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A report of *Pasteurella multocida* Type A infection in an organized poultry farm in Kerala

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Abstract

Fowl cholera is commonly occurring bacterial disease caused by *Pasteurella multocida* (Several serotypes) which affects most of the domestic and wild birds. A confirmatory diagnosis could be done by various biochemical test and polymerase chain reaction (PCR) for serotyping the organism. The main objective of the present study was to investigate the Fowl cholera infection in an organized poultry farm in Thrissur district of Kerala. The feacal swabs, tracheal swabs and blood samples were collected and subjected to microbiological and molecular investigations. Also, these isolates were exposed to antibiotic sensitivity test (ABST) to find out the drug choice for the treatment. Our research findings confirmed that out of 28 birds of native chicken about 21 birds were found to be positive for *Pasteurella multocida* Type A infection. As pasteurellosis has a great impact on economy, early diagnosis with effective treatment and vaccination strategies should be employed in poultry.

Keywords: Fowl cholera, Pasteurella, PCR, ABST

Introduction

Pasteurella multocida type A is the causative agent of fowl cholera, a highly contagious, systemic and fatal disease of chicken ^[13]. Fowl cholera also known as "avian pasteurellosis" and "avian haemorrhagic septicaemia" ^[9]. It's a septicaemic disease which is associated with high morbidity and mortality in poultry predominantly in chicken and ducks ^[1]. It has great impact on poultry economy and production. Fowl cholera usually occurs either acute or chronic forms. In acute cases the clinical symptoms of fowl cholera are often occurs few hours before death which includes fever, ruffled feathers, mucus discharge from mouth, nostrils and ears and cyanosis of comb and wattles [15]. Pasteurella multocida is Gram negative, nonmotile, cocco-bacillus, capsulated non spore forming bacteria occurring singly, in pairs or occasionally as chains or filaments belonging to Pasteurellaceae family ^[13]. In general, five capsular serotypes (A, B, D, E, and F) are found in *Pasteurella multocida* and each serotype is associated with specific host. For example, serotype A and D causes fowl cholera in avian species ^[7]. In Asian countries, fowl cholera is caused by *Pasteurella multocida* type A: 1, A: 3 and type D. Among different types of Pasteurella multocida, serotype A is the most pathogenic which causes 80% mortality whereas serotype D causes 20% mortality in chickens ^[12]. Precise and early diagnosis are the effective methods to rule out the fowl cholera infection. A conventional methods of diagnosis is not effective and less sensitive as compared to polymerase chain reaction (PCR) technique ^[1]. So far in Kerala, only limited studies were carried out on molecular identification of fowl cholera in chicken. As far as we know, there is no report on identification of *Pasteurella multocida* type A in chicken using molecular technique in Kerala. Hence, the present study was aimed at isolation, identification and molecular detection of Pasteurella multocida type A in naturally infected chickens. In addition, antibiotic sensitivity test (ABST) has been performed to find out the specific drug of choice for the effective treatment of infected chickens.

Materials and Methods Clinical examination

A total of 28 birds (one year old) of native chicken were showing the clinical signs of fowl cholera in an organized poultry farm, Thrissur. Upon clinical examination the infected birds were observed with pyrexia (107.6° F), greenish white diarrhoea, pale and hypertrophied comb, mucoid discharge from nostrils and swollen foot pad.

Collection of samples

The feacal swabs, tracheal swabs and blood samples (around 0.5 to 1 ml of blood from the wing vein) were collected aseptically from the infected birds in an organized poultry farm, Thrissur, Kerala. Blood smears were prepared according to the standard procedure. The collected samples were brought to laboratory for further investigation.

Isolation and identification of Pasteurella multocida

Pasteurella multocida organisms were cultured following the standard procedure described by ^[5]. The collected samples (Tracheal and feacal swabs) were streaked into different culture media namely, Blood agar (BA), Nutrient agar (NA) and Nutrient broth (NB) for the isolation of Pasteurella multocida. The inoculated culture plates were incubated at 37°C for the appearance of characteristic colony. To obtain the pure culture of Pasteurella multocida, a selective subculture media was prepared subsequently based on the morphological characteristics of the colony. For the morphological identification of bacteria, the isolated pure culture was subjected to Grams staining. The colonies obtained on the media was subjected to biochemical characterization using various tests viz., Methyl red (MR), Voges-Proskauer (VP), Indole production, catalase, oxidase and sugar fermentation test were carried out.

Antibiotic sensitivity test (ABST)

To find out the specific drug of choice for the treatment, the isolates were subjected to ABST using different antibiotic discs employing amoxicillin, tetracycline, cephalexin, cotrimaxazole, sulphadiazine and trimethoprim, enrofloxacin and gentamicin. The zone of inhibition was measured, recorded and interpreted.

DNA extraction and polymerase chain reaction (PCR) assay

The crude DNA was extracted from the isolates using boiling method given by ^[18]. Polymerase chain reaction was done using specific set of forward (5'-TCCGCAGAAAATTATTGACTC-3') and reverse primer (5'-GCTTGCTGCTTGATTTTGTC-3') to amplify the 511bp fragment of capE gene of Pasteurella multocida type A. Each diluted primer (10 pM/µl) was added to the template DNA [working solutions prepared from stock solution by diluting with sterile distilled water (Millipore) to get a final concentration of 100 ng/µl] and 2X PCR Smart Mix (origin) in a PCR tube and made up to the final volume of 20 µl using ultra filtered Millipore water. PCR was done in Bio-Rad thermal cycler and standardization was done for each reaction by mild adjustment of concentration of ingredients and annealing temperature with the following profile: initial denaturation of 5 min at 94°C; 35 cycles of 94°C for 30 s, annealing at 55°C for 30 s, and 72°C for 30 s with a final elongation of 5 min at 72°C. PCR amplicon was subjected to 2% agarose gel.

Agarose gel electrophoresis

The genomic DNA and PCR products were checked in agarose gels of 0.8 percent and 2%, respectively prepared using 1X TBE buffer. DNA and PCR products were mixed with loading dye (6X) and loaded in wells. The PCR products were loaded along with a molecular weight marker (50bp) for relative sizing. Electrophoresis was carried out at 5V/cm until the bromophenol blue dye migrated more than two by third length of the gel and was visualized in a Gel Doc System (Bio-Rad, USA). The amplicons of *capE* gene of *Pasteurella multocida* type A (511bp) were sequenced using respective forward and reverse primers in an automated sequencer using Sanger's dideoxy chain termination method at Agri Genome Labs Pvt. Ltd., Cochin.

Results and Discussion

In the present study out of 28 birds of infected native chickens, 21 birds were found to be positive for *Pasteurella multocida*. On Geimsa staining, blood smears revealed the presence of bipolar organisms (Fig.1). In comparison, ^[21] observed numerous bipolar organisms characteristic of *Pasteurella* spp. upon microscopic examination of Geimsa stained smears of heart blood in water fowl. The clinical samples collected during an outbreak of heamorrhagic septicaemia in a buffalo farm of West Bengal, India. On microscopic observation, peripheral, heart blood smear and tissue impressions revealed the presence of Gram negative, bipolar cocco – bacilli organisms alike of *Pasteurella* spp. ^[11]



Fig 1: Bipolar appearance of *Pasteurella multocida* in Geimsa staining

Cultural examination of tracheal and feacal swabs revealed the presence of small, glistening, mucoid, dew drop like colonies on blood agar (BA) after an overnight incubation (Fig.2). ^[16] Reported the same colony morphology in the samples which are collected from private poultry farm, Maharastra during the Fowl cholera outbreak. Similar findings was reported by ^[2] in cattle and buffalo of Karachi region of Pakistan. Also, ^[8, 13] reported the indistinguishable findings in native chicken of Bangladesh. In addition, the recovered bacteriological isolates were shown smooth, glistening and translucent colonies on blood agar and produced non-heamolytic dewdrop colonies on sheep blood agar. However, these isolates were failed to grow on McConkey's agar ^[11]. These findings were accordance with that of ^[6] who conducted a research on isolation, identification and antimicrobial susceptibility patterns of Pasteurella multocida isolated from diseased rabbits. As stated by ^[3], a few strains of Fowl cholera organisms may not grow in media without blood or blood serum. In the present study, the selected isolates were found to grow well in bovine blood agar media.

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Fig 2: Small, glistening, mucoid, dew drop like colonies on blood agar (BA) media

The selected isolates were found be to Gram-negative, coccobacillary shaped *Pasteurella multocida* on Gram staining (Fig. 3). Similarly, the microscopical examination of gram stained smears of suspected isolates revealed characteristic Gramnegative, non-spore forming and cocco-bacillary organisms arranging singly, in pairs or short in chains. Furthermore, a typical bipolar staining rods were evident in Giemsa stained smears of Pasteurella fresh cultures ^[6]. These results were accordance with that of ^[2, 8, 13, 1]. Also, ^[11] reported the indistinguishable findings from the collected samples suspected of Haemorrhagic septiceamia in an organized buffalo farm.



Fig 3: Gram negative, cocco bacillary or rod shaped *P. multocida* on Gram staining

On biochemical characterization, *Pasteurella multocida* organism were shown to be positive for catalase and oxidase test (Fig.4 and 5). On the contrary, Methyl red (MR) and Voges-Proskauer (VP) test did not show any reaction. Thus, *Pasteurella multocida* organism is negative for MR and VP tests. Sugars such as dextrose, sucrose and mannitol were fermented by *Pasteurella multocida* organism whereas maltose and lactose was not fermented. In comparison, these findings of biochemical reactions were similar with that of ^[18, 19, 8, 4 13]

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Fig 4: Catalase test of *Pasteurella multocida* showing positive reaction



Fig 5: Oxidase test of *Pasteurella multocida* showing positive reaction

The isolates were found to be sensitive to sulphadiazine and trimethoprim and cephalexin, and resistant to amoxicillin, tetracycline, cotrimaxazole, enrofloxacin and gentamicin (Fig.6). In contrast, ^[16, 14] were reported that isolates of *Pasteurella multocida* were sensitive to enrofloxacin, gentamycin and enroflaxacin, cephalexin in chicken and Pigeon, respectively. On the basis of ABST results, infected birds were treated with Injection Biotrim (Sulphadiazine and trimethoprim) at a dose rate of 25mg/Kg bodyweight intramuscularly and Cephalex powder (Cephalexin) at the rate of 0.3gm/ day orally for one week. During the course of treatment, about 3 birds were died out of 21. The remaining 18 birds were uneventfully recovered after one week of post treatment. Blood smear taken after a week was negative *Pasteurella* organisms.



Fig 6: ABST – Isolates were found sensitive to sulphadiazine and trimethoprim and cephalexin

addition, for the identification and molecular In characterization of Pasteurella spp., the isolated samples were subjected to polymerase chain reaction (PCR) with specific set of primers. The PCR assay revealed the 511bp amplicon fragment of capE gene of Pasteurella multocida type A (Fig. 7). In comparison, ^[13] carried a research on isolation and molecular detection of Pasteurella multocida type A in naturally infected chicken in Bangladesh. Also, ^[1] reported the similar findings in which out five dead chickens two were found to be positive for Pasteurella multocida type A infection in brother's poultry farm located at Konabari, Gazipur, Bangladesh. Similarly, ^[10] conducted a research on prevalent serotypes of Pasteurella multocida isolated from different avian species of India and found that majority of avian isolates were belonged to serotype A. On the other hand, ^[17] reported that prevalence of *Pasteurella multocida* capsular serotype B was found to be high in cattle and buffaloes of Iran. In case of ruminants, Pasteurella multocida serotype B causes Haemorrhagic septicaemia (HS) with high mortality and buffaloes are more susceptible than cattle. In Kerala, infection of Pasteurella multocida in poultry has been reported in many places but there were no reports have been made on molecular identification of Pasteurella multocida type A. The research findings of our present study reported the successful molecular diagnosis of Pasteurella multocida type A in naturally infected native chickens of Kerala. These results suggest that PCR assay could be an effective tool for the early diagnosis of disease. Also, PCR is comparatively more sensitive than conventional methods of diagnosis in order to identify the specific serotype of a particular organism. Hence PCR technique can be successfully applied to detect the different serotypes of Pasteurella multocida in poultry and other species.



Fig 7: Amplification of 511 fragment of capE gene of *Pasteurella multocida* type A. M- Marker, L1- Sample, L2- Negative control

Conclusion

In this study, a successful isolation and molecular characterization of *Pasteurella multocida* type A from naturally infected native chickens was carried out using PCR

assay. Since the bacterial isolation and identification are time consuming procedures, PCR could be an effective tool for timely diagnosis of the disease. Also, Pasteurellosis has great impact on poultry economy, early diagnosis with effective treatment and vaccination strategies should be implemented. Thus, molecular identification of fowl cholera could be an efficient method of diagnosis in future studies.

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