; B. K. and C. W. conducted LC-MS/MS and
629 corresponding data analysis. All authors contributed to drafting and revising the manuscript.

Data accessibility

632 All data associated with the study will be made available in Dryad.

Fig. 1: Effect of bacterial infection with *P. entomophila* (high load, low load), Ringer's solution or no manipulation (control) of *B. orientalis* males on: **A)** P:C ratio chosen, **B)** total consumption, **C)** P consumption, **D)** C consumption. Note different scales used for total P- and C- consumption. Area surrounding mean curves indicate 95% confidence intervals.

Fig. 2: Kaplan-Meier survival curves of: **A)** Unmanipulated *B. orientalis* males restricted to P-
rich (35 % protein and 7 % carbohydrate) or C-rich (7 % protein and 35 % carbohydrate)

diets. Survival data for three independent cohorts (1,2,3) for P- and C-rich diets are given in
blue and red respectively, with mean population survival on each diet indicated by a thick
bold line. Note the long period at the beginning of the experiment where no clear survival
differences between diets are observable. **B)** *B. orientalis* males restricted to P-rich (35 %
protein and 7 % carbohydrate) or C-rich (7 % protein and 35 % carbohydrate) diet and
infected with an LD_{50} of *P. entomophila* (infected), Ringer's solution (wounded) or
unmanipulated (control).

Fig. 3 Impact of diet on *in vitro* growth inhibition of *P. entomophila* in *B. orientalis*
hemolymph (1:4 and 1:8 dilution). A bacterial solution without hemolymph served as the
control.

Fig. 4 Effect of diet on abundance of male *B. orientalis* hemolymph proteins following
immune challenge (high dose). Points in blue and red reflect proteins that are significantly
(>2) more abundant in P- and C-rich diets respectively.

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