Molecular epidemiology of systemic *Salmonella enterica* serovar Typhimurium outbreak in canaries

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**ABSTRACT**

**Background and Objectives:** In May 2007, high mortality with severe septicemia was reported in 17 flocks of canaries in different regions of Tehran province. This study was designed to follow up and study a great outbreak of salmonellosis in these canaries.

**Materials and Methods:** Two carcasses from every flock, environment, food and water resources were examined. After isolating the bacteria, serotyping and multiplex PCR were performed to confirm the bacteria identified. The isolates within the same serovars were investigated by R-typing using 33 antibiotics and then subjected to RAPD-PCR with three primers. The genomic DNA from these isolates were digested with *Xba*I and the macro restriction fragment were separated by PFGE.

**Results:** *S. Typhimurium* was isolated from dead carcasses, visceral organs, stools and feed. Thirty-six isolated strains (35 isolates from canary carcasses and one isolate from feed) showed similar results in all of the tests, confirming the occurrence of an outbreak.

**Conclusion:** Canaries seem to be very susceptible to infection with *S. Typhimurium*. The clonality of isolated organisms and its characteristics is significantly important due to the severe septicemia and high mortalities in this outbreak and its public health threats. Environmental contamination within the cages, and food contaminated with stools of other canaries were the sources of infection. Inspection for food hygiene, daily cleaning of canary’s cage from stool and carrier insects and rodents are necessary to prevent such outbreaks. Combination of R-typing, RAPD-PCR and PFGE increase the differentiation power of isolates, however, they showed clonality of *S. Typhimurium* involved in this outbreak.

**Keywords:** *Salmonella* Typhimurium, PFGE, RAPD-PCR, R-typing, canary, Iran.

**INTRODUCTION**

*Salmonella* serovars are among the most frequent causes of bacterial infections in animals and human (1). *S. enterica* subsp *enterica* serovar Typhimurium is the most frequently isolated serovars and is one of the top three serotypes isolated worldwide over many decades from human, domestic and wild animals, feed, and the environment, with cattle and poultry being important source of infection for humans (2, 3). *Salmonella* can also be isolated from domestic birds, captive raptors, and other wild birds, especially gulls and corvids (4).

Effective public health monitoring of *Salmonella* infections by serotyping and/or other typing methods does interrupt transmission and reduce number of infections (4).

In May 2007, high mortality with severe septicemia was reported in 17 flocks of adult canaries in different regions of Tehran province. This study was performed to follow the infection, confirm an epidemic, test antibiotic susceptibility and fingerprinting characteristics of this highly virulent strain and finally interrupt transmission to humans.

**MATERIALS AND METHODS**

**Samples.** Carcasses of canaries were referred to the Department of Microbiology, Faculty of Veterinary Medicine, University of Tehran. Carcasses belonged to the seventeen flocks around the Molavi Street in downtown of Tehran. Number of canaries, number of deaths and mortality rate of every flock is shown in

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Table 1. Number of canaries and mortality rates in each flock in the outbreaks of salmonellosis.

<table>
<thead>
<tr>
<th>Number of flock</th>
<th>Number of canary in flock</th>
<th>Number of death in flock</th>
<th>Mortality rate (%)</th>
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</thead>
<tbody>
<tr>
<td>1</td>
<td>16</td>
<td>15</td>
<td>94</td>
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<tr>
<td>2</td>
<td>40</td>
<td>9</td>
<td>23</td>
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<td>3</td>
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<td>4</td>
<td>15</td>
<td>8</td>
<td>53</td>
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<td>5</td>
<td>26</td>
<td>16</td>
<td>62</td>
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<td>6</td>
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<td>4</td>
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<td>50</td>
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<td>9</td>
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<td>12</td>
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<td>18</td>
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<td>14</td>
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<tr>
<td>16</td>
<td>80</td>
<td>60</td>
<td>75</td>
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<tr>
<td>17</td>
<td>25</td>
<td>20</td>
<td>80</td>
</tr>
<tr>
<td><strong>Average</strong></td>
<td></td>
<td></td>
<td><strong>50.11%</strong></td>
</tr>
</tbody>
</table>

Primers selected for this study were P1254 5′-CCGCAGCCAAA-3′, 23L 5′-CCGAAGCTGC-3′ and OPA-4 5′-AATCGGGCTG-3′.

**PFGE.** Pulsed-field gel electrophoresis was performed according to the procedures developed by the CDC for molecular subtyping of Escherichia coli O157:H7, non-typhoidal Salmonella serovars and Shigella sonnei as previously described (10). Briefly, agarose-embedded DNA was digested with 50 U of XbaI (Fermentas, Vilnius, Lithuania) overnight in a water bath at 37°C. The restriction fragments were separated by electrophoresis in 0.5% TBE buffer at 14°C for 20 h in 6 V/cm using a CHEFF DR II electrophoresis system (Gene Navigator, Pharmacia, Sweden) with pulse times of 2.2 to 63.8 seconds. The gels were stained with ethidium bromide (1 μg/ml) and destained with the buffer remained in the electrophoresis apparatus for 60-90 min. A Gel Doc 2000 equipped with quantity one software (Bio-Rad, Hercules, CA) was used for image capture and conversion of gel images to Tiff file. Also isolates with DNA smear patterns were rechecked. The molecular weight standard used for all gels was XbaI-digested DNA from Salmonella Braenderup strain.
RESULTS

Pathology. Microscopic inspection revealed disseminated granulocyte and mononuclear leukocyte infiltration in liver and proventriculus. Macroscopic and microscopic examination indicated septicemia and focal necrosis with hepatitis in liver and proventriculitis. Mean mortality rate was calculated to be about 51% in the flocks.

Bacterial isolates and biotyping. Salmonella were isolated in biotyping by the conventional method in all of the 35 dead canaries aside from the feed. No bacterium was isolated from water sources.

Serotyping. Salmonella Typhimurium (4,5,12: i: 1, 2) resulted in serotyping while were tested by O:4, 5, 12; H; i and H; 1, 2 antisera (Difco, Detroit, MI, USA) in all of 35 dead canaries.

Multiplex PCR. Multiplex PCR confirmed the serotype of isolated Salmonella Typhimurium in all of 35 dead canaries. Results of multiplex PCR of some isolates are presented in fig. 1. Sizes of 183, 284, 526 and 663 bp indicate flIC, invA, fljB and rfbJ genes that represent H; i, invasion that indicates genus of Salmonella and virulence type, H; 1, 2 and O, respectively.

Antibiotic susceptibility testing. All of the isolated S. Typhimurium revealed the same R-type pattern. With the exceptions of sulphamethoxazole, tylosin, and cloxacillin, the isolates were susceptible to all antibiotics.

RAPD-PCR. RAPD-PCR with P1254, 23L and OPA-4 primers revealed identical patterns in all of the 36 S. Typhimurium (35 carcass isolates and one feed isolate) isolates. RAPD fingerprints of some isolates with OPA-4 primer are presented in figure 2 The similarity between S. Typhimurium ATCC 14028 and the outbreak strain was noticeable.

PFGE. Using XbaI endonuclease, PFGE revealed identical patterns in all of the 36 strains of isolated S. Typhimurium (figure 3). Interestingly, the type strain S. Typhimurium ATCC 14028 and the outbreak strain produce similar patterns in PFGE. They were different from an isolate cultured from pony in this study.

DISCUSSION

Diseases caused by non-host adapted Salmonella infection is uncommon and is usually seen in chicks, poultys or ducklings less than 2 weeks of age and rarely in birds over 4 weeks of age. The morbidity and mortality varies considerably and deaths usually occur in less than 20% of the affected group (11). In very susceptible young chicks and poultry, paratyphoid infection can sometimes lead to illness and death at high frequencies. Older birds are considerably less susceptible to the lethal effects of paratyphoid Salmonella and may experience intestinal colonization and even systemic dissemination without significant morbidity or mortality (12). Paratyphoid infection seldom causes acute systemic disease except in high subjected to stressful conditions (12).

This outbreak occurred in adult canaries with all of the systemic symptoms. Mortality rate varied from 20% to 94% with average of 51.1%. This indicates the high virulence of S. Typhimurium for a very susceptible host (canary) living in a stressful condition (cage). Keyvanfar et al. (1968) reported lethal systemic outbreak of S. Typhimurium in a flock in Tehran with 100 canaries with a mortality rate of more than 50%. Food contamination with other canary stool was the cause of that outbreak (13). Due to the low immune defense of canaries against S. Typhimurium, this organism is the causative agent of lethal infection in adult canaries. Moreover, breeding and environment conditions in the cage trigger the situation.

Death caused by non typhoid Salmonella occurs primarily in human adults with serious underlying disease such as HIV infection, cancer, leukemia and cirrhosis (4) Importance of infections such as salmonellosis of pet animals like canaries have
increased because of crowding and increasing rates of underlying type of infections in humans living in cities. It is obvious that following identification and reporting of these lethal strains for human health surveillance systems are very important.

Contamination of cages and sources of food with \textit{S. Typhimurium} observed during inspections in this epidemic and food contamination with other canary stools in the study of Keyvanfar \textit{et al.}, indicates importance of inspection of foodstuff and daily cleaning of canary cages from stool and carrier insects and rodents (13). However, distinguishing the source of infection and regions such as what is performed in this research is very important in campaign against these infections.

Madadgar \textit{et al.}, revealed that a combination of RAPD-PCR and R- typing is the method with high discriminatory ability for \textit{S. Typhimurium} strain differentiation (14). Practicality of this idea and discovery of identical isolates in an outbreak has been revealed in this study. In addition, similarity between \textit{S. Typhimurium} ATCC 14028 and this outbreak strain was observed in RAPD-PCR with all primers and PFGE with \textit{XbaI}. This is the first study in which molecular epidemiological methods were used to trace the source of outbreak with \textit{Salmonella Typhimurium} in canaries. This also showed the high susceptibility of canary to this serovar of \textit{Salmonella}.

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