Ineffective humoral anti-tick IgY-response in birds: reaction against pathogen constituents? [version 1; peer review: 1 approved, 1 approved with reservations]

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Abstract
Background: Variation in parasite burdens among hosts is typically related to differences in adaptive immunity. Comprehension of underlying mechanisms is hence necessary to gain better insights into endemic transmission cycles. Here we investigate whether wild songbirds that have never been exposed to ticks develop adaptive humoral immunity against endemic Ixodes ricinus ticks.

Methods: Blue tits were exposed three times in succession to wild Ixodes ricinus ticks. For each infestation, serum samples were obtained. An enzyme-linked immunosorbent assay was developed, using tick salivary antigens, in order to quantify the bird's IgY response against ticks. In addition, at every sampling occasion the birds' body weight (corrected for body size) and haematocrit level was determined.

Results: Individual IgY levels against the ticks' salivary proteins increased over three consecutive tick infestations, and large among-individual variation was observed. The responses were specifically directed against I. ricinus; cross-reactivity against the congeneric tree-hole tick Ixodes arboricola was negligibly low. IgY responses did not impinge on tick feeding success (engorgement weight and attachment success). Yet, those birds with the highest immune responses were more capable to reduce the acute harm (blood depletions) by compensating erythrocyte loss. Furthermore, at the end of the experiment, these birds had gained more body weight than birds with lower IgY levels.

Conclusions: Latter observations can be considered as an effect of host quality and/or tolerance mechanisms. Birds anticipate the (future) costs of the activation of the immune system by ticks and/or ongoing tick-borne pathogen infections. Furthermore, although
unsuccessful against tick feeding, the IgY responses may indirectly protect birds against tick-borne disease by acting against salivary protein secretions on which pathogens rely for transmission.

**Keywords**
Tick, Ixodes ricinus, bird, Borrelia burgdorferi s.l., IgY, antibody, constituent, immunity

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Songbirds are central elements in the ecological networks of ticks, but are heavily overlooked when it comes to elementary biological mechanisms like immune responses against ticks and tick-borne diseases. No studies so far have related individual variation in wild songbird’s adaptive immune responses to components of ectoparasite fitness. Although the immune response was specifically targeted against the tick *Ixodes ricinus*, tick feeding success was not reduced and thus birds clearly did not acquire resistance against the ticks after being exposed for a long time. Interestingly, birds with the highest immune responses were more capable to reduce the acute harm and had gained more body weight than birds with lower IgY levels. Latter observations can be considered as an effect of host quality and/or tolerance mechanisms: birds anticipate the (future) costs of the activation of the immune system by ticks and/or ongoing tick-borne pathogen infections. Although unsuccessful against tick feeding, the immune responses may indirectly protect birds against tick-borne disease by acting against salivary protein secretions on which pathogens rely for transmission. Our study can be considered as a primer for future work exploring tick epitopes that can be targeted by bird immune components.

**Methods**

**Ethical statement**

All procedures, including the tick infestation (for more details see below), were carried out in accordance with national environmental legislation and regulations, and were approved by the Ethical Committee for Animal Experiments of the University of Antwerp (Licenses N° 2009-32 and 2016-88). Wild birds were captured under licences N° 58/VERG/07-USR26 and ANB/BL/FF-V17-00029 of the Agency for Nature and Forests, Flemish Government, Belgium. Bird individuals were kept in optimal conditions at the University of Antwerp, with food and water *ad libitum* in large cages (surface floor 40 cm x 80 cm; height: 40 cm) and had the opportunity to take a bath in fresh water. Birds were monitored daily. Wild birds were released after a minimum time period in captivity. Manipulations of a bird (infestation, blood sampling, weighing, measurement of tarsus length) occurred in a separate section of the lab room, outside the view of the other birds. As manipulations (see below) cause mild distress or harm, the use of analesics was not necessary.

**Bird serum samples**

Sera from tick-exposed birds were obtained from 16 blue tits that were exposed three times in succession with 12 nymphs over a time span of 30 days in the summer of 2008 (see Figure 1 for schematic overview of study design). All of them made
part of a previous ethically approved experiment and were in good condition (Heylen et al., 2010). Birds were kept in tick-free aviaries since hatching, thus naïve to ticks at the start of the experimental exposure. A blood sample (maximum 65 μL) was taken from the ulnar vein collected into 75 μL heparinized capillary tubes and subsequently centrifuged for 10 min at 14,000 g, after which the serum was separated from the blood clot and stored in Eppendorf tubes at -80°C until further analysis. To this end, the vein was superficially punctured with a needle (27G). Due to the small body size of the songbirds under study, the sampled serum volumes were kept to a minimal. As the minimum requirement for biochemical analyses was approximately 30 μL serum/bird per sampling occasion, only a limited number of birds (16) of the original experiment (31, see Heylen et al., 2010) could enter the longitudinal analyses (i.e. enough volume in three consecutive infestation sessions).

Monitoring of repeatedly exposed blue tits for seroconversion and physiological changes

When blue tits were nine weeks old, individuals were infested with I. ricinus nymphs three times in succession (Infestation 1–3) (see Figure 1 for schematic overview of study design). Each infestation lasted 4–5 days, and the birds were kept free of ticks for a duration of 5–6 days between the consecutive infestations. We infested birds with tick loads corresponding to the maximum level found under natural conditions in our study population (Heylen et al., 2013). To this end, 12 randomly sampled I. ricinus nymphs were put underneath the feathers on the head of each bird in each infestation session using moistened tweezers (Heylen & Matthysen, 2008). To this end, for each bird, Eppendorf tubes containing a nymph each, were randomly picked out of a box containing the remaining tubes with ticks. Birds were then kept for 2 h in an air-permeable cotton bag (size: 20 cm × 15 cm) inside a darkened cage, which kept them inactive. After tick exposure, birds were placed in individual cages with a wire-mesh floor (40 cm × 80 cm). Below the wire-mesh was a plastic tray containing damp filter paper and edges were streaked with vaseline to prevent nymphs from escaping. The engorged nymphs that dropped through the mesh cage were collected each day with minimal disturbance to the host (Heylen et al., 2010).

We estimated the effects of the IgY-response (ELISA described below) on tick measures as the change between exposure 1

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**Figure 1. Schematic overview of the study design.** (A) Actions taken during an infestation session: “Δ Inf. 1” denotes the acute effect of the tick exposures (12 nymphs) on the birds’ health in session 1, i.e. the change in health parameters between the onset of infestation and the time point that the last tick detached. (B) Schedule of successive infestation sessions: “Δ Acute” denotes the change in acute responses between the first and the third infestation session, “Δ Chronic” denotes the change in (baseline) health parameters, comparing health measures taken just prior to the first infestation and just prior to infestation three, when birds had the opportunity to recover from the acute tick effects.
and exposure 3. We measured the effect of IgY-response on the health measures as the change in health status (described below) between the moments immediately before tick exposure 1 and immediately before tick exposure 3 (i.e. 5–6 days after Inf. 2). Furthermore, we studied the change in the acute responses (i.e. the change in health status immediately before and just after tick exposure; Figure 1A) between infestation 1 and infestation 3 (Figure 1B) in response to the cumulative IgY response (i.e. the summed OD’s; see further).

We measured two parameters reflecting the hosts’ health status immediately before and after the infestation (Figure 1A). (1) Haematocrit (Hct) level: anaemia, as indicated by low Hct (the volume percentage of erythrocytes in the blood), results in a reduced oxygen-carrying capacity of the blood and restricts oxygen-demanding processes (Dein, 1986). Reticulocytes (i.e. immature erythrocytes) are stored in bird bone marrow, and can be instantly released in the blood stream (Martinho, 2012) upon sudden erythrocyte reduction (e.g. injury). Heparinized capillary tubes containing blood samples were centrifuged for 10 min at 14,000 g, and the ratio of packed red blood cells to the total volume was measured with a digital calliper to the nearest 0.01 mm under optimal light conditions. (2) Body condition (mass/tarsus ratio): body mass was measured to the nearest 0.1 g using a digital balance. To this end, the bird was immobilised by gently placing it in a tube (6 cm length, diameter 2.5 cm).

We subsequently calculated the ratio between body mass and a skeletal measurement (tarsus length, measured with a digital calliper to the nearest 0.01 mm) as a measure of body condition (Yom-Tov, 2001). Metabolic processes, e.g. for the compensation of parasite harm and mounting immune responses are known to be energy demanding and may lead to a reduced body condition when anaabolic processes are hindered (e.g. restricted food conditions, constrained metabolic pathways, etc.) (Martin et al., 2003).

Ixodes ricinus collection and feeding success
Ixodes ricinus nymphs were caught by dragging a white flannel flag over suitable vegetation. The ticks were subsequently kept under sterile conditions in a climate room at >90% relative humidity, a 16 h:8 h (light:dark) photoperiod, and a 25°C:15°C temperature cycle until infestation. After feeding on the blue tits, the engorged nymphs were weighed to the nearest 0.01 mg. To investigate the influence of repeated infestations on feeding success, we estimated the following parameters: (1) the proportion of the administered ticks that successfully engorged (tick yield), (2) the total weight of engorged nymphs and (3) their feeding duration. If hosts acquire resistance, this is expected to result in the following observations compared to naïve hosts (Rechav, 1992): lower numbers of engorged ticks, smaller blood meals (lower weight of engorged ticks), and increased feeding durations. From these criteria, engorge-ment weight is considered as to be one of the most consistent indicators of resistance (Varma et al., 1990).

ELISA-based detection of immunoglobulins
Tick’s salivary gland extraction (SGE). Salivary glands were dissected from 32 semi-fled colony-reared infection-free adult female I. ricinus obtained from IS insect Services GmbH in Berlin, Germany. Before dissection, all ticks had fed on sheep for 5–7 days; feeding is known to significantly increase SGE concentrations (Mateos-Hernández et al., 2017; Ogden et al., 2002). The glands were washed four times in PBS to remove tick debris, pooled and homogenized. To test IgY-cross-reactivity with antigens of a congeneric tick species, we used material of 12 engorged adults I. arboricola ticks (Heylen et al., 2014) that had fed on great tit nestlings and that were dissected in the same way. A pool, containing the glands of 6–8 ticks in 60 μl PBS, was manually disrupted with a sterile pestle and the following steps were performed before storage at -80°C: sonication three times for five seconds with a treatment in ice (BRANSON 150), centrifugation at the maximum speed (10,000 rpm) for 10 minutes at 4°C, and filtering through a 0.2 μm filter (Chromafil AO-20/3 Macherey-Nagel GmbH, Düren, Germany) to remove contaminating bacteria. Total protein concentrations were estimated by Nanodrop (Cafiso et al., 2019) and equilibrated to 1 mg/mL prior to use in the assays.

IgY ELISA. Basing ourselves on Ogden et al. (2002), after optimization of the ELISA-protocol - including the binding capacities of the anti-chicken antibodies for passerine raised antibodies (sandwich ELISA in which plates are coated with bird sera) – the following volumes and concentrations yielded the most reliable and repeatable results for the indirect ELISA tests. To coat the 96-well microtiter plates (Nunc Maxisorp flat bottom, Thermo Fisher Scientific, Geel, Belgium), 150 μL PBS per well was used, containing a concentration of 1.8 μg/mL SGE. Negative controls were coated with 150 μL PBS only. Plates were incubated for 12 hours at 4°C. After coating, the plates were washed three times with 200 mL PBS to remove the unbound material prior to blocking, using 200 μL of 0.5% Bovine Serum Albumin in PBS per well, and incubation for 1h at 37°C. Subsequently, the solution was removed and the wells were rinsed with PBS. Primary antibodies were obtained from bird sera diluted 140 times in PBS. 150 μL of the diluted bird sera were added to appropriate wells and the plates were incubated for 1 hour at 37°C. Afterwards, the wells were washed four times with 200 mL of PBS and 150 μL of the labelled secondary antibody (Rabbit anti-Chicken IgG, FC specific-alkaline phosphatase antibody, Sigma-Aldrich, code SAB3700239, Overijse, Belgium, -15000 times diluted in PBS) was added. After one hour incubation at 37°C, the plates were washed three times with 200 mL of PBS and pre-washed once with 200 μL of alkaline phosphatase (AP) buffer (100 mM Tris, 2 mM MgCl2, pH 9.6 with HCl). The amount of secondary antibody bound to the primary antibodies is visualized through AP reaction after adding 150 μL of a 1 mg/mL 4-nitrofenylflosfaat dilution in AP reaction buffer and one hour incubation at 37°C. Plates are read with a plate reader (Biotek Synergy MX, BioTek, Winooski, VT, USA) measuring OD at 405nm.
Repeatability and qualitative discrimination infested vs. non-infested birds. Negative control serum samples were obtained from three adult (sex unknown) domesticated canaries (*Serinus canaria*, L. 1758), belonging to a captive population maintained for multiple generations (15 years) at the University of Antwerp. In addition, three 1st calendar year blue tits (*Cyanistes caeruleus*), and three 1st calendar year great tits (*Parus major*, L. 1758) that were kept in tick-free aviaries since hatching (sex unknown; see Heylen et al., 2010 for further details on origin and housing). Sera from tick-exposed birds were obtained from three free-living great tits (1st calendar male and female, one 2nd calendar year male) that showed to be *Ixodes ricinus* tick-infested upon capture with mist nets (early Autumn 2019, Antwerp, Belgium), three 1st calendar year great tits (sex unknown) that were three times experimentally exposed to 17 *Ixodes ricinus* nymphs over a time span of 30 days (Heylen et al., 2010), and three of the abovementioned blue tits.

The IgY levels in serum samples belonging to the same individuals showed to be highly repeatable within an ELISA-plate (Pearson’s Rho: 0.93; N= 17) as depicted by the scatterplot (Figure 2). One measurement was excluded (in the non-infested Sc 3) as a pipetting error had occurred. Considerable variation in IgY levels was observed among tick-exposed individuals of the same species (variance/mean in great tits: 10%; blue tits: 19%), but also in the naive blue tits (10%). Both naturally infested (caught in the wild and blood sampled once) birds and repeatedly infested birds (following scheme depicted in Figure 1) showed noticeably higher OD values than non-infested individuals.

Statistical analysis
For the qualitative comparisons between infested and non-infested birds (only three individuals per bird species, over the two groups) no statistical tests were performed (Figure 2), neither for the description of the IgY-profiles of three blue tits - for which sufficient amounts of serum and tick antigens allowed a quantification at each of the six time points (Figure 1 and Figure 3). Data of IgY levels at Day 1, 11 and 21 (Figure 1) of the latter three birds were combined with that of ten additional blue tits, followed by the parametric statistical analyses as described below:

generalized linear mixed effect models (GLMM’s) were fitted on health measures to model the acute infestation effects (Δ Inf. 1, 2 and 3; Figure 1) as a function of the bird’s IgY levels (OD value) within each infestation. To avoid collinearity problems and to adjust for differences in variation between infestations, the IgY levels were standardized (OD

\[ \text{Inf.x} - \text{mean}_{\text{Inf.x}} / \text{Standard deviation}_{\text{Inf.x}} \]

By adding a random bird individual effect, and using Kenward-Roger approximation for the denominator degrees of freedom, we took into account the correlation of observations within the same individuals.

In a second statistical analysis, we tested whether the changes in acute effects (‘Δ Acute’, Figure 1) were related to the summed IgY levels over the three infestations (SOD), which we consider as a proxy for the bird’s overall anti-tick IgY production over the course of the experiment. In a final analysis we modelled the change in initial values (Δ Inf. 3–1) (= chronic response) as a function of SOD. Effects of IgY levels on tick

**Figure 2.** ELISA’s optical density measures of two IgY-measurements on the same bird. IgY levels were obtained from non-infested and *Ixodes ricinus*-infested birds (great tit, blue tit, canary). Among the infested birds, three great tits were very recently infested when caught in the wild (filled squares). All other samples were obtained from birds that fledged in tick-free aviaries or cages (control), and which were three times experimentally infested with ticks (including bird 1–3 of the 16 birds in the blue tit experiment).
feeding parameters were modelled in a similar way, except for the fact that we only have one value per bird/infestation session (and not a difference). Before entering the analysis, the average tick measures (weight, feeding duration) were calculated for each bird/infestation. In all models, a stepwise selection procedure was used in which the model was iteratively refitted after exclusion of the least significant effect, until only significant factors and their lower order interactions terms were left. All data manipulations and statistical analyses were performed using SAS v 9.2 (SAS Institute, Cary, North Carolina, USA). Estimates are reported as mean ± standard error.

**Results**

**Sera I. ricinus-exposed birds and cross-reactivity with I. arboricola antigens**

In the three blue tits that were monitored at six time-points (Figure 3), we observed an OD-curve that tended to be bell-shaped, with the highest IgY levels around Day 15–21 (Heylen, 2021).

The ten blue tits (Figure 4), for which samples were analysed at three time points only (Day 1, 11, 21) showed OD-profiles that were monotonically increasing (0.014 ± 0.004 OD unit/Day unit, T-value = 3.49, df = 10.8, P = 0.0052, except for one (bird 3). Large individual variation was observed on each of the sampling days (variance/mean Day 1: 5%, Day 11: 17%, Day 21: 16%), and variation in slopes differed from zero (estimate: 0.14 ± 0.1 10⁻³; Likelihood ratio test: Z = 1.70; P = 0.044).

The antibodies against I. ricinus-antigens in the Day 15-samples showed almost no cross-reactivity against I. arboricola-antigens (Figure 4). In two birds (bird 2 and 4) the OD’s in the I. arboricola wells were slightly higher than those when birds were still naïve (Day 1).

**IgY correlations with anti-tick resistance**

Due to limitations in the amount of serum available, we were not able to provide standard positive samples over all plates. Although small plate differences in OD’s can occur,
the values of the three birds of Figure 3 were well situated within the variation of the 10 additional birds depicted in Figure 4 that were analysed in a separate assay, and therefore included in subsequent analyses.

Test statistics for the cross-sectional (Inf. 1, 2 and 3) and longitudinal analyses (Δ Inf. 3–Inf. 1) in relation to the IgY levels are presented in Table 1. Neither the average engorgement weight (Figure 5) nor the feeding duration showed significant associations with IgY level in any of the analyses (Table 1), despite the strong IgY-increase and large among-individual variation (Figure 3 and Figure 4). Furthermore, the proportion of ticks that successfully engorged did not covary with IgY levels. For absolute values of feeding parameters, we refer to Heylen et al. (2010). Conclusions did not change when restricting the analyses to the subset of 10 individuals that were simultaneously analysed on a single plate (Figure 4).

IgY correlations with tick virulence

Test statistics and model parameter estimates for the change in acute and chronic effects of the tick exposure in relation to the IgY levels are presented in Table 2. For absolute values of physiological variables we refer to Heylen et al. (2010).

Acute effects: while on average the Hct levels did not significantly change during the first two infestations (‘Acute’ Inf. 1: 1.97 ± 1.29; Inf. 2: -1.14 ± 1.08%), they decreased in the third infestation (Inf. 3: -6.24 ± 1.63%, T-value = -3.82, P = 0.0028). Birds with higher IgY levels prior to an infestation showed a less severe Hct decrease (2.23 ± 0.80%/OD unit, T-value = 2.92, df = 31, P = 0.0065; Figure 6). In the analysis, two statistical outliers belonging to the same bird were removed (see Figure 6). The difference in acute effects (‘Δ Acute’) between infestation 1 and 3 did not correlate with the summed IgY levels.

Body condition monotonically increased throughout the experiment for all birds (see below: ‘chronic effects’), but in none of the infestation sessions we found an acute tick effect.

The outcomes and interpretation of the above analyses did not change, when excluding the data of the three birds of Figure 3.

Chronic effects: We found that the increase in body condition over the infestation sessions (Δ Chronic: 0.027 ± 0.006 g/mm; T-value: 4.38, df = 11; P < 0.001) was higher in birds that had higher summed IgY levels (SOD) (0.033 ± 0.006 g/mm; T-value = 5.33, df = 11, P = 0.0002; Figure 7). Hct levels

Figure 4. IgY levels of sera from ten blue tits (bird 7–16) that were three times infested with I. ricinus (for design see Figure 1). Values of a second ELISA are included, in which the cross-reactivity of anti-I. ricinus IgY's was tested (Day 15 samples only) against I. arboricola salivary antigens.
Table 1. Type 3-test outcomes of GLMM’s for repeated measurements. IgY levels were measured in the serum samples taken at the beginning of each infestation session. In the analyses of the cross-sectional correlations (‘per infestation’) IgY levels have been standardized. For the longitudinal analysis (i.e. the cumulative response, ‘Inf. 3 minus Inf. 1’), the summed IgY levels over the three infestation sessions was calculated (See Figure 1). Test statistics before exclusion from the model are given, as well as the parameter estimates and statistics for the terms that remained in the model (P-value <0.05).

<table>
<thead>
<tr>
<th>Per infestation</th>
<th>Infestation</th>
<th>IgY (SD)</th>
<th>IgY x Infestation</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>(ndf,ddf)</td>
<td>F</td>
<td>(ndf,ddf)</td>
</tr>
<tr>
<td>Engorgement weights</td>
<td>(1,35)</td>
<td>0.15 NS</td>
<td>(1,33)</td>
</tr>
<tr>
<td>Feeding durations</td>
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<td>0.74 NS</td>
<td>(2,33)</td>
</tr>
<tr>
<td>Engorgement success</td>
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<td>0.63 NS</td>
<td>(2,32)</td>
</tr>
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</table>

<table>
<thead>
<tr>
<th>Inf. 3 minus Inf. 1</th>
<th>Summed IgY’s</th>
</tr>
</thead>
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<tr>
<td></td>
<td>(ndf,ddf)</td>
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<tr>
<td>Engorgement weights</td>
<td>(1,11)</td>
</tr>
<tr>
<td>Feeding durations</td>
<td>(1,10)</td>
</tr>
<tr>
<td>Engorgement succes</td>
<td>(1,11)</td>
</tr>
</tbody>
</table>

Infestation: early (Inf. 1), during sero-conversion (Inf. 2) and at maximum IgY levels (Inf. 3); NS: P-value >0.05.

Figure 5. Difference in the summed engorgement weights between the beginning of the experiment (Inf. 1, naïve bird) and the end (Inf. 3, previously exposed to 24 ticks) as function of the summed IgY levels in blue tit sera (bird 4–16). Total engorgement weight is a function of the average engorgement weight and the proportion of successfully fed ticks, none of which showed a significant association with IgY levels (Table 1).
**Figure 6.** Acute effects of ticks on Hct levels in response to the standardized IgY levels of sera samples in 13 blue tits (bird 4-16) presented in Figure 3 and Figure 4. Two outliers from the same individual were excluded from the statistical analyses.

**Table 2.** Outcomes of generalized linear mixed effect models (GLMMs) investigating the association between IgY levels and the health parameters in 13 birds repeatedly infested with *Ixodes ricinus* nymphs. See Figure 1 for the experimental design, and further explanation of 'Acute effect' ('after' minus 'before' exposure), Δ Acute (Change in acute effects Inf. 3 vs. Inf. 1) and Δ Chronic (Day 21 minus Day 1).

<table>
<thead>
<tr>
<th>Acute effect</th>
<th>Infestation</th>
<th>IgY (SD)</th>
<th>IgY x Infestation</th>
<th>Average Slope over</th>
</tr>
</thead>
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<tr>
<td>Haematocrit change</td>
<td>(2,31) 12.06</td>
<td>(1,31) 8.76</td>
<td>(2,29) 1.70</td>
<td>0.024±0.008</td>
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<tr>
<td>Condition change</td>
<td>(2,33) 0.06</td>
<td>(2,29) 0.69</td>
<td>NS</td>
<td></td>
</tr>
<tr>
<td>Δ Acute</td>
<td>Summed IgY's</td>
<td>(ndf,ddf) F</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Haematocrit</td>
<td>(1,11) 0.16</td>
<td>NS</td>
<td></td>
<td></td>
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<tr>
<td>Condition</td>
<td>(1,11) 4.42</td>
<td>0.064</td>
<td></td>
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<tr>
<td>Δ Chronic</td>
<td>Haematocrit</td>
<td>(1,11) 0.65</td>
<td>NS</td>
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<tr>
<td>Condition</td>
<td>(1,11) 28.41</td>
<td>0.0002</td>
<td>0.033±0.006</td>
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Infestation: early (Inf. 1), during sero-conversion (Inf. 2) and at maximum IgY levels (Inf. 3).

* Outcome without the bird giving outlying residuals. This bird led to violation of the normality assumption (Shapiro-Wilk W: 0.96); removing it improved the model fit (Shapiro-Wilk W: 0.98).

NS: P-value >0.05.
did not change as a function of summed IgY’s. The sum of acute health effects (sum over Inf. 1, 2 and 3) and SOD were not significantly correlated (all P-values > 0.05).

Discussion
This is the first study that investigates the interplay between the songbirds’ humoral immune response and natural blood-sucking ectoparasites (ticks), combining observational and experimental approaches. The acquisition of tick immunity has been an important topic in the field of tick biology, as patterns of immunity can be used to develop transmission blocking vaccines for humans or reservoir animals against tickborne pathogens. Immunity of mammals and non-mammalian hosts (such as birds) may differ in response to the same antigens (e.g. tick salivary proteins), which emphasizes the need of the current study. Furthermore, as wild birds can maintain and spread tick-borne pathogens, the relevance of the use of wild phenotype birds for endemic transmission cycles likely outweighs the limitation linked to the maintenance of wild animals in lab conditions. We put forward three questions (1) do birds develop a specific antibody response against *Ixodes ricinus* salivary antigens and how strong is the variation among birds? (2) Does the humoral immune response reduce tick feeding success (3) and virulence?

In order to address the first question, we developed an indirect ELISA, and succeeded to quantify the bird’s immunoglobulins (IgY) that bind to *I. ricinus* salivary gland antigens. We then could show that the level of tick-specific IgY was low at the beginning of the experiment, when birds were naïve, then steeply increased to peak at 15–20 days - the moment of seroconversion – and tended to decrease afterwards (Figure 3). This sero-conversion pattern is comparable to IgG-kinetics in mammals (Barriga et al., 1991). IgG is the mammalian analogue to the avian IgY, present in the chronic phase of parasite exposure, and is involved in the development of long-term resistance (Davison et al., 2008) against ticks (Ogden et al., 2002). In our study, the IgY-response turned out to be specifically targeted against *I. ricinus* salivary antigens, as the cross-reactivity against *I. arboricola*-antigens was shown to be negligibly low (Figure 4). We found significant individual variation among birds in immune profiles, as well as in initial values before being exposed for the first time. The most likely explanation for the latter finding is that maternal antibodies of mothers with anti-tick IgY-concentrations in their blood have been transferred via egg yolk to nestlings (Müller et al., 2004). This additional source of antibodies complicates the interpretation of IgY levels as signals of the individual’s previous tick exposure, especially in juveniles (Gasparini et al., 2001).
For our second question, we looked at pairwise relationships between the IgY levels and either tick feeding parameters or bird health measures. As observed in other natural hosts (Fielden et al., 1992; Ribeiro, 1989), the opportunistic I. ricinus turns out to be extremely efficient in circumventing the bird’s antibody response. In host types where tick resistance is acquired, strong decreases in engorgement weights are observed in subsequent tick exposures. We found that, despite the high among bird-individual variation in IgY-responses and tick feeding success, there was no significant association between them. We conclude that naïve juveniles do not acquire anti-tick resistance. The outcomes could be viewed as a mechanism of tolerance instead of resistance. The latter is the capacity to limit the parasite burden, while the first refers to the ability to limit the harm caused by a given burden by compensatory mechanisms (Råberg et al., 2009). The concept of tolerance is notoriously difficult to measure in animals, when measuring the slope of how host fitness decreases with parasite burden. Here, high burdens did not cause direct fitness effects (bird mortality), or gave rise to indirect fitness effects via physiological measure that link up with bird fitness; both findings are in favour of tolerance.

While on average the ticks performed equally well throughout the experiment (i.e. no significant difference between infestations 3 and 1), the harm (i.e. blood depletion) seemed to be better compensated for when birds had higher IgY levels. Massive amounts of reticulocytes (i.e. immature erythrocytes) are stored in bird bone marrow, and can be instantly released in the blood stream (Martinho, 2012). We point out that the net Hct difference in each infestation (Figure 1 – Figure 6) is the result of two processes: the immediate erythrocyte compensation (i.e. addition of erythrocytes in the blood stream) and acute erythrocyte depletion due to tick feeding. Although tick feeding did not decrease with IgY, the net effect of the abovementioned processes showed a correlation with IgY. We do not know of any role of IgY in these processes, but since both immune responses and harm compensation are energetically demanding, general health could simply be driving the observed correlation. Additionally, birds with a higher overall IgY-response gained more body weight (first 21 days). Metabolic processes for the compensation of the blood depletion by the ticks, the repair of skin lesions and blood vessels, and mounting immune responses are all energy demanding (Martin et al., 2003), and may lead to a reduced body condition. However, under lab conditions, with food ad libitum, those birds with the strongest IgY-response showed to be more successful in gaining body mass. The observed increase may relate to the gain in body mass for the regeneration of feathers (i.e. the post-juvenile moult) (Bojarinova et al., 1999) or other undefined seasonal physiological changes. Also, by gaining body mass, birds possibly anticipated the costs of (chronic) activation of the immune system due to tick infestations and/or ongoing tick-borne pathogen infections (as birds were exposed to ticks from the wild) (Heylen et al., 2015). In the end, this may benefit the fitness of both the ticks and micro-organism: future ticks could feed more successfully in those birds with a stronger body mass increase and it is conceivable that they may even induce such processes. We mention that our results are correlational, and do not prove causation; anti-tick IgY-response are not necessarily the cause of better health outcomes, but could be a correlated by-product of variation in quality among the birds. This quality (condition, health, vigour) could be affected by several factors, including good genetic constitution, higher quality maternal care, lower stress experience, fewer co-parasites, etc. In the ecological immunology literature, many studies have shown that variation in condition will drive associations between immunological traits (Sadd & Schmid-Hempel, 2009) but from our study it is clear that the measured IgY’s did not correlate with the tick’s feeding success, despite they targeted tick salivary antigens.

Birds are central elements in the ecological networks of ticks, but are heavily overlooked when it comes to elementary biological mechanisms like immune responses (De la Fuente et al., 2015). In fact, surprisingly few studies have related individual variation in host immunity to components of ectoparasite or even tick- fitness. Despite being unsuccessful in reducing tick feeding success via IgY’s, birds may benefit from the observed IgY-responses: by indirectly acting against vector-borne pathogen constituents (i.e. tick proteins functioning as carrier vehicles) tick-to-host transmission may become mitigated. As the transmission of pathogenic tick-borne agents heavily relies on salivary proteins (De la Fuente et al., 2015), it is worth studying this hypothesis for a variety of tick-borne pathogens in (in)competent natural reservoir songbird hosts. We may also wonder whether anti-tick immune responses are affected by the pathogens themselves, and whether these responses are comparable in different tick-pathogen-host systems. All these questions are heavily unexplored in birds but are crucial for understanding local transmission and life cycles. Answers could even inspire vaccine development, when mapping the tick epitopes that are effectively used by pathogens and could be targeted by host immune components.

Data availability

Data are available under the terms of the Creative Commons Attribution 4.0 International license (CC-BY 4.0).

Acknowledgements
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References


Maxime Madder
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Ineffective humoral anti-tick IgY-response in birds: reaction against pathogen constituents. Heyler et al.
The authors present an interesting study trying to define the effect of a humoral response in blue tits that are exposed to multiple infestations with wild caught *Ixodes ricinus* ticks.

General comments
The paper offers an interesting insight the way blue ticks react to multiple infestations of ticks, but some general information is missing as pointed out below.

Introduction
It might be interesting for the reader to explain what type of host immune responses are triggered by ticks (including birds), as other studies were referred to (Heylen et al., 2010, 2015), and this to explain why IgY was selected. Also why IgY was looked at and not IgA or M.

Methods
Having used wild caught ticks, and not knowing their infection status with viral, bacterial or protozoal disease agents, could the authors clarify if the use of infected ticks might have influenced the results and conclusions? Many articles describe the influence of infected ticks on the ticks' behaviour, but also on the host immune response. Could the authors specify where the ticks were collected, what the known pathogens are in this region and their prevalence, and where the birds originate from. Also the date the study was conducted is not specified.

Discussion
The authors state the the birds tolerated the ticks, and did not show anti-tick resistance. Could this be dose-dependent? Could higher infestation loads result in anti-tick resistance? It might be hypothesised that tolerance is seen as long as birds are not negatively impacted by infestation.

Is the work clearly and accurately presented and does it cite the current literature?
Yes

**Is the study design appropriate and does the work have academic merit?**
Yes

**Are sufficient details of methods and analysis provided to allow replication by others?**
Yes

**If applicable, is the statistical analysis and its interpretation appropriate?**
Yes

**Are all the source data underlying the results available to ensure full reproducibility?**
Yes

**Are the conclusions drawn adequately supported by the results?**
Yes

**Competing Interests:** No competing interests were disclosed.

**Reviewer Expertise:** Parasitology, acarology, ticks and tick-borne diseases, vector ecology, animal health

I confirm that I have read this submission and believe that I have an appropriate level of expertise to confirm that it is of an acceptable scientific standard.

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Michelle Wille
University of Sydney, Sydney, NSW, Australia

**Heylen: Ineffective humoral anti-tick IgY-response in birds: reaction against pathogen constituents**

The authors aimed to understand the immune response of Blue tits to tick infestation using an experimental approach. Overall, this is an interesting paper, but currently some shortcomings in data presentation (both text and writing) make it hard to understand the details and interpret of the findings.

**Some general questions:**
I think a bit more careful explanation is needed for the reader to understand the idea of “tick resistance”. Specifically, that it’s not the prevention of ticks infesting birds, but rather a decrease in the impact of ticks on birds (eg, lower tick engorgements and perhaps a more effective immune response).

Second, humoral immunity develops following the acute response (inflammation). How do these findings link into studies assessing inflammation following tick infestation? Would this variation in IgY perhaps be due to a variation in acute responses? I recognise that the authors did not measure this, but I wonder if there is data from previous studies?

Third, as these were field collected ticks, how do the authors control for things like viral infections which may affect Hct or body condition. These viral infections may also affect Engorgement weights and feeding durations of ticks (particularly for viruses infecting the ticks, not necessarily arboviruses which are transmitted to the birds). Theoretically they shouldn’t affect the IgY as this is specific. I recognise that the authors did not measure this, but it is worth point out in the discussion.

Specific comments:

Methods
○ I would defer to the expertise of other reviewers to appropriately assess the methods of this paper.

Results

“Sera I. Ricinus-exposed birds and...”:
○ I would encourage the authors to open the first sentence with the purpose of this experiment. For example, “In order to address XXXX, we three tits were monitored at six time points (as opposed to 3). In these birds we found....”

○ Similarly for the next paragraph. “Following this, 10 blue tits (including or excluding the 3 before?) were monitored at three time points. These time points corresponded to direction prior to infestation, and thus reflecting the IgY levels following the previous infestation event. In these birds, OD profiles were....”

“IgY correlations with anti tick resistance”:
○ The opening of the second paragraph could improve. Something like: “In order to investigate whether XX influenced YY, we utilised a GLMM. Neither age etc...”

○ Also, please help the reader a bit more. I'm not sure why “Inf 1” is cross-sectional and “delta Inf3-Inf 1” is longitudinal. Can you explain here what these values mean (especially given Figure 1 is not at all intuitive). So, based on this analysis, what did influence IgY levels? Perhaps add, “In sum, none of the factors we tested influenced IgY levels across either each infestation, or across all three infections”.

“IgY correlations with tick virulence”:
○ This section is not well written – it’s rather written like a list of bullet points rather than a “story” whereby the reader is walked through the analysis with care. For example, adding leader statements like “In order the assess the acute effects, i.e., the variation in Hct levels, XXX YY” are useful.
- Why were these data points deemed outliers – was this a technical problem?
- Why were the IgY levels summed?

**RE Body condition:**
- Perhaps rephrase to say something like “Body condition...for all birds, with the exception of infestation sessions wherein we found an acute effect (here explain which sessions those were).” I also don't understand why there is reference to chronic effect here?

**Paragraph 4:**
- Does this mean that you did the analysis twice – once including and once excluding the Birds in Figure 3 (i.e., the birds that were measured 6 times and therefore did not have enough sera remaining to be on all plates)?

**Paragraph 5:**
- Again, more common language interpretations would be useful here. Why did you use summed values? Why does Hct and Body condition refer to chronic? See comments above about including a leading sentence.

**Discussion**

**Paragraph 2 RE maternal antibodies:**
- Are there studies you can reference that have quantified how long maternal antibodies persist in birds. For example, this has been done in ducks for antibodies against influenza A, but I wonder if this has been done for any passerines for any virus/bacteria? This data is useful for future studies such that you may start experiments when birds are older, and you are sure that the effect of maternal antibodies has waned.

**Paragraph 3 RE tick resistance:**
- Can the authors please provide an example of hosts types that have been shown to acquire resistance to ticks?
- Also, you have a statement: “We conclude that naïve juveniles do not acquire anti-tick resistance”. Do adult passerines acquire this resistance? Are you sure you are using the correct measurements and experimental design for this as it is a strong statement to make?

**Paragraph 3 RE tolerance.**
- Given fitness wasn't really measured and the experimental design limited and controlled for tick infestations, I am not sure how convincing this is. Are there are examples that the authors could reference for animals that tolerate ticks?

**My suggestions for Figure improvement:**

I would think that a number of the figures that are related could be panelled into a larger figure? For example, Fig 3 and 4 together, perhaps Fig 5, 6, 7 together?

**Figure 1:**
- Concept figures should be immediately understandable without needing to read the legend, but unfortunately Figure 1 is very confusing. There is plenty of space so I suggestion you write out “Inf. 1” and “D1”. The legend isn't very clear either. The figure doesn't really explain well the different between “Acute” and “Chronic”. Overall, I suggest reconsidering this figure – there are lots of great concept figures out there in papers that rely on infection experiments for inspiration.

**Figure 2:**
- Based on the Figure and the legend, it is unclear to me the purpose of this figure. Perhaps a
more descriptive title such as: Correlation between first and second IgY measurement of
the same sample/bird”. Can you please add the Statistic onto the figure, and also the line.
I expect that you could leverage more colours to make it easier for the reader.

**Figure 3:**
- Again, I would change the figure legend so that this plot is easier to appreciate. For example: Change in IgY levels of 3 Blue Tits that were sampled on 6 occasions. Given Figure 1 isn’t intuitive, I still don’t really understand this plot. Perhaps add some extra metadata below the bars – lines or square brackets to indicate infestation periods, or different colours or arrows? Were the positive and negatives only for Day 1?

**Figure 4:**
- I would suggest that this plot be a boxplot or point graph, with day on the x-axis, and the different birds in different colours.
- You could, if you wanted, connect the individuals with lines. You could also then add a line or similar for the mean? Even better, you could model the change of IgY over time? An example of a suggestion: [link](https://stackoverflow.com/questions/59693411/r-boxplot-draw-lines-between-each-subject-in-case-of-repeated-measurements)

**Figure 5:**
- As with Figure 2 – can you add the test statistic and regression or correlation line to this graph? Also, please add the statistic to Figure 7.

**Is the work clearly and accurately presented and does it cite the current literature?**
Yes

**Is the study design appropriate and does the work have academic merit?**
Yes

**Are sufficient details of methods and analysis provided to allow replication by others?**
Yes

**If applicable, is the statistical analysis and its interpretation appropriate?**
I cannot comment. A qualified statistician is required.

**Are all the source data underlying the results available to ensure full reproducibility?**
Yes

**Are the conclusions drawn adequately supported by the results?**
Yes

**Competing Interests:** No competing interests were disclosed.

**Reviewer Expertise:** Host-pathogen interactions, virology, virus ecology, virus evolution, eco-immunology

I confirm that I have read this submission and believe that I have an appropriate level of expertise to confirm that it is of an acceptable scientific standard, however I have significant reservations, as outlined above.