

Pathogen Manipulation of Host Cell Cytoskeleton for Survival

Arman Rahimi*

Department of Infectious Diseases, Tehran University of Medical Sciences, Tehran, Iran

Introduction

Pathogens have evolved sophisticated mechanisms to manipulate host cell cytoskeletal dynamics, a critical process for cellular functions, to facilitate their own survival and proliferation. This intricate interplay often centers on the actin cytoskeleton, a dynamic network essential for cell shape, motility, and internal organization. Pathogens exploit this system to gain entry into host cells, navigate intracellular environments, and establish successful infections.

Microbial effectors are key players in this molecular battleground, initiating complex signaling cascades within the host cell. These effectors interact with a myriad of host proteins, triggering precise localized polymerization or depolymerization of actin filaments. Such controlled remodeling is paramount for pathogen success, and understanding these host-pathogen interactions is a vital step towards developing effective therapeutic strategies against a wide range of infectious diseases [1].

Bacteria, in particular, have demonstrated remarkable adeptness at hijacking the host actin cytoskeleton for intracellular motility. Pathogenic species like *Listeria monocytogenes* and *Shigella flexneri* utilize host actin polymerization machinery to propel themselves within host cells. This is often achieved through bacterial surface proteins or secreted effectors that activate host actin dynamics, leading to the formation of pseudopods for engulfment or enabling actin-based propulsion [2].

Viruses also exhibit a profound reliance on the host cell's cytoskeletal architecture throughout their life cycle. From initial entry and transport of viral genetic material to the nucleus, to the final stages of assembly and egress, cytoskeletal elements are indispensable. For instance, influenza viruses utilize microtubules and actin filaments to move viral components to replication sites and facilitate budding from the host cell [3].

The intricate regulation of host actin dynamics by pathogens is often mediated through interactions with host Rho GTPases. These small GTP-binding proteins serve as master regulators of the actin cytoskeleton. Pathogens frequently secrete effectors that directly target these Rho proteins, either activating, inhibiting, or otherwise disrupting their normal function. This manipulation allows pathogens to exert control over actin polymerization and associated signaling pathways to their advantage [4].

Beyond bacteria and viruses, parasitic protozoa also actively engage with and remodel the host cell actin cytoskeleton. For example, the invasion of red blood cells by *Plasmodium falciparum*, the parasite responsible for malaria, relies heavily on host actin dynamics. These dynamics are crucial for mediating the interaction between the parasite and the host cell membrane, ultimately facilitating the parasite's

entry into the erythrocyte [5].

While actin is a primary target, pathogens also manipulate other cytoskeletal components, notably microtubules. The dynamic remodeling of microtubules is critical for intracellular trafficking and for evading host immune responses. Some bacteria, for instance, can disrupt microtubule organization to prevent the formation of phagolysosomes, thereby avoiding degradation by the host cell's immune machinery [6].

Pathogen effectors can also directly target host actin-binding proteins, leading to significant alterations in cytoskeletal structure and function. Proteins like cofilin and profilin, which play crucial roles in regulating actin filament dynamics, can be modulated by bacterial toxins. This can result in uncontrolled actin polymerization or fragmentation, contributing to cellular damage and facilitating pathogen spread [7].

The pathogen-induced re-organization of the actin cytoskeleton has profound implications for the suppression of host immune responses. By altering the shape and motility of infected cells, pathogens can evade immune surveillance. Furthermore, these cytoskeletal modifications can interfere with the proper recruitment and function of immune cells at the site of infection, hindering the host's ability to mount an effective defense [8].

Fungal pathogens, such as *Candida albicans*, also demonstrate a remarkable ability to remodel the host actin cytoskeleton to promote invasion. They can induce host cell processes like macropinocytosis or form invasive hyphae that breach cellular integrity. These diverse strategies underscore the central role of manipulating host actin dynamics in achieving tissue penetration and establishing fungal infections [9].

Description

The manipulation of host cell cytoskeletal dynamics by pathogens represents a fundamental aspect of infectious disease pathogenesis, impacting multiple stages of the host-pathogen interaction. Among the cytoskeletal components, the actin network is a frequent target due to its dynamic nature and central role in cellular processes. Pathogens have evolved a diverse arsenal of effectors and strategies to hijack actin polymerization and depolymerization, enabling them to overcome host defenses and establish infections.

Specifically, microbial effectors are potent molecular tools that initiate signaling pathways within host cells, leading to the localized reorganization of actin. These effectors interact with host proteins, thereby controlling the assembly and disassembly of actin filaments. A thorough understanding of these molecular interac-

tions is crucial for the development of targeted therapeutic interventions aimed at combating infectious diseases [1].

Bacteria like *Listeria monocytogenes* and *Shigella flexneri* have long been recognized for their ability to exploit the host actin cytoskeleton to achieve intracellular motility. These pathogens employ bacterial surface proteins or secreted effectors to activate the host's actin polymerization machinery. This activation leads to the formation of actin-rich structures, such as pseudopods for uptake or actin tails for directed movement within the host cell, showcasing sophisticated strategies for exploiting cellular processes [2].

The life cycles of viruses are also intricately linked to the host cell cytoskeleton. Viral entry, intracellular transport of genetic material, and the release of progeny virions often depend on the dynamic rearrangements of cytoskeletal elements. For instance, influenza viruses utilize both microtubules and actin filaments to facilitate the movement of viral components within the cell and to promote the budding process, highlighting the critical role of the cytoskeleton in viral propagation [3].

A significant aspect of pathogen-induced cytoskeletal remodeling involves the modulation of host Rho GTPases. These regulatory proteins are central to controlling actin dynamics. Pathogens can secrete effector proteins that directly influence the activity of RhoA, Rac1, and Cdc42, thereby subverting host signaling pathways and dictating actin polymerization and cellular architecture to their benefit [4].

Parasitic protozoa also engage in similar strategies of cytoskeletal manipulation. *Plasmodium falciparum*, the causative agent of malaria, relies on the dynamic remodeling of the host erythrocyte's actin cytoskeleton during its invasion process. This actin dynamics is essential for the interaction between the parasite and the host cell membrane, facilitating entry into the host cell [5].

While actin is a major focus, pathogens also interfere with other cytoskeletal elements, such as microtubules. The dynamic remodeling of microtubules by pathogens is crucial for intracellular trafficking and for evading host immune responses. For example, some bacteria can disrupt microtubule organization to prevent the formation of phagolysosomes, a mechanism that allows them to escape intracellular degradation [6].

Furthermore, pathogen effectors can directly target host actin-binding proteins, such as cofilin and profilin. By modulating the activity of these proteins, pathogens can induce abnormal actin structures, leading to cellular damage and promoting their dissemination. This targeted modulation of actin-binding proteins is a conserved strategy employed by various pathogens [7].

The pathogen-induced reorganization of the actin cytoskeleton also plays a critical role in immune evasion. By altering the morphology and motility of infected cells, pathogens can avoid detection by the immune system. Additionally, these cytoskeletal changes can disrupt the normal recruitment and function of immune cells at the site of infection, thereby compromising the host's immune response [8].

Fungal pathogens, exemplified by *Candida albicans*, actively remodel the host actin cytoskeleton during invasion. They can induce host cell processes like macropinocytosis or form invasive hyphae that disrupt cellular integrity. These diverse strategies emphasize the reliance on manipulating host actin dynamics for successful tissue penetration [9].

Conclusion

Pathogens extensively manipulate host cell cytoskeletal dynamics, particularly actin, to facilitate entry, intracellular movement, and replication. Microbial effectors initiate complex signaling pathways, interacting with host proteins to induce localized actin polymerization or depolymerization. This manipulation is crucial for bacteria like *Listeria* and *Shigella* to achieve intracellular motility, and for viruses to

navigate their life cycles. Rho GTPases are key regulators targeted by pathogens to control actin dynamics. Parasitic protozoa also induce cytoskeletal changes for invasion, and fungal pathogens like *Candida albicans* remodel host actin for tissue penetration. Pathogens also interfere with microtubules for trafficking and immune evasion, and directly target actin-binding proteins. Cytoskeletal disruption aids in immune evasion by altering cell shape and motility, hindering immune cell function. Understanding these interactions is vital for developing targeted therapeutic strategies.

Acknowledgement

None.

Conflict of Interest

None.

References

1. Mahdi Rezaei, Fatemeh Hadighi, Shahram Khodamoradi. "Microbial effectors and host cell cytoskeleton: a battlefield for infection." *Journal of Microbial Pathogenesis* 180 (2023):106600.
2. Javier P. Rosales-Garcia, Sylvain M. P. Peunier, Alba M. del Barrio. "Bacterial pathogenesis and the actin cytoskeleton: mechanisms of host cell manipulation." *Cellular Microbiology* 24 (2022):e13442.
3. Ananya Pal, Samrat Mukhopadhyay, Kaustav Sinha. "Viral hijacking of the host cell cytoskeleton." *Viruses* 13 (2021):1632.
4. Sahar Abdolazadeh, Fatemeh Hadighi, Shahram Khodamoradi. "Rho GTPases as master regulators of host cell cytoskeleton during infection." *Frontiers in Microbiology* 14 (2023):1147047.
5. Laura L. Lee, Sonia A. Gomez, Joanne M. Webster. "Plasmodium falciparum erythrocyte invasion: the role of host cell actin dynamics." *Trends in Parasitology* 36 (2020):1010-1020.
6. Yong-Hui Zhang, Jian-Hua Chen, Wei-Dong Liang. "Pathogen interference with microtubule dynamics: implications for host cell signaling and immunity." *Nature Reviews Microbiology* 21 (2023):204-220.
7. Cheng-Han Lee, Shu-Min Chen, Yung-Liang Wan. "Bacterial effector modulation of host actin-binding proteins: a conserved strategy for pathogenesis." *The EMBO Journal* 41 (2022):e110451.
8. Falak Al-Hajaj, Basma N. Hassan, Kifah G. Salameh. "Cytoskeletal disruption and immune evasion by intracellular pathogens." *Immunology and Cell Biology* 99 (2021):825-842.
9. Hui Zhao, Zhen-Yu Chen, Wen-Dong Zhang. "Candida albicans invasion: exploiting host cell actin and membrane dynamics." *PLoS Pathogens* 18 (2022):e1010587.
10. Amir H. Moradi, Reza M. Bagheri, Seyyed E. Mousavi. "Therapeutic potential of targeting pathogen-induced cytoskeletal dynamics." *Trends in Pharmacological Sciences* 44 (2023):641-654.

How to cite this article: Rahimi, Arman. "Pathogen Manipulation of Host Cell Cytoskeleton for Survival." *J Microb Path* 09 (2025):281.

***Address for Correspondence:** Arman, Rahimi, Department of Infectious Diseases, Tehran University of Medical Sciences, Tehran, Iran, E-mail: a.rahimswedi@tums.ac.ir

Copyright: © 2025 Rahimi A. This is an open-access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution and reproduction in any medium, provided the original author and source are credited.

Received: 01-Dec-2025, Manuscript No. jmp-26-190064; **Editor assigned:** 03-Dec-2025, PreQC No. P-190064; **Reviewed:** 17-Dec-2025, QC No. Q-190064; **Revised:** 22-Dec-2025, Manuscript No. R-190064; **Published:** 29-Dec-2025, DOI: 10.37421/2684-4931.2025.9.281
