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SUBJECT: Understanding of multi-scale innate immune responses against bacterial pathogens

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DESCRIPTION: Infection triggers complex signalling events in cells, which aim to kill the pathogen directly or induce a protective cytokine response to activate bystander cells. Paradoxically, the immune cell activation exhibits extreme variability, for example protective cytokines are produced by a small subset of genetically identical macrophages. Consequently, the interactions between host and pathogen are inherently heterogeneous and result in different and “seemingly” probabilistic outcomes at the single cell level. How a robust immune response emerges from the heterogeneous single cell behaviour remains one of the fundamental questions in immunology and infection biology. The main aim of the project is to quantitatively understand mechanisms involved in the multi-scale coordination of innate immune responses against bacterial pathogens at the single cell level. We will focus on events occurring at low physiological multiplicities of infection and physiological febrile temperature range, which give rise to heterogeneous and stochastic effects in the infected population. We will study direct interactions between innate immune macrophages and the important food-borne bacteria *Listeria monocytogenes* that causes significant levels of human morbidity and mortality. Using live-cell microscopy approaches and mathematical modelling to elucidate how pathogen threats are encoded in dynamical responses of NF- κ B/STAT/IRF systems upon *L. monocytogenes* infection and how they relate to different (and seemingly probabilistic) outcomes of single-cell host-pathogen interactions. The goal of this proposal is to identify key regulatory mechanisms and understand how to change the course of infection in single cells and cellular populations.

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