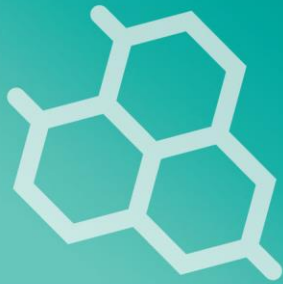


ISSN: 2663-9513 (Online)

ISSN: 2663-9505 (Print)



South Asian Journal of **BIOLOGICAL RESEARCH**



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To cite the article: Mst. Deloara Begum, Sujon Kumer Ray, Md. Mostafizer Rahman, Md. Fakhruzzaman, Md. Nurnoby Islam, and Md. Tazul Islam Sarker. Isolation, characterization and antibiogram study of *salmonella* isolated from apparently healthy Japanese quail, *South Asian Journal of Biological Research*, 4 (2): 84-94.

Link to this article: <http://aiipub.com/journals/sajbr-220323-1003/>

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ISOLATION, CHARACTERIZATION AND ANTIBIOGRAM STUDY OF *SALMONELLA* ISOLATED FROM APPARENTLY HEALTHY JAPANESE QUAIL

*Mst. Deloara Begum*¹, *Sujon Kumer Ray*², *Md. Mostafizer Rahman*³, *Md. Fakhruzzaman*⁴, *Md. Nurnoby Islam*⁵, and *Md. Tazul Islam Sarker*⁶

Corresponding Author E-mail: fakhrul.hstu@gmail.com

¹Mst. Deloara Begum, Assistant Professor, Department of Microbiology, Faculty of Veterinary and Animal Science, Hajee Mohammad Danesh Science and Technology University, Dinajpur-5200, Bangladesh

²Sujon Kumer Ray, MS student, Department of Microbiology, Faculty of Veterinary and Animal Science, Hajee Mohammad Danesh Science and Technology University, Dinajpur-5200, Bangladesh

³Dr. Md. Mostafizer Rahman, Professor, Department of Microbiology, Faculty of Veterinary and Animal Science, Hajee Mohammad Danesh Science and Technology University, Dinajpur-5200, Bangladesh

⁴Dr. Md. Fakhruzzaman, Professor, Department of Microbiology, Faculty of Veterinary and Animal Science, Hajee Mohammad Danesh Science and Technology University, Dinajpur-5200, Bangladesh

⁵Md. Nurnoby Islam, Assistant Professor, Department of Medicine, Surgery and Obstetrics, Faculty of Veterinary and Animal Science, Hajee Mohammad Danesh Science and Technology University, Dinajpur-5200. Bangladesh

⁶Md. Tazul Islam Sarker, MS student, Department of Microbiology, Faculty of Veterinary and Animal Science, Hajee Mohammad Danesh Science and Technology University, Dinajpur-5200, Bangladesh

ARTICLE INFO

Article Type: Research

Received: 27, March. 2022.

Accepted: 06, April. 2022.

Published: 06, April. 2022.

Keywords:

Salmonellosis, Public health, Antibiogram, cloacal swabs, Multidrug resistant

ABSTRACT

Salmonellosis is a major public health problem around the world affecting both animals and humans caused by various species of *Salmonella*. The present study was aimed at isolation, identification and antibiogram profiling of *Salmonella* spp. isolated from cloacal swabs of apparently healthy Japanese quail of selected live bird markets at Dinajpur Sadar Upazila under Dinajpur district of Bangladesh. The isolates were identified based on their morphological, cultural and biochemical characteristics and motility test. While the antibiogram profiling of the *Salmonella* isolates were performed with standard disc diffusion methods. Among the collected samples (n=53), 16.98% samples (n=9/53) were found positive for *Salmonella* spp. Interestingly, all isolates were found to be motile while tested with Motility Indole Urease

(MIU) test. The antibiogram study revealed that all the isolates were completely (100%) resistant to Amoxicillin, Gentamicin, Erythromycin, Tetracycline and Ceftriaxone, whereas the sensitivity rate of the isolates to Ciprofloxacin, Amikacin and Streptomycin were 77.8%, 55.6% and 33.3%, respectively. While, among the resistant *Salmonella* isolates, all were multidrug resistant (MDR). Therefore, findings of the present research work suggested that presence of multidrug resistant *Salmonella* spp. in quails could be a potential threat for public health.



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