

Isolation of Pathogenic Bacteria from Poultry Wastages at Chennai Suburban

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Abstract : India is one of the largest broiler producers in the world. Huge quantities of poultry wastages are generated and dumping of these wastes causes several environmental pollution problems. One of the serious problems is, it also serves as a nutritional source for the growth of several pathogenic bacteria and thereby causes disease outbreaks. This work addresses this environmental issue through enumeration, isolation and identification of microorganisms from the dumping sites of poultry wastages. Samples were collected from five different regions and bacterial isolates obtained were then purified into pure culture and identified based on their morphological, cultural and biochemical tests by standard microbiological procedures. The isolated pathogenic organisms were *Vibrio cholerae*, *Vibrio parahaemolyticus*, *Salmonella typhi*, *Shigella dysenteriae*, *Shigella sonnei* and *Klebsiella pneumonia*. The colonization of pathogenic bacteria in poultry waste dumping sites is highly hazardous to the environment and these poultry wastages need a prompt disposal system.

Keywords: Poultry waste, Environmental pollution, Total Bacterial Count, IMViC, Enteric pathogens, Opportunistic pathogens.

I. Introduction

Poultry is one of the fastest growing agricultural sectors in India [1]. This leads to significant raise in the production of broilers at the rate of 8 to 10% per annum [2] with an annual turnover of 30,000 crores [3]. The advantages of low investment and the requirement of small area have increased the number of poultry shops and creating employment opportunities [4]. This also leads to the generation of huge quantities of poultry wastes usually composed of broiler and layers, feathers, bones, blood, hatchery debris and dead birds. These wastes pose serious environmental pollution problems through microbial infection, offensive odours, promotion of flies and rodent breeding [5]. There are no proper disposing units for these wastages in developing countries like India but attempts are in process to reuse these materials as beneficiary ones such as fertilizers and animal feed supplements [6]. As these wastes are composed of tissues and blood, we have hypothesized that these deposited wastes may serve as a reservoir for the multiplication of several pathogenic microorganisms that can cause severe disease outbreaks. Hence, the present study was carried out to identify the potential pathogens that can survive in these poultry wastages which can be hazardous.

II. Materials And Methods

Study Area

The poultry waste samples were collected from the poultry waste dumping sites in and around Poonamallee town, a suburban situated in the west side of Chennai, Tamil Nadu state, India. Samples were collected from five different regions namely Iyappanthangal, Kumananchavadi, Senneerkuppam, Chembarambakkam and Kuthambakkam. All the sampling sites were densely dumped with the poultry wastages as shown in fig.1 (Kumananchavadi).



Fig 1. Kumananchavadi Poultry wastage dumping site

Sample Collection

The samples were collected according standard microbiological procedures [7,8] during pre-summer season in the month of January 2013. In brief, about 2 gram of samples were collected in sterile containers using sterile spatula (as shown in fig.2) and immediately transported to the Microbiology laboratory of Sree Sastha Institute of Engineering and Technology for further analysis.



Fig 2. Aseptic collection of poultry wastes

Media and Reagents used

The media and reagents used for the study such as Nutrient Broth, Nutrient agar, Eosin-Methylene-Blue agar, Blood agar, Salmonella-Shigella agar, Thiosulfate-Citrate-BileSalt-Sucrose agar, Mannitol Salt agar, MacConkey agar, Methyl red and Voges Proskauer were procured from Himedia, India.

Sample Processing

The collected samples were aseptically processed in the microbiology laboratory. 500 mg of each sample was inoculated into 100 ml of freshly prepared nutrient broth and incubated for 24 hrs at 37⁰C. After incubation, the total viable count of the bacteria and screening for the potential pathogens were carried out.

Total Bacterial Count

The total bacterial count was performed to all the samples by Lazy Susan plating method in triplicate on solid Nutrient agar plates after serial dilution of samples at 1 in 10 concentrations. The seeded plates were sealed and incubated at 37⁰C. The colony forming units were counted after 24 hours using colony counter (I.L.E, India) and expressed in CFU/mg.

Isolation of Pathogens

Isolation and identification of bacterial pathogens were carried out based on their growth pattern and colony characteristics on selective and differential media. In addition, Grams staining, IMViC, Catalase, Oxidase, Urease and motility tests were carried out according to the standard test procedures [7,8] to confirm the findings.

III. Results And Discussion

The total bacterial count was assessed to all the samples and described in fig 3. Bacterial counts are presented as the average of the actual triplicate values. All the study regions have shown significant load of bacteria. Among the five different regions Kumananchavadi has the highest bacterial load with 1088×10^9 CFU/mg of sample. This region had a huge dumping of fresh poultry wastages and debris in sacs when compared to other study regions. Moreover this dumping site usually founds with fresh blood and other body fluids that supports the growth and multiplication of most of the pathogenic bacteria [9].

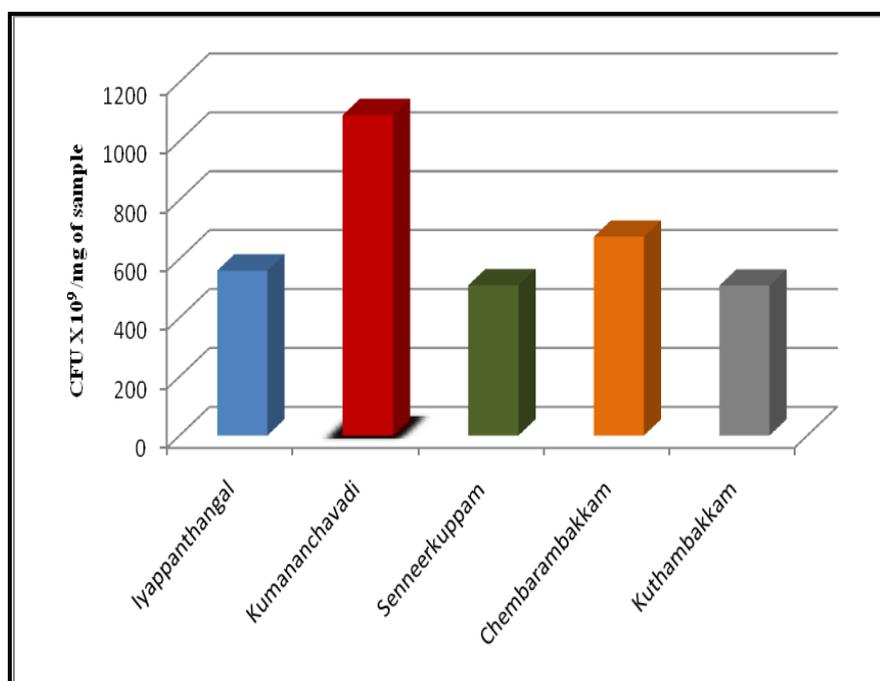


Fig 3. Total Bacterial Count at Study Regions

The pathogenic organisms were identified and confirmed through the standard biochemical tests and the region-wise distribution is given in Table 1. Present study has revealed the presence of potential pathogens such as *Vibrio cholerae*, *Vibrio parahaemolyticus*, *Salmonella typhi*, *Klebsiella pneumoniae*, *Shigella dysenteriae* and *Shigella sonnei* in the study regions. The opportunistic pathogens such as *E.coli* and *Staphylococcus sp.* were also isolated. The typical growth characteristics of *Salmonella typhi* and *Vibrio cholerae* on SS agar and TCBS agar respectively are given in fig 4 and fig 5.

Table 1. Distribution of pathogens

S. No	Samples Sites	Bacterial pathogens	Opportunistic Bacterial pathogens
1	Iyappanthangal	<i>Vibrio cholerae</i> <i>Salmonella typhi</i> <i>Shigella dysenteriae</i> <i>Shigella sonnei</i>	<i>E. coli</i> <i>Staphylococcus sp.</i>
2	Kumananchavadi	<i>Vibrio cholerae</i> <i>Vibrio parahaemolyticus</i> <i>Salmonella typhi</i> <i>Shigella dysenteriae</i> <i>Shigella sonnei</i>	<i>E. coli</i> <i>Staphylococcus sp.</i>
3	Senneerkuppam	<i>Klebsiella pneumoniae</i> <i>Salmonella typhi</i> <i>Shigella dysenteriae</i> <i>Shigella sonnei</i> <i>Vibrio parahaemolyticus</i>	<i>E. coli</i> <i>Staphylococcus sp.</i>
4	Chembarambakkam	<i>Klebsiella pneumoniae</i> <i>Salmonella typhi</i> <i>Shigella sonnei</i>	<i>E. coli</i> <i>Staphylococcus sp.</i>
5	Kuthambakkam	<i>Salmonella typhi</i> <i>Shigella sonnei</i>	<i>E. coli</i> <i>Staphylococcus sp.</i>



Fig.4 Growth of *Salmonella typhi* on SS agar with H₂S production

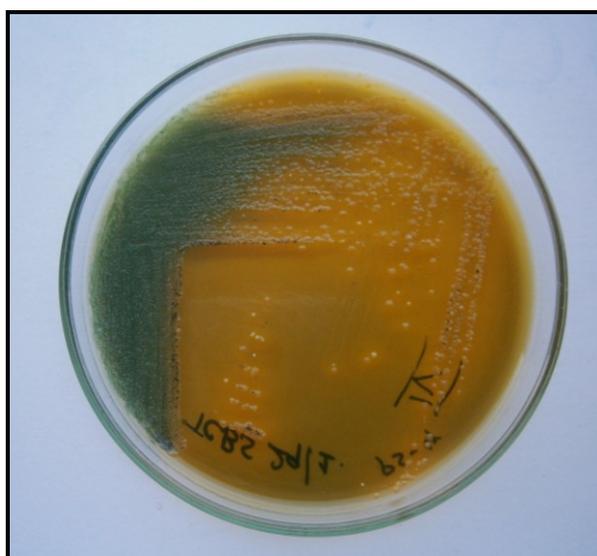


Fig.5 Growth of *Vibrio cholerae* (Golden yellow coloured) on TCBS agar

India is the ninth largest broiler producer in the world today [10]. The production and processing of poultry involve killing, defeathering, eviscerating and cleaning that lead to the generation of wastes. Coker *et al* [9] showed that these wastes can affect water, land and air qualities if proper practices of management are not followed. Hence it is important to study the prevalence and distribution of different bacteria in these poultry wastes.

The findings of the present study support the study hypothesis that the deposited poultry wastes serve as a reservoir for the growth and multiplication of bacteria. All the samples have showed significant bacterial growth. The isolated organism *Vibrio cholerae* is the causative agent of cholera that spreads through the contaminated water. Epidemic outbreaks are more common as it spreads through convalescent carriers and the incubation period for these bacteria is of 1-5 days [11]. Another species *V. parahaemolyticus* is capable of causing gastroenteritis through contaminated water and food. Presence of these organisms is a big threat to the community living in these regions.

The samples also revealed the presence of *Shigella dysenteriae* and *Salmonella typhi* which are also serious enteric pathogens. *Shigella dysenteriae* can cause severe dysentery and prostration to the infected people [12]. The major threats with this organism are; it remains viable in moist environment for several days and a small infective dose such as 20-100 bacilli is enough to cause the infection. *Salmonella typhi* is the causative agent of typhoid fever in which the organism invades the intestinal mucosa and multiplies there. Occasionally they manage to enter the lymphatic and cardio vascular systems to cause serious illness [13]. Presence of other bacteria such as *K. pneumoniae*, *E. coli* and *S. aureus* is also a major environmental health problem. They can cause urinary tract infections, respiratory tract infections and gastrointestinal tract infections [14].

The colonization of pathogenic bacteria in these dumping sites is highly hazardous to the environment. If these infectious poultry wastes are discharged into lakes, ponds and/or other drinking water resources it can harm aquatic life and jeopardise the quality of our drinking water [15]. In addition, the nuisance odour arises

from these materials has also an impact on greenhouse gas emissions associated with global climate change [16]. These waste materials need prompt disposal system or should be utilized as an alternative source. Several ways of disposing methods are followed worldwide such as burial, rendering, incineration, composting, feed for livestock, fertilizer or source of energy [6] as recommended by Hazard Analysis Critical Control Point (HACCP) guidelines [17] and Environmental, Health, and Safety Guidelines for Poultry Production Guidelines [18]. Poultry waste materials also contain organic matter that can be converted into bioenergy under anaerobic conditions which yields biogas, a gas mixture with varying concentrations of combustible methane [19]. But, before proceeding with these processes an effective screening procedures like the present study are important.

Present study has some limitations. It has focussed to screen only the aerobic bacteria and not other microorganisms such as anaerobes and fungus. The reason is the limited availability of resources and laboratory facilities. Special infrastructure and culture conditions are required to screen for more number of other pathogens. Investigations with advanced techniques like PCR, molecular sequencing and flow cytometry may reveal more scientific information.

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