

Darkling Beetles (*Alphitobius diaperinus*) and Their Larvae as Potential Vectors for the Transfer of *Campylobacter jejuni* and *Salmonella enterica* Serovar Paratyphi B Variant Java between Successive Broiler Flocks[∇]

Wilma C. Hazeleger,^{1*} Nico M. Bolder,² Rijkelt R. Beumer,¹ and Wilma F. Jacobs-Reitsma^{2†}
Laboratory of Food Microbiology, Wageningen University, Wageningen, The Netherlands,¹ and Animal Sciences Group, Wageningen University and Research Centre, Lelystad, The Netherlands²

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Broiler flocks often become infected with *Campylobacter* and *Salmonella*, and the exact contamination routes are still not fully understood. Insects like darkling beetles and their larvae may play a role in transfer of the pathogens between consecutive cycles. In this study, several groups of beetles and their larvae were artificially contaminated with a mixture of *Salmonella enterica* serovar Paratyphi B Variant Java and three *C. jejuni* strains and kept for different time intervals before they were fed to individually housed chicks. Most inoculated insects were positive for *Salmonella* and *Campylobacter* just before they were fed to the chicks. However, *Campylobacter* could not be isolated from insects that were kept for 1 week before they were used to mimic an empty week between rearing cycles. All broilers fed insects that were inoculated with pathogens on the day of feeding showed colonization with *Campylobacter* and *Salmonella* at levels of 50 to 100%. Transfer of both pathogens by groups of insects that were kept for 1 week before feeding to the chicks was also observed, but at lower levels. Naturally contaminated insects that were collected at a commercial broiler farm colonized broilers at low levels as well. In conclusion, the fact that *Salmonella* and *Campylobacter* can be transmitted via beetles and their larvae to flocks in successive rearing cycles indicates that there should be intensive control programs for exclusion of these insects from broiler houses.

Salmonella and *Campylobacter* are responsible for many cases of human food-borne disease. Many of these cases can be related to the handling or consumption of contaminated chicken meat, and these pathogens occur frequently in broiler husbandry. Due to the introduction of pathogen control programs in The Netherlands, the prevalence of *Salmonella*-positive broiler flocks (cecal carriage) has been reduced from 22% in 1997 to approximately 5% in 2006. At the retail level, this resulted in a contamination rate of about 8% for poultry meat in 2006. For *Campylobacter*, however, the prevalence at the cecal level remained quite stable at 30% of the broiler flocks positive, leading to a prevalence of 16% for poultry meat products in 2006 (38).

So far, how broilers become infected with these microorganisms is not fully understood, but it is assumed that there are horizontal transmission routes with multiple sources of infection (40). Insects like flies, beetles, and larvae are some of the potential sources, as these animals are reported to be frequent carriers of pathogens such as *Escherichia coli*, *Shigella*, *Salmonella*, and *Campylobacter* around the world, especially near animal-rearing facilities (3, 13, 14, 25, 26, 27, 29, 36, 42). The insects that frequently occur in poultry houses include *Alphitobius diaperinus*, the darkling beetle, and its larvae, the lesser

mealworm (19, 30, 34). These insects are persistent in poultry houses and can be carriers of zoonotic bacteria, such as *Salmonella* and *Campylobacter*, and viruses (10, 27, 41), and they are considered to be a risk factor for *Campylobacter* contamination in broilers (28). Beetles and their larvae can survive in empty broiler houses between rearing cycles and may cause damage to the building's infrastructure by eating their way into insulation material in ceilings and walls, or they may hide under floors and in cracks and joints. When litter is taken out of the houses, the beetles may spread into the neighborhood (20).

It is not clear whether darkling beetles can be a vector for transmission of *Salmonella* and *Campylobacter* in consecutive broiler rearing cycles. The time spans between cycles are important for pathogen survival in these insects. These time spans may be different in different countries; e.g., in The Netherlands new rotations start after 1 week, but the time span is up to 6 weeks in Scandinavian countries (6, 32). Workers have described contradictory results regarding the survival of pathogens in insects, and the survival time ranges from 1 to 20 days (2, 16, 35, 37). In a study of broiler houses (32), *Salmonella* survived in beetles in the period (up to 2 weeks) between rearing cycles in broiler houses; however, during the same period no *Campylobacter* was found in beetles even after a first rotation with positive chickens. On the other hand, this could be explained by the fact that stressed *Campylobacter* cells may not grow on selective media. In the case of a less selective growth environment, such as the intestines of young chickens, sublethally damaged bacteria may still grow and colonize the chickens (9).

* Corresponding author. Mailing address: Wageningen UR, Agrotechnology & Food Science, Laboratory of Food Microbiology, P.O. Box 8129, 6700 EV Wageningen, The Netherlands. Phone: 31-317-482887. Fax: 31-317-484978. E-mail: wilma.hazeleger@wur.nl.

† Present address: RIKILT Institute of Food Safety, Wageningen University and Research Centre, Wageningen, The Netherlands.

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TABLE 1. Isolation of *Campylobacter* and *Salmonella* from beetles and their larvae at the start of the chicken feeding experiment and total numbers of chickens positive for *Campylobacter* and *Salmonella* at any time

Groups ^a	Treatment	Isolation of pathogens from beetles and larvae				No. of positive chickens/total no. tested			
		<i>Campylobacter</i>		<i>Salmonella</i>		<i>Campylobacter</i>		<i>Salmonella</i>	
		Beetles	Larvae	Beetles	Larvae	Beetles	Larvae	Beetles	Larvae
A-b and A-l	Beetles and larvae, respectively, inoculated on the day of feeding to chicks	+	+	+	+	11/12	10/12	12/12	10/12
B-b and B-l	Beetles and larvae, respectively, inoculated daily for 4 wk but not 1 wk prior to feeding to chicks	–	–	+	+	1/15	2/15	3/15	4/15
C-b and C-l	Beetles and larvae, respectively, daily inoculated for 5 wk prior to feeding to chicks	+ ^b	+ ^b	+	+	13/15	8/14	15/15	7/14
D-b and D-l	Beetles and larvae, respectively, inoculated daily for 5 to 6 wk until feeding to chicks and fed daily to chicks for 7 days	+ ^b	+ ^b	+	+	13/15	15/15	14/15	14/15
E-b and E-l	Beetles and larvae, respectively, collected at a commercial broiler farm 2 days before feeding to chicks	–	+ ^b	+	+	1/15	1/16	3/15	1/16

^a b, beetles; l, larvae.

^b Positive only after enrichment.

Therefore, in this study, different groups of darkling beetles (*A. diaperinus*) and their larvae that were either artificially contaminated with *Salmonella* and *Campylobacter* or caught in commercial broiler houses that were regularly positive for both pathogens were fed to young chickens to examine the possibility that the insects were a vector for pathogen transmission. Cecal droppings of the chickens were examined for the presence of the pathogens regularly, and isolated strains were characterized by antibiotic resistance testing (*Salmonella*) or amplified fragment length polymorphism (AFLP) typing (*Campylobacter*).

MATERIALS AND METHODS

Bacterial strains. Stock cultures of *Campylobacter jejuni* strains C224 (isolated from darkling beetles), C356 (isolated from broilers), and C81116 (often used in colonization experiments) and *Salmonella enterica* serovar Paratyphi B variant Java (nalidixic acid-resistant strain) were kept in cryovials with beads in heart infusion broth supplemented with 20% (vol/vol) glycerol at –80°C.

Insects. *A. diaperinus* beetles and their larvae were purchased from a commercial supplier (Kreca, Ermelo, The Netherlands) or were collected from a commercial broiler farm in The Netherlands known to commonly have *Salmonella*- and *Campylobacter*-positive broiler flocks. The insects were kept in glass containers and were fed rat feed and water.

Inoculation of insects with bacteria. Equal amounts of freshly prepared cultures of the *Salmonella* strain and the three *Campylobacter* strains in heart infusion broth were mixed, and the resulting bacterial suspension (containing about 10⁷ CFU/ml of *Salmonella* and 10⁸ CFU/ml of *Campylobacter*) was fed as droplets to the beetles and their larvae. The amount of the cocktail consumed was at least 1 µl, resulting in infection levels of 10⁴ to 10⁵ CFU of each of the pathogens per insect. Five groups of beetles and their larvae were used (Table 1): (i) insects inoculated with pathogens on the day of feeding to the chicks, simulating incidental contamination (groups A-b and A-l); (ii) insects inoculated daily for 4 weeks but not 1 week prior to feeding to the chicks, simulating the Dutch situation in broiler houses between rearing cycles (groups B-b and B-l); (iii) insects inoculated daily for 5 weeks until they were fed to the chicks, simulating a contaminated broiler house during rearing (groups C-b and C-l); (iv) insects inoculated daily for 5 to 6 weeks until they were fed to the chicks daily for 7 days, simulating repeated exposure of the chicks to contaminated insects during rearing (groups D-b and D-l); and (v) insects collected at a commercial broiler farm

2 days before feeding to the chicks to study possibly naturally contaminated insects (groups E-b and E-l).

***Salmonella* and *Campylobacter* isolation.** The presence of pathogens in the insects was checked by crushing the insects in a mortar and subsequent specific enrichment and/or plating using Preston broth (Oxoid CM067, SR0117, and 5% lysed horse blood) or cefoperazone charcoal desoxycholate broth and agar (Oxoid CM0963 and CM0739, each with SR0155) for *Campylobacter* and using buffered peptone water (Oxoid CM0509), brilliant green agar (Oxoid CM0329 with 100 mg/liter nalidixic acid), or modified semisolid Rappaport-Vassiliadis agar (Oxoid CM0910 with SR0161 incubated for 24 h at 41.5°C) for *Salmonella*. *Campylobacter* preparations were incubated microaerobically at 42°C for 2 days, and *Salmonella* preparations were incubated for 1 day at 37°C unless indicated otherwise. The identity of *Campylobacter* was confirmed microscopically and serologically (Microscreen *Campylobacter*; Microgen Bioproducts Ltd., Camberley, United Kingdom). The identity of *Salmonella* was confirmed by using the standard biochemical test and serological agglutination (Pro-Lab Diagnostics, BioTrading, Mijdrecht, The Netherlands).

Chicken experiments. A previous feeding trial showed that, once accustomed to the insects, chicks ate the beetles and their larvae eagerly. One-day-old chicks obtained from a commercial *Salmonella*-free breeder flock were placed on a litter floor for 5 days. On day 5, broilers were individually transferred to battery cages that were separated from each other by one empty cage and then had ad libitum access to feed and water in a strictly controlled broiler facility. Upon arrival, chicks were screened for the presence of *Salmonella* and *Campylobacter* by testing the paper liners of the chicken transport crates. Furthermore, eight pooled samples were taken from fecal droppings on day 7 just before insect feeding started, and all samples were negative. At the age of 7 days, each broiler was fed three beetles or three larvae after feed deprivation for 2 h. The number of chicks in groups A-b and A-l was 12, the number of chicks in group C-l was 14, the number of chicks in group E-l was 16, and all other groups contained 15 chicks. The broilers that received insects belonging to the different groups were distributed randomly in the cages.

Sampling of the chickens. Trays covered with clean paper sheets were placed below the cages in order to sample the cecal (or, if not present, fecal) droppings since the cecum is reported to be the primary site of colonization (1, 5). After each sampling, trays were cleaned and new paper was applied. Swab samples were taken from the droppings of all chickens on days 1, 2, 5, 7, 9, and 12 after the first insects were fed to the chickens. Chickens were sacrificed after 14 days, ceca were removed, and swab samples of the contents were taken. All swabs were streaked on cefoperazone charcoal desoxycholate agar and subsequently mixed with buffered peptone water, incubated for 16 to 20 h at 37°C, and streaked onto brilliant green agar. The incubation and confirmation procedures used were the

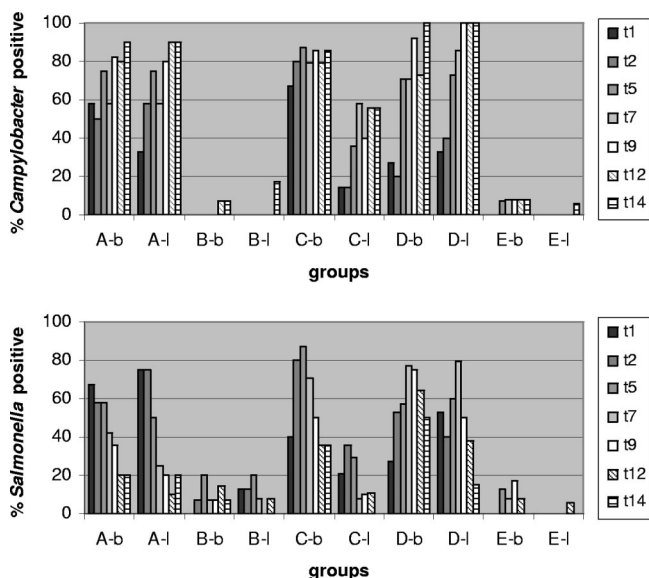


FIG. 1. Percentages of *Campylobacter*-positive (upper panel) and *Salmonella*-positive (lower panel) chickens, arranged by feeding group. The different bars for each group indicate the results for samples taken 1, 2, 5, 7, 9, 12, and 14 days after the insects were fed to the chickens. For a description of the groups see Table 1.

procedures described above. Isolated strains were collected and kept at -80°C until they were characterized further.

Antibiotic resistance. A random selection of the isolated *Salmonella* strains was tested for antibiotic resistance as described previously (18). In brief, *Salmonella* (10^8 CFU) was plated on Mueller-Hinton agar plates, and four antibiotic-containing filter disks (Neo-sensitabs; Rosco Diagnostics A/S, Taastrup, Denmark) were applied to each plate. The plates were incubated for 24 h at 37°C , and inhibition zones were measured. The antibiotics used were ampicillin (33 μg per disk), nalidixic acid (130 μg), amoxicillin (30 μg), gentamicin (40 μg), ciprofloxacin (5 μg), doxycycline (80 μg), trimethoprim (5.2 μg), spectinomycin (200 μg), norfloxacin (40 μg), chloramphenicol (60 μg), enrofloxacin (10 μg), flumequinone (30 μg), streptomycin (100 μg), and tetracycline (80 μg).

AFLP typing and data processing. A range of the *Campylobacter* strains isolated from beetles, larvae, and chicken ceca and the three strains used for inoculation were typed using AFLP genotyping, essentially as described previously (12). In short, chromosomal DNA was digested with HindIII and HhaI and ligated to restriction site-specific adapters. A preselective PCR and then a selective PCR were performed using a D4-labeled HindIII selective primer (Biologio, Malden, The Netherlands). The final products were mixed with an internal standard (Size Standard-600; Beckman) and separated using a CEQ 8000 capillary sequencer (Beckman). Data in SCF 3.00 format were imported into BioNumerics 3.5 (Applied Maths, Sint-Martens-Latem, Belgium). An AFLP analysis was performed for fragments ranging from 60 to 500 nucleotides long using BioNumerics 3.5. The similarity of patterns was calculated using the Pearson product-moment correlation coefficient. The unweighted pair group method using arithmetic averages and 1% optimization for position tolerance was used for cluster analysis.

RESULTS

Insects ($n = 50$) from all groups were examined for the presence of the pathogens just before they were fed to the chickens on day 0, and all of them were positive for *Salmonella*. Most groups were positive for *Campylobacter* as well; however, this pathogen could not be isolated from group B-b, B-l, and E-b insects (Table 1). For groups C, D, and E-l, *Campylobacter* was found only after enrichment and not by direct plating.

All chicken groups fed insects that were inoculated with the pathogens on the day of feeding (groups A, C, and D) showed

colonization with *Campylobacter* and *Salmonella* at levels of 50 to 100% (Table 1). Insects from groups B and E, however, colonized chickens at levels of 10 and 25% for *Campylobacter* and *Salmonella*, respectively.

In general, with groups A, C, and D the number of *Campylobacter*-carrying chicks increased during the first 5 to 9 days after feeding and remained stable at a high value until the end of the experiment (Fig. 1). In contrast, the colonization percentages for *Salmonella* were initially high and decreased gradually toward the end of the experiment most clearly for groups A, C, and D. *Salmonella*-positive birds fed group B insects were found from the beginning of the experiment, but *Campylobacter* colonization for this group was found only after 12 to 14 days. With group E (insects collected from a commercial broiler farm), low numbers of chicks were colonized with *Campylobacter* and *Salmonella* from day 5 onward.

The AFLP types of *Campylobacter* strains C81116, C224, and C356 used for inoculation were designated types R, S, and T, respectively. Seven of the strains isolated from the insects directly were identical to the type R strain, and the type of one strain was designated type T2 (similar to type T with minor differences). A total of 31 strains isolated from chickens were typed; type R was the predominant type found, and some type T-related types were observed as well (Table 2). Furthermore, a new AFLP type was identified, which was designated type P. No type S strains were recovered from insects or chickens.

A total of 19 *Salmonella* isolates from insects and chickens were characterized further, and all strains exhibited antibiotic resistance patterns similar to those of the *Salmonella* variant Java strain used for inoculation (data not shown).

DISCUSSION

The easy transfer of *Salmonella* and *Campylobacter* from beetles and their larvae to chickens in groups A, C, and D confirmed that these insects may have a role as vectors in one rearing cycle. There were no major differences between incidental contact (groups A and C) and repeated exposure to the inoculated insects (group D), except for group C larvae, which

TABLE 2. AFLP typing results for *Campylobacter* strains used for inoculation of the insects and for isolates from beetles, larvae, and chickens

Strain or source	Group	No. of strains for AFLP type:						
		R	S	T	T1	T2	T3	P
C81116		1						
C224			1					
C356				1				
Beetles		4				1		
Larvae		3						
Chicken	A-b	2						
Chicken	A-l	5						
Chicken	B-b	1						
Chicken	B-l				1			1
Chicken	C-b	5					1	1
Chicken	C-l	1					1	
Chicken	D-b	7						
Chicken	D-l	1			2			
Chicken	E-b	1						
Chicken	E-l	1						

showed an inexplicably lower level of transfer of both pathogens. Apparently, just a single exposure of chicks to contaminated insects may be sufficient for colonization of the intestines, which was also observed for *Campylobacter* in a previous study (35). In general, no differences between beetles and their larvae were observed, indicating that the two types of insects are equally important in transmission of the pathogens.

In The Netherlands, an empty 1-week period between rearing cycles is a common practice to clean out broiler houses. Removal of the litter results in removal of many of the insects, but some insects remain and hide in insulation material or in crevices in the building material or beneath the floor. Transmission to the next rearing cycle occurs if the microorganisms grow in the insects or just survive, which is most likely the case for *Campylobacter* due to its high minimal growth temperature (30°C). The empty week between rearing cycles was mimicked with group B, where inoculated beetles and their larvae were isolated for 1 week before they were fed to the chicks. Just before feeding, all insects were positive for *Salmonella*, but *Campylobacter* could not be isolated from group B insects. The rapid decrease in the number of campylobacters in insects may be explained by normal die-off kinetics due to temperature or humidity levels, which are season dependent, and also by the presence of antimicrobials in insects, which has been described for flies and unicorn beetles (17, 21). However, although the levels of colonization were low, with group B transfer of both pathogens to the chicks was observed. Stressed *Campylobacter* cells may not grow on selective media under these conditions (9), but in a less selective growth environment, such as the intestines of young chickens, sublethally damaged bacteria may still grow and colonize the birds. Slow recovery of damaged campylobacters could explain the fact that these pathogens were isolated from the chickens only at the end of the trial. The possibility of cross-contamination between different birds cannot be completely excluded, but cross-contamination was unlikely since empty cages were placed between the separate chickens and no direct contact was possible. Even though the numbers of positive chicks fed groups B and E insects were low, one positive chicken in a broiler house soon results in most chickens becoming colonized quite quickly and stably until slaughter (19). Furthermore, once a chick was found to be *Campylobacter* positive, it was also found to be positive postmortem, which also confirms that there was stable colonization.

On the other hand, shedding of *Salmonella* by the birds was intermittent and decreased toward the end of the experiment, similar to results described previously (23). *Salmonella* was isolated in all broiler groups throughout the trial, indicating that *A. diaperinus* can indeed be a vector for this organism in broiler houses, even after an empty week between rearing cycles. This is in accordance with the results of a previous study (32), in which similar genotypes of *S. enterica* serovar Indiana were found in broilers in two successive cycles and in beetles in a 2-week empty period between flocks. *Salmonella* variant Java was chosen for this study since this pathogen is known to be a cause of gastroenteritis (8) and it is an increasingly common organism found in broiler houses in The Netherlands (39), which was confirmed by the fact that it was found in insects from the commercial broiler farm (group E).

Some chickens carried one of the pathogens, but concomitant infection was also observed frequently for all groups (data

not shown), which is consistent with previous data showing no differences in single-colonization and cocolonization levels in chicks for both *Salmonella* and *Campylobacter* (33).

AFLP typing of the *Campylobacter* strains showed that mainly type R was isolated from beetles, larvae, and chickens, which indicates that strain C81116 either survives best in insects or colonizes chickens best or both. C81116 showed enhanced colonization of chickens after a single passage in vivo, which may be an explanation for the rapid spread of *Campylobacter* in broiler houses once a chicken is infected (7). Furthermore, some type T-related strains were also isolated, indicating colonization with strain C356. Compared to this original strain, there were minor differences in the genotypes, which is not uncommon since it is known that *Campylobacter* is genetically unstable and intragenomic alterations may occur, especially after chicken passage (11). Type S was not recovered, which is remarkable since strain C224 was originally isolated from beetles, which suggests that there would be some level of survival in the insects. However, considering the fact that not all isolates were typed and the fact that only one colony per plate was isolated, some types might have been missed, especially if low numbers were present. A new AFLP type (type P) was also found in the cecum of a group C-b chicken, which could be explained by the genetic rearrangement within or between the inoculated strains mentioned above. A great variety of *Campylobacter* genotypes in broiler flocks and less variation in beetles were reported previously, which was explained by preferential survival of certain clones in beetles or by different levels of recovery via enrichment (beetles) and direct plating (chickens) (4).

Type R was also isolated from group E-b and E-l chickens, which was unexpected, since these groups were fed insects that were not artificially contaminated but were caught at a commercial broiler farm. Cross-contamination with the insects from the other groups was not likely, since even at day 0 group E-l larvae were shown to carry a strain whose type was similar to type R, indicating that there was coincidental resemblance to C81116. This is plausible since this type was previously isolated from the majority of the samples in a human outbreak and chicken isolates from geographically distant sources were type R (24).

In conclusion, the finding that *Salmonella* and, to a lesser extent, *Campylobacter* may be transmitted via beetles or their larvae to successive rearing cycles indicates that there should be intensive control programs for elimination of insects from broiler houses. Depending on the type of house, controlling insect populations in broiler houses with physical barriers (15) and insecticides is possible (31), but only when carefully planned strategies are used (22). Another option might be to increase the time between rearing cycles, but, at least in countries where this type of husbandry is used, this is not economically feasible, especially in the case of *Salmonella*, which might survive for several weeks. Such measures should contribute to reducing the levels of *Campylobacter* and *Salmonella* in poultry meat products and thereby improve public health.

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