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The role of *invG* in *Salmonella enterica* adherence to and invasion of human intestinal epithelial cells

Nontyphoidal *Salmonella enterica* are primary foodborne pathogens infecting both humans and animals worldwide. Among nontyphoidal *Salmonella*, the serovar Typhimurium (*S. Typhimurium*) and serovar Enteritidis (*S. Enteritidis*) are the most common *Salmonella* serovars implicated in foodborne illness in the United States and Europe. Recently, we identified an immunogenic protein of *S. Enteritidis* called *InvG*, which is currently being investigated as a vaccine candidate to reduce *Salmonella* colonization and shedding in layer chickens. The *invG* was shown to be involved in *S. Enteritidis* adherence to and invasion of chicken intestinal epithelial tissues. The long-term goal of the proposed study is to assess the suitability of *InvG* as a vaccine candidate against human foodborne infection. As the first step, in this preliminary study, we plan to evaluate if *invG* is involved in the adherence and invasion of *S. Enteritidis* and *S. Typhimurium* to the murine gut similar to what we observed in the chicken gut.

Specific aims:

1. Study the role of *invG* in *S. Enteritidis* and *S. Typhimurium* adherence and invasion of human intestinal epithelial cells using an in vitro cell invasion assay established.
2. Perform a time course analysis of *invG* expression using a reverse transcription quantitative PCR (RT-qPCR) during interaction of *Salmonella* with intestinal epithelial cells.

Student's role:

1. Create an isogenic mutant of *S. Typhimurium* lacking the *invG* and its complemented strain (we have already constructed the  $\Delta invG$  *S. Enteritidis* strain and the corresponding complemented strain). Student will begin the experiments with *S. Enteritidis* while the mutant strain of *S. Typhimurium* is being created.
2. Establish the monolayers of Caco-2 cells (ATCC HTB-37) and perform adherence and invasion assays using wild-type, mutant, and complemented strains.
3. Bacterial RNA purification from cell culture assays and perform RT-qPCR to measure *invG* expression.
4. Statistical analysis of data and drafting the manuscript.