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An exploration on animal and public health significance of salmonella from major meat sources in Puducherry, India

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Abstract

A study was framed to isolate the predominant strains of Salmonella prevalence in Puducherry. A total of 150 (Chicken, Chevon and Beef each 50) raw meat samples procured from different retail outlets in and around Puducherry. The study objective included Salmonella identification and PCR confirmation, serotyping, biofilm formation assay and antimicrobial resistance testing with selected antibiotics. Conventionally confirmed isolates were subjected for PCR against *invA* gene. Overall prevalence was 7.4 per cent and chicken was found to harbour highest isolates. On serotyping *Salmonella typhimurium* was the only strain recorded in all meat samples. Recovered *Salmonella typhimurium* were biofilm producers. High antimicrobial resistance were observed to Amoxycillin, Co- trimaxazole and Metronidazole whereas, maximum sensitivity to Azithromycin, Chloramphenicol, Cephalexin, Cefotaxime and Tetracycline. This study indicated the need to take up foodborne disease surveillance in Puducherry under the Indian context and to identify the common high-risk food commodities for microbial contamination and antibiotic resistance.

Keywords: Foodborne illness, raw meat, salmonella, biofilm, antibiotic resistance

Introduction

Meat, in general considered of major protein sources in India. Meat production in India estimated at 6.18 million tones, standing 5th of world meat production [29]. Though India stands at 5th position, export in the livestock area is not up to the standards. The expected meat production in 2030 is 3.3 times more than present situation [16]. Over the last 25 years, the global incidence of foodborne infections has markedly increased, with nearly a quarter of the population at a high risk of foodborne diseases [31]. World Health Organization [44] estimated that foodborne and waterborne diarrheal diseases together kill around 2.2 million people annually. The contaminated meat and dairy products are the most common cause of human Salmonellosis outbreaks in worldwide [20]. In India, majority of food borne outbreaks remains underreported or non-investigated and may only be noticed after major health effects or economic damage has occurred. Nevertheless few peoples already have been reported on Salmonella food poisoning [8, 5, 1, 12]. Salmonella survivability also increasing in other side by means of microbial biofilm on the contact surfaces in the food industry or small intestine. The direct contact with raw materials or foodstuffs can cause contamination and due to which the product become unsafe [39]. The biofilm forming isolates are considered an emerging menace to public mainly of resistant to antimicrobial agents [15].

Even though studies on the microbial quality of fresh meat sold in Puducherry, was carried out in Rajiv Gandhi Institute of Veterinary Education and Research, no systematic study has been conducted on the Public and animal health significance of Salmonella. Therefore the present work framed with the objectives of isolation and identification Salmonella from raw meat (chicken, chevon and beef), Serotyping, PCR based confirmation of the *invA* gene, biofilms production and antimicrobial resistance profiling of the isolates.

Materials and methods**Collection and transportation of meat samples**

A 100 gram of raw meat samples were collected from different retail meat shops in and around Puducherry (chicken, chevon and beef each 50) and subjected to this study. All the samples

were collected in sterile polyethylene bags and immediately transported to laboratory in ice box for further analysis.

In this study all the chemicals, primers, reagents and culture media were procured from the Merck Limited, Hi media from Mumbai and Bangalore genei. The PCR master mix was procured from Gene Technologies, Chennai

Primary and Selective enrichment

Twenty five gram of each meat sample (chicken, chevon and beef) was homogenized in 225 ml of Peptone Water (PW) and incubated at 37 °C overnight for the metabolic recovery and proliferation of the microbes. One milliliter of pre-enrichment culture was inoculated into tubes containing 9 ml of Selenite F Broth (SB) and Rappaport Vassiliadis Soya (RVS) Broth and incubated for 24 hours at 37 °C for secondary enrichment.

Isolation on selective and differential media

Bacteria growing in SB and RVS broth were streaked onto Mac Conkey's agar for differentiation of lactose and non-lactose fermenters. Non lactose fermenters were streaked onto Xylose Lysine Deoxycholate (XLD) and Bismuth Sulfate agar (BSA) incubated at 37 °C overnight. Agar plates were examined for the presence of colorless colonies on Mac Conkey's agar, pink colonies with or without black centers on XLD and black colonies surrounded by a black or brownish zone with a metallic sheen on BSA. All the remaining negative samples were decontaminated as per standard procedures.

Identification of Salmonella

The presumptive colonies were subjected to cultural, morphological and biochemical characters as per [9]. Primary identification tests include Gram's staining, Catalase test (Slide test), Oxidase test and Motility test were performed. Secondary identification tests like Indole production, Methyl Red (MR) reaction, Voges-Proskauer (VP) Reaction, Citrate utilization test, Urease activity, Triple Sugar Iron test, Lysine Decarboxylase and Carbohydrate Utilization tests proceeded as per the standard procedures. All biochemical tests were performed at 37 °C for 24 hours.

Preservation and revival of isolates

A loopful of the isolated organism was added to the sterile DMSO-glycerol broth vials and mixed well in the vortex mixture. The vials were then labeled and stored at - 40 °C. The isolates were revived once in three months. The vials stored at - 40°C was thawed and streaked on Nutrient agar and incubated at 37 °C overnight. The isolated organism was also streaked onto MH slants in screw capped test tubes and incubated at 37 °C for 24 hours. Once the growth was obtained the slants were stored at 4 °C.

Molecular confirmation by PCR

Extraction of DNA from isolates

The reference strain of *Salmonella* spp was obtained from Institute of Microbial Technology, Chandigar. Template preparation and PCR was carried out for the detection of *invA* gene as per the method described by [34]. Primer, Reaction mixture and PCR conditions details are given in table 5, 6 & 7.

Serotyping

Isolates of Salmonella were sent to National Salmonella and Escherichia Centre (Central Research Institute, Kasauli,

Himachal Pradesh, India) on nutrient agar slants.

Assessment of biofilm production of the isolates

The confirmed Salmonella colonies were further purified on Muller Hinton agar before being subjected for biofilm production assay. Slime production assay was performed as per [41]. Briefly, Brain heart infusion agar supplemented with 5 per cent sucrose and Congo red (0.08 g/l) was prepared and autoclaved. The isolates were inoculated and incubated aerobically for 24 to 48 hours. The ability of the isolates to produce bio-films was indicated by black colonies with a dry crystalline consistency and red colour colonies indicates negative. Procedure was done with the comparison of positive and negative controls.

Antimicrobial resistance profiling of the isolates

The drug susceptibility of Salmonella isolates was performed on Mueller Hinton agar plates by disc diffusion method in accordance with the recommendations of the Clinical and Laboratory Standards Institute [10]. List of antibiotics showed in Table 2.

Results

Isolation of Salmonella

In the present study, peptone water was used as a non-selective pre enrichment medium. The pre enrichments were sub cultured in selective enrichment media Rappaport Vassiliadis Soy (RVS) broth and Selenite F broth (SB). A total of 150 raw meat samples were processed and screened for *Salmonella* spp. Eleven (7.4 per cent) samples positive by culture and all the 11 samples were confirmed by polymerase chain reaction. In total isolates, eight (72.7 per cent) were from Rappaport Vassiliadis Soy (RVS) broth, three (27.2 per cent) were isolated from Selenite F broth.

Out of 11 (7.4 per cent) isolates, five (45.4 per cent) from chicken, four (36.6 per cent) from beef and two (18.1 per cent) from chevon were isolated only by RVS. Among the five isolates from chicken, three (60 per cent) from RVS and two (40 per cent) from SB were recovered (Table 3). Chicken being a natural host for Salmonella the prevalence was high compared to the other meat. The rest of the all samples were negative by culture.

Identification of Salmonella

Out of 150 samples processed, 11 (7.4 per cent) samples were found to show pale colonies on Mac Conkey's agar, pink colonies with black centre in XLD agar and black colonies with metallic sheen on BSA (Plate - 1,2 &3). On Gram's staining, all were found Gram-negative slender rods. Catalase positive cultures were further identified up to subspecies level as per [9]. Results showed in Table 4.

Serotyping of Salmonella

The serotype isolated Salmonella was *S. typhimurium*. All the isolates were found to have the somatic antigens as 4,12 and flagellar phase I antigen was 'I' and the phase II antigen was 1,2.

PCR for primers targeting *invA* gene

All the 11 positive meat isolates of Salmonella by cultural and biochemical tests were subjected for PCR with Salmonella genus virulence specific primers (*invA* gene). DNA was prepared by the boiling-lysis method from the suspected Salmonella culture. PCR products analysed on 1.5 per cent

Agarose gel electrophoresis (along with positive and negative control and 100 bp ladder) revealed the presence of expected 244 bp products under UV transilluminator. The primers targeting the *invA* gene successfully amplified the DNA prepared by the boiling-lysis method from all Salmonella culture. The details of results of PCR are depicted in Picture 1.

Assessment of biofilm production

PCR confirmed isolates of Salmonella are subjected for biofilm production test and all were possess the ability to produce black colonies with dry crystalline consistency on modified Congo Red Agar. The results indicated isolates were able to produce biofilm (Plates - 4&5).

Antimicrobial resistance profiles

The details of the results and consolidated antimicrobial sensitivity pattern of 15 antibiotics depicted in Table 8. Salmonella isolates from chicken, were 100 per cent sensitive to azithromycin, chloramphenicol and cefotaxime, 100 per cent resistant to amoxicillin and metronidazole, 60 per cent resistant to tetracycline and norfloxacin, 40 per cent resistant to nalidixic acid and 20 per cent resistant to ciprofloxacin.

Out of two isolates of Salmonella from chevon, 100 per cent were sensitive to azithromycin, cephalexin, chloramphenicol, timethoprim, tetracycline and cefotaxime, 100 per cent resistant to amoxicillin, co-trimoxazole and metronidazole and 50 per cent resistant to gentamicin, nalidixic acid, ciprofloxacin and norfloxacin. Out of four Salmonella isolates from beef 100 per cent were sensitive to amoxicillin, cephalexin, cefotaxime, perfloxacin and sulphamethizole, 100 per cent resistant to metrinadazole, 75 per cent resistant to norfloxacin and tetracycline, 50 per cent resistant to gentamicin, nalidixic acid and 25 per cent resistant to ciprofloxacin.

Discussion

Food borne diseases always follow the consumption of contaminated food especially from animal products such as meat or carcasses. In developing countries most of the foodborne outbreaks are mainly associated with Salmonella spp. These organisms are subsequently transferred to human through production, handling and consumption of meat and meat products. Food is an important vehicle for human infection of *Salmonella* spp but the number of Salmonella shedding from food is usually low. It is necessary to use pre enrichment media such as peptone water, buffered peptone water or universal pre enrichment broth to assist isolation. This may allow the small number of food or contaminated Salmonella to grow, that may otherwise be killed by the toxic effects of different enrichment media. Pre enrichment normally helps to multiply or resuscitate Salmonella that have been sub lethally damaged. Many studies have documented the use of peptone water for pre enrichment. In the present study peptone water was used as a non- selective pre enrichment media. Peptone water was a preferred pre enrichment media for Enterobacteriaceae family especially Salmonella as it favoured the repair and growth of stressed or sub lethally injured isolates arising from exposure to heat, desiccation, high osmotic pressure or wide temperature fluctuations [2, 13, 47, 26].

Enrichment media contain additives that selectively permit Salmonella to grow while inhibiting the growth of other contaminated bacteria. Some however, are also relatively

toxic to certain serovars of Salmonella, e.g. Selenite inhibits *S. cholerae suis*. Elevated temperature has also been used in some laboratories especially with RVS broth, where the recommended temperature for incubation is around 41.5°C [40]. Rappaport Vassiliadis Soy (RVS) broth and Selenite F broth (SB) were used as selective enrichment media in the present study and better isolation was observed with RVS broth compared to SB. RVS broth also was found to be an effective enrichment media for preparation of template DNA for PCR examination when compared to SB which did not produce good quality of DNA. Authors [4] used buffered peptone water for non - selective enrichment and RVS broth for selective enrichment of Salmonella isolates. [13] Used SB, RVS broth for secondary enrichment of Salmonella. Solid, selective agars permit differential growth to varying degrees. They inhibit the growth of other unwanted bacteria other than Salmonella, which gives the information on some of the principal differential biochemical characteristics usually non lactose fermentation and hydrogen sulphide production. The results were recorded after 24 to 48 hours of culture at 37°C. Salmonella form characteristic colonies on such media that are usually distinguishable from the colonies of other bacteria with the possible exceptions of Proteus, Pseudomonas and Citrobacter. In this study XLD and BSA were used as selective and differential media for the isolation of Salmonella from the selectively enriched cultures. Mac Conkey's agar was used as a selective and differential media at initial stage for streaking of enriched culture from selective broth and also used to purify the Salmonella suspected colonies from XLD, BSA and Salmonella-Shigella agar. [35] Recommended the Selenite F enrichment broth for selective isolation and Mac Conkey's agar and XLD agar for purification.

Out of 150 raw meat samples processed and screened for the presence of Salmonella in meat, 11 (7.4 per cent) isolates were recovered. Among the 11 isolates five (45.4 per cent) from chicken, two (18.1%) from chevon and four (36.6 per cent) from beef were recorded. This result was found to be higher than [22] who reported a total of ten isolates of *Salmonella* spp from raw meat with an overall prevalence of 5 per cent among food animals in Maharashtra, India. An overall prevalence of Salmonella in 7.4 per cent food animals in current study is comparable with the findings [23]. In their study 5 per cent (n=300) Salmonella incidence was reported from the municipal slaughter houses and the retail meat shops in Hyderabad and Karnataka, India. This higher and lower incidence variation may also relay the suitable environment persistence for Salmonella. In present study increased isolates shows the presence of suitable ecology for Salmonella.

In a total of 50 samples screened 5 (10 per cent) were positive in chicken. Similarly, [11] reported the 8 per cent (n=100) incidence in Puducherry. In an earlier study [35] on poultry carcasses reported 8.3 per cent (n=60) incidence in Tamil Nadu. [37] Reported, a total of 40 (n=80) Salmonella isolates from poultry in Assam, India. Chicken is considered as a natural host for *Salmonella* spp hence, the prevalence of Salmonella in the present study is high compared to other meat. However, variations observed between the reported Salmonella prevalence in previous investigations around the world may be due to several factors, including the initial Salmonellosis in live birds, sanitation within the slaughterhouse, possible contamination during poultry processing steps (e.g., the amount of cross-contamination of chicken carcasses by contact with intestinal tracts during slaughter or processing) and differences among isolation

methods applied to detect *Salmonella*.

Four percent incidence in chevon was recorded in the present study (n= 50). This observation differs from the works done in the other parts of India wherein, 17.6 per cent and 38.7 per cent prevalence of *Salmonella* were recorded in goats slaughtered at Bareilly and Wardha district of Maharashtra, respectively [11, 28]. The differences in the prevalence reported could be associated with the sampling procedures, type of sample, transportation of samples, isolation techniques or the actual difference in the occurrence and distribution of *Salmonella* in the study population itself. This increased prevalence than earlier studies clearly indicates the presence of health hazards in chevon.

Out of 50 beef samples processed and screened, 8 per cent isolates were recovered. An overall 8 per cent *Salmonella* prevalence among beef samples in the current work is higher than a report [22] from 6 per cent (n=50) in Maharashtra, India and also higher than those reported 4 per cent (n=50) in faecal samples from diarrhoeic young animal, from Telangana, Chennai, Maharashtra, Goa, Uttar Pradesh, and Rajasthan [30]. The workers attributed the reason for lower prevalence in young animals and absences of *Salmonella* in bovine to the limited number of samples included in the study. The higher prevalence in the present study could be due to the actual prevalence of *Salmonella* in cattle itself or it might be attributed to the unhygienic conditions in slaughtering, dressing, washing and handling the carcasses leading to the cross contamination of the meat samples.

Isolates of *Salmonella* were sent to National *Salmonella* and *Escherichia* Centre (Central Research Institute, Kasauli, Himachal Pradesh, India) on nutrient agar slants for serotyping. The serotype isolated from the meat samples was *S. typhimurium*. All the isolates were found to have the somatic antigens as 4,12 and flagellar phase I antigen was 'I' and the phase II antigen was 1,2. This is in accordance with that of Chaiba [7], who had reported that *Salmonella typhimurium* was the predominant and most frequently isolated *Salmonella* serotypes among the 57 *Salmonella* isolated from meat samples. The isolation of the serovar *Salmonella typhimurium* in the present study is an indication of a close association of man and meat animals during any of the farm operations or consumption of contaminated meat. Isolation of *Salmonella typhimurium* alone from the meat samples might be attributed to the fact that only a small geographical area was covered in the present study.

A total of 11 (7.4 per cent) isolates of *Salmonella* were subjected for PCR with *invA* gene (genus specific for *Salmonella* genus). All the isolates were confirmed to carry the *invA* gene. [35] Recommended PCR for the detection of *Salmonella* from broiler meat and the incidence was 8.3 per cent (n=60) in Tamil Nadu. In this study primer targeting the *invA* gene was found to be an accurate for the identification of *Salmonella*. [21] Examined the presence of *Salmonella* by polymerase chain reaction in poultry carcasses. A total of 60 neck skin swab samples were taken from 12 different broiler flocks in Iran. PCR amplification of *invA* gene was the method for detection of *Salmonella*. In their study *Salmonella* was isolated from 11.6 percent of samples by conventional culture method. All positive isolates from conventional culture method were confirmed by PCR amplification and *invA* gene was superior for the confirmation of *Salmonella*.

A total of 11 confirmed isolates of were subjected for biofilm production assay. All were found to produce black colonies with dry crystalline consistency colonies on modified Congo

Red Agar. The result indicates that all the isolates were able to produce biofilm. [42] Studied the biofilm formation ability of *Salmonella* isolates from a chicken slaughter plant in China. The isolates have exhibited variation in biofilm forming behaviour with a relatively high biofilm production was observed for *Salmonella agona* compared to *Salmonella typhimurium* isolates.

[27] Assessed the biofilm forming ability of *Salmonella enterica* isolated from meat products in USA. Twenty *Salmonella enterica* subsp *enterica* strains were isolated from meat (n=11) and meat products (n=7). Two (10 percent) *Salmonella enterica* subsp *enterica* strains from ground chicken meat and chicken sausage were defined as biofilm forming but others were considered non biofilm forming. Different studies have been conducted to see the biofilm formation of *Salmonella* isolates on different surfaces [3]. Various factors (growth medium, incubation period, fixation of adhered cells and staining) affect development of *Salmonella* biofilm [28]. According to those studies, the source of isolates (from humans, animals or food) did not affect the biofilm [18]. The evidence of biofilm shows the existing survivability of *Salmonella* in meat samples which indicates the persisting threat to public.

Out of five *Salmonella* isolates from chicken, 100 per cent were sensitive to azithromycin, chloramphenical and cefotaxime, 100 per cent were resistant to amoxicillin and metronidazole, 60 per cent were resistant to tetracycline and norfloxacin, 40 per cent were resistant to nalidixic acid and 20 per cent were resistant to ciprofloxacin. [14] Screened the raw meat to find out the prevalence of multi drug resistance *Salmonella* isolates. Among the variety of antibiotics tested, the highest resistance was observed with nalidixic acid followed by tetracycline, trimethoprim, and streptomycin. The isolates of chicken resistant to nalidixic acid, tetracycline, trimethoprim, and streptomycin were 90.6 per cent, 71.9 per cent, 56.6 per cent and 25 per cent respectively.

Out of two isolates of *Salmonella* from chevon, 100 per cent were sensitive to azithromycin, cephalixin, chloramphenical, trimethoprim, tetracycline and cefotaxime, 100 per cent were resistant to amoxicillin, co-trimoxazole and metronidazole and 50 per cent were resistant to gentamicin, nalidixic acid, ciprofloxacin and norfloxacin. A cross sectional study was conducted on 249 apparently healthy slaughtered goats at the municipal abattoir of Dire Dawa to determine the antimicrobial susceptibility pattern of the isolates. Of all the isolates, 41 (93.2 per cent) were multiple antimicrobial resistant and the highest level of resistant was observed for tetracycline (100 per cent). In their study, isolates of *Salmonella* were showing highest sensitivity to ciprofloxacin [17].

Out of four *Salmonella* isolates from beef 100 per cent sensitive to amoxicillin, cephalixin, cefotaxime, perfloracin and sulphamethizole, 100 per cent were resistant to metrinadazole, 75 per cent were resistant to norfloxacin and tetracycline, 50 per cent were resistant to gentamicin, nalidixic acid and 25 per cent were resistant to ciprofloxacin.

[6] Author cultured the *Salmonella* from beef samples in Nigeria, the isolates were moderately sensitive to ciprofloxacin, sulphademetoxazole, chloramphenical. The highest sensitivity was observed for gentamicin and the isolates were completely resistant to tetracycline. Several studies have documented the high level of resistance of *Salmonella* to the tetracycline [45, 24] which is in agreement with the result obtained in this study. The high resistance of

tetracycline observed in the study could be as a result of the use of tetracycline in food animal production which has led to worldwide spread of tetracycline resistance observed in *Salmonella* isolates [43, 25, 32].

In this study the resistance of ciprofloxacin, nalidixic acid and cephalosporin has also been reported because these antibiotics are commonly used in veterinary and human medicine. *Salmonella* resistance to the fluoroquinolones (ciprofloxacin) is of great concern to public health as invasive forms of *Salmonellosis* are treated with these compounds [19, 43, 24]. The absence of resistance to the fluoroquinolones by *Salmonella* serovars from foods in Nigeria has been documented [33]. Even though the sensitivity with cefotaxime was noticed, it is quite costlier than other commonly available antibiotics. Chloramphenicol is not frequently used nowadays due to many side effects but in this study complete sensitivity of chloramphenicol is noticed. The occurrence of multi-drug resistance *Salmonella* from food is of global health concern as this could lead to major healthcare challenge since multidrug resistance hinders the possibility of therapeutic treatments. Nowadays most of the human strains are resistant to chloramphenicol, sulphonamides and tetracycline. Recent resistant additions include resistance to trimethoprim and also particular concern to the fluoroquinolones [46]. Equally the use or misuse of antibiotics in human for example also leads to the development of antibiotic resistant. Drug resistant *Salmonella* emerge in response to antimicrobial usage in humans and in food animals. So, selective pressure from the use of antimicrobials is a major driving force behind the emergence of resistance.

Conclusion

Contamination of meat with *Salmonella* observed in this study might be attributed to the unhygienic and improper handling of meat during slaughter, dressing and evisceration. The usual practice of washing the carcass in the same water, in which the intestines and offal had been washed, was considered as one of the predominant reasons for the increased prevalence of *Salmonella* in the carcasses. The knowledge on hygienic

meat production among the butchers was found poor. Henceforth, there is an urgent need for educating the butchers regarding the significance of *Salmonellosis* and its public health significance and also regarding clean meat production. Proper cooking and hygienic measures need to be followed by the consumer to avoid these organisms. There is also a need for a comprehensive epidemiological study and control of *Salmonella* contamination at various levels of meat production and slaughter points in Puducherry.

Table 1: Sources of raw meat samples

S. No	Place of sample collection	Number of Samples		
		Chicken	Chevon	Beef
1	Mettupalayam	2	3	-
2	Muthirapalayam	2	3	-
3	Kurumbapet	2	-	-
4	Dharmapuri	2	-	-
5	Kathirkamam	2	2	-
6	Thilasupet	1	2	-
7	Thattanchavady	2	3	-
8	Kalapet	1	3	3
9	Lawspet	2	3	-
10	Iyankuttipalayam	2	-	-
11	Reddiyaralayam	2	2	1
12	Moolakulam	1	1	1
13	Manaveli	2	2	2
14	Arumbathapuram	2	2	2
15	Ariyankuppam	2	4	1
16	Thavalakkuppam	-	3	6
17	Othianpet	2	1	4
18	Koonichampet	2	2	4
19	Pillaiyarkuppam	2	1	1
20	Karikkalampakkam	2	1	4
21	Bahour	3	3	1
22	Thirukkanur	3	3	4
23	Thirubuvanai	2	1	4
24	Vazhuthavur	2	1	-
25	Sedarapet	2	1	2
26	Sulthanpet	3	3	10
	Total	50	50	50

Table 2: List of antibiotics used in this study.

Sr.no	List of antibiotics	Content/disk
01	Amoxycillin	10µg
02	Azithromycin	15µg
03	Co – trimoxazole	30µg
04	Cephalexin	30µg
05	Ciprofloxacin	30µg
06	Chloramphenical	30µg
07	Cefotaxime	10µg
08	Gentamicin	120µg
09	Metronidazole	50µg
10	Norfloxacin	15µg
11	Nalidixic acid	30µg
12	Perfloxacin	5µg
13	Sulphamethizole	30µg
14	Trimethoprim	25µg
15	Tetracycline	30µg

Table 3: Comparison of RVS and SB broth

Samples (n=150)	Salmonella		
	Total isolates	RVS	SB
Chicken (50)	5 (7.4%)	3 (60%)	2 (40%)
Chevon (50)	2 (4%)	2 (4%)	-
Beef (50)	4 (8%)	4 (8%)	-

Table 4: Primary and secondary biochemical test results

Characteristics	Sample No.											
	Chicken					Chevon			Beef			
	1	6	7	14	21	1	16	17	23	26	25	
Primary and secondary biochemical tests												
Shape	R	R	R	R	R	R	R	R	R	R	R	R
Motility	+	+	+	+	+	+	+	+	+	+	+	+
Catalase	+	+	+	+	+	+	+	+	+	+	+	+
Oxidase	-	-	-	-	-	-	-	-	-	-	-	-
Urease	-	-	-	-	-	-	-	-	-	-	-	-
Simmon's citrate	+	+	+	+	+	+	+	+	+	+	+	+
Indole	-	-	-	-	-	-	-	-	-	-	-	-
MR Test	+	+	+	+	+	+	+	+	+	+	+	+
VP Test	-	-	-	-	-	-	-	-	-	-	-	-
H2S production in TSI slant	+	+	+	+	+	+	+	+	+	+	+	+
Lysine decarboxylase	+	+	+	+	+	+	+	+	+	+	+	+
Ornithine decarboxylase	+	+	+	+	+	+	+	+	+	+	+	+
ONPG	-	-	-	-	-	-	-	-	-	-	-	-
Sugar fermentation tests												
Dulcitol	+	+	+	+	+	+	+	+	+	+	+	+
Dextrose	+	+	+	+	+	+	+	+	+	+	+	+
Maltose	+	+	+	+	+	+	+	+	+	+	+	+
Xylose	+	+	+	+	+	+	+	+	+	+	+	+
Trehalose	+	+	+	+	+	+	+	+	+	+	+	+
Arabinose	+	+	+	+	+	+	+	+	+	+	+	+

+ : Positive; - : Negative

Table 5: Primer sequence for *invA* gene

Organisms	Target gene	Primer sequence (5'- 3')	Expected Product size (bp)	References
<i>Salmonella</i> spp	<i>invA</i>	F:GTGAAATTATCGCCACGTTCCGGCAA R:TCATCGCACCGTCAAAGGAACC	244	Rahn <i>et al.</i> (1992)

Table 6: Details of the Reaction mixture

Sr. No	Reaction mixture	Salmonella (<i>invA</i>)
1	Template DNA	5 µl
2	Primer	20 pmol each primer
3	Master mix	12.5 µl
4	Triple distilled water	5.5 µl

Table 7: Details of the PCR conditons

Sr. No	PCR Programme	Salmonella (<i>invA</i>)
1	Initial Denaturation	95 °C for 5 minutes
2	Denaturaation	94 °C for 30 seconds
3	Annealing	56 °C for 30 seconds
4	Extension	72 °C for 2 minutes
5	Final extension	72 °C for 5 minutes

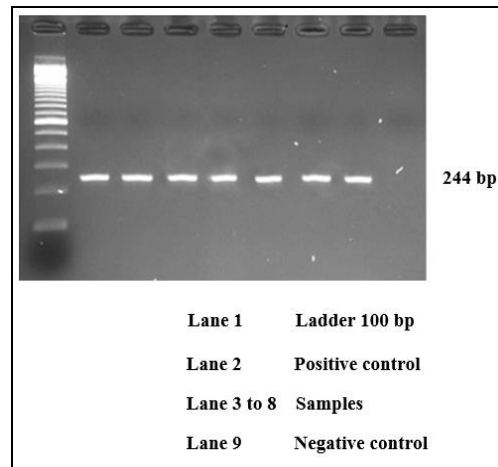


Table 8: Consolidated antimicrobial sensitivity pattern (n=11)

List of antibiotics	Chicken (%)			Chevon (%)			Beef (%)		
	R	I	S	R	I	S	R	I	S
Amoxycillin	100	-	-	100	-	-	100	-	-
Azithromycin	-	-	100	-	-	100	-	25	75
Co – trimoxazole	80	20	-	100	-	-	100	-	-
Cephalexin	-	20	80	-	-	100	-	-	100
Ciprofloxacin	20	20	60	50	-	50	25	75	-
Chloramphenical	-	20	100	-	-	100	-	25	75
Cefotaxime	-	-	100	-	-	100	-	-	100
Gentamicin	20	20	60	50	-	50	25	25	50
Metronidazole	100	-	-	100	-	-	100	-	-
Norfloxacin	60	20	20	50	50	-	75	25	-
Nalidixic acid	40	60	-	50	50	-	25	-	75
Perfloxacin	-	40	60	-	50	50	-	-	100
Sulphamethizole	20	20	60	50	-	50	-	-	100
Trimethoprim	20	-	80	-	-	100	50	-	50
Tetracycline	60	-	40	-	-	100	50	-	50

R – Resistant; S- Sensitive and I – Intermediate

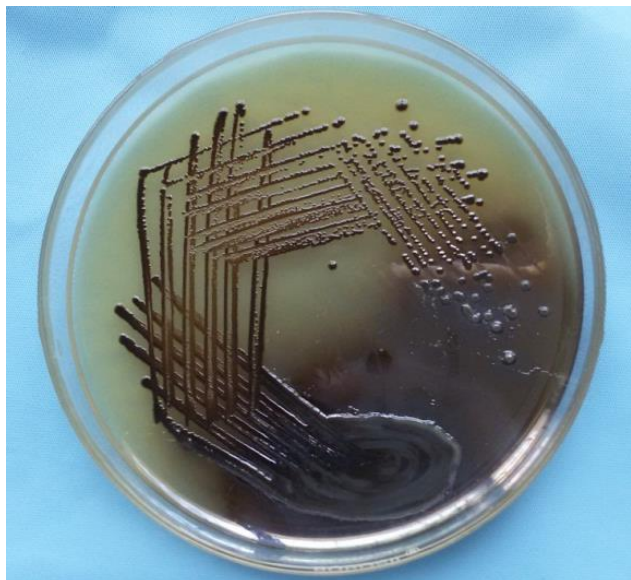


Plate 1: Salmonella – BSA

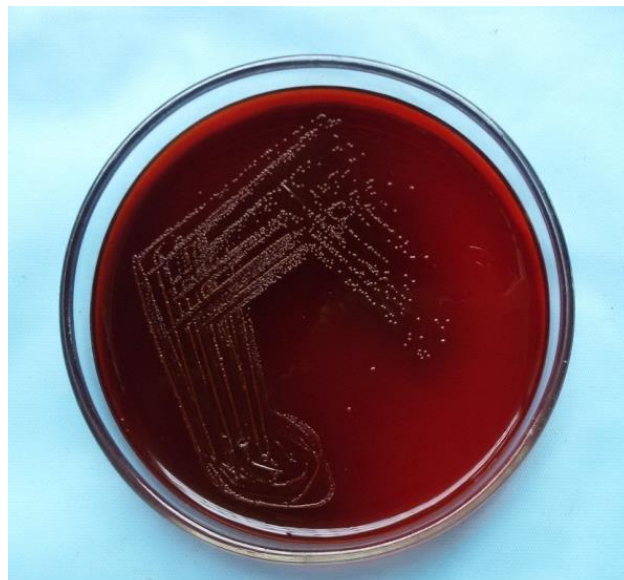


Plate 4: Biofilm positive

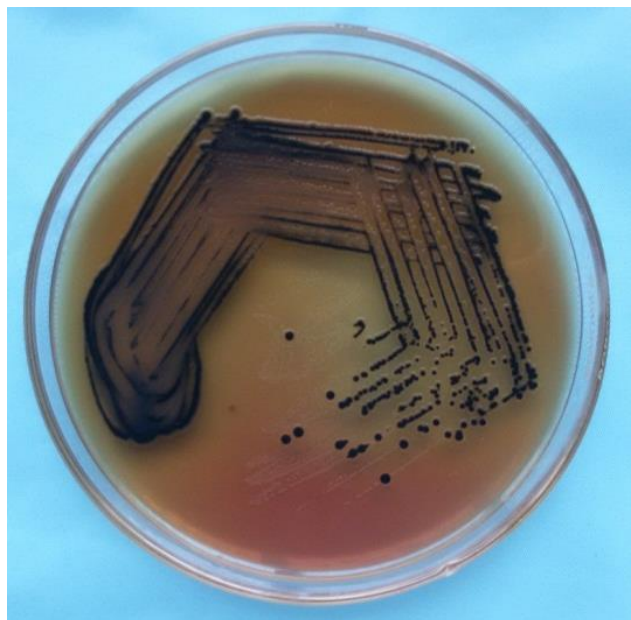


Plate 2: Salmonella – SSA



Plate 5: Biofilm negative



Plate 3: Salmonella – XLD

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