



## Original Research Article

# Molecular analysis of Virulence of field isolates of *Salmonella typhimurium* isolated from poultry in Tarai region (Uttarakhand State) of India

B. H. Jagadish and M. K. Saxena\*

College of Veterinary and Animal Sciences, G.B. Pant University of A & T,  
Pantnagar, (U.A), India

\*Corresponding author

## ABSTRACT

### Keywords

Plasmid,  
Salmonella,  
*vir* gene

In the pathogenic life cycle of a number of microbes invasion of eukaryotic cells is an essential step. In case of enteric bacteria invasion is the critical step for systemic infection. For *Salmonella typhimurium* invasion decides whether it is less or more pathogenic. We here report variation in the virulence of *Salmonella typhimurium* isolates in terms of their ability to invade the chick embryo fibroblast CEF. Further the role of plasmid in invasion and the presence of *vir* gene and its value in detection of virulent strains of *Salmonella typhimurium*. We also report that, CEF can be very well used as an *in vitro* model for the invasion studies.

## Introduction

Virulence is defined as the relative capacity of microbes to cause damage in host. Virulence is dependent on different independent variables that includes the microbial aggressiveness, invasiveness, infectivity, toxigenicity and communicability (Lipsitch and Moxon, 1997). Many of the pathogens' virulence exclusively depend on single most critical virulent factor. In case of *Salmonella typhimurium* invasion of gastrointestinal mucosa is an essential and crucial step for pathogenesis and strains unable to invade animal cells are avirulent (Giannella *et al.*, 1973).

Characterization of virulence of different isolates of *Salmonella typhimurium* in terms of their ability to invade eukaryotic

cells is one approach that may enable a better understanding of their pathogenicity. Cultured eukaryotic cell lines are commonly used to study bacterial invasion since they provide reproducible and uncomplicated infection models (Tang *et al.*, 1993)

In the present study, nine isolates of *Salmonella typhimurium* which were isolated from poultry farm of Tarai region were studied for their virulence status.

These isolates were screened for plasmid and their invasion in chicken embryo fibroblast cells. Gene was amplified and findings were correlated between invasion rate, plasmid profile and "vir" gene amplification.

## Materials and Methods

### Bacterial isolates

Nine isolates of *Salmonella typhimurium* used in this study are maintained at the Animal biotechnology Centre, Department of Veterinary Biochemistry, College of Veterinary and Animal Sciences, G.B. Pant University of Agri. & Tech. Pantnagar, (U.A), India. These isolates were isolated from poultry farms of Tarai region of Uttarakhand state. The isolates were antigenically characterized at National *Salmonella* Centre, Division of Bacteriology, IVRI, Izatnagar, UP, India.

The isolates were grown overnight at 37°C in Luria Bertani broth (LB broth) and were sub cultured into shaking L.B broth culture until the late-logarithmic phase of growth (Wavelength 600nm).

### Preparation of Chicken Embryo Fibroblast (CEF) culture

Monolayer required for bacterial invasion study were prepared from 10-11 day old chicken embryos procured from flocks free of *Salmonella* from Instructional Poultry Farm (IPF), GBPUA&T, Pantnagar. Primary CEF culture was prepared as per the method described by Tang *et al.*, 1993 with some modifications. The embryos were collected aseptically and placed in Hank's Balances Salt Solution (HBSS, Himedia, India). After removal of head, extremities and visceral organs, the remaining tissue was washed and minced using a sterile scissors. Then the tissue was digested with 0.25% trypsin solution (Himedia, India) for 20 min. with constant stirring. The cells were filtered through sterilized muslin cloth and centrifuged at 800 g for 5min. at 4°C. The cells were then washed in HBSS and finally in minimum essential medium (MEM,

Himedia, India). The washed cells were re-suspended and counted using Neubauer's chamber under a microscope. Finally the cell concentration was adjusted to 10<sup>7</sup> cells per ml in MEM containing 10% fetal calf serum (Himedia, India) and distributed in to 6-Well culture plates.

### Bacterial infection of fibroblast

Infection of chicken embryo fibroblasts with *Salmonella typhimurium* isolates was carried out according to the protocol of Cano *et al.*, 2001 with some modifications. Briefly, CEF were grown in MEM medium containing 10% fetal calf serum to 30% confluence at the time of infection. Bacteria were grown overnight in the LB broth at 37°C under static conditions without shaking. The multiplicity of infection was 10 bacteria per fibroblast cell, and the infection time and pH were standardized for maximal invasion. Infection time were 15min, 30min, 45min, 1hr 15min and 2hr, and the pH selected were 4.5, 6.5, 7.5 and 8.5. Infected cells were washed three times with phosphate buffer saline (PBS pH 7.4) and incubated in fresh medium containing 100µg of Gentamicin/ml. The antibiotic concentration was decreased to 10µg/ml at 2 h post infection. After 2 h the media was removed and the monolayer was washed with PBS and was exposed to 1% Triton-X to lyse the cells and to recover the bacteria that were in side the cells, the lysates were poured on BGA plates for finding the viable count which gave the estimate of invasion of given isolate.

### Isolation of genomic DNA and 'vir' PCR

For isolation of genomic DNA, the bacteria were grown overnight at 37°C for 18hr in 25 ml LB Broth with constant shaking in shaker cum incubator. Genomic DNA was isolated by the C-TAB method (Wilson (1987)).The

protocol for 'vir' PCR was carried out as per the method of Rexach *et al.* (1994) and Saxena *et al.* (2004) with some modifications.

Bacterial DNA was amplified with a programmable Thermal cycler (Biometra). Taq. polymerase (1.5U) was added to 25µl of a reaction mix containing 1-X assay buffer (1.5mM MgCl<sub>2</sub>), 200µM of each dNTPs, 20 pico mole of each primer and 40ng template of target DNA. PCR was performed with initial denaturation at 94°C for five minutes, followed by 30 cycles of denaturation for one minute at 94°C, annealing at 65°C for 1min, polymerization at 72°C for 1min, and final polymerization for 5min. Negative control reaction was set up in each replica, containing all reagents except for the template DNA.

### **Plasmid isolation**

The plasmid DNA was isolated and electrophoresed in 0.8% agarose gel as described in Sambrook *et al.* (1990).

The molecular weight of plasmid was calculated by comparing with the Marker plasmid (*E. coli* MTCCV517).

## **Results and Discussion**

### **The invasiveness of *S. typhimurium* in CEF**

For this experiment ML36, a representative isolate of *Salmonella typhimurium* was taken and the invasiveness was checked at different time of exposure, viz, 15min, 30min, 45min and 1hr 15min (Table 1). The results of this experiment indicate that *Salmonella typhimurium* need 45min for maximum invasion.

For determining the optimum pH for

invasion time was kept at 45 min and pH was varied (Table 2). It indicates that pH 6.5 is optimum for invasion of CEF by *Salmonella typhimurium*.

### **Invasiveness among the isolates of *Salmonella typhimurium* varies significantly:**

In present study we have differentiated different isolates of *Salmonella typhimurium* based on their ability to invade CEF (Table 3).

Comparative study of invasiveness was made for nine different isolates of *Salmonella typhimurium* (Table 3). As per the observations, the isolate ML7 showed highest invasion and ML25 least invasion. The degree of invasiveness between least and maximum invading isolates was about 200 times.

### **Plasmid isolation**

The plasmid of (32.2kD) was isolated from six isolates ML4, ML6, ML7, ML14, ML23, and ML36. ML11 has small plasmid of approximately 2 kDa. In remaining two cultures there was no plasmid.

### **'vir' gene PCR**

All the nine isolates were tested for the presence of 'vir' gene by PCR which on amplification with specific primers produced a product of approximately 530bp in 7 of the 9 isolates. The isolates ML5 and ML25 were not positive for 'vir' gene PCR (Fig. 1).

Invasion of eukaryotic cells is an essential step in the pathogenic life cycle of a number of microbial pathogens. Different organism has evolved different strategies to enter and survive inside the eukaryotic host cells (Falkow, 1991). A number of enteric

bacteria including *Salmonella* have developed the ability to penetrate non phagocytic cells. This ability of *Salmonella* constitutes a very important pathogenic/virulent property. The optimal pH and time of pH 6.5 and 45 min obtained for maximum invasiveness in this study is in the agreement with earlier reported findings. Similar study in bovine suggested bovine halt of bacterial interaction with follicle-associated epithelium by 60 minutes. Time dependence invasiveness has greater relevance for deciding the incubation period of a pathogen in host organism. *Salmonella* invasion is receptor mediated; therefore a fixed number of receptor on membrane allows a maximum number of organisms to enter in to host cells at a given point of time

The first major stressful environment that *Salmonella* encounters after an oral infection is exposure to acidic gastric contents. Acid tolerance may contribute to virulence of *S. typhimurium* (Humphrey *et al.*, 1996). Adaptation to specific pH range is likely to be an important variable in *Salmonella* pathogenicity, since pH has been identified as a regulator of virulence gene expression in animal pathogens. In slightly different study was also carried out in *S. Enteritidis* in which different isolates were tested for acid tolerance at pH 4 and they found wide variation in their ability to tolerate this pH and they also reported that the isolates with higher acid tolerance were more virulent than those with acid intolerance isolates (Lu *et al.*, 1999). In *S. Typhi* invasion at pH of 5 and 6.5 in human intestinal epithelial cell culture reveal that the *S. Typhi* has higher invasion rate at 6.5 pH than at pH 5. This variation is mainly because, the expression of gene responsible for invasion are pH dependent. For example, the expression of the *invF*, *pagC*, *prgH*, *prgK*, *orgA*, *sspA*, and *sspC* genes of *S. typhimurium* (Bajaj *et al.*, 1996) and the *inv*

gene of *Y. enterocolitica* (Pepe *et al.*, 1994) are regulated by the pH of the immediate environment. Our study supports these findings.

Invasion of epithelial cells is a characteristic of *Salmonella* associated with virulence. The comparative study of invasiveness of different isolates of *S. typhimurium* in CEF monolayer revealed that, within a given serovar there is great variation in the invasiveness. We found almost 200 times difference in degree of invasiveness between lowest and highest invasiveness obtained. Similar experiment performed earlier in *S. typhimurium* reported variance in invasiveness from 51.80 to 36.80 percent with the different isolates in CEF cells. Difference in invasion between different isolates was also found in *S. Cholerasuis* in MDCK epithelial cells monolayer and *S. Typhi* in HeLa cell monolayer. Similarly the variation in the invasion ability of different isolates was also reported by Lu *et al.*, 1999 in *Salmonella* Enteritides. The variation in the ability to invade the cultured mammalian cells is accounted for the presence of virulence plasmid (Galan *et al.*, 1989) and SPI1. An operon (*spvRABCD*), containing five genes, is present on plasmids commonly associated with some serovars. The *spv* genes possibly have the ability to increase the severity of enteritis and allow infection and persistence at extra-intestinal sites (Libby *et al.*, 2000). The chromosomally located invasion gene *invA* is thought to trigger the invasion of *Salmonellae* in to cultured epithelial cells. The locus of this gene is in SPI1. Hence the presence or absence of plasmid and SPI1 is considered responsible for the variation found in the invasion rate. In our study we found the isolates without the large plasmid had low invasion rate was found less compared to the isolates with the plasmid. The earlier findings also support these observation viz.

ability of virulent *Salmonella typhimurium* strains to adhere to and invade HeLa cells, and to kill mice following oral administration, depends on the presence of 92kbp large plasmid (Jones *et al.*, 1982).

PCR has been used for detection of pathogen (Shivchandra *et al.*, 2006, Shivshankar *et al.*, 2001) for molecular epidemiology (Saxena *et al.*, 2006) and for detection of virulence (Saxena *et al.*, 2006) in bacteria. Virulence associated gene in the plasmid is detected by the amplification of approximately 530bp product with vir gene specific primers. The presence of gene linked to virulence in the isolates of *Salmonella typhimurium*. All the isolates were tested for the presence of ‘vir’ gene, out of 9 isolates 7 isolates (ML4, ML11,

ML6, ML7, ML14, ML23, ML36) showed amplification of a single product of approximately 530 bp. The amplification was observed only in those isolates in which plasmid was present. Virulence associated gene has been reported in Indian isolates of *S. Gallinarum* and British isolates of *S. typhimurium* (Jones *et al.*, 1982). Similarly, plasmids encoding ‘vir’ gene in isolates of *Salmonella* was also reported (Saxena *et al.*, 2004). However, it has been reported that every plasmid bearing serovars are not necessarily the carriers of virulence gene (Boyd *et al.*, 1998). In ML 11 plasmid of low molecular weight was obtained but “vir” gene was present. This isolate also exhibit low invasion, so we conclude that there is a positive correlation between large plasmid and invasion rate.

**Table.1** The invasiveness of *S. typhimurium* in CEF at different times of infection

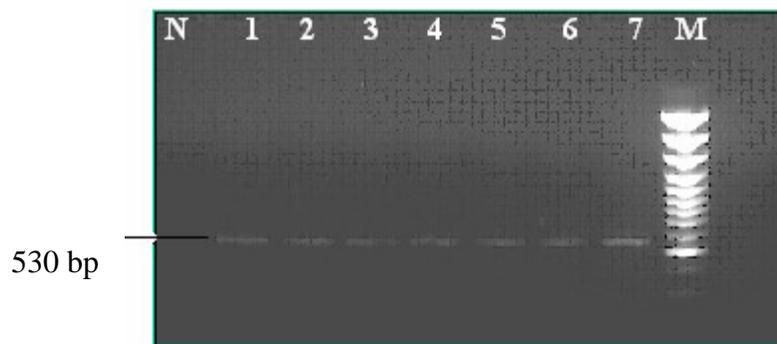
| Isolate | Number of CEF cells. | Number of bacteria inoculated | Recovered intracellular viable bacteria at different exposure period |                   |                   |                   |
|---------|----------------------|-------------------------------|--|-------------------|-------------------|-------------------|
|         |                      |                               | 15min  | 30min             | 45min             | 75min             |
| ML-36   | $4 \times 10^7$      | $40 \times 10^7$              | $1.3 \times 10^3$  | $9.3 \times 10^3$ | $2.1 \times 10^4$ | $1.4 \times 10^4$ |

**Table.2** The effect of pH on the invasion rate of *S. typhimurium* in CEF

| Isolate | Number of CEF cells | NO, of bacteria inoculated | Colonies on BGA (colonies/100µl of lysate) |                    |                   |                   |
|---------|---------------------|----------------------------|--|--------------------|-------------------|-------------------|
|         |                     |                            | 4.5  | 6.5                | 7.5               | 8.5               |
| ML36    | $4 \times 10^7$     | $40 \times 10^7$           | $1.8 \times 10^2$                          | $2.19 \times 10^3$ | $7.5 \times 10^2$ | $1.6 \times 10^3$ |

**Table.3** Comparative invasion rate of different isolates of *S. typhimurium* in CEF

| Isolate | Number of CEF cells. | Number of bacteria infected. | Recovered intracellular viable bacteria(CFU/100µl) | Recovered intracellular viable bacteria (CFU/ml) |
|---------|----------------------|------------------------------|--|--|
| ML4     | $4 \times 10^7$      | $40 \times 10^7$             | 500  | $5.0 \times 10^3$                                |
| ML6     | $4 \times 10^7$      | $40 \times 10^7$             | 350  | $3.5 \times 10^3$                                |
| ML5     | $4 \times 10^7$      | $40 \times 10^7$             | 30   | $3.0 \times 10^2$                                |
| ML7     | $4 \times 10^7$      | $40 \times 10^7$             | 3300   | $3.3 \times 10^4$                                |
| ML11    | $4 \times 10^7$      | $40 \times 10^7$             | 20   | $2.0 \times 10^2$                                |
| ML14    | $4 \times 10^7$      | $40 \times 10^7$             | 1200   | $1.2 \times 10^4$                                |
| ML23    | $4 \times 10^7$      | $40 \times 10^7$             | 440  | $4.4 \times 10^3$                                |
| ML25    | $4 \times 10^7$      | $40 \times 10^7$             | 18   | $1.8 \times 10^2$                                |
| ML36    | $4 \times 10^7$      | $40 \times 10^7$             | 2100   | $2.1 \times 10^4$                                |
| Control | $4 \times 10^7$      | $40 \times 10^7$             | No Colonies  | No Colonies                                      |



N-Negative, 1-ML4, 2-ML36, 3-ML6, 4-ML7,  
5-ML11, 6-ML14, 7-ML23,  
M- marker 100 bp plus ( Fermentas)

Fig.1

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