



Identification of the infective dose of recovered multi-drug resistant non-typhoidal *Salmonella* spp. and the intestinal histomorphometric changes in broiler chicken

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ABSTRACT

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The study is aimed to determine the ID₅₀ of the two recovered *Salmonella* isolates from various broiler farms in Wayanad district, Kerala. The isolates were confirmed as *Salmonella* Typhimurium through cultural and molecular characterisation. On antibiotic susceptibility testing both the isolates were found to be resistant to more than six classes of antibiotics. Vencobb 430Y chicks (n=24) were equally distributed into four groups (T1, T2, T3 and T4) with six birds in each group. All the birds tested negative for *Salmonella* on the fifth day. On the seventh day, the birds of treatment T2, T3 and T4 were orally inoculated with 10⁸, 10⁹ and 10¹⁰ cfu/mL of MDR-S. Typhimurium. The birds were examined closely for clinical symptoms and changes in faecal matter consistency. The body weight of the birds was taken before infection and also on weekly basis post-infection. The clinical symptoms started showing from the fifth-day post-infection in T4 and the seventh day post-infection in T2 and T3. Fifty percentage of the birds from T4 exhibited progressive weakness with diarrhoea after the first-week post-infection without causing mortality. The body weight difference of birds from different treatment was non-significant during pre-infection and during the first and second week post-infection body weight was in the order T1>T2>T3>T4. The caecal *Salmonella* load was highest in T4 and it was found to increase with time. When the histomorphometry of the different segments of small intestine were compared, a significantly lower villi-to-crypt ratio was exhibited by T4, indicating impaired intestinal function. The ID₅₀ of recovered S. Typhimurium was established as 10¹⁰ cfu/mL based on the clinical findings, faecal matter consistency pattern, body weight variations and intestinal histomorphometry parameters.

Keywords: Broiler chicken, MDR-NTS; ID₅₀, Intestinal histomorphometry, *Salmonella*

INTRODUCTION

Poultry is one of the most crucial elements of the agricultural sector worldwide with a growth rate of 16.81 percent in its population in India alone from 2012 to 2019 (BAHS, 2022). The fastest-growing segment of worldwide meat production and export is poultrymeat (Aswathi *et al.*, 2013). Poultry, constituting approximately 50 percent of India's total meat production, stands as the predominant species among meat producers (Varunkumar *et al.*, 2022). The poultry sector is affected regularly by the introduction of bacterial infections. *Salmonella* infection is one of the most significant bacterial illnesses in chickens, resulting in significant economic loss due to death and decreased production (Moultotou *et al.*, 2017; Jubeda *et al.*, 2022). The genus *Salmonella* is a member of the bacterial family Enterobacteriaceae and consists of more than 2500 serologically distinguishable variants. Among them, non-typhoidal *Salmonella* serotypes (*S. enterica* subspecies *enterica* serovar Enteritidis and *S. enterica* subspecies *enterica* serovar Typhimurium) are virtually invariably present in both wild and domestic animals, are primarily of concern as a source of foodborne illness in humans

(Saif, 2009). Poultry products are frequently identified as important sources of *Salmonellae* that cause human illness, although most non-typhoidal *Salmonella* infections are associated with self-limiting gastroenteritis, they have the potential to cause fatal infections among infants, young children, older adults and immunocompromised individuals (Pandey and Saxena, 2015; Wen *et al.*, 2017; Balasubramanian *et al.*, 2019).

The infective Dose 50 (ID₅₀) is a quantitative measurement to describe the number of microorganisms or particles required to cause an infection in 50 percent of the individuals or experimental subjects exposed to the pathogen (Ramesh *et al.*, 2020). The infective dose of *Salmonella* in broiler chickens can be influenced by a range of factors including, strain variability, host susceptibility, gut microbiota composition, stress and many other factors (Cosby *et al.*, 2015). The present study was conducted to estimate the ID₅₀ of MDR-S. Typhimurium in broiler chicken of one week age.

MATERIALS AND METHODS

Confirmation of *Salmonella* isolates

The study utilised the two *Salmonella* isolates obtained from distinct broiler farms in Wayanad district, Kerala. Isolates were selectively plated on Xylose Lysine

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Deoxycholate (XLD) agar, and molecular confirmation was achieved through polymerase chain reaction employing the *invA* and *spy* genes (Nair et al., 2015). Antibiotic susceptibility test was done by disc diffusion method using commonly used antibiotic discs as per guidelines (Bauer et al., 1966). One of the multi-drug resistant non-typhoidal *Salmonella* isolates among the tested samples was used further in the study.

Experimental design

Day-old Vencobb-430 Y broiler chicks (n=24) were distributed equally on body weight (43 g) basis into four groups (T1-T4) each having 6 birds. All birds were fed commercial broiler chicken feed. On the fifth day, cloacal swabs were obtained from each group and tested for the existence of *Salmonella* by streaking onto XLD agar. The birds of treatments T2, T3 and T4 which tested negative for *Salmonella* were orally inoculated with multi-drug resistant *S. Typhimurium* at the concentration of 10^8 , 10^9 and 10^{10} cfu/ml respectively on the seventh day. Faecal matter consistency and other clinical changes were monitored on daily basis and cloacal swab examination was done on weekly basis. The caecal and liver samples were collected in sterile PBS broth to ascertain the existence of the organism by spread plate method on XLD agar plates supplemented with 100 µg of ampicillin. The body weights of each bird were measured at weekly intervals. All the birds were slaughtered humanely on the third week and the duodenal, jejunal and ileal segments were collected in 10 percent formalin for histomorphometric study.

Intestinal histomorphometry

The representative pieces of the duodenum, jejunum and ileum kept in 10 per cent neutral buffered formalin were used for histomorphometric examination. After seven days the 0.5 cm long tissue was cut from the duodenum, jejunum and ileum part and placed in a cassette and processing was done by standard tissue processing protocols (Luna, 1968). Sections were taken at 5µm thickness by using a semi-automatic M-TECH microtome. Haematoxylin and Eosin staining method was used for histological studies. The histomorphometric measurements were taken at 40x magnification for intestinal villus and crypt depth using image analysing software (MicapsMicroview).

RESULTS AND DISCUSSION

On cultural characterisation, red colonies with black centre confirmative for *Salmonella* were obtained and these isolates were tested positive for the *invA* and *spy* gene on PCR. On antibiotic susceptibility testing both the isolates were resistant to more than six classes of antibiotics and hence they were multidrug-resistant *S. Typhimurium*. The study reports published by Salehi et al. (2005), Alcaine et al. (2007), McDermott et al. (2016), Liljebjelke et al. (2017), Sharma et al. (2019),

Jayaweera et al. (2020), Gayathridevi et al. (2021) and Raghavendra (2022) also stated that the isolated NTS strains from broilers, carcass and poultry farm environment exhibited resistance to more than three classes of antibiotics. Michael and Schwarz (2016) reviewed the existence of resistance genes, mutations, and genetic modifications associated with *Salmonella* spp. leading to the emergence of MDR conditions.

The clinical symptoms started showing from the fifth day post-infection in T4 and the seventh day post-infection in T2 and T3. The clinical symptoms included dullness, depression and reduced feed intake. Fifty percentage of the birds from T4 exhibited progressive weakness with diarrhoea after first-week post-infection without causing mortality. The results obtained were in agreement with Dra et al. (2019), who reported the development of clinical symptoms from the first week in broiler chicken infected with *S. Typhimurium* at the dose of 2×10^8 cfu/ml. On caecal and liver sample examination the *Salmonella* load was found to be higher in T4, the birds that were infected with 10^{10} cfu/ml. Based on these findings and through the application of the probit regression model the ID_{50} for the recovered MDR-*Salmonella* spp. challenge was conclusively set at 1×10^{10} CFU/ml. Akil and Ahmad (2019) reported that the consumption of at least 1.46×10^4 CFU/g for *Salmonella enterica* serovar Enteritidis or 6.4×10^3 CFU/g for *S. Typhimurium* is required to develop an infection in 50 per cent of the population. The obtained infective dose 50 was different from the previous studies and that might be attributed to the changes in the *Salmonella*'s virulence pattern and other factors which affect the colonization of bacteria such as survival of *Salmonella* through the gastric barrier and genetic background of the chicks (Cosby et al., 2015).

From the third day post-infection, the body weight of birds up to the third week was analysed and a significant difference ($P < 0.001$) could be observed during the second and third weeks (Table 1). Significantly higher body weight observed in T1 and T4 exhibited lower body weight. The body weight of the different treatments after post-infection was as follows $T1 > T2 > T3 > T4$. *Salmonella* infection in poultry is linked to severe morbidity, malabsorption, slowed development, and ineffective feed utilisation, especially in young birds.

The villi height, crypt depth and villi-to-crypt ratio showed significant differences between the four treatments. In duodenum, villi height and crypt depth were significantly higher in T1 ($P < 0.001$) and villi to crypt ratio was significantly lower in T4 ($P < 0.001$). The villi height and villi to crypt depth were significantly higher in T1 ($P < 0.001$) and the crypt depth was significantly higher in T4. The ileal villi to crypt ratio was significantly higher in T1 and the villi height was lower in T4 ($P < 0.001$). The villi height to crypt depth

Table 1: Body weight upto third week (g)

Week	T1	T2	T3	T4	P-value
Week 1	211.50 ± 1.54	208.83 ± 0.60	209.83 ± 0.60	207.50 ± 0.76	0.051 ^{ns}
Week 2	321.33 ± 1.05 ^a	305.33 ± 1.11 ^b	290.83 ± 0.94 ^c	282.66 ± 0.66 ^d	< 0.001 ^{**}
Week 3	752.33 ± 0.98 ^a	730.33 ± 0.66 ^b	712.50 ± 0.99 ^c	706.33 ± 0.76 ^d	< 0.001 ^{**}

Table 2: Histomorphometric parameters of duodenum, jejunum and ileum

Group	Duodenum			Jejunum			Ileum		
	Villi height (µm)	Crypt depth (µm)	Villi: Crypt	Villi height (µm)	Crypt depth (µm)	Villi: Crypt	Villi height (µm)	Crypt depth (µm)	Villi: Crypt
T1	1398±14 ^a	164±3 ^a	8.51±0.17 ^b	833±8 ^a	110±5 ^b	7.68±0.41 ^a	738±17 ^{ab}	104.1±1.13 ^b	7.09±0.18 ^a
T2	788±16 ^c	109±2 ^c	7.22±0.22 ^c	565±16 ^c	110±1 ^b	5.15±0.17 ^b	765±14 ^a	130.2±6.60 ^a	3.08±0.14 ^c
T3	964±4 ^b	101±3 ^d	9.55±0.28 ^a	591±31 ^c	112±2 ^b	5.27±0.25 ^b	697±36 ^b	105.3±4.69 ^b	6.61±0.19 ^{ab}
T4	708±4 ^d	151±3 ^b	4.71 ± 0.07 ^d	702±15 ^b	131±6 ^a	5.37±0.20 ^b	398±5 ^c	129.0±10.25 ^a	6.06±0.42 ^b
P-value	<0.001	<0.001	<0.001	<0.001	0.004	<0.001	<0.001	0.014	<0.001

ratio is utilised as a predictor of the small intestine's potential capability for digestion. This ratio rises in proportion to increased digestion and absorption as per Montagne *et al.* (2003). Consistent with the present investigation, Sieo *et al.* (2005) observed that an enhanced intestinal villi height and an increased ratio of villi height and crypt depth indicate a larger surface area for nutrient absorption and improved absorption function. The findings are indicative of the impaired nutrient absorption and utilization in MDR-NTS infected birds especially in birds orally infected at a dose of 10¹⁰cfu/ml.

CONCLUSION

The present study investigated the ID₅₀ of recovered MDR-S. Typhimurium in broiler chicken. The infective dose of recovered MDR-S. Typhimurium is determined as 10¹⁰ cfu/mL as per clinical symptoms, faecal matter consistency and intestinal histomorphometry.

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