Host responses to pathogenic *Salmonella* infection

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Exposure to bacteria, fungi, and protozoa is an inescapable part of life from the moment that we are born until we die. Our contact with many of these microorganisms is very brief, whereas others stay behind and establish a more stable relationship. These organisms become part of our normal flora and are found, among other places, on the skin, in the mouth, and in the gastrointestinal tract. Typically these relationships are beneficial to both the host and the microorganism. Although the commensal flora appears to provide several benefits, it can also, under certain conditions, represent a significant threat to our health. The immune system has the primary responsibility of ensuring that the growth and location of commensal organisms is optimal for the host. If defensive immunity is compromised or fails in any way then benign commensals quickly become opportunistic pathogens.

Some of the microorganisms that we encounter rely upon their ability to cause cellular damage to a host for survival and replication. Upon entering a suitable host, these pathogens initiate a series of events that test innate host defense mechanisms and the capabilities of the immune system. From the perspective of the microorganism, these interactions challenge the ability of the bacteria to respond to constantly changing obstacles within the host environment. This review will focus on the interactions between virulent *Salmonella* strains and the hosts that they infect as an example of how selective pressure helps to shape host-pathogen relationships. Efforts will be made to highlight the various immunological and cellular host responses that occur in response to *Salmonella* infection and invasion.

*Salmonella* infections are initiated when a pathogenic strain is ingested into a suitable host via contaminated food or water. Ingested organisms move from the mouth to the extremely acidic environment of the stomach. Organisms that survive this low pH environment then move into the small intestine where they begin to compete with the normal enteric flora of the host for nutrients and space. The first pathogenic step of the bacteria is taken when organisms breach the intestinal epithelium of lymphoid follicles and enter the lymphatic system, sequentially moving from the Peyer's patch tissue to the lymph nodes and then to the spleen and liver. At each step of infection these pathogens respond to changing host environments by expressing appropriate factors that ensure their survival. The ability of these sophisticated pathogens to respond to changing host conditions places a heavy burden on the host as it must also adapt to ensure its own well being. Host cell responses to bacteria provide clues to the mechanisms used by the bacteria to infect the host and provide insight into possible mechanisms that the host might invoke to engage and eliminate the invading pathogens.

**Tissue culture cell responses to Salmonella**

Understanding the virulence mechanisms of bacterial pathogens depends upon the ability to directly examine host–bacterial interactions. Often, the lack of suitable models has made research on bacterial pathogens difficult. Fortunately, the development of an in vitro tissue culture model (Jones et al. 1981) has largely overcome this problem for the study of the role of *Salmonella* invasion in pathogenesis. Importantly, many of the interactions that occur between invasive *Salmonellae* and mammalian tissue culture cells duplicate the changes that have been observed by microscopy in *Salmonella*-infected host tissue (Takeuchi 1967; Finlay and Falkow 1990; Francis et al. 1992). In vitro tissue culture studies have identified some of the host cell responses that lead to *Salmonella* entry (or occur as a result of entry) and include actin rearrangement and polymerization at the host cell membrane, accumulation of cytoskeletal proteins at the site of bacterial entry, and calcium fluxes. A prominent, visible feature of the *Salmonella* entry process is the induction of membrane rearrangements or ruffles at the site of bacterial attachment (Ginocchio et al. 1992; Francis et al. 1993; Jones et al. 1993) [Fig. 1]. These protrusions occur on the membrane at the site of bacterial attachment and are presumed to occur in direct response to a bacterial signal. Time-lapse video microscopy has demonstrated that these membrane ruffles erupt from the host cell within seconds of bacterial attachment to the cell. Subsequently, the columns of polymerized actin surround and enclose the organism. Within a short period of time, the entry signal peaks and, as it dissipates, the membrane ruffles return to the cell.
along with the enveloped bacterium (Francis et al. 1992). The formation of the membrane ruffles is dependent on actin monomer polymerization into microfilaments, as specific inhibitors [cytochalasins] [Tanenbaum 1978; Forscher and Smith 1988] of actin polymerization prevent membrane rearrangements and block Salmonella entry [Kihlström and Nilsson 1977; Finlay and Falkow 1988; Mroczenski-Wildey et al. 1989].

Efforts have been made to identify the signal transduction pathway of the cell that leads to uptake of the bacteria by using specific inhibitors. The protein kinase inhibitor genistein [Akiyama et al. 1987] blocks entry of Yersinia and enteropathogenic Escherichia coli into cells but has no effect on the invasion of Salmonella typhimurium into a variety of different cell lines [Rosenshine et al. 1992, 1994]. Recent work has demonstrated that Listeria entry can be blocked by wortmannin and LY294002, inhibitors of phosphoinositide 3-kinase, but no effect was observed on S. typhimurium internalization [It et al. 1996]. These experiments have helped to identify signal transduction pathways that are not involved in Salmonella entry but have provided only limited information on the actual pathway that is stimulated.

Increased quantities of cytoskeletal proteins, in addition to actin, are also found at the site of bacterial entry. These proteins include α-actinin, ezrin, talin, tropomysin, and tubulin, although the mechanism of accumulation of these proteins is unknown [Finlay et al. 1991]. The induction of intracellular calcium fluxes has also been reported in cells that have been exposed to invasive S. typhimurium [Ruschkowski et al. 1992; Pace et al. 1993]. It is unknown whether these calcium changes are an integral part of the signal pathway to actin or foils that are scattered throughout the small intestine. This lymphoid tissue, also called Peyer’s patches, has a specialized epithelium and an underlying dome that contains mature lymphocytes as well as developing lymphoblasts. The primary function of this tissue is believed to be immune surveillance of the gut. Invasive Salmonella strains specifically target these Peyer’s patches for the initial penetration of the host small intestinal tissue. Early infection studies with mice demonstrated that virulent S. typhimurium, administered by an oral route, associated almost exclusively with Peyer’s patch tissue of the terminal ileum [Carter and Collins 1974] as quickly as 3 hr postinoculation [Hohmann et al. 1978]. M cells are specialized residents of the epithelium of lymphoid follicles and are interspersed within the epithelial layer of an intestinal lymphoid follicle at a ratio of approximately 1 per 10–20 enterocytes [Owen and Er PAUK 1990]. M cells have several distinctive characteristics that allow them to be identified easily [Owen and Jones 1974; Bockman 1983]. These cells possess microvilli that are visibly shorter than those of enterocytes. They also lack a rigid cytoskeleton that allows migratory lymphocytes to deform and distort their cytoplasm. In addition, they have an elevated pinocytic activity that facilitates the uptake of intestinal microorganisms and particles. This suggests that a primary function of these cells is to transport and process luminal antigens to prime local intestinal immunity in the Peyer’s patches [Jones et al. 1995]. M cells also share similarities with other epithelial cells in that they form tight junctions with adjacent enterocytes and align along the basal lamina surface.

Because M cells are unique to follicle-associated epithelia and S. typhimurium strains target lymphoid follicles for the initial penetration of host tissue, one strategy to study early interactions between Salmonella and host cells has been to look for interactions of the bacteria with M cells at the earliest stages of infections. These studies typically utilize an intestinal ligated loop model system. In this system, a section of intestine of an anesthetized animal is tied off and injected with a pathogenic bacteria, which are then allowed to interact with the host tissue. Experiments using a murine-ligated model demonstrated that invasive S. typhimurium is able to invade M cells within the epithelium of the murine lymphoid follicles within 30 min of infection.

**Intestinal epithelium responses to Salmonella**

The first specific interactions during natural infection between pathogenic Salmonella and a host occur at the epithelial surface of organized patches of lymphoid tis-
Salmonella-host cell interactions (Clark et al. 1994; Jones et al. 1994) (Fig. 2). The bacteria display a clear preference for M cells, as no interactions can be detected with enterocytes at these early time points. Importantly, the entry of Salmonella into murine M cells occurs by a membrane-ruffling mechanism that appears to be identical to Salmonella entry into tissue culture cells. The pathogenic effect of the bacteria on the tissue was increased at 60 min postinfection (Jones et al. 1994). Dying and disintegrating M cells that contained bacteria were observed detaching from the mucosal surface. This damage allowed organisms that had been confined to the lumen of the small intestine access to underlying tissue. At later time points (120 and 180 min), significant damage to the intestinal epithelium could be observed that included invasion of enterocytes, sloughing of large sections of epithelial cells, and penetration of significant numbers of bacteria into deeper tissues.

Experiments performed to determine whether the presence of invasive S. typhimurium in the murine intestine has an effect on the cellular composition of Peyer’s patches showed that the presence of the invasive pathogen caused a number of different host responses in the Peyer’s patches (Savidge et al. 1991). The total number of M cells increased in the follicle-associated epithelium as compared to uninfected mice, the average crypt depth lengthened, and the rate of enterocyte migration from the intestinal crypts increased. In addition, the numbers of CD4+ cells increased and CD8+ cells decreased. These results indicate that the damage elicited by invasive Salmonella induces a host response in the Peyer’s patch tissue. Although the pathogen causes significant damage to the epithelium of lymphoid follicles at early points of infection, the findings of this study indicate that the host quickly replaces cells that have been damaged and/or destroyed. In addition, it is clear that the immune system is activated by and responds to the presence of Salmonella by activating both humoral and cellular immunity (Mastroeni et al. 1993).

Figure 2. Scanning electron microscopy of the apical membrane of murine lymphoid follicles. (A) Uninfected Peyer’s patch tissue. The M cells have visibly different morphologies compared to the enterocytes, owing to the difference in the height of their microvilli. Two M cells are indicated by the arrows, although several other M cells are also visible. Magnification, 2250x. (B) Murine intestinal tissue infected for 30 min with invasive S. typhimurium. Substantial membrane rearrangements are visible at the surface of the Peyer’s patch. The cells with membrane rearrangements are M cells, as interactions with enterocytes are not detected at this time point. The cell indicated by the arrow to the far left is clearly an M cell and has a small, round ruffle protruding from the surface. Magnification, 2250x.
These studies highlight the importance of correlating in vitro findings with in vivo observations. *Salmonella* entry into either M cells or tissue culture cells occurs by a membrane-ruffling mechanism. However, the bacterial entry event typically does not kill tissue culture cells but clearly induces the death of M cells that reside in the epithelium of Peyer’s patches. Furthermore, in vivo studies of infection have provided first-hand observations of the progression of *Salmonella* infections that can result in systemic disease and the host cell response that leads to regeneration of host tissue and activation of immune defenses. Current research efforts are focused on understanding how and why invasive *Salmonella* selectively enter M cells and determining the molecular mechanism of *Salmonella*-induced M cell death.

**Phagocytic cell responses and interactions**

Following passage through the intestinal epithelium of the Peyer’s patch, invading organisms quickly enter the lymphatic system where interactions with professional killing cells determine the ultimate fate of the infection. These cells possess both oxygen-dependent and -independent killing mechanisms to kill internalized bacteria [Falkow et al. 1992]. The production of toxic oxygen molecules such as superoxide, hydrogen peroxide, and hydroxy radicals that are pumped into the phagolysosome are the primary oxygen-dependent killing molecules of macrophages. The oxygen-independent killing mechanisms of the macrophages include acidification of the phagolysosome as well as secretion of small bactericidal peptides into the compartment.

Because the oxygen-dependent killing mechanisms of macrophages appear to have little effect on pathogenic *Salmonella*, the host is more dependent on oxygen-independent killing mechanisms. However, experimental evidence suggests that *Salmonella* has evolved mechanisms to circumvent or delay the killing activity of these mechanisms. For example, upon entry into macrophages, *S. typhimurium* delays significantly the fusion of the phagosome to the lysosome [Buchmeier and Heffron 1991]. The majority of viable and dividing organisms observed by microscopy were found within unfused vacuoles, whereas nonpathogenic *E. coli* and dead *Salmonella* organisms were predominantly within fused phagolysosomes. Acidification of vacuoles containing live *Salmonella* organisms was delayed significantly [4-5 hr] compared to vacuoles that contained dead organisms [Alpuche et al. 1992]. The appearance of vacuole membrane markers, such as lysosomal membrane glycoprotein (lgp), Rab7, Rab9, cathepsin D, and mannose-6-phosphate receptor (M6PR), on normal and *Salmonella*-containing vacuoles has been monitored as a way to examine the effect of pathogenic *Salmonella* on phagolysosome maturation. These protein markers sequentially appear in (and disappear from) the membrane of an uninfected intracellular compartment as it matures from an endosome into a phagosome [Griffiths et al. 1988; Tooze et al. 1990; Chavrier et al. 1991]. The early marker proteins lgp, Rab7, and Rab9 can be found on *Salmo-

nella*-containing vesicles, but two later markers, cathepsin D and M6PR, are excluded [Jones and Falkow 1996]. These results indicate that *Salmonella* begins to modify the vacuole in which it resides after early stages of vesicle maturation but before the appearance of the late-stage markers cathepsin D and M6PR.

The growth state of *S. typhimurium* seems to change rapidly after entry into macrophages [Abshire and Neidhardt 1993]. Immediately after entry of bacteria into phagocytic cells, the intracellular bacteria can be killed by treatment with chloramphenicol and ampicillin, which indicates that the bacteria are growing rapidly and require protein synthesis for survival. However, if the bacteria are allowed to reside within the cells for 2 hr before treatment with the same antibiotics, the bacteria are no longer killed, although net growth is inhibited. This suggests that as infection time increases, the bacteria adapt to the intracellular environment of the macrophage and switch from a rapidly growing state to a survival mode that does not require protein synthesis. Collectively, the data indicate that upon entering the phagocytic vacuole, pathogenic *Salmonella* are intent on altering their environment by delaying and modifying the formation of the phagolysosome. Perhaps, these phagolysosome alterations are necessary to provide the bacteria sufficient time to alter their own physiological state in response to the intracellular environment of the macrophage. Future work is aimed at identifying the bacterial factors, and their mechanism of action, that leads to disruption of normal endosome maturation.

The PhoP/PhoQ is a two-component system that *Salmonella* encodes, which regulates its ability to cause systemic disease and to survive within the intracellular environment of phagocytic cells [Fields et al. 1989; Groisman et al. 1989; Miller et al. 1989]. The PhoP protein is a transcriptional regulator and is a member of the OmpR subgroup of this protein family [Volz 1993]. The PhoQ protein is a sensor protein that has two transmembrane domains and a periplasmic region in its predicted structure that may play a role in the sensing of environmental signals [Miller et al. 1989]. The PhoP/PhoQ transcriptional regulon controls ~40 genes [Garcia-Vescovi et al. 1994] that are essential for various stages of *Salmonella* virulence, including resistance to host antimicrobial peptides [Groisman et al. 1992], passage or invasion through a host cell membrane [Behlau and Miller 1993; Bajaj et al. 1996], and the ability to adapt to and/or modify the phagolysosome [Alpuche et al. 1992]. In fact, it appears that the majority of the virulence functions of *Salmonella* strains is regulated by the PhoP/PhoQ system. It seems fair to speculate that a primary role of this regulatory system is to monitor environmental signals from the host during infection to orchestrate a variety of bacterial survival strategies. In turn, it is likely that the host is responding in a similar fashion to the presence of the bacteria by activating tissue repair mechanisms and antibacterial killing mechanisms.

Recent work has indicated that virulent *Salmonella* induces the death of cultured macrophages [Lindgren et al. 1996]. Moreover, it has been reported that death in
cultured and bone marrow-derived macrophages occurs by an apoptotic mechanism based on nuclear morphology, cytoplasmic vacuolization, and cell DNA fragmentation (Chen et al. 1996; Monack et al. 1996). Programmed cell death was dependent on an intact bacterial invasion phenotype, as noninvasive S. typhimurium strains were unable to induce death. Currently it is not clear whether apoptosis benefits the pathogen by inactivating the professional killing cell or whether the host benefits by reducing the capacity of the bacteria to spread. Answers to those questions await future experimentation.

**Host immune responses to Salmonella infection**

Exposure to the human pathogen S. typhi usually results in acquired immunity to future infections. However, immunity may not develop if antibiotic treatment is received during the early stages of infection, suggesting that the immune system may require a prolonged exposure to the bacteria before a substantial defense can be mounted. Many studies have been performed to identify the principle components of the human immune response to typhoid fever. Adult volunteers who were administered the oral vaccine strain S. typhi Ty21a developed significant levels of circulating IgA and IgG antibodies that were directed against the lipopolysaccharide (LPS) of the organism (Forrest et al. 1991). In a Nepalese S. typhi vaccine field trial, subjects that developed serum antibody to the Vi capsular polysaccharide had a typhoid attack rate of only 4.1% compared to 16.2% for the control group, suggesting that circulating antibody levels participate in the initial resistance to infection (Acharya et al. 1987). Cell-mediated immunity is also observed in S. typhi-infected individuals (Taglia­bue et al. 1985; Nencioni et al. 1987; Murphy et al. 1989). Although specific antigens that trigger the development of immunity have not been identified, it is believed that CD4+ cells, stimulated in response to the bacterial infection, activate professional killing cells. Thus, the primary line of defense against systemic Salmonella infection is presumed to be the activation of phagocytic cells.

Animal infection models have also been used extensively to study the immune response to Salmonella strains. The majority of this work has focused on the immune response to systemic infection of mice caused by S. typhimurium. Clearly, host resistance to Salmonella infection depends on a number of different factors. One gene that has received considerable attention has been designated Ity [also Lsh and Bcg]. Mice that are Ity-defective [Itya] are significantly more susceptible to pathogenic Salmonella than are Ity+ animals. The molecular mechanism for this difference in resistance to the bacteria is unknown, although Itya mice permit low numbers of bacteria to proliferate rapidly to high levels in the spleen and liver while much higher inocula are required to allow net growth of Salmonella in Ity+ mice. One possible explanation for these growth differences in mice is provided by in vitro studies. Resident phagocytic cells from Itya and Ity+ mice were isolated and examined for their ability to kill intracellular bacteria (van Dissel et al. 1985, 1986). Phagocytic cells from the Ity+ mice killed intracellular Salmonellae at a much higher rate than the cells from the Itya mice. The mechanism of bacterial killing has not yet been elucidated, although macrophages from the Itya mice had enhanced expression of the activated macrophage markers la and Acg-1 (Buschman et al. 1989). In addition, the cells produced increased levels of oxygen- and nitrogen-free radicals in response to the activator l interferon (IFN-γ) [Skamene and Pietrangeli 1991].

The gene responsible for the Ity phenotype recently has been cloned and sequenced from the mouse and designated Nramp (natural resistance-associated macrophage protein) [Vidal et al. 1993; Malo et al. 1994]. Structural analysis of the Nramp protein sequence suggests that the molecule is a membrane transporter, although no transport function has been demonstrated yet [Cellier et al. 1996]. Another member of the Nramp gene family, the Saccharomyces cerevisiae protein SMFl, has been shown to be a manganese transporter, which suggests that this protein family may be a new class of membrane transporters of channels [Supek et al. 1996]. Northern blot analysis has demonstrated that the Nramp mRNA is expressed exclusively in macrophages derived from lymphoid organs as well as from macrophage tissue culture lines such as J774 (Vidal et al. 1993). Transfection of the Nramp gene into RAW 264.7 tissue culture macrophages resulted in increases in the respiratory burst, nitrite release, and uptake of l-arginine, all of which are indicators of macrophage activation [Barton et al. 1995]. The Nramp gene, which plays a role in regulating expression of interleukin-1β, inducible nitric oxide synthase, maphistocomputability [MHC] class II molecules, and tumor necrosis factor α (TNFα) levels is thought to exert its activity early in the macrophage activation pathway [Blackwell et al. 1994; Skamene 1994; Blackwell 1996]. Future experimentation should elucidate the function of the Nramp protein and how this function translates into macrophage activation and immunity. However, it is clear that the Nramp protein is important in the activation of macrophages at an early stage and that this activation is important in the elimination of intracellular pathogens such as Salmonella, Mycobacteria, and Leishmania from an infected host.

Mice with an Itya background have been useful in the study of the specific host immune response to virulent Salmonella infection. Antibody-mediated and cell-mediated immunity are both required for long-term protection against infection with pathogenic Salmonella strains in Itya mice, whereas innately resistant mice respond to Salmonella infection with a nonspecific immune response [Collins 1974; Hormaeche et al. 1990; Mastroeni et al. 1993]. Presumably, a part of these different immune responses are attributable to different levels of IFN-γ that are produced. Two different groups have found that Itya mice produce significantly lower levels of IFN-γ than Ity+ mice [Freudenberg et al. 1991];
which cells might be responsible for the difference in
membrane rearrangements and entry (Degre et al. 1989). Pre-
IFN-7 production. They demonstrated that murine natu­
from proliferating and inducing septic shock. In
phagocytes, as evidenced by increased fusion of phago­
response to virulent infection. Treatment of
Salmonella-induced membrane rearrangements and entry [Degre et al. 1989]. Pre-
treatment of cultured macrophages with IFN-7 has been
shown to activate killing mechanisms of professional
phagocytes, as evidenced by increased fusion of phago­
S. typhimurium early in the infection of mice [Muotiala and Makela 1990; Nauciel and Espinasse 1992]. Mice injected with
exogenous IFN-7 at early stages of infection with patho-
genic S. typhimurium had fewer numbers of bacteria in
the lymphatic organs than control mice. However, when
IFN-7 was administered at later stages, it had no detect-
able effect on the course of the infection. Reciprocally,
when circulating levels of IFN-7 were depleted in mice by
the injection of anti-IFN-7 antibody, mice were unable
to prevent even very low numbers of S. typhimu-
from proliferating and inducing septic shock. In
addition, mice that have a mutation in the gene encoding
IFN-7 [IFN-7 knockout or GKO mice], and are therefore
unable to produce IFN-7, are hypersusceptible to infec-
tion with virulent S. typhimurium [Hess et al. 1996]. The
observation that the lymphoid cells of Peyer's patches are stimulated to produce large quantities of IFN-7 in
response to the presence of Salmonella is indicative that the host depends on IFN-7-mediated effects early in in-
fection for activation of nonspecific defensive mecha-
nisms [Ramarathinam et al. 1991].

Additional experiments have established that TNFα
plays a significant role in the host cell response to S. typhimurium [Nakano et al. 1990]. Administration of
TNFα to mice inoculated with S. typhimurium reduced the
numbers of bacteria that could be recovered from the
animals compared to untreated controls. In contrast, in-
jection of TNFα improved the survival rate of mice re-
ceiving pathogenic Salmonella. As observed for IFN-7,
depletion of TNFα in mice with anti-TNFα antibodies
decreased the dose of bacteria required to achieve a lethal
systemic infection in mice. Also, treatment of phago-
cytes with anti-TNFα antibodies blocked LPS-stimu-
lated activation. These data suggest that TNFα is pro-
duced as a result of two separate signals; the LPS of the
infecting Salmonella organisms and tissue damage that
results from invasion and destruction of host cells. Clearly, the ability of the host to produce IFN-7 and
TNFα in response to the presence of Salmonella is criti-
cal. Presumably, these two lymphokines function by ac-
tivating early nonspecific killing mechanisms of profes-
sional phagocytes, as their activity is most important at
the initial stages of Salmonella infection. A specific ef-
effect of IFN-7 on macrophages has been identified, which
is to increase the level of phagosome-lysosome fusion in
the cell. Increased levels of endosome maturation would
protect the host by neutralizing efforts by Salmonella
increase survival by delaying fusion. Research efforts
continue to define these processes more clearly as well
to identify other activities of IFN-7 and TNFα in host
immunity.

The specific immune response to pathogenic S. typhi-
murium appears to result in a general host immuno-
pression [Lee et al. 1985]. Mice that have been immu-
nized previously with S. typhimurium have been exam-
ined for their ability to mount immune responses to other antigens or pathogens [Lee et al. 1985]. Those ex-
periments revealed that immunized mice are unable to
respond to new antigens for up to 3 weeks after immu-
nization with S. typhimurium. Apparently, macrophages
are primarily responsible for this inability to respond to
new antigenic challenge, as in vitro depletion of phago-
cytes eliminated the observed immunosuppression. Fur-
ther work revealed that large quantities of nitric oxide,
produced by activated macrophages, inactivated lympho-
cytes, thereby reducing the capacity of the immune sys-
tem to mount a cellular response to antigens. Treatment
of mice with IL-4 or anti-IFN-7 antibody prevented Sal-
monella-induced immunosuppression by preventing ni-
tric oxide poisoning of lymphocytes [al-Ramadi et al.
1992].

Conclusion
The focus of this review has been the host response to
Salmonella infection and virulence mechanisms. The
first significant interactions between the bacteria and
the host occur at the lymphoid follicles of the intestine.
This action would seem to play into the hands of the
host because the primary function of lymphoid follicles
is to protect the host from intestinal invasion. However,
Salmonella displays a highly evolved ability not only to
initiate contact with the host at Peyer's patches but to
benefit from the interaction. Subsequently, the bacteria
move into the lymphatic system where the survival of
the organisms requires the ability to productively inter-
act with professional killing cells. Again, these patho-
gens display a remarkable adaption to an environment
that is considered to be very hostile. Thus, it is apparent
that the bacteria have discovered that survival within
the host depends on their ability to directly engage the
host immune system and alter the host response in their
favor. It seems certain that the pressure being exerted by
this and other bacterial pathogens is honing the host
response to infection. It is also clear that the bacteria
will not be left behind in the evolutionary battle. From a
scientific point of view, these interactions offer the op-
portunity to gain insights both into cell biology and bac-
terial pathogenicity.
Salmonella-host cell interactions

Acknowledgments

Thanks go to Nitin Mathur for providing the scanning electron micrographs. B. Jones is supported by U.S. Public Health Service grant AI38268 and a grant from the Roy J. Carver Charitable Trust.

References


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*Genes Dev.* 1997, 11:
Access the most recent version at doi:10.1101/gad.11.6.679

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