

Importance of Virulence Factors in Bacterial Pathogenicity: A Review

Sura A. Abdulateef¹, Hasan A. Aal Owaif², Mohanad H. Hussein³

¹Department of Applied Sciences, University of Technology-Iraq

²Department of Plant Biotechnology, College of Biotechnology, Al-Nahrain University, Baghdad, Iraq

³Department of Molecular and Medical Biotechnology, College of Biotechnology, Al-Nahrain University, Baghdad, Iraq.

ABSTRACT

Understanding the strategies of bacterial infections requires an extensive knowledge of the essential role virulence factors play in bacterial pathogenicity. These components are essential in spreading of the infection, the capacity of the bacteria to avoid host defenses and the start of disease. The absence or presence of certain virulence factors has significant influence on the rate of infection dissemination. Virulence factors are divided into four groups: adhesion factors, invasion factors, toxin factors and immune evasion factors. The review here, addresses how virulence parameters affect host cells and tissues, how they may influence disease development and how they may be utilized to identify and treat bacterial infections. Controlling virulence parameters is also an important aspect of this. Virulence parameters expression can be influenced by variety of genetic and environmental variables, including quorum sensing which is not always present. Finally developing efficient prevention and therapeutic techniques for bacterial infections requires detailed understanding of the control of virulence parameters. Indicating the important role of virulence factors in bacterial pathogenicity is vital in the development of powerful treatment methods.

KEYWORDS: Virulence factors, adhesion, toxins, immune evasion, bacterial pathogenicity, regulation.

ARTICLE DETAILS

Published On:
22 April 2023

Available on:
<https://ijmscr.org/>

I. INTRODUCTION

Virulence factors are essential components that enable bacteria to establish infection, overcome host defenses and cause disease. These elements can be generally divided into two categories: those that assist tissue injury and those that encourage colonization. Adhesins, pili, and capsules are colonization factors, whereas toxins, proteases and lipases are factors that cause tissue injury.^{1,2} The severity of the infection and the progression of the disease are influenced by the presence or absence of particular virulence factors. For instance, while other strains of *Escherichia coli* lack these virulence factors and only produce mild gastroenteritis, some strains do carry virulence factors that allow them to induce severe diarrhea and kidney damage.³

Developing strategies to fight bacterial infections requires an understanding of the molecular mechanisms that control virulence factors. The understanding of virulence factors and the signaling pathways that regulate their expression has advanced significantly over the years.^{1,2} Virulence factors, their significance in bacterial pathogenicity, and the most

recent developments in our understanding of their regulation will all be covered in this review.

II. TYPES OF VIRULENCE FACTORS

Virulence factors are classified into toxins, immune evasion factors, adhesion factors and invasion factors. Understanding these virulence determinants, is critical for effectively preventing and treating bacterial infections.⁴

Adhesion Factors

Adhesion factors are substances that allow bacteria to adhere to host cells or elements of the extracellular matrix, promoting colonization and infection. Examples of protein-containing adhesion parameters, are pili and fimbriae while lipopolysaccharides and capsules, are non-protein-containing adhesion factors. Fimbriae and pili are surface features that protrude from the bacterial surface and help the bacterium attach to host cells. The proteins that make up fimbriae and pili can differ between bacterial species, enabling various bacteria to bind to various host cells. For instance, *Pseudomonas aeruginosa* causes pneumonia by attaching to

Importance of Virulence Factors in Bacterial Pathogenicity: A Review

pulmonary epithelial cells with its type IV pili.⁵ Non-proteinaceous adhesion components called capsules surround bacteria in a protective coating, shielding it from the host immune system's detection. A typical bacterium called *Streptococcus pneumoniae* employs a capsule to escape the host's immune system and spread infections.⁶

Invasion Factors

These components include effector proteins that change host cell signaling pathways to aid bacterial survival as well as secretion systems that deliver bacterial proteins into host cells to affect cellular processes. For instance, *Salmonella enterica* uses a type III secretion system to enter host cells and release effector proteins that alter host cell signaling pathways to aid in bacterial replication and survival.⁷ Motility, which permits bacteria to move through host tissues and acquire fresh niches for colonization, is another role in invasion. It is possible for bacteria to migrate through liquids and into host tissues thanks to their surface features called flagella. For instance, gastritis is brought on by *Helicobacter pylori*, which utilizes its flagella to pierce the mucous layer of the stomach.⁸

Toxins

Toxins are virulence components that harm host tissues and advance the development of bacterial diseases. Endotoxins and exotoxins are the two main types of bacterial toxins. Endotoxins which are essential factors of the bacterial cell wall, are released during bacterial lysis. Lipopolysaccharide (LPS) as endotoxin, causes inflammatory reaction in the host tissue. When host immune cells known as macrophages encounter LPS molecules, they release cytokines that stimulate inflammation and draw other immune cells to the infection site.⁹ Bacteria release exotoxins into the extracellular environment. Exotoxins can be further divided into groups according to how they work, including cytotoxins, superantigens and toxins that damage membranes. Hemolysin which destroy red blood cells and leukocidin which destroy white blood cells, are examples of membrane-damaging toxins. Superantigen toxin can cause systemic inflammation and shock by activating an overactive immune response. While cytotoxin can cause host cells to die by rupturing or impairing their metabolic systems.¹⁰ The cholera toxin produced by *Vibrio cholerae* is an illustration of a cytotoxin, it results in diarrhea by interfering with ion transport in the gut.¹¹

Immune Evasion Factors

Immune evasion factors as a virulence factors allow bacteria to subvert or control the immunological response of the host. Proteins that prevent complement activation, a crucial part of the host innate immune system, are among these factors. The complement system is a collection of proteins that can be triggered by elements of the bacterial cell wall or antibodies, which results in the bacterial cells being destroyed. Bacteria can avoid complement activation by creating proteins that bind to and deactivate complement proteins. *Staphylococcus aureus* as an example, produces staphylococcal complement

inhibitor protein that binds to C3 complement protein and prevents its activation.¹²

By producing proteins that prevent phagocytosis, the process by which immune cells engulf and kill bacterial cells, bacteria can also elude the host immunological response. For instance, the bacteria *Streptococcus pyogenes* makes the M protein, which binds to complement factor H, a regulator of the alternative complement pathway. M protein inhibits the opsonization and phagocytosis of *S. pyogenes* by host immune cells by binding to complement factor H.¹³

III. IMPORTANCE OF VIRULENCE FACTORS IN BACTERIAL PATHOGENESIS

Although most species of bacterial are not hazardous to humans, several can cause serious diseases that include simple infections to severe problems. The presence of virulence factors, which are molecular parameters, that enable bacterial colonization, penetration and harm host tissues is highly responsible for initiation of disease caused by bacteria. In order to develop successful prevention and treatment methods for bacterial infections, it is important to comprehend the significance of virulence factors in bacterial pathogenesis.¹⁴

Impact of Virulence Factors on Host Cells and Tissues

Virulence factors of bacteria have important role in bacterial colonization and invasion of host cells and tissues. The infection process and the development of disease depend on the bacteria capacity to adhere to and infiltrate host cells. When bacteria adhere to host cells, the infection process begins,¹⁵ and this process is controlled by a variety of adhesins or fimbriae on the bacterial surface. Adhesins attach to particular receptors on the host cell surface, enabling the bacteria to adhere to the host and colonize it. For instance, it has been reported that Type 1 fimbriae of *E. coli* increase bacterial pathogenicity for the urinary system by encouraging adherence to uroepithelial cells.¹⁵

Invasion factors enhance Bacterial pathogenicity by allowing bacteria to enter and multiply inside host cells. *Salmonella's* secretion system is one such; it is utilized to introduce bacterial proteins into host cells, interfering with normal cellular operations and promoting bacterial proliferation.¹⁶ A different example is the lipoarabinomannan (LAM) generated by *Mycobacterium TB*, which causes phagosome maturation arrest and enables the bacteria to multiply inside macrophages while thwarting the host immune response.¹⁷

Role of Virulence Factors in Disease Progression

Infectious disease progression depends heavily on bacterial virulence factors. Colonization of the host tissue, the first step of bacterial infection, necessitates bacterial attachment to host cells, which is mediated by virulence proteins such fimbriae and adhesins. After effectively invading host tissues, bacteria can multiply and create toxins and other virulence factors that can harm host cells and tissues and cause illness symptoms. As an illustration, *Streptococcus pyogenes* creates

Importance of Virulence Factors in Bacterial Pathogenicity: A Review

the streptolysin O (SLO) toxin, which harms host cells and tissues and causes the signs and symptoms of streptococcal pharyngitis.¹⁸

Bacterial virulence factors can interfere with the immune response of the host causing direct harm to host cells and tissues and lead to support bacterial survival and persistence. For instance, altering surface features of certain bacteria make them unrecognized by the host immune system. In order to avoid being recognized by antibodies created by the host immune system, a process known as antigenic variation modifies surface proteins.¹⁹

The role of virulence factors in biofilm formation represents another way in which they can affect disease progression. Biofilms are bacterial assemblages that attach to surfaces and are encased in a matrix of extracellular polymeric molecules. By forming biofilms, bacteria can resist the host immune responses and antibiotic causing chronic infections that are difficult to treat. Adhesins and exopolysaccharides which enable bacterial attachment to the surfaces and build the extracellular matrix, are important virulence factors in biofilm formation.²⁰

Bacterial Infection Diagnosis and Treatment Using Virulence Factors

Bacterial virulence factors are also utilized to diagnose and treat infections caused by bacteria. Bacterial pathogenicity can be detected by using virulence factors as diagnostic tool. The presence of the *S. aureus* virulence component Pantone-Valentine leukocidin, for example, is associated with severe skin and soft tissue infections.²¹ Some virulence factors can also be used to differentiate between bacterial pathogenic strains and track the disease progress.²²

Identification of virulence factors is important because they represent potential targets for generating new antibiotics for bacterial infections. Traditional antibiotic therapy focuses on important bacterial functions including the formation of cell wall and the synthesis protein. However, the need for generating new treatment techniques is resulted from the increasing incidence of bacterial antibiotic-resistant. Targeting virulence factors offers an alternate method of treating bacterial infections by interfering with bacterial ability to cause disease without necessarily killing the bacterium.²³ The use of small molecule inhibitors that interfere with virulence factor function is another promising strategy to addressing virulence factors. The chemical 2-aminobenzimidazole, as an example, was reported in animal models of infection to prevent the development of virulence factors in *P. aeruginosa* and decrease pathogenicity. This technique aids to provide new antimicrobial agents for treatment of bacterial infections, especially those caused by antibiotic-resistant bacteria.²⁴

IV. REGULATION OF VIRULENCE FACTORS

The virulence factors expression is regulated by genetic and environmental parameters.

Genetic Regulation of Virulence Factors

Bacterial virulence factors are encoded by genes that are usually organized in operons. The expression of such genes is often regulated by transcriptional factors that respond to environmental signals. The *fim* operon, as an example, regulates the type 1 fimbriae expression which are essential adhesins in *E. coli*.²⁵ This operon contains the genes required for fimbriae synthesis, assembly and export. As well as FimB, a transcriptional regulator, that regulates the switch between the type 1 fimbriae expression and the expression of type P fimbriae is another type of fimbriae that produced by *E. coli*.²⁶ Other transcriptional regulators regulate virulence factor expression in response to host signals.²⁷ The *agr* (accessory gene regulator) system, for example, regulates the expression of various virulence factors, including alpha-toxin, in *S. aureus*.²⁸ Post-transcriptional processes, in addition to transcriptional regulation, can influence virulence gene expression. In *P. aeruginosa*, for example, the expression of the exotoxin A (ETA) gene is regulated by RsmY, a short RNA that binds to and stabilizes the ETA gene's mRNA. In contrast, another short RNA termed RsmZ binds to RsmY and sequesters it, lowering ETA expression.²⁹

Environmental Factors that Influence Virulence Expression

Bacterial virulence factors can also be influenced by environmental parameters such as temperature, pH, nutrition availability, and oxygen levels. For example, low pH and high temperature promote the production of the cholera toxin in *V. cholerae*, mimicking the conditions encountered in the human small intestine.¹¹ In a similar way, high pH and low Mg²⁺ which are intestinal lumen cues, promote type III secretion system production in *S. enterica*.³⁰

Moreover, the expression of virulence factors can be stimulated by host parameters. In *H. pylori*, for instance, the acidic conditions in stomach and interaction with epithelial cells stimulate the expression of the *cag* pathogenicity island.^{31,32} The iron-depleted medium can increase the type III secretion system expression in *Yersinia pseudotuberculosis*, which is essential for survival in the host.³³

The Role of Quorum Sensing in Virulence Factors Expression

Quorum sensing is a communication system between cells that allows bacteria to monitor their population density and adjust their activity accordingly. Quorum sensing is mediated by small signaling molecules known as autoinducers, which are produced by bacteria and diffuse across the cell membrane. When autoinducers reach a certain concentration, they bind to transcriptional regulators known as LuxR-type proteins, which activate or repress the expression of target genes including virulence genes.³⁴

Quorum sensing is found throughout bacteria and is known to regulate a number of bacterial characteristics such as biofilm formation, motility, and virulence expression.^{34,35}

Importance of Virulence Factors in Bacterial Pathogenicity: A Review

Quorum sensing is usually involved in the regulation of virulence factors in pathogenic bacteria. In *P. aeruginosa*, for example, it regulates the expression of many virulence factors such as pyocyanin, elastase and rhamnolipids. Quorum sensing is mediated in *P. aeruginosa* by 3-oxo-C12-HSL, C4-HSL and PQS (2-heptyl-3-hydroxy-4-quinolone), as well as two LuxR-type transcriptional regulators, LasR and RhIR. LasR binds to 3-oxo-C12-HSL and activates the expression of genes involved in pyocyanin and elastase production, whereas RhIR attaches to C4-HSL and increases the expression of genes involved in rhamnolipid formation.^{36,37,38} The agr system mediates quorum sensing in *S. aureus* and regulates the synthesis of surface proteins such as protein A, which is important for immune evasion, also agr system regulates virulence factors expression.³⁹

V. CONCLUSION

The role of virulence factors in bacterial pathogenicity cannot be emphasized. These factors are essential for bacterial ability to cause diseases and their absence or presence can affect the severity of infection and progression of disease. Virulence factors are divided into four categories, adhesion, invasion, toxins and immune evasion factors. Understanding these characteristics is essential for developing successful treatment techniques, and prevent bacterial infection. Moreover; the regulation of virulence factors is an important topic with current studies shedding insight on the genetic and environmental variables that regulate virulence factor expression. This information can be utilized to generate new treatments and also prevent the spread of antibiotic-resistant bacteria.

Finally, studying virulence factor of pathogenic bacteria is an essential field of research that has significant implications for human health. It is important to carry on research about the role of virulence factors in bacterial infections, and to generate new techniques to prevent diseases.

REFERENCES

- I. Finlay, B. B. and Falkow, S. (1997). Common themes in microbial pathogenicity revisited. *Microbiology and Molecular Biology Reviews*, 61(2): 136-169.
- II. Pizarro-Cerdá, J. and Cossart, P. (2006). Bacterial adhesion and entry into host cells. *Cell*, 124(4): 715-727.
- III. Pakbin, B., Brück, W. M. and Rossen, J. W. A. (2021). Virulence Factors of Enteric Pathogenic *Escherichia coli*: A Review. *Int J Mol Sci*. 22(18): 9922.
- IV. Casadevall, A. and Pirofski, L. A. (2009). Virulence factors and their mechanisms of action: the view from a damage-response framework. *J Water Health*. 7: S2-S18.
- V. Giltner, C. L., Nguyen, Y. and Burrows, L. L. (2012). Type IV pilin proteins: versatile molecular modules. *Microbiology and molecular biology reviews*, 76(4): 740-772.
- VI. Weiser, J. N., Ferreira, D. M. and Paton, J. C. (2018). *Streptococcus pneumoniae*: transmission, colonization and invasion. *Nat Rev Microbiol*. 16(11): 355-367.
- VII. Figueira, R. and Holden, D. W. (2012). Functions of the *Salmonella* pathogenicity island 2 (SPI-2) type III secretion system effectors. *Microbiology*, 158(5): 1147-1161.
- VIII. Lertpiriyapong, K., Whary, M. T., Muthupalani, S., Lofgren, J. L., Gamazon, E. R., Feng, Y. and Fox, J. G. (2014). Gastric colonisation with a restricted commensal microbiota replicates the promotion of neoplastic lesions by diverse intestinal microbiota in the *Helicobacter pylori* INS-GAS mouse model of gastric carcinogenesis. *Gut*, 63(1): 54-63.
- IX. Raetz, C. R. and Whitfield, C. (2002). Lipopolysaccharide endotoxins. *Annu Rev Biochem*. 71: 635-700.
- X. Bae, J., Jin, H., Kim, J., Park, M., Lee, J. and Kim, S. (2021). Molecular Characteristics and Exotoxins of Methicillin-Resistant *Staphylococcus aureus*. *Biomedical Science Letters*, 27(4): 195-207.
- XI. Kaper, J. B., Morris, J. G. and Levine, M. M. Cholera. (1995). *Clin Microbiol Rev*. 8(1): 48-86.
- XII. Hair, P. S., Echague, C. G., Sholl, A. M., Watkins, J. A., Geoghegan, J. A., Foster, T. J. and Cunnion, K. M. (2008). *Staphylococcus aureus* clumping factor A binds to complement regulator factor I and increases factor I cleavage of C3b. *J Infect Dis*. 198(1):125-133.
- XIII. Fieber, C. and Kovarik, P. (2014). Responses of innate immune cells to group A *Streptococcus*. *Front. Cell. Infect. Microbiol*. 4:140.
- XIV. Depluvere, S., Devos, S. and Devreese, B. (2016). The Role of Bacterial Secretion Systems in the Virulence of Gram-Negative Airway Pathogens Associated with Cystic Fibrosis. *Front Microbiol*. 7: 1336.
- XV. Mulvey, M.A., Schilling, J. D., Martinez, J. J. and Hultgren, S. J. (2000). Bad bugs and beleaguered bladders: interplay between uropathogenic *Escherichia coli* and innate host defenses. *Proc Natl Acad Sci U S A*. 97(16): 8829-8835.
- XVI. McGhie, E. J., Brawn, L. C., Hume, P. J., Humphreys, D. and Koronakis, V. (2009). *Salmonella* takes control: effector-driven manipulation of the host. *Curr Opin Microbiol*. 12(1): 117-124.
- XVII. Cambier, C. J., Falkow, S. and Ramakrishnan, L. (2014). Host evasion and exploitation schemes of *Mycobacterium tuberculosis*. *Cell*. 159(7): 1497-1509.

Importance of Virulence Factors in Bacterial Pathogenicity: A Review

- XXVIII. Cunningham, M. W. (2000). Pathogenesis of group A streptococcal infections. *Clinical Microbiology Reviews*, 13(3): 470-511.
- XIX. Moxon, E. R., Rainey, P. B., Nowak, M. A. and Lenski, R. E. (1994). Adaptive evolution of highly mutable loci in pathogenic bacteria. *Current Biology*, 4(1): 24-33.
- XX. Costerton, J. W., Stewart, P. S. and Greenberg, E. P. (1999). Bacterial biofilms: a common cause of persistent infections. *Science*, 284(5418): 1318-1322.
- XXI. Lina, G., Piémont, Y., Godail-Gamot, F., Bes, M., Peter, M. O., Gauduchon, V., Vandenesch, F. and Etienne, J. (1999). Involvement of Panton-Valentine leukocidin-producing *Staphylococcus aureus* in primary skin infections and pneumonia. *Clinical Infectious Diseases*, 29(5): 1128-1132.
- XXII. Grundmann, H., Schouls, L. M., Aanensen, D. M. A., Pluister, G. N., Tami, A., Chlebowicz, M., Glasner, C., Sabat, A. J., Weist, K., Heuer, O. and Friedrich, A. W. (2014). The dynamic changes of dominant clones of *Staphylococcus aureus* causing bloodstream infections in the European region: results of a second structured survey. *Euro Surveill*. 19(49): 20987.
- XXIII. Dehbanipour, R. and Ghalavand, Z. (2022). Anti-virulence therapeutic strategies against bacterial infections: recent advances. *Germs*. 12(2): 262-275.
- XXIV. Hentzer, M., Wu, H., Andersen, J. B., Riedel, K., Rasmussen, T. B., Bagge, N., Kumar, N., Schembri, M. A., Song, Z., Kristoffersen, P., Manefield, M., Costerton, J. W., Molin, S., Eberl, L., Steinberg, P., Kjelleberg, S., Høiby, N. and Givskov, M. (2003). Attenuation of *Pseudomonas aeruginosa* virulence by quorum sensing inhibitors. *EMBO J*. 22(15): 3803-3815.
- XXV. Sivick, K. E. and Mobley, H. L. T. (2010). Waging War against Uropathogenic *Escherichia coli*: Winning Back the Urinary Tract. *Infect. Immun*. 78(2): 568-585.
- XXVI. Gahlot, D.K., Taheri, N. and MacIntyre, S. (2023). Diversity in Genetic Regulation of Bacterial Fimbriae Assembled by the Chaperone Usher Pathway. *Int. J. Mol. Sci*. 24(1): 161.
- XXVII. Aal Owaif, H. A., Mhawesh, A. A. and Abdulateef, S. A. (2019). The role of BipA in the regulation of K1 capsular polysaccharide production of uropathogenic *Escherichia coli*. *Ann Trop Med Public Health*. 22: S254.
- XXVIII. Novick, R. P., Ross, H. F., Projan, S. J., Kornblum, J., Kreiswirth, B. and Moghazeh, S. (1993). Synthesis of staphylococcal virulence factors is controlled by a regulatory RNA molecule. *The EMBO Journal*, 12(10): 3967-3975.
- XXIX. Brencic, A., McFarland, K. A., McManus, H. R., Castang, S., Mogno, I., Dove, S. L. and Lory, S. (2009). The GacS/GacA signal transduction system of *Pseudomonas aeruginosa* acts exclusively through its control over the transcription of the RsmY and RsmZ regulatory small RNAs. *Molecular Microbiology*, 73(3): 434-445.
- XXX. Eriksson, S., Lucchini, S., Thompson, A., Rhen, M. and Hinton, J. C. (2003). Unravelling the biology of macrophage infection by gene expression profiling of intracellular *Salmonella enterica*. *Mol Microbiol*. 47(1): 103-118.
- XXXI. Odenbreit, S., Puls, J., Sedlmaier, B., Gerland, E., Fischer, W. and Haas, R. (2000). Translocation of *Helicobacter pylori* CagA into gastric epithelial cells by type IV secretion. *Science*, 287: 1497-1500.
- XXXII. Amieva, M. R., and El-Omar, E. M. (2008). Host-bacterial interactions in *Helicobacter pylori* infection. *Gastroenterology*, 134(2): 306-323.
- XXXIII. Zhang, Y., Romanov, G. and Bliska, J. B. (2011). Type III Secretion System-Dependent Translocation of Ectopically Expressed Yop Effectors into Macrophages by Intracellular *Yersinia pseudotuberculosis*. *Infect Immun*, 79(11): 4322-4331.
- XXXIV. Miller, M. B. and Bassler, B. L. (2001). Quorum sensing in bacteria. *Annu Rev Microbiol*, 55: 165-199.
- XXXV. Fuqua, W. C., Winans, S. C. and Greenberg, E. P. (1994). Quorum sensing in bacteria: the LuxR-LuxI family of cell density-responsive transcriptional regulators. *J Bacteriol*, 176(2): 269-275.
- XXXVI. Passador, L., Cook, J. M., Gambello, M. J., Rust, L. and Iglewski, B. H. (1993). Expression of *Pseudomonas aeruginosa* virulence genes requires cell-to-cell communication. *Science*, 260(5111): 1127-1130.
- XXXVII. Pearson, J. P., Pesci, E. C. and Iglewski, B. H. (1997). Roles of *Pseudomonas aeruginosa* las and rhl quorum-sensing systems in control of elastase and rhamnolipid biosynthesis genes. *Journal of bacteriology*, 179(18): 5756-5767.
- XXXVIII. Latifi, A., Foglino, M., Tanaka, K., Williams, P., and Lazdunski, A. (1996). A hierarchical quorum-sensing cascade in *Pseudomonas aeruginosa* links the transcriptional activators LasR and RhIR (VsmR) to expression of the stationary-phase sigma factor RpoS. *Molecular microbiology*, 21(6): 1137-1146.
- XXXIX. Le, K. Y. and Otto, M. (2015). Quorum-sensing regulation in staphylococci-an overview. *Frontiers in microbiology*, 6: 1174.