

# How to "Hijack" the Host Cells

——The manipulation of host-cell pathways by bacterial pathogens

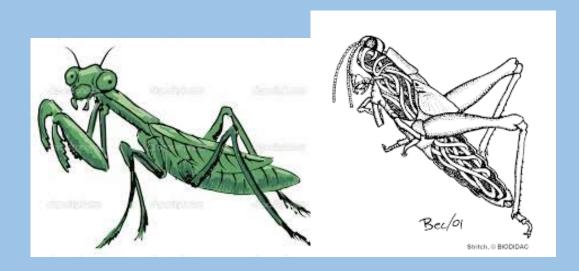
YE Yu

Supervisor: Prof. Guoping Zhao

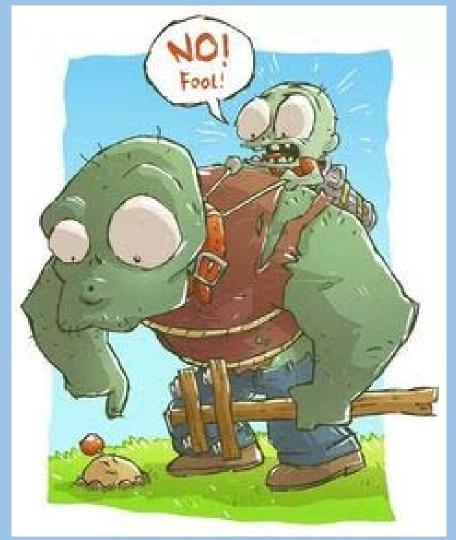
Microbiology

Date: Dec. 1 2016

• Virus "Hijack" zombie's brain



• Gordiacea "Hijack" Mantis



#### Outline

- Introduction
- Research Objectives
- "Hijack" targets
  - During invading
    - Cytoskeleton(Actin Filaments)
    - Microtubules
    - Phagosomes
    - Signaling Pathway
  - During preservation
    - Inflammation reaction
    - antigen presentation
- A typical "hijacking" pathogen: Mycobacterium tuberculosis
- Summary

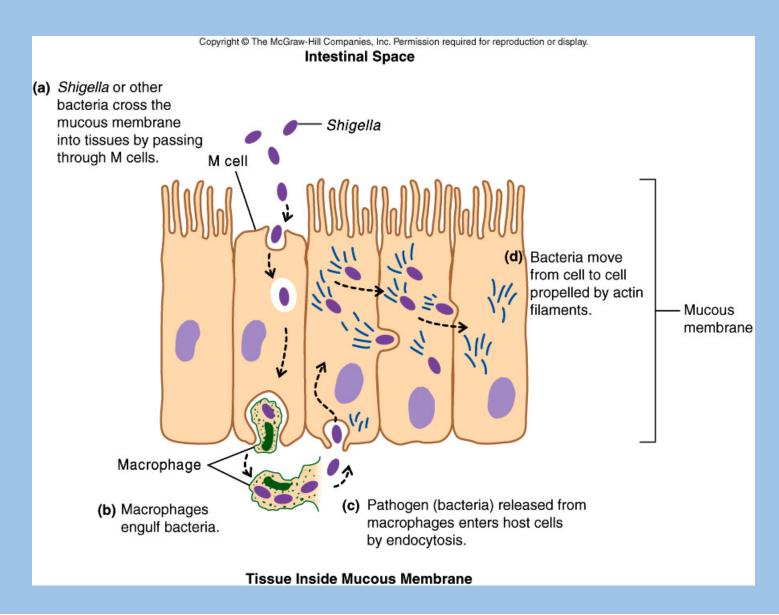
#### Relationships between bacteria and host

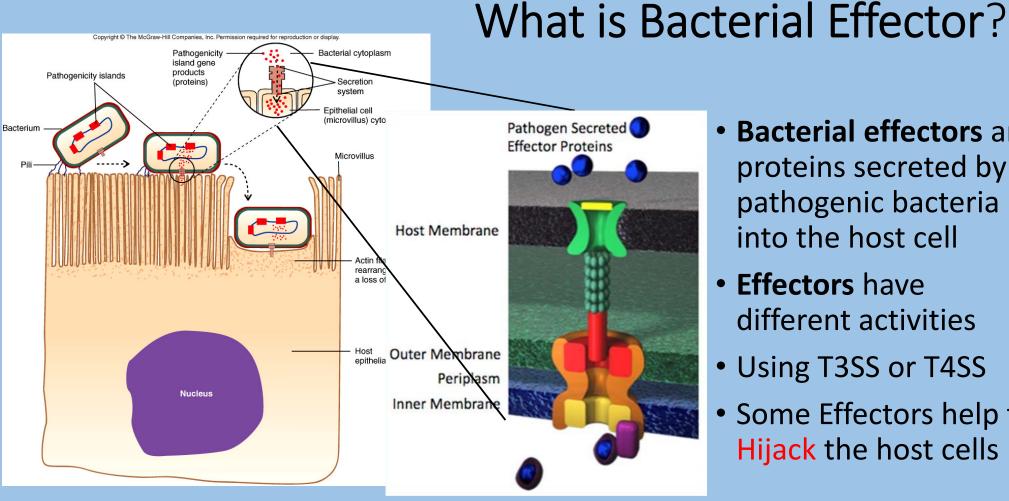
- mutualistic relationship
- both the bacteria and the host benefit
- The host provides the microbiota a niche with a stable nutrient supply
- Intestinal flora

- parasitic relationship
- the bacteria benefit while the host is harmed
- The bacteria seize the a niche and nutrient from host
- Pathogens
- Infectious disease

## Struggle during infection

- Establishing an Infection
- Invasion
- Avoiding the Host Defenses
- Survival Strategies within Phagocytes
- The effector proteins may much helped





- Bacterial effectors are proteins secreted by pathogenic bacteria
- Effectors have different activities

into the host cell

- Using T3SS or T4SS
- Some Effectors help to Hijack the host cells

### Why bacteria hijacking is worth to study?



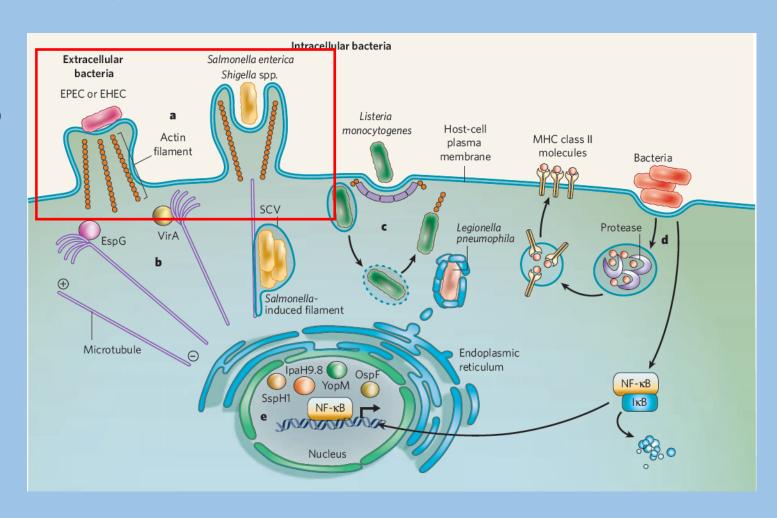
- To demonstrate how pathogens overcome host defenses
- To understand the bacterial virulence at a molecular level
- To find new method to prevent, block or eliminate infectious pathogens

#### Periods for pathogen to Hijack host cells

- During invading and replicating
  - Targets: cytoskeleton, membranous structures and key signalling pathways
- During preservation
  - Target: immune response

#### Cytoskeleton

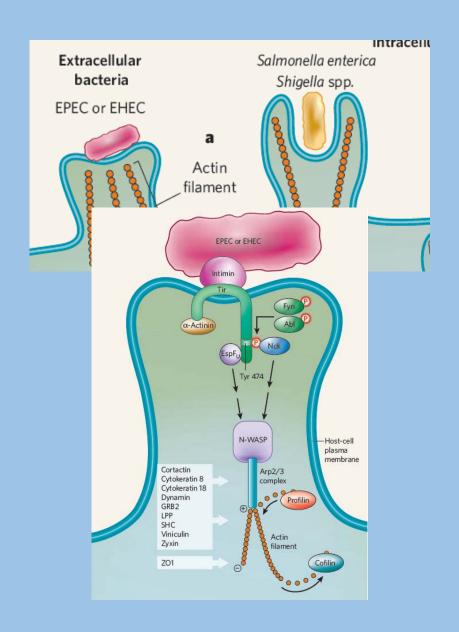
- Bacterial pathogens manipulate the cytoskeleton to help invade a host cell and/or to gain motility in the cell
- They often interact with actin filaments in particular, by modulating G proteins.



#### Cytoskeleton

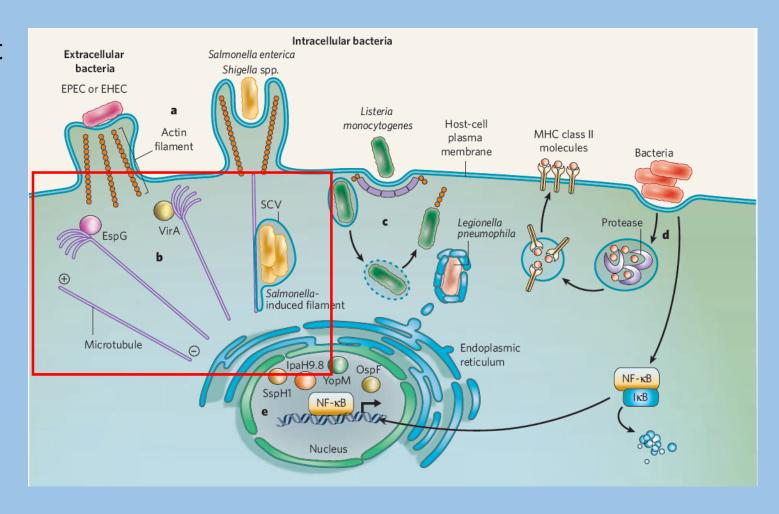
Salmonella enterica

- Enterohaemorrhagic *Escherichia coli* (EHEC)
  - Having an elaborate actin recruiting process as an extracellular pathogen



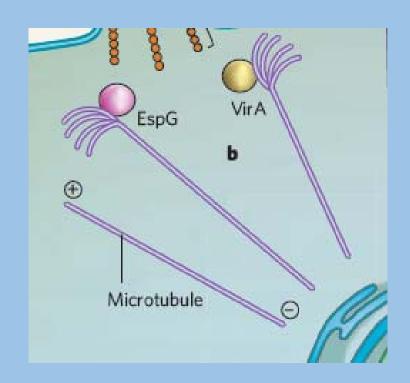
#### Microtubules

- Guide and transport intracellular cargo
- Transport and the microtubule assembly and/or disassembly dynamics can be modified and controlled by the pathogen



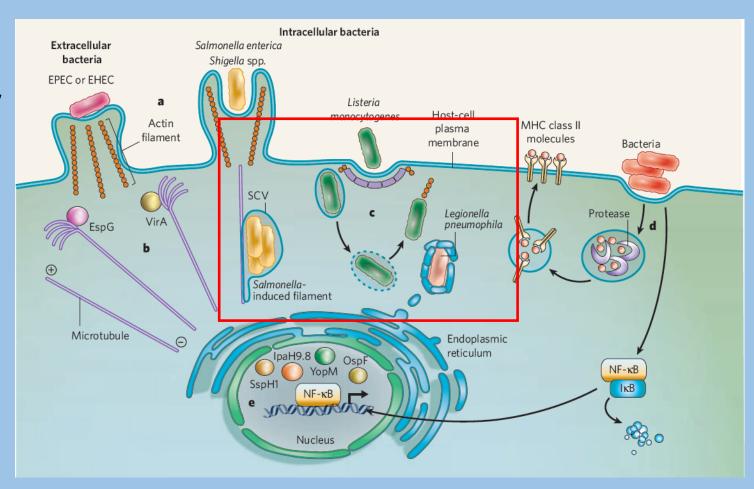
#### Microtubules

- Shigella spp
- Effector: VirA protein
- This results in invasion by Shigella spp.



#### Vacuoles and modified phagosomes

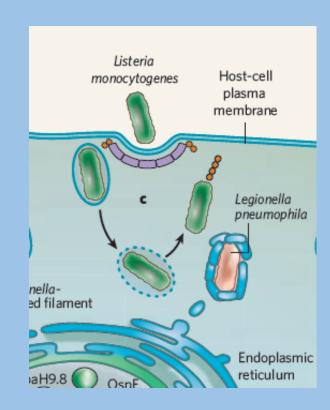
- Pathogens have adopted various strategies to multiply in, or escape from, these structures before surviving in the cytosol
- They will able to live in vacuoles and modified phagosomes



#### Phagosomes

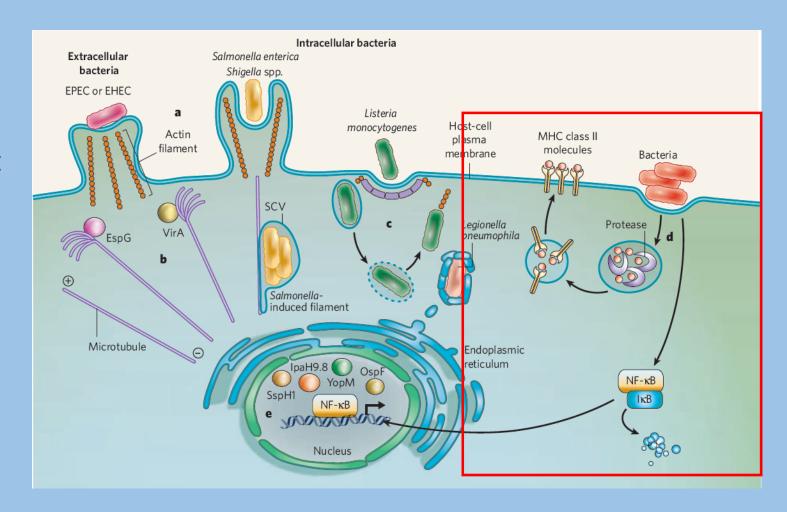
• Legionella spp.

Providing a nutrient-rich resource for the bacteria



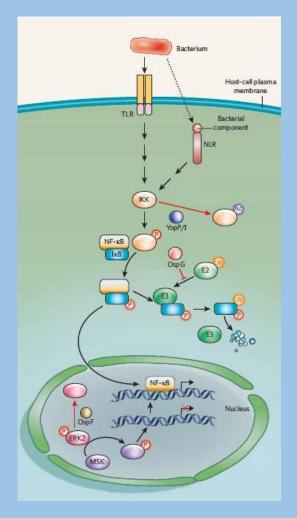
#### Signaling pathways

- Many pathogens interfere with the phosphorylation cascades of the host cell.
- bacterial effector proteins mimicked host kinases and phosphatases



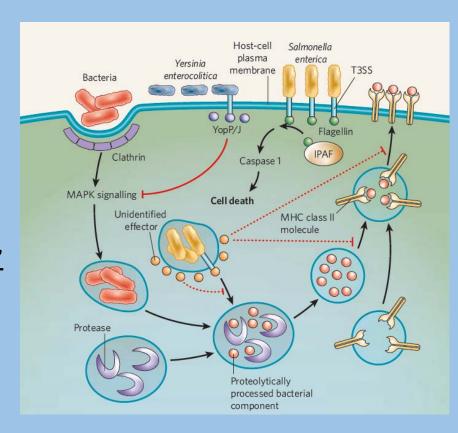
#### Inflammation and nuclear factor-kB

- Factor nuclear factor-κB (NF-κB) plays a role in inflammation reactivation
- S. flexneri have come to 'understand' the NF-κBactivation pathway
- Sprevent IkB from being ubiquitylated and remain inactive
- This bacteria effect this through the T3SS effector proteins OspG



#### Antigen presentation

- Bacteria are recognized and internalized by specialized cells known as antigenpresenting cells
- Several pathogens able to subvert the immune responses.
- S. enterica can block antigen presentation, resulting in the lower activation of fewer T cells



#### Other targets

Apoptosis pathways

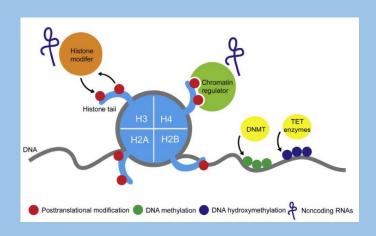
### Bacterial pathogens modulate an apoptosis differentiation program in human neutrophils

Scott D. Kobayashi, Kevin R. Braughton, Adeline R. Whitney, Jovanka M. Voyich, Tom G. Schwan, James M. Musser, and Frank R. DeLeo\*

Laboratory of Human Bacterial Pathogenesis, Rocky Mountain Laboratories, National Institute of Allergy and Infectious Diseases, National Institutes of Health. Hamilton. MT 59840

Edited by Richard M. Krause, National Institutes of Health, Bethesda, MD, and approved July 15, 2003 (received for review June 3, 2003)

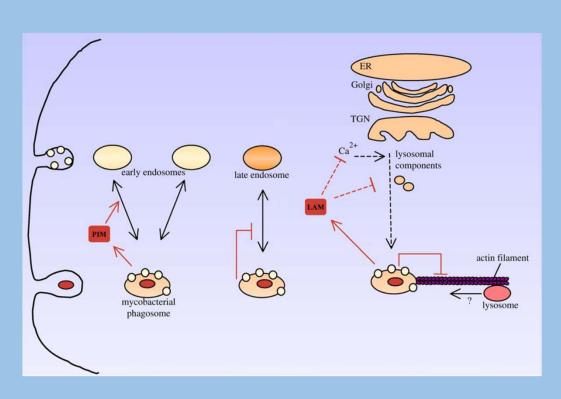
Epigenome



## A typical "hijacking" pathogen: *Mycobacterium* tuberculosis

- A most "successful" pathogen in our planet
- The macrophage represents a highly specialized cell of the immune system, yet fails to eliminate the M. tuberculosis bacillus
- *M. tuberculosis* has managed to establish itself within the macrophage, avoiding the host killing mechanisms.
- Forming granuloma and living in extreme hypoxia and high lipid environment for decades as latent infection

#### Mtb hijacks the macrophages



- M.tb interference with macrophage trafficking events
- M.tb suppress MHC class II expression in macrophages
- M.tb reside within phagosomes that retain early endosomal properties and resist fusion with later stages of the endosomal/lysosomal pathway.

#### New mechanisms?

• The nucleic acid secretion from *M.tb* has been found in macrophage during the infection.

#### Review article

Molecular requirements for sensing of intracellular microbial nucleic acids by the innate immune system

Stefanie Luecke<sup>a</sup>, Søren R. Paludan<sup>a,b,\*</sup>

<sup>a</sup> Department of Biomedicine, Aarhus University, Wilhelm Meyers Allé 4, 8000 Aarhus C, Denmark

<sup>b</sup> Aarhus Research Center for Innate Immunology, Aarhus University, Aarhus, Denmark

#### ARTICLE INFO

Article history: Received 5 October 2016 Accepted 11 October 2016 Available online xxxx

Keywords: Innate immunology Intracellular nucleic acid sensing PAMP Interferon response

#### ABSTRACT

Nucleic acids sensors of the innate immune system recognize various RNA and DNA structures during infection to induce transcription of interferon and pro-inflammatory cytokines and activation of inflammasomes. Cytosolic RNA is recognized by RIG-1 and MDA5, while intracellular DNA is sensed among others by cGAS, AIM2, IF116 and RNA polymerase III. The diversity of nucleic acid species produced during infection in the cytosol and nucleus and the limited chemical differences between self and non-self nucleic acids challenge the host's innate pattern recognition system to ensure reliable sensing while avoiding immune activation by self nucleic acids. We review the molecular characteristics of intracellular nucleic acid sensor ligands, the structural basis of the binding preferences of the sensors, the identity and origin of immunostimulatory nucleic acid species during infection, the influence of intracellular localization of the sensor on immune activation, and the ability of viruses to use the ligand specificity of the sensors to evade recognition.

© 2016 Published by Elsevier Ltd.

#### Summary

- The manipulation of host-cell pathways by bacterial pathogens is extensively existing in infections, might be much more than we can imagine.
- Knowledge of how pathogens target distinct cytoskeletal components and immune-cell signalling pathways is rapidly advancing, together with the understanding of bacterial virulence at a molecular level.
- Studying how these bacterial pathogens subvert host-cell pathways is central to understanding infectious disease.

#### Reference

- 1. Amit P. Bhavsar, Julian A. Guttman B. Brett Finlay. Manipulation of host-cell pathways bybacterial pathogens. NATURE (2016)
- 2. Dennis J Murphy, James R Brown\*. Identification of gene targets against dormant phase Mycobacterium tuberculosis infections. BMC Infectious Diseases (2014).
- 3. Calvin Boon & Thomas Dick\*. How *Mycobacterium tuberculosis* goes to sleep: the dormancy survival regulator DosR a decade later. *Future Microbiol* (2016). 7(4), 513–518
- 4. Martin Gengenbacher & Stefan H.E. Kaufmann. Mycobacterium tuberculosis: success through dormancy. *FEMS Microbiol Rev* 36 (2012) 514–532
- 5. Shiloh MU, Manzanillo P, Cox JS. *Mycobacterium tuberculosis* senses host-derived carbon monoxide during macrophage infection. *Cell Host Microbe* 3(5), 323–330 (2015).
- 6. Chao JD, Papavinasasundaram KG, Zheng X *et al.* Convergence of Ser/Thr and two-component signaling to coordinate expression of the dormancy regulon in *Mycobacterium tuberculosis. J. Biol. Chem.* 285(38), 29239–29246 (2014).
- 7. Leistikow RL, Morton RA, Bartek IL, Frimpong I, Wagner K, Voskuil MI. The *Mycobacterium tuberculosis* DosR regulon assists in metabolic homeostasis and enables rapid recovery from nonrespiring dormancy. *J. Bacteriol.* 192(6), 1662–1670 (2010).
- 8. Watanabe S, Zimmermann M, Goodwin MB, Sauer U, Barry CE 3rd, Boshoff HI. Fumarate reductase activity maintains an energized membrane in anaerobic *Mycobacterium tuberculosis. PLoS Pathog.* 7(10), E1002287 (2011).
- 9. Cole ST, Brosch R, Parkhill J et al. Deciphering the biology of Mycobacterium tuberculosis from the complete genome sequence. Nature 393(6685), 537–544 (1998).
- 10. Dasgupta N, Kapur V, Singh KK *et al.* Characterization of a two-component system, *devR*–*devS*, of *Mycobacterium tuberculosis*. *Tuber. Lung Dis.* 80(3), 141–159 (2000).
- 11. Kumar A, Deshane JS, Crossman DK *et al.* Heme oxygenase-1-derived carbon monoxide induces the *Mycobacterium tuberculosis* dormancy regulon. *J. Biol. Chem.* 283(26), 18032–18039 (2008).
- 12. Commandeur S, Lin MY, Van Meijgaarden KE *et al.* Double- and monofunctional CD4 and CD8 T-cell responses to *Mycobacterium tuberculosis* DosR antigens and peptides in long-term latently infected individuals. *Eur. J. Immunol.* 41(10), 2925–2936 (2011).
- 13. David R. Sherman and Gary K. Schoolnik et al. Inhibition of Respiration by Nitric Oxide Induces a Mycobacterium tuberculosis Dormancy Program. J. Exp. Med. (2003)
- 14. Santhosh Sivaramakrishnan and Paul R. Ortiz de Montellano. The DosS-DosT/DosR Mycobacterial Sensor System. *Biosensors* 2013, *3*, 259-282; doi:10.3390

## THANK YOU!