ABSTRACT

Over the years, *Salmonella* Heidelberg (SH) has gained prominence in North America poultry production and in the poultry production of other countries. *Salmonella* Heidelberg has been isolated and reported from poultry and poultry products in Brazil since 1962, whereas *Salmonella* Enteritidis (SE) has only emerged as a serious problem in poultry and public health since 1993. These strains of *Salmonella* can cause intestinal problems in newly hatched chicks, and infection may persist until adulthood. Upon slaughter of chickens, *Salmonella* can contaminate carcasses, a condition that poses a threat to human health. The aim of this study was to compare the fecal excretion of *Salmonella* Enteritidis and *Salmonella* Heidelberg in newly hatched chicks (orally inoculated with $10^5$ UFC/mL each) until 20 days of age. In addition, the ratio of cecal villus height: crypt depth (morphometry) and liver and cecum cell counts was analyzed in chicks ranging from 0 to 3 days of age and infected with these two *Salmonella* strains. One hundred seventeen chicks were separated into one of three experimental groups: a control group, an SE-infected group and an SH-infected group. Eight chicks per group were euthanized at 6, 12 and 72 hours post-inoculation (pi) to allow for *Salmonella* isolation from the liver and cecum and for the collection of the cecum for villi and crypt analysis. Other birds were allowed to mature to 20 days of age and cloacal swabs were taken at 2, 6, 13 and 20 days pi to compare the fecal excretion of inoculated strains. The *Salmonella* Enteritidis group had a higher number of cells excreted during the trial. Both strains were isolated from the liver and cecum by 6h pi. At 12h pi the *Salmonella* Heidelberg group had high cell counts in the cecum. No difference was found in liver cell counts. Both strains showed lower villus height: crypt depth ratio than the control group post-infection.

Key words: *Salmonella* Heidelberg, *Salmonella* Enteritidis, Fecal Excretion, villus: crypt ratio, Broiler
INTRODUCTION

Many strains of Salmonella are zoonotic agents, infecting humans through contaminated animal origin food products. In humans, salmonellosis is one of the most common causes of food poisoning. The most common serovars causing disease in humans are Salmonella Enteritidis and Salmonella Typhimurium. In recent years, Salmonella Heidelberg has gained prominence in North America and has also been found in the poultry production of other countries. Salmonella is commercially important in laying hens, turkeys and broiler chickens because of increase in meat and eggs production (3, 9, 20). Salmonella Heidelberg has been isolated and reported from poultry and poultry products in Brazil since 1962 (14), while Salmonella Enteritidis emerged as a serious problem in poultry and public health in 1993. The growth of the Brazilian poultry industry in the 1990’s and the increasing number of birds reared in high density environments has created favorable conditions for the maintenance of Salmonella positive flocks (24). The control of Salmonellae in Brazil is regulated by the Poultry National Health Plan developed by the Ministry of Agriculture, Livestock and Food Supply. In 2003, a vaccination for broiler breeders using killed vaccine against Salmonella Enteritidis was approved (23). Some authors have suggested that the presence of one serovar in a flock will reduce the accumulation of others; e.g., such as the situation with the reduction of Gallinarum and the increase of Enteritidis, or the increased presence of Enteritidis displacing other serovars (24). In broiler production, when cecal tonsil colonization is established by Salmonellae bacteria are consistently excreted in the feces (1). Salmonella Heidelberg is present in Brazilian broiler flocks and in an effort to better understand the behavior of this strain, this study was performed to compare the intestinal and liver presence of Salmonella Heidelberg and ATCC Salmonella Enteritidis strains following chick inoculation.

MATERIALS AND METHODS

Chicks: Recently hatched commercial Ross chicks were obtained from a Brazilian poultry company. All chicks were Salmonella free, hatched from breeders, unvaccinated against Salmonellae and were housed in battery cages. All cages were placed in the same room in three different sectors, and management of the sectors was performed separately to avoid cross-contamination. Water and feed free from antibiotics and animal protein were given ad libitum, and the environmental temperature was set to ensure chick physiological comfort.

Bacteria: Salmonella Heidelberg (SH) was isolated from broiler carcasses in Southern, and Salmonella Enteritidis (SE) ATCC (American Type Culture Collection) 1980 was kindly provided by Dr. Paul Barrow (AFRC Institute for Animal Health, Berkshire, UK). The organisms were retrieved from frozen culture stocks and cultured overnight at 37ºC in brain heart infusion broth (BHI; Merck, Germany). The culture was streaked on xylose lysine deoxycholate agar (XLD, Difco, UK) and incubated at 37ºC for 24 hours. Five colony forming units (CFU) of each strain were selected and transferred to BHI and incubated at 37ºC for 24 hours. CFU was determined by tenfold dilutions in 0.1% buffered peptone water (BPW, Oxoid, UK) and streaked on XLD agar. The suspension with $10^5$ CFU/mL was selected for inoculation.

Experimental design: One hundred seventeen chicks were separated into three groups designated as Control, SE and SH. Two groups were inoculated orally with 1 mL containing $10^5$ CFU/mL of SE and SH, and another group was inoculated with saline solution as a control. From each group during the trial, 24 birds were killed, and 15 were kept for Salmonella fecal excretion analysis. All birds were individually numbered by a metal ring placed on the wing. Eight chicks per group were killed by cervical dislocation at 6, 12 and 72 hours post-inoculation (pi). At necropsy, the cecum and the liver were collected for Salmonella bacteriological enumeration. Ceca were collected 3 centimeters away from the colon and taken for histological examination and submission for electron microscopy analysis. Cloacal swabs of 15 birds from each group were collected for bacteriological examination at 2, 6, 13 and 20 days pi. At necropsy (21 days pi), cecum and liver were collected for bacteriological examination. The trial was
approved and performed according to the recommendations of the Faculty Ethics Committee of Veterinary Medicine, Federal University of Parana, Brazil.

**Bacterial identification in chick liver and cecum:** Two pools of four birds per group had their organs homogenized in 2% BPW (ten times the volume of organs by weight). BPW cultures were diluted tenfold in BPW up until a dilution of $10^{-3}$, followed by streaking in duplicate on XLD agar plates, incubation for 24 hours at 37°C and CFU counting, adapted from Desmidt et al. (6). All organ samples were also incubated for 16-18 hours at 37°C, and 100 µL from each BPW were inoculated in 10 mL of Rappaport Vassiliadis (RV; Merck, Germany) and placed at 42°C for 24 hours. The RV samples were streaked on XLD agar (24 hours at 37°C). *Salmonella* CFU were counted and the results expressed according to the Plate Count Proceedings recommended by Normative Instruction Nº6 from the Brazilian Ministry of Livestock and Food Supply (18). One typical colony per plate was biochemically and antigenically screened to confirm *Salmonella* identity.

**Bacteriological examination for cloacal swabs:** Some cloacal swabs were directly inoculated on XLD agar, and others were placed in 2 mL of 2% BPW. BPW cultures were diluted tenfold in BPW until a dilution of $10^{-3}$ was reached, and 100 µL of the diluted suspension was plated out in duplicate on XLD agar, adapted from Desmidt et al. (6). *Salmonella* colonies were counted, and the results were expressed according to the Plate Count Proceedings recommended by Normative Instruction Nº62 published August 26, 2003 and recommended by the Brazilian Ministry of Agriculture, Livestock and Food Supply (18). One typical colony per plate was biochemically and antigenically screened to confirm *Salmonella* identity.

**Histological examination and villus and crypt analysis:** Cecum samples were fixed in 4% phosphate-buffered formaldehyde, paraffin-embedded, sectioned at 5 µm and stained with hematoxylin and eosin (HE). The cecum samples processed for histological examination were measured for villi height and crypt depth. These data were taken from the tip of the villus to the valley between individual villi, and measurements for crypt depth were taken from the valley between individual villi to the baso-lateral membrane (Thompson, Applegate, 2006). Villi and crypts (50 of each) were measured for each time per group (21). The villus and crypt study was carried out using an image-analyzing system (Motic Images Plus 2.0-Motic China Group Co.2006), coupled to a microscope (Olympus BX41 Olympus America INC., NY, USA).

**Statistical analysis:** Data were submitted for analysis of variance using StatView SAS software (1992-98-SAS Institute Inc. USA) and means were compared using the Tukey test at a 5% significance level (P<0.05).

**RESULTS**

*Salmonella* isolation and villi height-crypt depth analysis. Quantification of *Salmonella* per gram of cecum and liver and the relationship between villi height and crypt depth were analyzed for 72 hours pi with two different *Salmonella* strains, serovars Enteritidis (SE) and Heidelberg (SH). The control group was not infected with *Salmonella*. At 6 hours pi, the two strains were present in the cecum and liver. SH had a higher CFU/g count in the cecum at 12 hours pi than did SE, but no
difference was observed in number of cells in the liver between the two strains (Table 1). For the villus height:crypt depth ratio, the SE and SH groups were significantly different from the control group (Table 2).

**Table 1.** Bacteriological counting of *Salmonella* (S) Heidelberg and *Salmonella* Enteritidis per gram of caeca and liver tissue from newly hatched chicks inoculated with $10^5$ CFU/mL. Data are the means of organs collected from eight animals at each time point.

<table>
<thead>
<tr>
<th>Time post - inoculation</th>
<th>Control cecum</th>
<th>S. Enteritidis cecum</th>
<th>S. Heidelberg cecum</th>
<th>Control liver</th>
<th>S. Enteritidis liver</th>
<th>S. Heidelberg liver</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>CFU/g</td>
<td>CFU/g log$_{10}$</td>
<td>CFU/g log$_{10}$</td>
<td>CFU/g</td>
<td>CFU/g log$_{10}$</td>
<td>CFU/g log$_{10}$</td>
</tr>
<tr>
<td>0h</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
</tr>
<tr>
<td>6h</td>
<td>-</td>
<td>3.22 ± 2.16</td>
<td>3.11 ± 2.32</td>
<td>-</td>
<td>2.20 ± 2.12</td>
<td>2.69 ± 2.81</td>
</tr>
<tr>
<td>12h</td>
<td>-</td>
<td>3.99 ± 0.06$^b*$</td>
<td>4.70 ± 0.03$^a$</td>
<td>-</td>
<td>0.69 ± 0.00</td>
<td>0.69 ± 0.00</td>
</tr>
<tr>
<td>72h</td>
<td>-</td>
<td>4.10 ± 0.64</td>
<td>4.37 ± 0.36</td>
<td>-</td>
<td>1.74 ± 0.29</td>
<td>1.08 ± 0.55</td>
</tr>
</tbody>
</table>

CFU/g: Colony Forming Units per gram. Original counts transformed into log$_{10}$.

$^*$Different letters in the same line denote means that are significantly different according to a Tukey test at a 95% level of confidence (p<0.05).

**Table 2.** Cecal villus height:crypt depth ratio (μm) measurements from cecal samples of chicks inoculated with *Salmonella* (S) Heidelberg and *Salmonella* Enteritidis at different time points post-inoculation.

<table>
<thead>
<tr>
<th>Time post-inoculation</th>
<th>Control Villus:crypt</th>
<th>S. Enteritidis Villus:crypt</th>
<th>S. Heidelberg Villus:crypt</th>
</tr>
</thead>
<tbody>
<tr>
<td>6h</td>
<td>$^a$5.21 ± 2.28</td>
<td>$^b$4.49 ± 1.62</td>
<td>$^b$5.09 ± 1.56</td>
</tr>
<tr>
<td>12h</td>
<td>$^b$6.59 ± 9.44</td>
<td>$^b$4.28 ± 1.10</td>
<td>$^b$5.21 ± 1.87</td>
</tr>
<tr>
<td>72h</td>
<td>$^a$7.62 ± 2.36</td>
<td>$^b$5.63 ± 1.89</td>
<td>$^a$7.14 ± 2.20</td>
</tr>
</tbody>
</table>

**Salmonella Enteritidis and Salmonella Heidelberg Fecal Excretion:** The SH strain at 2 days pi showed a higher cell count than the SE strain, which could only be counted after selective enrichment. At 6 days pi, the strains showed no difference in the number of cells excreted in the cloacal swabs. 6 days pi, the SE strain showed higher levels of excretion than the SH strain (Table 3 and Figure 4). From the direct agar plate (DAP), it was possible to determine that the SH group had a high number of positive birds at 2 days pi, whereas the SH positive birds showed a gradual increase in total numbers according to the DAP. Due to the individual identification of birds in the groups, it was possible to observe that all birds were positive for *Salmonella* at some point during the experiment. At the necropsy performed 21 days pi, all birds were found to be positive for SH or SE in the liver, while the cecum bacterial counts were higher in the SH group than in the SE group. Means and standard deviation to bacterial counts into the cecum were 1.34 ± 1.37 for SE and 2.96 ± 1.52 for SH, with p<0.05 between the two groups. Data obtained from analyses of the cecal contents did not correlate with the results from the last cloacal swab taken one day before euthanasia.

**Electron microscopy:** Figure 1 shows the cecum scanning photomicroscopy of birds from the control group, and some injured areas are apparent. Figures 2 and 3 from birds infected with *Salmonella* Heidelberg ($10^5$ CFU/mL) show villi with aspects of edema, rough aspect, mucosal erosion, degenerate cells and attached bacteria.
Table 3. Percentage of positive samples in direct agar plating (DAP) and fecal excretion counting of *Salmonella* Enteritidis (SE) and Heidelberg (SH) ($10^5$CFU/mL). Original number transformed in log_{10}.

<table>
<thead>
<tr>
<th>Group</th>
<th>DAP+ (%) 2 days pi</th>
<th>Excretion (log_{10}) 2 days pi</th>
<th>DAP+ (%) Excretion (log_{10}) 6 days pi</th>
<th>DAP+ (%) Excretion (log_{10}) 13 days pi</th>
<th>Excretion (log_{10}) DAP+ (%) 20 days pi</th>
</tr>
</thead>
<tbody>
<tr>
<td>Control</td>
<td>0</td>
<td>-</td>
<td>0</td>
<td>-</td>
<td>0</td>
</tr>
<tr>
<td>SE</td>
<td>0</td>
<td>0.70±0.00b</td>
<td>20</td>
<td>1.23±1.12</td>
<td>73</td>
</tr>
<tr>
<td>SH</td>
<td>53</td>
<td>1.52±0.13a</td>
<td>13</td>
<td>0.93±0.61</td>
<td>20</td>
</tr>
</tbody>
</table>

P value: <0.01 = 0.36 <0.01 <0.0001

* Different letters in the same column denote means that are significantly different according to a Tukey test with a 5% probability of error (p<0.05). pi: post-inoculation; DAP+: direct agar plating positive; pi: post-inoculation.

**Figure 1.** Electron microphotograph of the cecum of chicks from the Control group, 3 days post-inoculation with a saline solution (X100). In the circles, the apical region of some villi with injured areas.

**Figure 2.** Electron microphotograph of the cecum of a chick at 3 days post-inoculation with $10^5$ CFU/mL of *S. Heidelberg* (X100). Villi have an aspect of edema and surfaces with rough aspect or mucosal erosion. Space between the villi shows structures similar to bacilli and cellular debris.
DISCUSSION

Previous studies performed with *Salmonella* Enteritidis in different broiler lines have demonstrated an inverse relationship between the severity of cecal infection and the colonization of systemic organs in animals of similar ages and challenge (16, 22). In the present study, we also found cell counts from the cecum to be higher than in the liver for chicks infected with SE and SH (Table 1). The different cecum and liver colonization levels may be due to *Salmonella* colonization virulence systems that are based on two *Salmonella* pathogenic island systems (SPI 1 and 2) responsible for intestinal (SPI 1) and systemic (SPI 2) host infection. The expression of these island systems are regulated in response to different environmental factors. For example, SPI 1 responds to low oxygen levels and high osmolarity in the gut lumen, whereas SPI 2 responds to low magnesium and phosphorus levels inside host cells (4).

According to Nabburs (19), under optimal circumstances, intestinal villi should be high and crypts should be shallow, resulting in a high villus height to crypt depth ratio. In the present work, from 0 to 3 days pi, SE and SH groups had low villus height:crypt depth ratios when compared to the control group, possibly because of SE and SH damage caused to the...
intestinal mucosa. Reductions in the villi height:crypt depth ratios can occur due to intestinal pathogen infection and are seen under other certain conditions in poultry, including after feed withdrawal (25) and in response to some feed additives that are used during growout (12, 8, 21). These changes have been attributed to reduced cell migration and proliferation rates, along with increased rates of cell loss and programmed cell death (7).

Intestinal mucosa alterations found in this work may have occurred because immediately after hatching, chicks have an immature immune system, and the protection of intestinal mucosa depends on barrier cells (macrophages, natural killer cells, heterophils), chemical and physical factors, and the presence of Paneth cells. Paneth cells are located in the crypt and secrete defensins and phospholipase that lyses bacteria, fungi and enveloped viruses (15).

Infection of newly hatched chicks with Salmonella Enteritidis by oral inoculation or by horizontal contact exposure can lead to the establishment of intestinal colonization that persists into adulthood (10) because of the ability of this pathogen to replicate inside host cells, most prominently in phagocytic cells such as macrophages (13). In our study, SE and SH strains were inoculated in chicks, and cloacal swabs were taken until 20 days of age. As with previous trials utilizing these two strains (2), the birds infected with SE excreted a very low Salmonellae number at 2 days pi (Table 3).

Initially, the SH group had the highest bacterial counts from the cloacal swabs, but this number decreased over time until 20 days pi. In contrast, birds with SE showed an increase in the number of positive birds and cell counts throughout the trial. Table 3 shows a comparison between isolation in direct agar plates (DAP) and the number of cells at each time point, showing that the number of birds DAP positive in the SE group increased along with the number of cells isolated. From the measurements, it could be determine that when the excretion level per bird was above 10 CFU/mL, it was detectable in DAP. When individual birds were identified it was also determined that all birds were positive at a minimum of one time point analyzed and often the same bird was positive for more than one time point.

Scanning photomicroscopy showed that the apical region of some villi was affected in the control group, which can be ascribed to natural cell turnover (Figure 1). Figures 2 and 3 show the changes in the intestinal villi of the SH group. Aspects of edema, rough surface, mucosal erosion, epithelial cell degeneration, and the presence of cellular debris in the space between villi were suggested by scanning photomicroscopy. These results are in agreement with Desmidt et al. (5), who observed adhesion and colonization of Salmonella in the cecal lumen.

The importance of Salmonella Enteritidis is well established in the poultry industry and human health. Similarly, in the last few years, the presence of Salmonella Heidelberg in laying hens producing contaminated eggs (7, 8, 9, 26) and as a cause of human illness (3) has been documented. The present work demonstrates that changes in the intestinal mucosa caused by Salmonella Heidelberg are similar to those caused by Salmonella Enteritidis; specifically, Salmonella is present in the liver of newly hatched chicks and persists in the fecal excretion of broiler chickens. These findings suggest that Salmonella Heidelberg may have importance as a pathogen to newly hatched chicks and is a potential broiler carcass contaminant.

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