Two functionally distinct Type III Secretion Systems for *Salmonella* pathogenesis

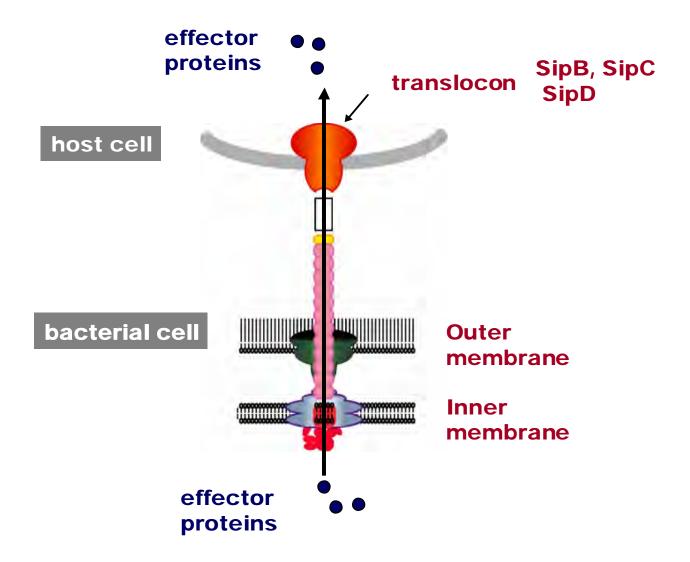
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Type III protein secretion system (TTSS)

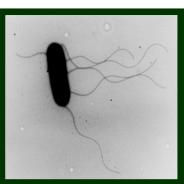


Salmonella spp

important pathogens of humans and animals

•Cause a wide variety of diseases ranging from mild diarrhoea to severe systemic infections like typhoid fever

•Estimated 16 million cases of typhoid fever per year occur with about 600,000 fatal outcomes



an interesting model organism

•For the study of host-pathogen interaction

-able to enter into non-phagocytic cells (e.g. epithelial cells) and grow within phagocytic cells (e.g. macrophages)

Salmonella Pathogenicity island (SPI): SPI1 and SPI2 encoding Type III protein secretion system

Outline

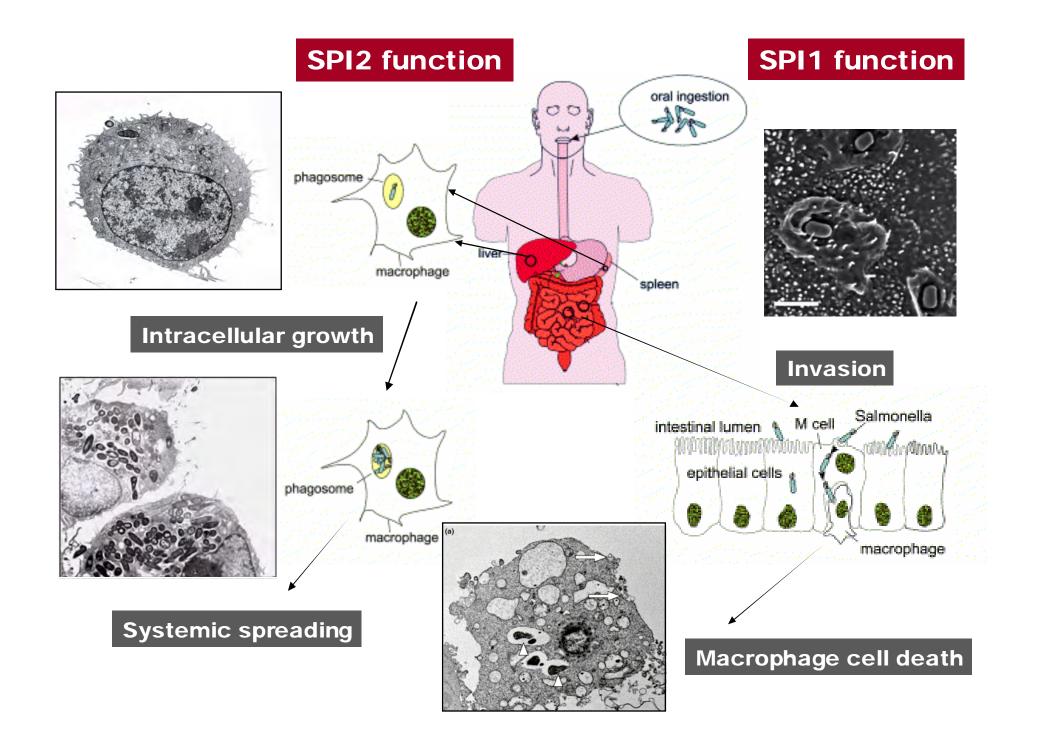
Functions of SPI-1 and SPI-2 Type III secretion systems (TTSSs) on Salmonella pathogenesis

Inverse regulation of SPI1-TTSS and SPI2-TTSS within macrophages

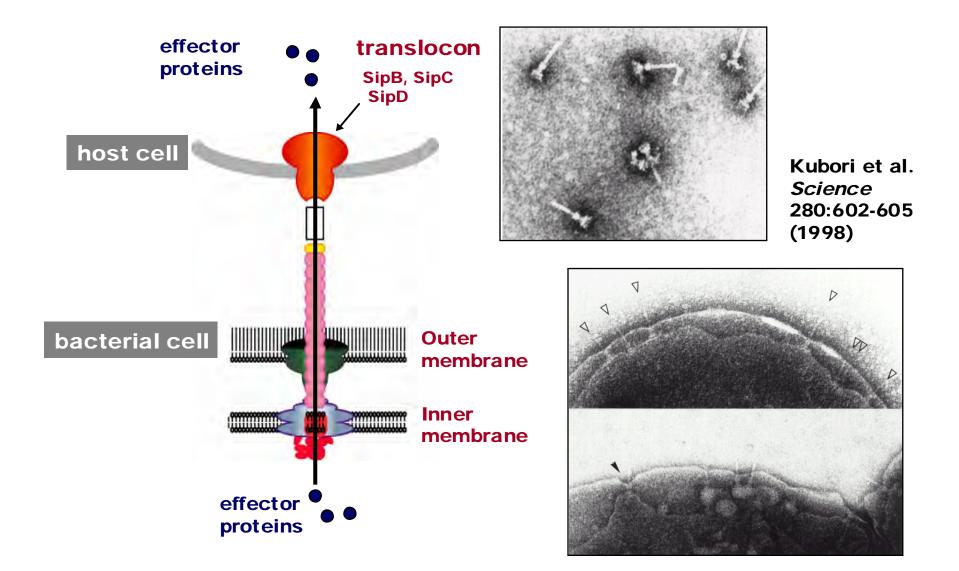
Control of host macrophage cell death, pyroptosis and apoptosis, by SPI-1 effectors



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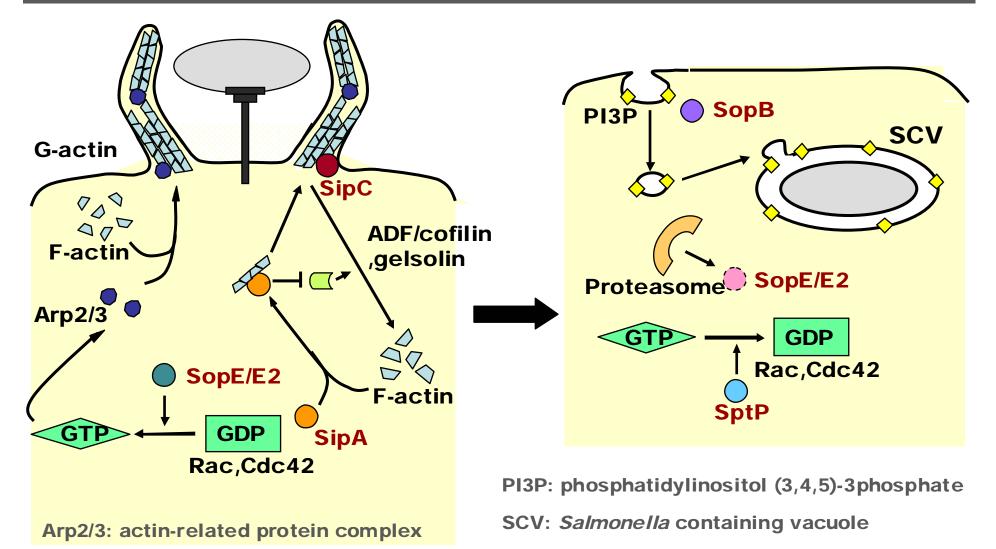


SPI1-Type III secretion system

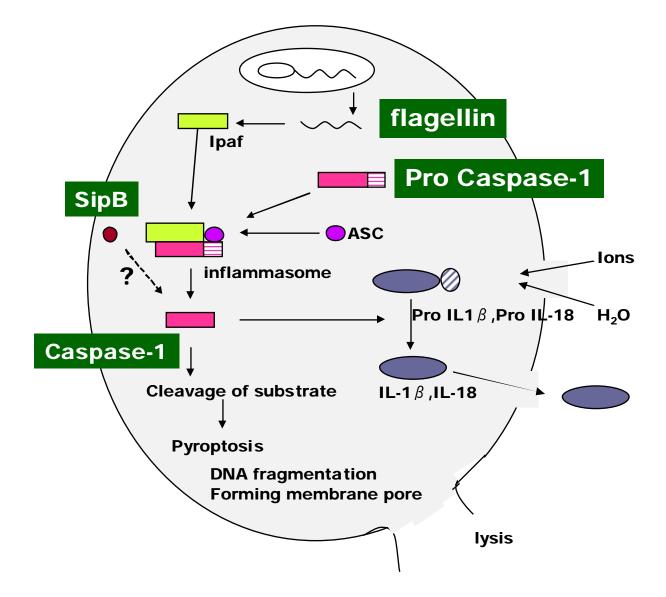


SPI1-TTSS promotes invasion

Effectors induce actin rearrangements and alter vacuole trafficking to trigger invasion, without causing cellular damage

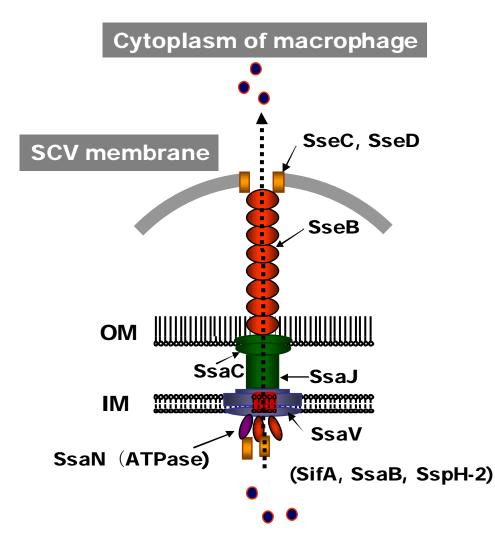


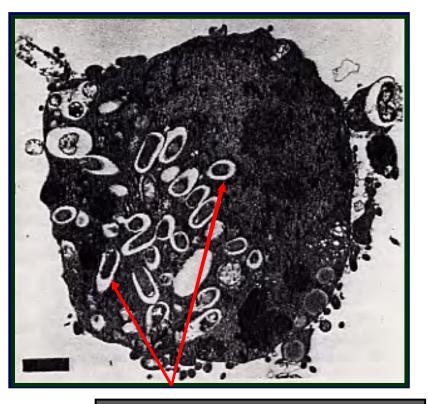
SPI1-TTSS-dependent activation of caspase-1, leading to macrophage cell death, pyroptosis



SPI2-Type III secretion system

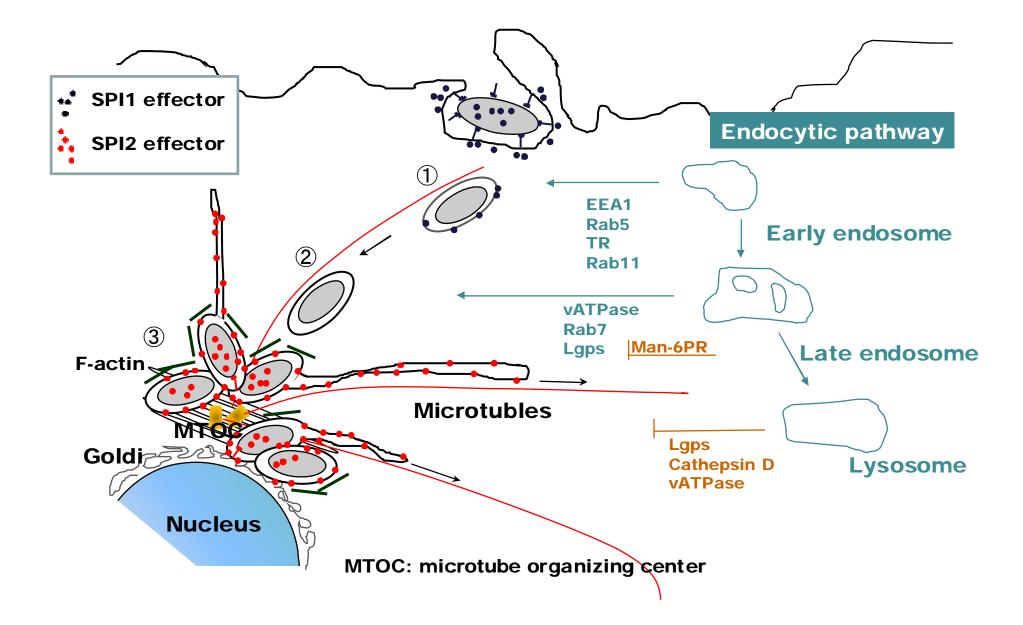
*SPI2: Salmonella pathogenicity Island at cs31



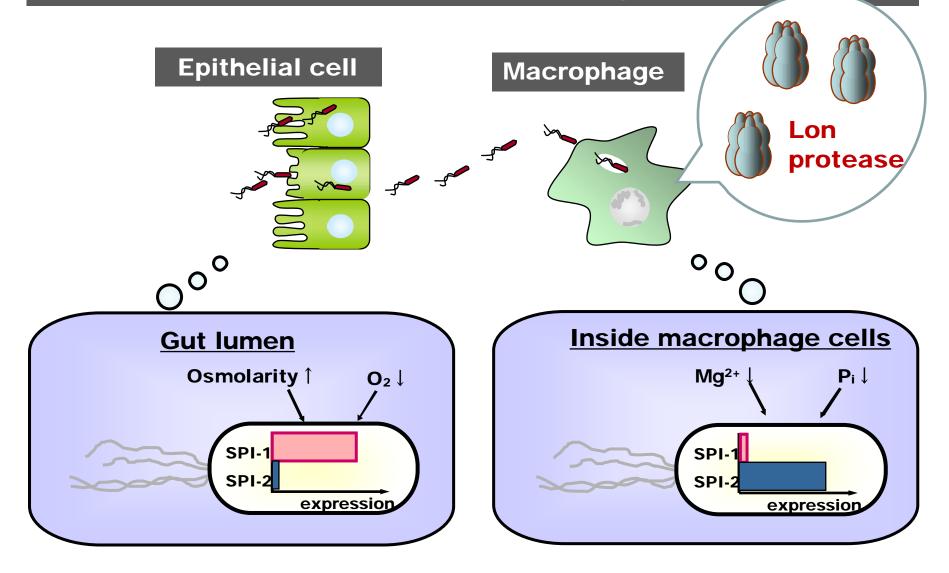


SCV :*Salmonella* containing vacuole

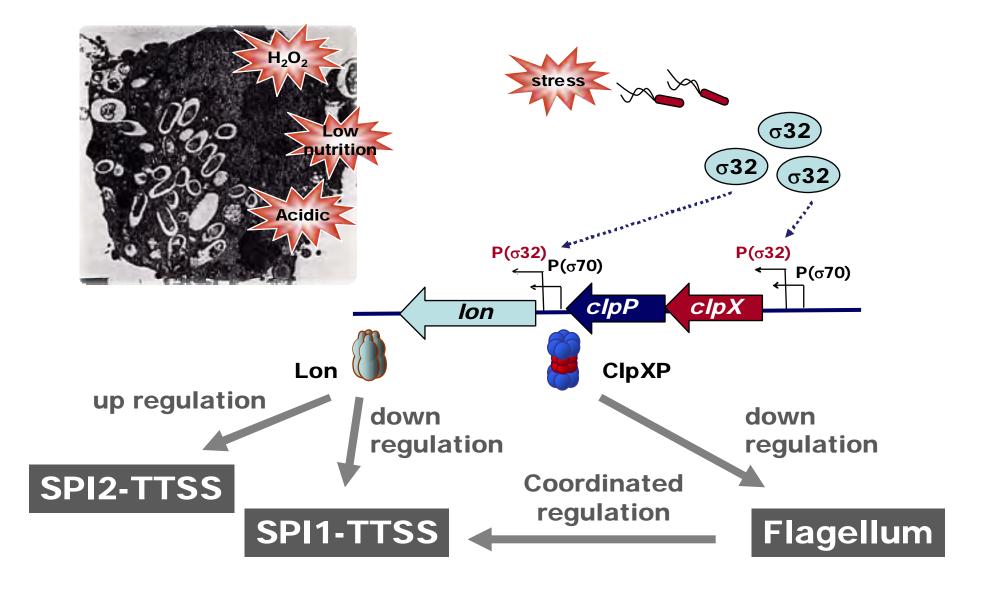
SPI2-mediated SCV membrane dynamics, leading to *Salmonella* multiplication within host cells

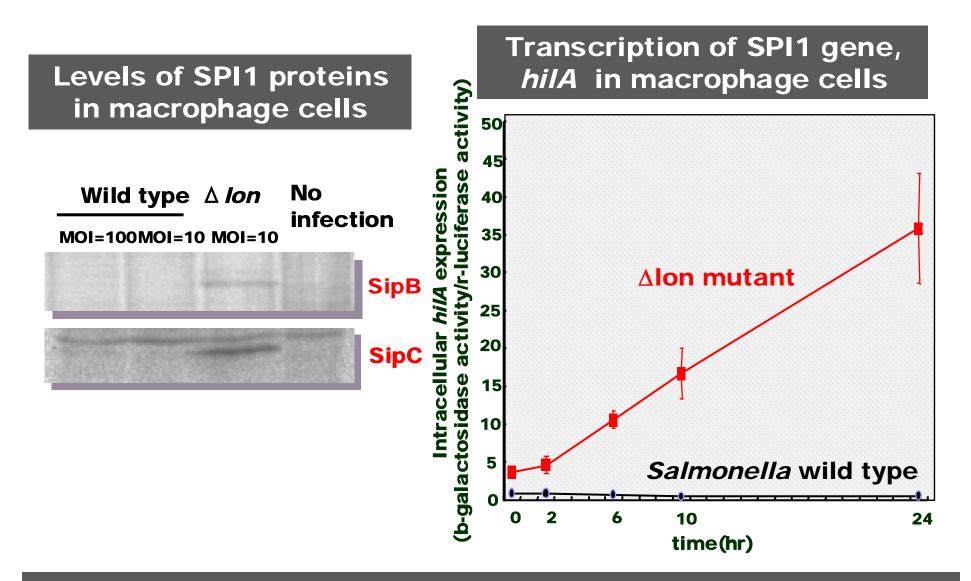


Inverse regulation of expression of SPI-1 and SPI-2 genes



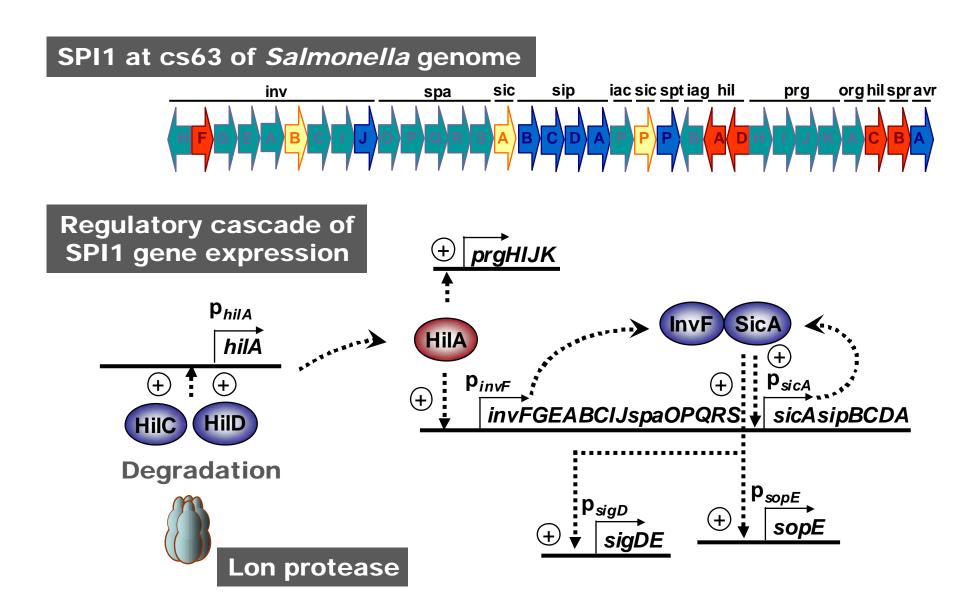
Lon is induced as a stress response by *Salmonella* to hostile environment in macrophages





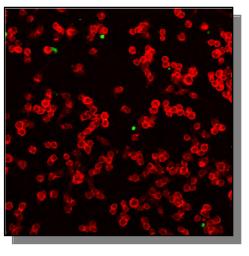
Lon is essential for down-regulation of SPI1expression in macrophage cells after phagocytosis

Lon degrades HilC and HilD to down-regulate the expression of SPI1 genes

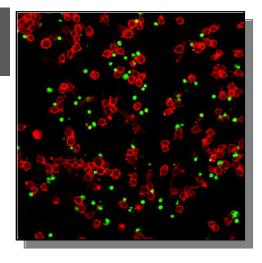


Salmonella ∆lon mutant induces massive apoptosis in macrophages

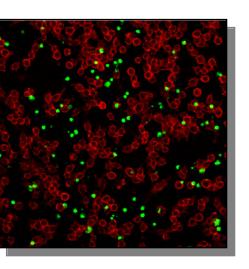
wild type MOI=10



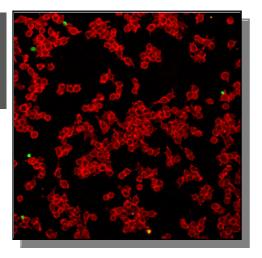
∆*Ion* MOI=10



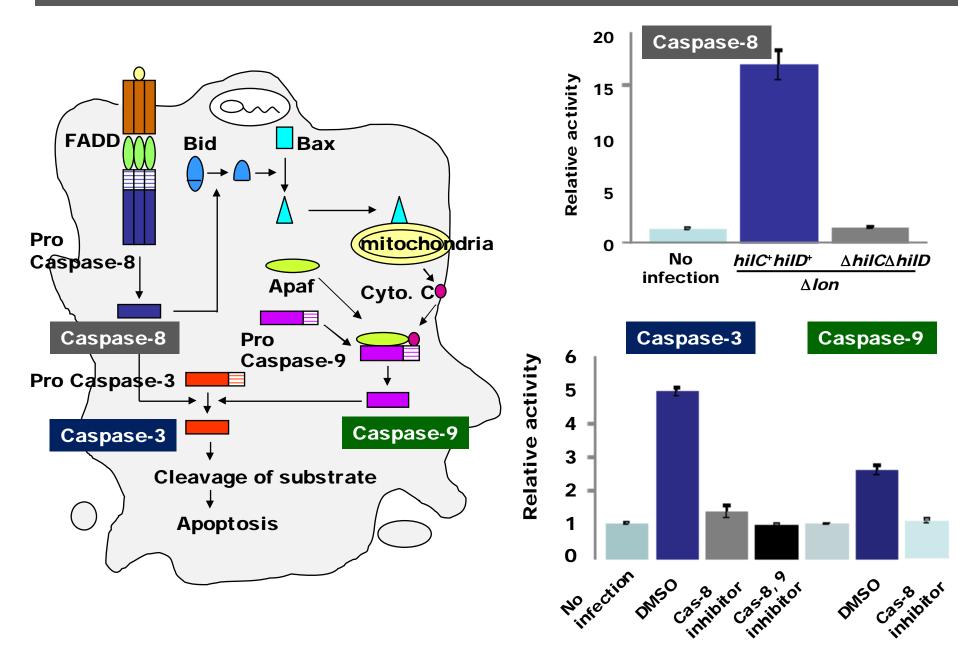
∆*lon* MOI=1



∆*Ion* ∆SPI1 MOI=10



Over-expression of SPI1 activates caspase-8-dependent procaspase-3 activation pathway



Physiological significance of control of cell death via negative regulation of SPI1 expression

At the initial stage of infection (intestinal phase of infection), Salmonella escape the macrophage killing mechanism by induction of flagellin-dependent and SPI1-dependent cell death, pyroptosis.

>Once Salmonella has established a systemic infection, excess cell death like apoptosis would be detrimental to the pathogen because Salmonella resides in macrophage cells.

>It would be required to suppress apoptosis to allow time for the bacterium to replicate, escape and invade new macrophages for systemic infection.

>Therefore, negative regulation of SPI1-TTSS expression by Lon which is induced in response to the hostile environment in macrophage cells would be essential for the suppression of apoptosis through the control of caspase-8 activity in the macrophage cells after *Salmonella* infection.

Summary

 Two functionally distinct TTSS
SPI1 • invasion of epithelial cells
• release of inflammatory cytokines
• induction of cell death, pyroptosis
• induction of caspase-8, leading to apoptosis
SPI-2 • SCV membrane dynamics leading to replication and spatial distribution

Inverse regulation of Two TTSSs Lon which is induced in Salmonella growing in macrophage cells after phagocytosis controls negatively SPI1 expression and positively SPI2 expression





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