Coupling between tolerance and resistance for two related *Eimeria*

parasite species

Short title: Resistance-tolerance coupling for two *Eimeria*

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Abstract

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Resistance (host capacity to reduce parasite burden) and tolerance (host capacity to 2 3 reduce impact on its health for a given parasite burden) manifest two different lines of defence. Tolerance can be independent from resistance, traded-off against it, or the 4 two can be positively correlated because of redundancy in underlying (immune) 5 processes. We here tested whether this coupling between tolerance and resistance 6 could differ upon infection with closely related parasite species. We tested this in 7 8 experimental infections with two parasite species of the genus *Eimeria*. We measured 9 proxies for resistance (the (inverse of) number of parasite transmission stages (oocysts) per gram of feces at the day of maximal shedding) and tolerance (the slope of 10 11 maximum relative weight loss compared to day of infection on number of oocysts per gram of feces at the day of maximal shedding for each host strain) in four inbred mouse 12 strains and four groups of F1 hybrids belonging to two mouse subspecies, 13 Mus musculus domesticus and M. m. musculus. We found a negative correlation 14 15 between resistance and tolerance against E. falciformis, while the two are uncoupled 16 against E. ferrisi. We conclude that resistance and tolerance against the first parasite 17 species might be traded off, but evolve more independently in different mouse genotypes against the latter. We argue that evolution of the host immune defences can 18 19 be studied largely irrespective of parasite isolates if resistance-tolerance coupling is 20 absent or weak (E. ferrisi) but host-parasite coevolution is more likely observable and best studied in a system with negatively correlated tolerance and resistance 21 22 (E. falciformis).

Keywords: Resistance, Tolerance, *Eimeria*, Coevolution

Introduction

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Host defence mechanisms evolve to alleviate the detrimental effect of parasites. They can be categorised into two components: resistance and tolerance (Råberg et al. 2009). Resistance is the ability of a host to reduce parasite burden, resulting from defence against parasite infection or proliferation early after infection (Schmid-Hempel 2013). The negative effect of resistance on parasite fitness can lead to antagonistic coevolution. According to theoretical models, fluctuating host and parasite genotypes arise, and balancing selection maintains resistance alleles polymorphic (Boots et al. 2008; Roy & Kirchner 2000). Resistance has been the classical "catch all" measure for host-parasite systems, but recently it has been shown to be incomplete, especially with respect to potential fitness effects on the host (Kutzer & Armitage 2016; Råberg et al. 2009). Disease tolerance be confused from "immunological (not to unresponsiveness to self antigens; Medzhitov et al. 2012) is the ability of the host to limit the impact of parasite on its fitness (Råberg et al. 2009; Vale & Little 2012; Kutzer & Armitage 2016). By potentially providing a longer-living niche, this defence mechanism improves, or at least does not deteriorate, the fitness of the parasite. Tolerance alleles are thus predicted by theoretical models to evolve to fixation due to positive feedback loops (Boots et al. 2008; Restif & Koella 2004; Roy & Kirchner 2000). From a mechanistic perspective tolerance alleviates direct or indirect damage (e.g. immune response underlying resistance against parasites, excessive immunopathology; Graham et al. 2005) caused by parasites (Råberg et al. 2009). Tolerance mechanisms include modulation of inflammatory response (Ayres & Schneider 2012), tissue repair (stress response, damage repair and cellular

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regeneration mechanisms; Soares et al. 2017), and compensation of parasite-induced damage by increase of reproductive effort (Baucom & Roode 2011). Even in the absence of parasite infection, the maintenance of tolerance mechanisms can be detrimental to other functions, ultimately affecting host fitness (Stowe et al. 2000; Råberg et al. 2009). The resulting costs of resistance and tolerance determine the optimal (steady state and infection inducible) extent of both immune defences (Sheldon & Verhulst 1996). Resistance and tolerance can be positively associated if they involve the same metabolic pathway, as was shown in the plant model *Arabidopsis thaliana* in response against herbivory (Mesa et al. 2017). In animals, genetic association studies of resistance and tolerance of *Drosophila melanogaster* against the bacterium Providencia rettgeri have shown positively correlated genetic effects, as the same loci were associated with changes of both traits in the same direction (Howick & Lazzaro 2017). Nevertheless, resistance and tolerance can also be genetically and physiologically independent, involving different proximate mechanisms. Lack of correlation between both defences was shown for example in monarch butterflies (Danaus plexippus) infected by the protozoan parasite Ophryocystis elektroscirrha. This study found genetic variation in resistance between butterflies families, but a fixed tolerance (Lefèvre et al. 2010). Similarly, no correlation could be found between resistance and tolerance for the fish Leuciscus burdigalensis in response to infection with its parasite Tracheliastes polycolpus. The authors explain the decoupling of both defences by the fact that, in this system, tolerance likely involves wound repair rather than immune

71 regulation, making resistance and tolerance mechanisms independent (Mazé-Guilmo et 72 al. 2014). 73 In other systems, resistance and tolerance have been found negatively correlated. For 74 example, inbred laboratory mouse strains lose weight upon infection with *Plasmodium* 75 chabaudi. The extent of this impact on host health is negatively correlated with the peak number of parasites found in the blood (Råberg et al 2007), meaning that mouse 76 strains with higher resistance present lower tolerance. Similarly, infections of sea trout 77 78 (Salmo trutta trutta) and Atlantic salmon (Salmo salar) with the trematode Diplostomum 79 pseudospathaceum showed that resistance and tolerance were negatively correlated when assessing mean levels of both traits in different host populations (Klemme & 80 81 Karvonen 2016). This is interpreted as a result of trade-off between resistance and tolerance (Sheldon & Verhulst 1996; Restif & Koella 2004; Råberg et al. 2009). 82 We have seen that depending on the system studied, resistance and tolerance can be 83 84 (1) uncoupled (independent), (2) positively correlated (involving same genes and 85 mechanisms), or (3) negatively correlated (traded-off). Theoretical models show that coupling between resistance and tolerance (or absence thereof) could depend not only 86 on the host but also on the parasite (Carval & Ferriere 2010). Here we tested this 87 hypothesis. More precisely, we asked whether there could be differences in the 88 89 resistance-tolerance coupling upon infection of one host type with two closely related parasite species. To answer this question, we infected four inbred mouse strains and 90 four groups of F1 hybrids representative of two house mouse subspecies, 91 M. m. domesticus and M. m. musculus, with two parasite isolates representative of two 92 93 naturally occurring parasite species, the protozoan parasites Eimeria ferrisi and E. falciformis (Jarquín-Díaz et al. 2019). Eimeria spp. are monoxenous parasites that 94

expand asexually and reproduce sexually in intestinal epithelial cells, leading to malabsorption of nutrients, tissue damage and weight loss (Chapman et al. 2013). The evolutionary history of these different *Eimeria* species in the two house mouse subspecies is unknown and it is unclear whether subspecies-specific adaptation exists in one or the other. We tested if coupling between resistance and tolerance differs between both parasite species and discussed the implication for parasite-host coevolution.

Material and methods

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1. Parasite isolates

The three parasite isolates used in this study were isolated from feces of three different 104 M. m. domesticus/M. m. musculus hybrid mice captured in Brandenburg, Germany, in 105 106 2016 (capture permit No. 2347/35/2014). The parasite isolates belong to both the most 107 prevalent Eimeria species in this area, namely E. ferrisi (isolate Brandenburg64) and 108 E. falciformis (isolate Brandenburg88)(Jarquín-Díaz et al. 2019). Isolate Brandenburg64 was isolated in a 92% M. m. domesticus individual (hybrid index (HI) = 109 110 0.08: Proportion of M. m. musculus alleles in a set of 14 diagnostic markers, see 111 Balard et al. (2020)) and isolate Brandenburg88 in a 80% M. m. domesticus (HI=0.2). 112 Pre-patency and the peak day of parasite shedding for these isolates were estimated 113 during infection in NMRI laboratory mice (Al-khlifeh et al. 2019) which were also used 114 for serial passaging of the isolates. Previous to the experiment, the isolates had been 115 passaged respectively 3 and 4 times in NMRI laboratory mice. Parasite infective forms 116 (oocysts) were recovered by flotation in saturated NaCl solution followed by washing 117 and observation under light microscope (following the protocol described in Clerc et al. 118 (2019)) and stored at room temperature in 1mL of 2% potassium dichromate for a

maximum of 2 months before infection of the wild-derived mice. Oocysts were allowed to sporulate 10 days before infection in a water bath at 30°C.

2. Mouse groups

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122 We used four wild-derived inbred mouse strains from which we generated four groups 123 of F1 hybrids. Hybrids between M. m. domesticus and M. m. musculus are used in the present study solely to increase statistical power for comparisons among strains (such 124 125 as resistance-tolerance correlations). In the future, analyses of a hybrid effect (Balard 126 et al. 2020) could investigate tolerance and resistance employing a larger panel of such 127 hybrid strains allowing statistical analysis of an outbreeding effect. Two parental strains 128 represented M. m. domesticus: SCHUNT (Locality: Schweben, Hessen, Germany [N: 129 5°0 26', E: 9° 36'] (Martincová et al. 2019)) and STRA (Locality: Straas, Bavaria, 130 Germany [N: 50° 11', E: 11° 46'] (Piálek et al. 2008), and two derived from M. m. musculus: BUSNA (Locality: Buškovice, Bohemia, Czech Republic [N: 5°0 14', 131 E: 1°3 22'] (Piálek et al. 2008)) and PWD (Locality: Kunratice, Bohemia, Czech 132 133 Republic [N: 5°0 01', E: 14 2°9'] (Gregorová and Forejt 2000)). These four strains were fully inbred, i.e. passing more than 20 generations of brother-sister mating. The four 134 135 groups of F1 hybrids consisted of two intrasubspecific hybrids (SCHUNTxSTRA and PWDxBUSNA) and two intersubspecific hybrids (STRAxBUSNA and SCHUNTxPWD) 136 (Figure 1). Age of the mice at the time of infection ranged between 5.6 and 21.4 137 weeks, with the mean for each eight mouse group ranging between 10.5 and 14.7 138 weeks. All mouse strains and F1 hybrids were obtained from the Institute of Vertebrate 139 140 Biology of the Czech Academy of Sciences in Studenec (license number 61974/2017-141 MZE-17214; for further details on strains see https://housemice.cz/en).

Parasites of the *Eimeria* genus are known to induce host immune protection against reinfection (Rose, Hesketh, and Wakelin 1992; Smith and Hayday 2000). To ensure that our mice were *Eimeria*-naive, mouse fecal samples were tested before infection for the presence of *Eimeria* spp. oocysts by flotation in saturated NaCl solution followed by washing and observation under light microscope.

3. Experimental infection

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Mice were kept in individual cages during infection. Water and food (SNIFF, Rat/Mouse maintenance feed 10 mm) were provided ad libitum supplemented with 1 g of sunflower and barley seeds per day. Mice were orally infected with 150 sporulated oocysts of one Eimeria isolate suspended in 100 μl phosphate-buffer saline (PBS) and monitored daily until their sacrifice by cervical dislocation at time of regression of infection (reduction of oocyst output). Individuals presenting severe health deficiency and/or a weight loss approaching 18% relative to their starting weight were sacrificed earlier at defined humane end points (experiment license Reg. 0431/17). Weight was recorded and feces collected on a daily basis. Fecal pellets were collected every day from each individual cage and suspended in 2% potassium dichromate. Parasite oocysts were recovered using NaCl flotation (see above). All individuals were negative for Eimeria at the beginning of our experiment (before infection of first batch, as described in the next paragraph). In total, 143 mice were infected. Mice were randomly allocated to experimental groups ensuring homogeneous distribution of ages and sexes between groups. Our experiments were conducted in four (partially overlapping) consecutive batches for logistical reasons. The first two batches were infected with E. ferrisi isolates (Brandenburg64), the third and fourth by one E. ferrisi isolate (Brandenburg64) and one E. falciformis isolate (Brandenburg88). Our experimental design is summarized in **Table 1** (chronology of experimental batches can be scrutinized in **Appendix 1**).

Nematode infection is common in breeding facilities (Baker, 1998) and could interact with Eimeria (Clerc et al. 2019). We surveyed for their presence and nematode eggs (Syphacia sp. and Aspiculuris sp.) were observed in flotated feces of mice belonging to all genotypes before the experiment. Despite treatment of the first infection batch of mice (B1, 12 mice) with anthelminthics (Profender®, Bayer AG, Levekusen, Germany) following the protocol of Mehlhorn et al. (2005), nematodes were still detected with PCR (following the protocol of (Floyd et al. 2005)) in randomly sampled fecal samples a week later. We therefore decided not to treat mice of the following infection batches. Moreover, we observed Eimeria oocysts in the feces of 28 mice belonging to the last experimental batch (batch B4) at the day of infection, likely due to cross-contamination between batches. For following statistical analyses, we considered along with the full data set (N=143) a conservative data set in which cross-contaminated animals and animals treated by anthelminthic were removed (N=103). Results obtained on the conservative data set can be found in Appendix 2 and 3. Despite differences in significance due to a lower statistical power, the main conclusions of our analyses were consistent with those obtained on the main data set.

4. Statistical analyses

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4.1. Choice of proxies for resistance, impact of parasite on host and tolerance

As resistance is the capacity of a host to reduce its parasite burden, it is usually estimated by the inverse of infection intensity (Råberg et al. 2009). Pre-patency (the time to shedding of infectious stages, so called oocysts) is longer for *E. falciformis* (7

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days) than for E. ferrisi (5 days) (Al-khlifeh et al. 2019). Therefore, as a proxy of (inverse of) resistance we used the number of oocysts per gram of feces (OPG) at the day of maximal shedding. Using the Spearman's non-parametric rank correlation test, we found this measurement to be tightly correlated with the sum of oocysts shed throughout the experiment (Spearman's ρ =0.93, N=168, P<0.001). Due to the aggregation characteristic of parasites (Shaw and Dobson 1995), the appropriate distribution for maximum number of OPG was found to be the negative binomial distribution. This was confirmed based on log likelihood, AIC criteria and goodness-offits plots (density, CDF, Q-Q, P-P plots; R packages MASS (Venables & Ripley 2002) and fitdistrplus (Delignette-Muller & Dutang 2015)). We confirmed the fit of our models by assessing the uniformity of the distribution of model residuals. Both parasite species provoke inflammation, cellular infiltration, enteric lesions, diarrhea, and ultimately weight loss (Ankrom, Chobotar, and Ernst 1975; Ehret et al. 2017; Schito, Barta, and Chobotar 1996; Al-khlifeh et al. 2019). Therefore, the impact of parasites on host health was measured as the maximum relative weight loss compared to day 0 (body weight measured at the start of the experimental infection). For mice sacrificed at humane end points before the end of the experiment, the last weight of the living animal was used. This weight (loss) can be expected to be a very conservative estimate for our analyses (rendering tolerance conservatively low for these animals, which might have lost more weight if not sacrificed). Tolerance is usually defined as a reaction norm, i.e. the regression slope of host fitness (or health condition if that is the parameter of interest) on infection intensity per host genotype (Simms 2000; Råberg et al. 2009). Thus tolerance was assessed as the slope of maximum relative weight loss compared to day 0 on number of OPG at the 213 day of maximal shedding, within each mouse group and for each parasite isolate. A
214 steep slope indicates a low tolerance (high weight lost for a given parasite burden).

4.2. Statistical comparison of resistance, impact on health and tolerance in

E. ferrisi and E. falciformis

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(parasite isolate-mouse strain).

The comparison between *E. ferrisi* and *E. falciformis* was performed using respectively the isolates Brandenburg64 and Brandenburg88 with which we infected all our eight mouse groups (see Table 1). Maximum OPG and relative weight loss were modelled separately as a response of mouse group, parasite isolate and their interaction. We used a negative binomial generalised linear model for maximum OPG, and a linear model for relative weight loss. Tolerance was assessed by modelling relative weight loss as a response of maximum OPG interacting with mouse group, parasite isolate and the interaction of the two latter. As each mouse was controlled against itself at the start of the experiment, before losing weight or shedding parasites, we performed a linear regression with null intercept. To test the significance of the marginal contribution of each parameter to the full model, each parameter was removed from the full model, and the difference between full and reduced model was assessed using likelihood ratio tests (G). For each of our models that showed a significant interaction term, we also asked within each parasite isolate if the response differed between mouse groups using likelihood ratio tests (G) as described above. In the case of a non-significant interaction term, we performed post-hoc tests corrected for multiple testing (Tukey Honest Significant Differences (HSD)) to compare within all pairwise comparisons between groups Of note, four mice infected with *E. falciformis* isolate Brandenburg88 did not shed any oocysts as death occurred at or one day before the peak of oocysts shedding in other mice. For this reason, we modelled maximum OPG when mice infected with this parasite were included using a zero-inflated negative binomial (ZINB) generalised linear model, after verifying that it provided a better fit than the simple negative binomial based on log likelihood and AIC criteria.

4.3. Test of coupling between resistance and tolerance

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We tested coupling between resistance and tolerance for E. ferrisi and E. falciformis using the isolates Brandenburg64 and Brandenburg88 and our eight mouse groups. To test such coupling, one can assess the strength of correlation between measure of resistance and measure of tolerance (Råberg et al 2007). Of note, tolerance (in absolute value) is measured as the slope α of the linear regression of parasite load (x) on maximum relative weight loss (y) of equation $y = \alpha x + \beta$ (α being the slope and β the intercept, 0 in our case). Therefore, tolerance is expressed as $\alpha = y/x - \beta/x$. As x and y/x are by definition not independent, testing the correlation between resistance and tolerance can lead to spurious correlation (Brett 2004). To alleviate the dangers of this statistical artifact, we additionally tested differences in resistance, impact on health and tolerance between mouse groups separately (as described before, see 4.2) and also the underlying correlation between mean parasite load (x) and mean relative weight loss (y). We use the terminology "coupling" (between resistance and tolerance) to describe genotype-level correlation between tolerance and resistance additionally supported by the absence of positive correlation between health-effect and resistance. Correlations were tested using Spearman's rank correlation.

After testing the resistance-tolerance coupling separately in both parasites, we tested the statistical difference in the relationship between (1) health-effect and resistance and (2) tolerance and resistance in the two *Eimeria* species infections. To achieve this aim, we used the mean values predicted by our three models (see 4.2) for each eight mouse groups to perform first a linear regression of the mean predicted relative weight loss as a response of the mean predicted OPG, parasite isolate and their interaction, and second a linear regression of the mean predicted tolerance value as a response of the mean predicted OPG, parasite isolate and their interaction. The significance of the marginal contribution of each parameter to the full model was assessed by removing each parameter from the full model, and the difference between full and reduced model was assessed using likelihood ratio tests (G). All analyses were performed using R version 3.5.2 (R Development Core Team 2013) (negative binomial: function glm.nb from R package MASS (Venables and Ripley 2002); ZIBN: function zeroinfl from R package pscl (Jackman 2020; Zeileis, Kleiber, and Jackman 2008); linear model: function Im from R core package stats; mean and 95% confidence intervals: function gapredict from R package ggeffect (Lüdecke 2018)). Graphics were produced using the R package ggplot2 (Wickham 2016) and compiled

Results

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1. General

Parasites of all isolates successfully infected all mouse groups (at the exception of 5 individuals infected with the *E. falciformis* isolate Brandenburg88 that died or had to be sacrificed due to a strong weight loss before the peak of shedding for this parasite),

using the free software inkscape (https://inkscape.org).

meaning that no "qualitative infection resistance" (*sensu* (Gandon and Michalakis 2000)) was detected. For *E. ferrisi* isolate Brandenburg64, the pre-patent period was 5 days post infection (dpi) and the median day of maximal oocyst shedding was 6 dpi (standard deviation sd=0.9). The median day of maximum weight loss was 5 dpi for both isolates (sd=1.7). For *E. falciformis* isolate Brandenburg88 pre-patency was 7 dpi, median day of maximal shedding was 8 dpi (sd=1.3) and median day of maximal weight loss 9 dpi (sd=1.6)(**Figure 2**). Of note a considerable number of mice infected with this isolate (13 out of 56 = 23%) died or had to be sacrificed at humane end points less than 3 days after the oocysts shedding peak for the group, all belonging to *M. m. musculus* subspecies (PWD, BUSNA, or their F1 PWDxBUSNA; 5 died at dpi 8, 5 at dpi 9, 3 at dpi 10). *E. falciformis* isolate Brandenburg88 was more lethal for the *M. m. musculus* mice strains than for the other strains (χ_7^2 = 31.96, P<0.001; **Table 2**).

- 294 2. Comparison of resistance-tolerance coupling between *E. ferrisi* and
- 295 E. falciformis

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- 296 2.1. Differences in resistance and tolerance between mouse groups depends on
- 297 the parasite
- Considering all mice infected with either E. ferrisi isolate Brandenburg 64 and 298 E. falciformis isolate Brandenburg 88, we found our proxy for (inverse of) resistance 299 300 (maximum number of OPG) to be statistically different between mouse groups, parasite 301 isolates and their interaction (LRT: mouse groups: G=55.5, df=28, P<0.01; parasite 302 isolates: G=40.5, df=16, P<0.001; interaction: G=27.9, df=14, P=0.015). Results were similar for our proxy for tolerance (LRT: mouse groups: G=28.4, df=14. P=0.01; 303 304 parasite isolates: G=20.1 df=8, P=0.01; interaction: G=18.8, df=7, P<0.01). Our proxy 305 for impact on weight (maximum relative weight loss) was significantly different between

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mouse groups and parasite isolates, but not for their interaction (LRT: mouse groups: G=44.9, df=14, P<0.001; parasite isolates: G=33, df=8, P<0.001; interaction: G=7.5, df=7, P=0.38). For the latter model, impact on weight, post-hoc tests showed that the only statistical differences between two mouse groups within a parasite infection were found in E. falciformis infection, between PWD and STRA (Tukey HSD test, p-value = 0.02), PWD and STRAxBUSNA (Tukey HSD test, p-value = 0.03) and PWD and SCHUNTxPWD (Tukey HSD test, p-value = 0.02). No difference was found within one mouse group between the two parasite isolates at the 0.05 significance threshold. We found that the mean predicted number of OPG varies with the mean predicted relative weight loss (LRT: G=10, df=2, P<0.01), differs between both parasites (LRT: G=8.9, df=2, P=0.012), and more importantly we found a significant interaction term (LRT: G=8.3, df=1, P<0.01). This means that the relationship between mean healtheffect and mean resistance differs between the two *Eimeria* species infections. Then, we performed a linear regression of the mean predicted tolerance for each eight mouse groups as a response of the mean predicted OPG, parasite isolate and their interaction. In this case we found that the mean number of OPG varies along with tolerance (LRT: G=8.5, df=2, P=0.01) but does not statistically differ between both parasites (LRT: G=1.1, df=2, P=0.57), and the interaction term was not found significant (LRT: G=0.03, df=1, P=0.86). In this respect, the correlation between resistance and tolerance was not found to significantly differ between both parasites. Following these results, we looked at the coupling of resistance and tolerance within each of the two isolates. 2.2. Resistance and tolerance to E. ferrisi isolate Brandenburg64 are uncoupled We tested coupling between resistance and tolerance for *E. ferrisi* isolate Brandenburg64 in our eight mouse groups. First, we tested whether our proxies for 330 resistance and tolerance were different between the mouse groups. We found the 331 maximum number of OPG to be statistically different between mouse groups (LRT: 332 G=26.6, df=7, P<0.001; Figure 3A). Tolerance was not found to significantly differ 333 between mouse groups for this parasite isolate (LRT: G=6.8, df=7, P=0.45; Figure 3B). 334 We found a non significant positive correlation between resistance (inverse of maximum number of OPG) and impact on health (maximum weight loss) (Spearman's 335 336 ρ =0.69, P=0.07, N=8; **Figure 3C**). Moreover, we did not find a correlation between 337 resistance (inverse of maximum number of OPG) and tolerance (inverse of slope of 338 maximum weight loss on maximum OPG) (Spearman's ρ =0, P=1, N=8; **Figure 3D**). 339 In conclusion, we did not find indications of resistance-tolerance coupling for E. ferrisi 340 isolate Brandenburg64, the different mouse groups infected by this parasite presenting 341 a similar level of tolerance while showing an effect of quantitative resistance on health. 342 2.3. Coupling between resistance and tolerance to *E. falciformis* We then tested coupling between resistance and tolerance for E. falciformis isolate 343 344 Brandenburg88 in our eight mouse groups. First, we tested if our proxies for resistance and tolerance were different between the mouse groups. We found the maximum 345 346 number of OPG to be statistically different between mouse groups (LRT: G=28.6, 347 df=14, P=0.012; Figure 4A). Contrary to our results on E. ferrisi isolate 348 Brandenburg64, the tolerance slopes for E. falciformis isolate Brandenburg88 were 349 different between mouse groups (LRT: G=13.9, df=7, P=0.05; Figure 4B). 350 We detected a strong negative correlation between (inverse of) resistance (maximum 351 number of OPG) and tolerance (inverse of slope of maximum weight loss on maximum OPG) (Spearman's ρ =-0.95, P=0.001; **Figure 4D**). This result was robust to the 352

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exclusion of the extreme point corresponding to mouse strain PWD (point 8 in Figure 4D; Spearman's ρ =-0.93, P<0.01). We conclude that this correlation is unlikely a statistical artifact, as (1) mouse groups present statistically different values of resistance and tolerance (see 2.1) and (2) we found a (non significant) negative correlation between resistance (inverse of maximum number of OPG) and impact on health (maximum weight loss) (Spearman's ρ =-0.5, P=0.22; Figure 4C), indicating that mouse groups losing more weight also shed less parasites. We conclude that our results indicate the presence of negative resistance-tolerance coupling for *E. falciformis* isolate Brandenburg88. Discussion In this study, we assessed resistance and tolerance to two closely related parasites, E. ferrisi and E. falciformis, in four mouse strains and their intra- and intersubspecific hybrids. Understanding this coupling has two major implications: From a practical "measurement" perspective we can ask whether tolerance can be predicted from resistance, as the latter is easier to measure (e.g. in field sampling). Many studies assess the impact of parasites on host fitness based on resistance. If, as we found in the present study, resistance and tolerance are decoupled this can be misleading. In our host system, the house mice, for example, it has been shown that hybrids between M. m. domesticus and M. m. musculus are more resistant to parasites (Baird et al. 2012; Balard et al. 2020), including Eimeria, but tolerance could not be measured under natural conditions (Balard et al. 2020). The effect of parasites on host fitness in the evolution of the house mouse hybrid zone is thus still rather

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tolerance, likely implying a trade-off.

ambiguous (Baird and Goüy de Bellocq 2019). We show that careful distinction between parasite species is necessary when analysing parasite host interaction (see also Jarquín-Díaz et al. 2019) and that it is indispensable to measure both resistance and tolerance in Eimeria infections of house mice. In this work we used the concept of tolerance as used originally in the plant literature and later on transferred to animal studies (Fineblum and Rausher 1995). This concept of tolerance can be criticised, as it links tolerance mathematically to resistance. Nevertheless, we argue that this view is biologically meaningful considering resistance and tolerance as a double defence system, one step limiting the parasite multiplication, the other limiting the impact of this multiplication on fitness-related traits. To limit the possibility of statistical artifact, our approach did not only consist in calculating correlations between resistance and tolerance, but also in testing differences in resistance, impact on health and tolerance. Of note, a positive correlation between mean health-effect and mean resistance of each host strains could indicate some host strains having few parasites-few effects on health, and others more parasites-more effects on health; This configuration would limit the possibility of detecting an actual resistance-tolerance trade-off by lack of a full range of resistance values. For this reason, our approach consisted in testing the "coupling" between resistance and tolerance, that is (1) a genotype-level correlation between tolerance and resistance additionally supported by (2) the absence of positive correlation between health-effect and resistance. We argue that this additional step increases the confidence in the presence of a biologically meaningful negative correlation between resistance and

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Differences between parasite species could explain the evolution of different strategies: E. ferrisi commits to sexual reproduction after a relatively short time with few cycles of asexual expansion (Al-khlifeh et al. 2019; Ankrom, Chobotar, and Ernst 1975), while E. falciformis has a relatively longer life cycle (Al-khlifeh et al. 2019; Haberkorn 1970). As E. ferrisi infections do not reach extremely high intensities, high tolerance might be the optimal strategy for both house mouse subspecies. Resistance could then evolve relatively freely without any major impact of the parasite on the hosts' health. In the case of *E. falciformis*, the long life cycle might lead to high tissue load. Tissue damage is observed during sexual reproduction for this parasite (Ehret et al. 2017) and might mean that a certain level of resistance is required. On the other hand, immunopathology has been observed in advanced E. falciformis infections (Stange et al. 2012). These intrinsic characteristics of *E. falciformis* might lead to multiple different optima for resistance and tolerance, leading to a trade-off. More generally, from an evolutionary perspective, coupling between resistance and tolerance might help determine if coevolution between host and parasite can be expected: a host-parasite system in which one finds negative coupling between tolerance and resistance would be an especially promising system for studies of hostparasite coevolution. Indeed, coevolution in host-parasite systems is often assumed but rarely proven (Woolhouse et al. 2002). Janzen (1980) notes that not all parasite-host systems are coevolving. The presence of efficient host defences against a given parasite is not necessarily produced in response to this parasite specifically and the parasite does not necessarily respond specifically. In the mouse-E. ferrisi system, where resistance and tolerance are decoupled, host and parasite fitness might be decoupled as a result, making host-parasite coevolution less likely. In the mouseE. falciformis system we found a negative coupling between tolerance and resistance,
 making coevolution between host and parasite more likely.
 In conclusion, we show that the coupling between resistance and tolerance can differ
 between closely related parasite species and we argue that this trait of a host-parasite

system determines the questions to be best approached with a particular parasite.

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References

429 Al-khlifeh, E., Balard, A., Jarquín-Díaz, V. H., Weyrich, A., Wibbelt, G. & Heitlinger, E. (2019). 430 Eimeria falciformis BayerHaberkorn1970 and novel wild derived isolates from house 431 mice: Differences in parasite lifecycle, pathogenicity and host immune reactions. bioRxiv, 432 611277. doi:10.1101/611277 433 Ankrom, S. L., Chobotar, B. & Ernst, J. V. (1975). Life cycle of *Eimeria ferrisi* Levine & Ivens, 434 1965 in the mouse, Mus musculus. The Journal of Protozoology, 22, 317–323. 435 doi:10.1111/j .1550-7408.1975.tb05177.x 436 Ayres, J. S. & Schneider, D. S. (2012). Tolerance of infections. Annual Review of Immunology, 437 30, 271-294. doi:10.1146/annurev-immunol-020711-075030 438 Baird, S. J. E. & Goüy de Bellocg, J. (2019). Shifting paradigms for studying parasitism in 439 hybridising hosts: Response to Theodosopoulos, Hund, and Taylor. Trends in ecology & 440 evolution, 34, 387-389. doi:doi.org/10.1016/j.tree.2019.01.011 441 Baird, S. J. E., Ribas, A., Macholán, M., Albrecht, T., Piálek, J. & Goüy de Bellocq, J. (2012). 442 Where are the wormy mice? A reexamination of hybrid parasitism in the European house 443 mouse hybrid zone. Evolution, 66, 2757–2772. doi:10.1111/j.1558-5646.2012.01633.x 444 Balard, A., Jarquín-Díaz, V. H., Jost, J., Martincová, I., Ďureje, Ľ., Piálek, J., Macholán, M., de 445 Bellocq, J. G., Baird, S. J. E. & Heitlinger, E. (2020). Intensity of infection with intracellular 446 Eimeria spp. and pinworms is reduced in hybrid mice compared to parental subspecies. 447 Journal of Evolutionary Biology, 33, 435–448. doi:10.1111/jeb.13578 Baucom, R. S. & de Roode, J. C. (2011). Ecological immunology and tolerance in plants and 448

animals. Functional Ecology, 25, 18-28. doi:10.1111/j.1365-2435.2010.01742.x

450 Boots, M., Best, A., Miller, M. R. & White, A. (2008). The role of ecological feedbacks in the 451 evolution of host defence: What does theory tell us? Philosophical Transactions of the 452 Royal Society B: Biological Sciences, 364, 27–36. doi:10.1098/rstb.2008.0160 453 Brett, M. T. (2004). When is a correlation between non-independent variables "spurious"? 454 Oikos, 105, 647-656. doi:10.1111/j.0030-1299.2004.12777.x 455 Carval, D. & Ferriere, R. (2010). A unified model for the coevolution of resistance, tolerance, 456 and virulence. Evolution, 64, 2988-3009. doi:10.1111/j.1558-5646.2010.01035.x 457 Chapman, H. D., Barta, J. R., Blake, D., Gruber, A., Jenkins, M., Smith, N. C., Suo, X. & 458 Tomley, F. M. (2013). Chapter two - a selective review of advances in coccidiosis 459 research. 83, 93-171. doi:10.1016/B978-0-12-407705-8.00002-1 460 Clerc, M., Fenton, A., Babayan, S. A. & Pedersen, A. B. (2019). Parasitic nematodes 461 simultaneously suppress and benefit from coccidian coinfection in their natural mouse 462 host. Parasitology, 146, 1096-1106. doi:10.1017/S0031182019000192 463 Delignette-Muller, M. L. & Dutang, C. (2015). Fitdistributions: An r package for fitting distributions. 464 Journal of Statistical Software, 64, 1-34. doi:10.18637/jss.v064.i04 465 Ďureje, Ľ., Macholán, M., Baird, S. J. E. & Piálek, J. (2012). The mouse hybrid zone in Central 466 Europe: From morphology to molecules. Journal of Vertebrate Biology, 61, 308–318. 467 doi:10.25225/fozo.v61.i3.a13.2012 468 Ehret, T., Spork, S., Dieterich, C., Lucius, R. & Heitlinger, E. (2017). Dual RNA-seg reveals no 469 plastic transcriptional response of the coccidian parasite Eimeria falciformis to host 470 immune defenses. BMC Genomics, 18, 686. doi:10.1186/s12864-017-4095-6 471 Fineblum, W. L. & Rausher, M. D. (1995). Tradeoff between resistance and tolerance to 472 herbivore damage in a morning glory. Nature, 377, 517–520. doi:10.1038/377517a0

Floyd, R. M., Rogers, A. D., Lambshead, P. J. D. & Smith, C. R. (2005). Nematode-specific 473 474 PCR primers for the 18S small subunit rRNA gene. Molecular Ecology Notes, 5, 611–612. 475 doi:10.1111/j.1471-8286.2005.01009.x Gandon, S. & Michalakis, Y. (2000). Evolution of parasite virulence against qualitative or 476 477 quantitative host resistance. Proceedings of the Royal Society of London. Series B: 478 Biological Sciences, 267, 985-990. doi:10.1098/rspb.2000.1100 479 Graham, A. L., Allen, J. E. & Read, A. F. (2005). Evolutionary causes and consequences of 480 immunopathology. Annual Review of Ecology, Evolution, and Systematics, 36, 373–397. 481 doi:10.1146/annurev.ecolsys.36.102003.152622 482 Gregorová, S. & Foreit, J. (2000). PWD/Ph and PWK/Ph inbred mouse strains of Mus m. 483 musculus subspecies—a valuable resource of phenotypic variations and genomic 484 polymorphisms. Folia Biologica, 46, 31–41. 485 Haberkorn, A. (1970). Die Entwicklung von Eimeria falciformis (Eimer 1870) in der weißen 486 Parasitenkunde, Maus (Mus musculus). Zeitschrift für 34, 49–67. 487 doi:10.1007/BF00629179 Howick, V. M. & Lazzaro, B. P. (2017). The genetic architecture of defence as resistance to and 488 489 tolerance of bacterial infection in *Drosophila melanogaster*. Molecular Ecology, 26, 1533-490 1546. doi:10.1111/mec.14017 491 Jackman, S. (2020). pscl: Classes and methods for R developed in the political science 492 computational laboratory. United States Studies Centre, University of Sydney, Sydney, 493 New South Wales, Australia. 494 Janzen, D. H. (1980). When is it coevolution? Evolution, 34, 611-612. doi:10.1111/j.1558-495 5646.1980.tb04849.x

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Jarquín-Díaz, V. H., Balard, A., Jost, J., Kraft, J., Dikmen, M. N., Kvičerová, J. & Heitlinger, E. (2019). Detection and quantification of house mouse Eimeria at the species level -Challenges and solutions for the assessment of coccidia in wildlife. International Journal for Parasitology: Parasites and Wildlife, 10, 29-40. doi:10.1016/j.ijppaw.2019.07.004 Klemme, I. & Karvonen, A. (2016). Vertebrate defense against parasites: Interactions between avoidance. resistance, Ecology and 7. and tolerance. Evolution, doi:10.1002/ece3.2645 Kutzer, M. A. M. & Armitage, S. A. O. (2016). Maximising fitness in the face of parasites: A review of host tolerance. Zoology, 119, 281-289. doi:10.1016/j.zool.2016.05.011 Lefèvre, T., Williams, A. J. & de Roode, J. C. (2010). Genetic variation in resistance, but not tolerance, to a protozoan parasite in the monarch butterfly. Proceedings of the Royal Society B: Biological Sciences, 278, 751–759. doi:10.1098/rspb.2010.1479 Lüdecke, D. (2018). Ggeffects: Tidy data frames of marginal effects from regression models. Journal of Open Source Software, 3, 772. doi:10.21105/joss.00772 Macholán, M., Baird, S. J. E., Fornůsková, A., Martincová, I., Rubík, P., Ďureje, Ľ., Heitlinger, E. & Piálek, J. (2019). Widespread introgression of the Mus musculus Y chromosome in Central Europe. bioRxiv. doi:10.1101/2019.12.23.887471 Martincová, I., Ďureje, Ľ., Kreisinger, J., Macholán, M. & Piálek, J. (2019). Phenotypic effects of the Y chromosome are variable and structured in hybrids among house mouse recombinant lines. Ecology and Evolution, 9, 6124-6137. doi:10.1002/ece3.5196 Mazé-Guilmo, E., Loot, G., Páez, D. J., Lefèvre, T. & Blanchet, S. (2014). Heritable variation in host tolerance and resistance inferred from a wild host-parasite system. Proceedings of the Royal Society B: Biological Sciences, 281, 20132567. doi:10.1098/rspb.2013.2567

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Medzhitov, R., Schneider, D. S. & Soares, M. P. (2012). Disease tolerance as a defense strategy. Science, 335, 936–941. doi:10.1126/science.1214935 Mesa, J. M., Scholes, D. R., Juvik, J. A. & Paige, K. N. (2017). Molecular constraints on resistance-tolerance trade-offs. Ecology, 98, 2528-2537. doi:10.1002/ecy.1948 Piálek, J., Vyskočilová, M., Bímová, B., Havelková, D., Piálková, J., Dufková, P., Bencová, V., Ďureje, L., Albrecht, T., Hauffe, H. C., Macholán, M., Munclinger, P., Storchová, R., Zajícová, A., Holáň, V., Gregorová, S. & Forejt, J. (2008). Development of unique house mouse resources suitable for evolutionary studies of speciation. Journal of Heredity, 99, 34-44. doi:10.1093/jhered/esm083 R Development Core Team. (2013). R: A language and environment for statistical computing. http://www.R-project.org/. R Foundation for Statistical Computing. Vienna, Austria. Råberg, L., Graham, A. L. & Read, A. F. (2009). Decomposing health: Tolerance and resistance to parasites in animals. Philosophical Transactions of the Royal Society B: Biological Sciences, 364, 37-49. doi:10.1098/rstb.2008.0184 Råberg, L., Sim, D. & Read, A. F. (2007). Disentangling genetic variation for resistance and tolerance to infectious diseases in animals. Science, 318, 812-814. doi:10.1126/science.1148526 Restif, O. & Koella, J. C. (2004). Concurrent evolution of resistance and tolerance to pathogens. The American Naturalist, 164, E90–E102. doi:10.1086/423713 Rose, M. E., Hesketh, P. & Wakelin, D. (1992). Immune control of murine coccidiosis: CD4+ and CD8+ T lymphocytes contribute differentially in resistance to primary and secondary infections. Parasitology, 105, 349-354. doi:10.1017/S0031182000074515

541 Roy, B. A. & Kirchner, J. W. (2000). Evolutionary dynamics of pathogen resistance and 542 tolerance. Evolution, 54, 51-63. doi:10.1111/j.0014-3820.2000.tb00007.x 543 Schito, M. L., Barta, J. R. & Chobotar, B. (1996). Comparison of four murine *Eimeria* species in 544 immunocompetent and immunodeficient mice. The Journal of Parasitology, 82, 255–262. 545 doi:10.2307/3284157 546 Schmid-Hempel, P. (2013). Evolutionary parasitology: The integrated study of infections, 547 immunology, ecology, and genetics. Oxford University Press. doi:10.1093/acprof:oso/ 548 9780199229482 .001.0001 549 Shaw, D. J. & Dobson, A. P. (1995). Patterns of macroparasite abundance and aggregation in 550 wildlife populations: quantitative Parasitology. 111, review. S111-S133. 551 doi:10.1017/S0031182000075855 552 Sheldon, B. C. & Verhulst, S. (1996), Ecological immunology: Costly parasite defences and 553 trade-offs in evolutionary ecology. Trends in ecology & evolution, 11, 317–321. 554 Simms, E. L. (2000). Defining tolerance as a norm of reaction. Evolutionary Ecology, 14, 563-555 570. doi:10.1023/a:1010956716539 556 Smith, A. L. & Hayday, A. C. (2000). Genetic Dissection of primary and secondary responses to 557 a widespread natural pathogen of the gut, Eimeria vermiformis. Infection and Immunity, 558 68, 6273-6280. doi:10.1128/IAI.68.11.6273-6280.2000 559 Soares, M. P., Teixeira, L. & Moita, L. F. (2017). Disease tolerance and immunity in host 560 protection against infection. Nature Immunology, 17, 83-96. Reviews 561 doi:10.1038/nri.2016.136 562 Stange, J., Hepworth, M. R., Rausch, S., Zajic, L., Kühl, A. A., Uyttenhove, C., Renauld, J.-C., 563 Hartmann, S. & Lucius, R. (2012). IL-22 mediates host defense against an intestinal

564 intracellular parasite in the absence of IFN-y at the cost of Th17-driven immunopathology. 565 Journal of Immunology, 188, 2410–2418. doi:10.4049/jimmunol.1102062 566 Stowe K., Marguis R., Hochwender C., Simms E.L. (2000). The evolutionary ecology of 567 tolerance to consumer damage. Annual Review of Ecology, Evolution, and Systematics, 568 31, 565-595. doi:10.1146/annurev.ecolsys.31.1.565 569 Vale, P. F. & Little, T. J. (2012). Fecundity compensation and tolerance to a sterilizing pathogen 570 in Daphnia. Journal of Evolutionary Biology, 25, 1888-1896. doi:10.1111/j.1420-571 9101.2012.02579.x 572 Venables, W. N. & Ripley, B. D. (2002). Modern Applied Statistics with S (4th ed.). New York, 573 NY: Springer. doi:10.1007/978-0-387-21706-2 574 Wickham, H. (2016). Ggplot2: Elegant graphics for data analysis (second edition). New York, 575 NY: Springer. doi:10.1007/978-0-387-98141-3 576 Woolhouse, M. E. J., Webster, J. P., Domingo, E., Charlesworth, B. & Levin, B. R. (2002). 577 Biological and biomedical implications of the co-evolution of pathogens and their hosts. 578 Nature Genetics, 32, 569-577. doi:10.1038/ng1202-569 579 Zeileis, A., Kleiber, C. & Jackman, S. (2008). Regression models for count data in R. Journal of 580 Statistical Software, 27. doi:10.18637/jss.v027.i08

581 Tables

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Но	st	Parasite			
Mouse strains	Mouse subspecies	E. ferrisi Brandenburg64	E. falciformis Brandenburg88		
SCHUNT	F0 M.m.domesticus	14 (6M / 8F)	6 (3M / 3F)		
STRA	F0 M.m.domesticus	15 (8M / 7F)	7 (4M /3F)		
SCHUNTxSTRA	F1 M.m.domesticus	6 (2M / 4F)	8 (5M / 3F)		
STRAXBUSNA	F1 Hybrid	8 (5M / 3F)	8 (3M /5F)		
SCHUNTxPWD	F1 Hybrid	8 (3M / 5F)	6 (4M / 2F)		
PWDxBUSNA	F1 M.m.musculus	9 (4M / 5F)	7 (4M /3F)		
BUSNA	F0 M.m.musculus	14 (8M / 6F)	7 (3M /4F)		
PWD	F0 M.m.musculus	13 (10M / 3F)	7 (1M / 6F)		

Table 1. Infection experiment design.

Mous	status at dpi 11			
subspecies	group		alive	dead
Mmd	SCHUNT		6	0
Mmd	STRA		7	0
Mmd	SCHUNTxSTRA		8	0
Mmd-Mmm	STRAxBUSNA		8	0
Mmd-Mmm	SCHUNTxPWD		6	0
Mmm	PWDxBUSNA		4	3
Mmm	BUSNA		3	4
Mmm	PWD		1	6
		total	43	13

Table 2. Contingency table: number of mice and status at dpi 11 for each mouse group upon infection with E. falciformis isolate Brandenburg88.

Figures

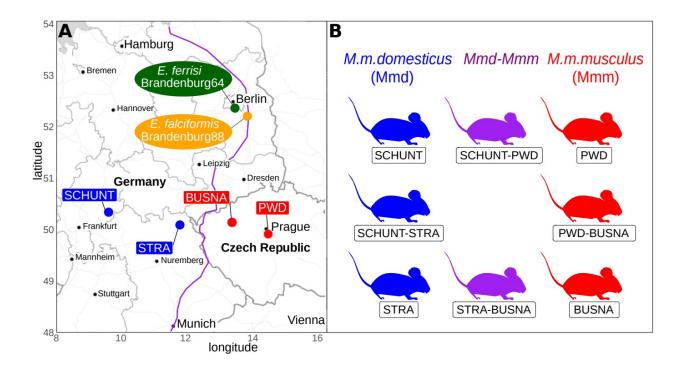


Figure 1. Parasite isolates and mouse wild-derived strains. (A) Map showing locations at which mice were collected for breeding of mouse strains and isolation of parasites. The purple line is an estimation of the center of the house mouse hybrid zone between *M. m. domesticus* and *M. m. musculus* based on sampling and genotyping of mice in this area (Balard et al. 2020; Ďureje et al. 2012; Macholán et al. 2019). (B) The eight mouse groups (parents and F1s) used in our experimental infections.

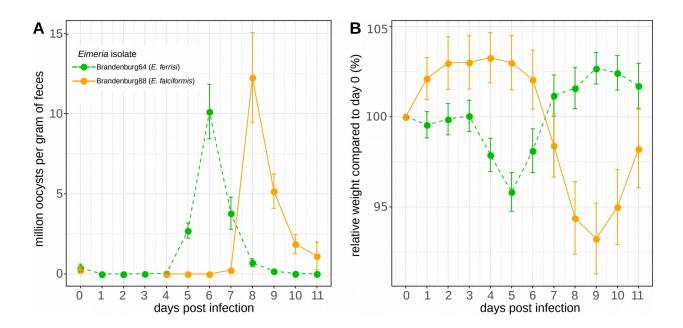


Figure 2. Parasite density (A) and host relative weight (B) during *Eimeria* infection. Parasite density is calculated as number of oocysts detected (in millions) per gram of feces, host relative weight is calculated as the percentage of weight compared to day 0. Mean and 95% CI are plotted for each parasite isolate. All mouse groups are pooled for each parasite isolate.

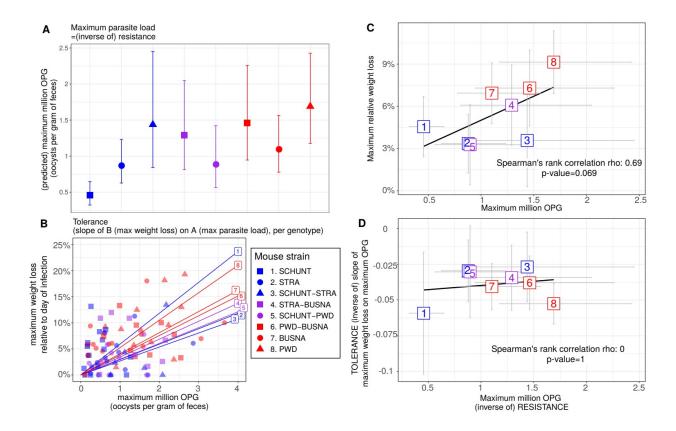


Figure 3. No indication of resistance-tolerance coupling for *E. ferrisi* isolate Brandenburg64. Colors represent mouse subspecies (blue: *M. m. domesticus*, red: *M. m. musculus*, purple: Mmd-Mmm). Left side: comparison of maximum oocysts per gram of feces used as a proxy for (inverse of) resistance (A) and tolerance (B) between mouse groups estimated by the slope of the linear regression with null intercept modelling maximum relative weight loss as a response of maximum oocysts per gram of feces, a steep slope corresponding to a low tolerance. Maximum number of OPG differs between mouse groups, but tolerance is similar. Right side: non significant positive correlation between mean maximum oocysts per gram of feces and mean relative weight loss (C) and absence of correlation between maximum oocysts per gram of feces used as a proxy for (inverse of) resistance and tolerance (D); Grey error bars represent 95% confidence intervals. Our results do not support coupling between resistance and tolerance *E. ferrisi* isolate Brandenburg64.

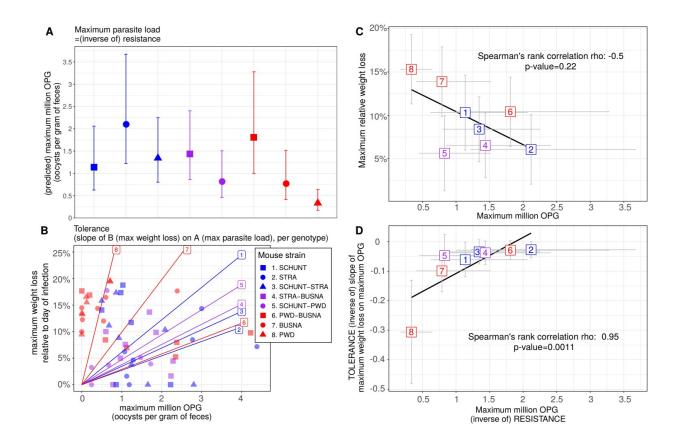


Figure 4. Coupling between resistance and tolerance for *E. falciformis* isolate Brandenburg88. Colors represent mouse subspecies (blue: *M. m. domesticus*, red: *M. m. musculus*, purple: Mmd-Mmm). Left side: comparison of maximum oocysts per gram of feces used as a proxy for (inverse of) resistance (A) and tolerance between mouse groups estimated by the slope of the linear regression with null intercept modelling maximum relative weight loss as a response of maximum oocysts per gram of feces, a steep slope corresponding to a low tolerance (B). Maximum number of OPG and tolerance differ between mouse groups. Right side: non significant negative correlation between mean maximum oocysts per gram of feces and mean relative weight loss (C) and strong positive correlation between maximum oocysts per gram of feces used as a proxy for inverse of resistance and tolerance (corresponding to a negative correlation between resistance and tolerance) (D); Grey error bars represent 95% confidence intervals. Our results support coupling between resistance and tolerance *E. falciformis* isolate Brandenburg88.

625 Appendix:

				Mouse strai	n (species)				
Batch	SCHUNT (Mmd)	STRA (Mmd)	SCHUNTX STRA (Mmd)	STRAX BUSNA (Mmd-Mmm)	SCHUNTX PWD (Mmd-Mmm)	PWDx BUSNA (Mmd)	BUSNA (Mmm)	PWD (Mmm)	Eimeria isolate (species)
B1	3	4					2	3	Brandenburg64 (E. ferrisi)
B2	4	4					5	3	Brandenburg64 (E. ferrisi)
B3	3	3	2	2	3	3	3	3	Brandenburg64 (E. ferrisi)
БЗ	3	3	3	4	3	3	3	3	Brandenburg88 (E. falciformis)
В4	4	4	4	6	5	6	4	4	Brandenburg64 (E. ferrisi)
	3	4	5	4	3	4	4	4	Brandenburg88 (E. falciformis)

626 Appendix 1. Chronology of experimental infections.

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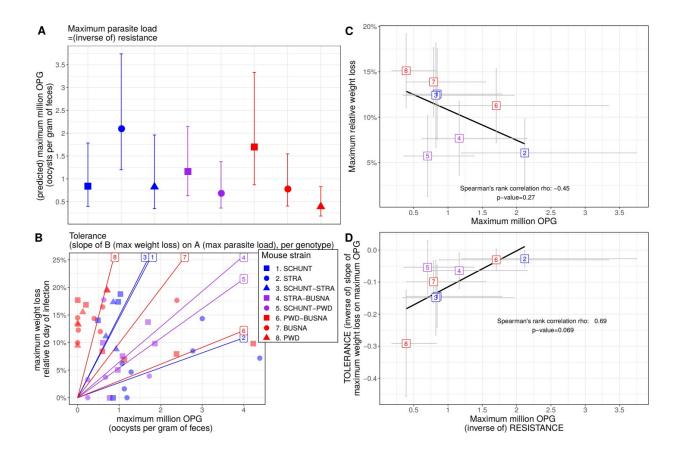
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(oocysts per gram of feces)

(inverse of) RESISTANCE

Appendix 2. No indication of resistance-tolerance coupling for *E. ferrisi* isolate Brandenburg64 in the conservative dataset. Colors represent mouse subspecies (blue: *M. m. domesticus*, red: *M. m. musculus*, purple: Mmd-Mmm). Left side: comparison of maximum oocysts per gram of feces used as a proxy for (inverse of) resistance (A) and tolerance (B) between mouse groups estimated by the slope of the linear regression with null intercept modelling maximum relative weight loss as a response of maximum oocysts per gram of feces, a steep slope corresponding to a low tolerance. Maximum number of OPG differs between mouse groups, but tolerance is similar. Right side: positive correlation between mean maximum oocysts per gram of feces and mean relative weight loss (C) and absence of correlation between maximum oocysts per gram of feces used as a proxy for (inverse of) resistance and tolerance (D); Grey error bars represent 95% confidence intervals. Our results do not support coupling between resistance and tolerance *E. ferrisi* isolate Brandenburg64.



Appendix 3. Coupling between resistance and tolerance for *E. falciformis* isolate Brandenburg88 in the conservative dataset. Colors represent mouse subspecies (blue: *M. m. domesticus*, red: *M. m. musculus*, purple: Mmd-Mmm). Left side: comparison of maximum oocysts per gram of feces used as a proxy for (inverse of) resistance (A) and tolerance (B) between mouse groups estimated by the slope of the linear regression with null intercept modelling maximum relative weight loss as a response of maximum oocysts per gram of feces, a steep slope corresponding to a low tolerance. Maximum number of OPG and tolerance differ between mouse groups. Right side: non significant negative correlation between mean maximum oocysts per gram of feces and mean relative weight loss (C) and strong positive correlation between maximum oocysts per gram of feces used as a proxy for inverse of resistance and tolerance (corresponding to a negative correlation between resistance and tolerance) (D); Grey error bars represent 95% confidence intervals. Our results support coupling between resistance and tolerance *E. falciformis* isolate Brandenburg88.