

Review Article

Microbial Science Archives (ISSN: 2583-1666) An International Research Journal Journal homepage: www.microbial.sciencearchives.org

https://doi.org/10.47587/MSA.2024.4201



# Virulence and Pathogenicity of Clostridium perfringens

Shler Ali Khorseed<sup>™</sup>

College of sciences, Kirkuk university, Iraq Received: April 22, 2024/ Revised: May 20, 2024/Accepted: May 29, 2024 (☑) Corresponding Author: <u>shler-ali@uokirkuk.edu.iq</u>

## Abstract

*Clostridium perfringens* is anaerobic bacterium, exhibits a broad spectrum of pathogenicity with significant implications for human and animal health. Its virulence is intricately linked to the production of various toxins, notably alpha toxin, beta toxin, and epsilon toxin, which play pivotal roles in tissue damage and disease manifestation. The pathogen makes convoluted host-pathogen interactions requiring some kind of adhesion, colonization, and evasion of the host immune response Epidemiologically, *C. perfringens* could be a main reason for foodborne diseases, with Enterotoxigenic strains making gastroenteritis. Improper food handling practices play part in outbreaks. The field of veterinary epidemiology links *C. perfringens* to necrotic enteritis in poultry, which affects animal welfare and agricultural productivity. The significance of making collaborations with a One Health perspective when coping with issues related with *C. perfringens* is important. Surveillance, preventive measures, and outbreak examinations are seen to be included within interventions of public health. Understanding the epidemiology of *C. perfringens* could be essential for sake of implementing targeted strategies aiming at diminishing foodborne infections.

Keywords: Clostridium perfringens, Pathogenicity, Virulence.

# Introduction

The Gram-positive, spore-generating bacterium Clostridium perfringens can be broadly renowned for its clinical significance just because of its severe pathogenicity. As far as the organism is anaerobic (Robertson, 2013), soil and the human digestive system could be better instances of lowoxygen environments where Candida perfringens flourish. Too many poisons and pili made by this kind of bacterium assist its pathogenic qualities as well as cause various infections in both humans and animals (Adams Vroom et al., 2018). However, the ability of C. perfringens to generate some spores, last in a variety of conditions, leading to a diverse variety of toxins with unique mechanisms of action are the basic factors that have a kind of contribution to its pathogenicity. From minor gastrointestinal disorders to severe, fatal diseases, such toxins are considered as key factors leading to a variety of diseases. (Craig & Melville, 2013). By stressing on the factors of virulence and complex interactions that C. perfringens has with its hosts, this introduction helps in

a more in-depth examination of the cellular and molecular aspects of the bacteria's pathogenicity. Clostridium *perfringens* is a pathogen which could basically be a leading threat for the health of both human and animal (Bio Number Details Page, 105474). C. perfringens is a spore-forming, Gram-positive anaerobe that employs its versatility to flourish in a range of niches, from soil to the intestines of its hosts (Beingesser et al., 2013). Furthermore, the pathogenicity related to C. perfringens can definitely and increasingly be related to its capacity to generate numerous toxins such as beta, epsilon, iota, and alpha toxins, where each one of those kinds of toxins holds a unique effect on host cells. Therefore, such toxins can extremely grant C. perfringens the capability to disrupt cellular processes, which undoubtedly causing tissue damage and leading multiple diseases to appear. The process of detecting the mechanisms laid beyond C. perfringens' virulence could indefinitely be a core part for creating effective treatments and preventative strategies to fight infections that are originally caused by this virulent bacteria.(Navarro, et al., 2021).. In this study of C. perfringens

virulence, the molecular mechanisms are initially examined in front of toxin production, host-pathogen interactions, and the repercussions of such kind of processes on disease outcomes. By addressing the enigma ambient to the virulence of C. perfringens, it is hoped that it may facilitate novel methods to appear in order to diminsh the detrimental effects on public health. (Fernandez-Miyakawa and Redondo, 2016). Beta, epsilon, iota, and alpha toxins are considered as basic tools in C. perfringens' armory. Each of them targets a different area of the cell through a unique mechanism of action. Collectively, the cells could extremely be damaged by such toxins, causing inflammation, and facilitate the spread of infections in both humans and animals..(Duncan et al., 1978). Recognizing the complexity of *C. perfringens* pathogenicity requires that knowing the nuances of host-pathogen dynamics are imperative. The bacteria evades immune surveillance and makes use of the mechanisms of its host cell to promulgate and stay alive as it circumnavigates the complex surroundings of the host.( Hall et al., 2018). The research goal is to provide up-to-date knowledge to the complex interactions among C. perfringens and its host organisms, so that targeted handlings can be improved to decrease the damaging effects of the bacterium on human health and welfare. Contextually, the mechanisms of action and host cell defenses were dissected and employed by C. perfringens toxins to examine their molecular makeup.(Li et al., 2011; Brown et al., 1993). In epidemiology, Clostridium perfringens, a bacterium wellknown for its association with foodborne infections, is critical. The epidemiological landscape of the bacteria is shaped by the ability of C. perfringens to make toxins that cause gastroenteritis in humans, produce robust spores, and survive in a variety of conditions. Animal and public health are at risk from the multifarious pathogen Clostridium perfringens, which has to do with wound infections and necrotic enteritis. It undoubtedly has a kind of considerable effects on both human and animal health. Toxins and spores made by C. perfringens take privilege of wound infections to cause gas gangrene and other severe tissue-related diseases(Kio & Hall, 2018). These diseases require a sophisticated understanding of host-pathogen interactions and bacterial virulence factors for effective management due to their significant clinical problems. C. perfringens frequently causes necrotic enteritis in animals, which has detrimental financial effects on the poultry industry (Revit-Mills, 2015).

# Clostridium perfringens

The round or subterminal spore found in the rod-shaped bacterium *Clostridium perfringens* gives it a boxcar-like appearance. Gram Stain: The violet stain from the Gram staining procedure is kept because it is Gram-positive (Hall & Kio, 2018). *Clostridium perfringens* is an anaerobic bacteria that grows in environments with little to no oxygen (Rood et al., 2018).Formation of Spores: It can produce spores, resilient structures that aid in the bacterium's survival in harsh environments. (Hassan et al., 2015). One of the main virulence factors that contribute to *Clostridium perfringens'* pathogenicity is its production of alpha toxin, one of the toxins

that the bacterium is known for producing. Because of its peritrichous flagella, which are uniformly distributed across its surface, the bacterium is typically motile (Wampum, Joseph; et al., 2021). It is a common cause of food poisoning, usually linked to the consumption of contaminated food, and is widely distributed in nature, especially in soil and the intestinal tracts of animals and humans. Infections and gas gangrene can also result from it. It can ferment a wide range of carbohydrates, yielding organic acids and gas as byproducts. Recall that *Clostridium perfringens* can be harmful in some situations but can also be helpful in others (such as the gut microbiota) (McClane, 2007).

# Pathogenicity and transmission

Through several different mechanisms, *Clostridium perfringens* can cause diseases in both humans and animals. *Clostridium perfringens* capacity to generate toxins and form spores is intimately linked to both its pathogenesis and transmission.

# Transmission

Foodborne Transmission: Eating food contaminated with *Clostridium perfringens* is the most frequent way for humans to contract the infection (Kawai et al., 2014; Suzuki et al., 2015). Common sources include undercooked or improperly stored food, as well as tainted meat and poultry products (Korajkic et al., 2014). Environmental Transmission: The bacterium is extensively dispersed throughout the environment, particularly in soil and in animals' and humans' digestive systems. Transmission may be aided by contact with contaminated surfaces or soil (Korajkic et al., 2014). Person-to-Person Transmission: Although less frequent, direct transmission between individuals can still happen, particularly in medical environments. In light of nosocomial infections, this is more pertinent.(Sawyer, 2021).

# Pathogenicity

Toxin Production: The primary cause of Clostridium perfringens' pathogenicity is its ability to produce a variety of toxins. Alpha toxin, a phospholipase that ruptures cell membranes and causes cell lysis, is the most significant toxin (Welch & Nutgall, 2023). Food Poisoning: When bacteria in food become contaminated, it can produce toxins in cases of foodborne transmission. The poisons in the contaminated food induce gastroenteritis, which manifests as diarrhea and cramping in the abdomen. The sickness usually goes away on its own in a day or two (Singer et al., 2018). Gas Gangrene: When a wound becomes infected, Clostridium perfringens spores can enter the body through contaminated wounds. Once inside, the bacteria can proliferate and produce the toxins that result in gas gangrene. Gas gangrene is characterized by rapid tissue destruction, gas production, and the potential for systemic toxicity (Unbolts et al., 2023). Clostridium perfringens Necrotic Enteritis: can induce necrotic enteritis in the intestines, particularly in animals (Wade et al., 2020).

Toxins produced by the bacteria cause damage to the intestinal mucosa. Nosocomial Infections: In hospitals, Clostridium perfringens have the ability to make in nosocomial infections, principally in surgical persons and those with compromised immune systems. Knowing the Clostridium perfringens pathogenesis and transmission is a prerequisite for taking precautionary measures in healthcare environments, such as appropriate food handling and hygiene practices, wound care, control measures. Understanding and infection the pathogenicity and mode of transmission of Clostridium perfringens is vital before making precautionary measures in healthcare conditions, such as better food treatment, hygiene practices, caring of wound, and control of infection (Alemayehu et al., 2019).

# Virulence

The ability of *Clostridium perfringens* to produce toxins and severe spores is considered as one of numerous factors leading to the virulence of bacterium's, or ability to be infected with disease. Nonetheless, the main cause for C. perfringens virulence is phenololipase, also known as alpha toxin. Cell lysis is the result of damage to cell membranes produced by a phospholipase known as alpha toxin. It increases the damaging effects of multiple C. perfringens infections, such as gas gangrene (Ode et al., 2012). Beta toxins can make necrotic enteritis in animals. It damages cell membranes and aids in the disintegration of the intestinal mucosa (Ma et al., 2014). Epsilon toxin has been connected to certain types of enterotoxaemia in animals. It could highly affect blood vessels and can cause edema and necrosis in a range of tissues (Wadley et al., 2019). C. perfringens produces an enterotoxin linked to food poisoning. It could lead to gastrointestinal symptoms by acting on the intestines.( Sheathe et al., 2016). Creation of Spores: It produces spores that can endure harsh environmental conditions. The bacteria has the possibility of spreading to novel hosts thanks to the spores' assistance in letting it withstand and flourish in a range of environments (Ammon et al., 2022). Enzymatic Activities: It produces a variety of enzymes, including collagenase and hyaluronidase, which aid in tissue degradation and facilitate the bacteria's spread throughout the host tissues (Comrade et al., 2010). Some strains of C. perfringens can form capsules, which may help them avoid detection by the host immune system (Eunsu et al., 2019) The motile bacteria C. perfringens is able to move around because of its peritrichous flagella, which are flagella dispersed throughout its surface

# Clostridium perfringens and its hosts' interactions

Numerous significant interactions occur between *Clostridium perfringens* and both its host and pathogen. Adhesion and Colonization: Adhering to intestinal epithelial cells is the first way *Clostridium perfringens* interacts with its host. Adhesins that the bacterium produces could help it adhere to the mucosal surface (Luis et al., 2017). Production of Toxins: *Clostridium perfringens* is known to be pathogenic due to its production of several toxins, such as beta toxin, epsilon toxin,

iota toxin, and alpha toxin (phospholipase C). These toxins cause tissue damage and are accountable for the symptoms linked to the various diseases that C. perfringens causes (Ode et al., 2012). Food Poisoning: When enterotoxin-producing strains of *Clostridium perfringens* are consumed through tainted food, the enterotoxin acts in the intestines, causing symptoms like diarrhea and cramping in the abdomen (Rood, 2018). Gas Gangrene: This type of necrotic infection, marked by gas production and the breakdown of surrounding tissues, is caused by C. perfringens infecting wounds or damaged tissues. Alpha toxin and other toxins work together to cause systemic effects and tissue destruction (Unbolts et al., 2023). Immune Response: To phagocytosis and eradicate the bacteria, neutrophils and other immune cells are brought to the infection site by the host, which is an important part of the immune response against C. perfringens infections. Both the amount of tissue damage and the bacterial toxins can affect how well the immune system responds (Xiao et al., 2019). Host Factors: Variations exist in the susceptibility of hosts to C. perfringens infections. The dysfunction of immune system or underlying medical conditions might probably lead to more vulnerable hosts (Luis et al., 2017).

#### Epidemiology of Clostridium perfringens

The research of prevailing bacterium, dispersion, and causal factors in communities is called epidemiology of *Clostridium perfringens* infections. The following are important facets of *C. perfringens* epidemiology: Foodborne illness risk factors, incidence and prevalence, surveillance, and outbreak investigations. In order to lower the burden of *C. perfringens*-associated diseases, enhance food safety procedures, and implement effective preventive measures, it is imperative to understand the epidemiology of *Clostridium perfringens* (Hama & Gonium, 2015).

# Risks to Animal Welfare and Public Health as posed by *Clostridium perfringens*

Foodborne illnesses, toxins, and tissue damage have an impact on public health. Animal Welfare Issues: Environmental Reservoirs, Necrotic Enteritis in Poultry (Li et al., 2011). Public health interventions, surveillance, and outbreak response are part of the One Health Perspective and Interconnected Health.

The pathogenicity of *C. perfringens* is noteworthy, mainly because it produces a range of toxins. The virulence factors play a role in the bacterium's capacity to cause severe tissue damage, especially alpha toxin to (Rood et al., 2018). Comprehending the molecular processes underlying pathogenicity is essential for creating focused interventions as in (Gunosewoyo et al., 2023).. The wide range of poisons that *C. perfringens* produces has a major impact on how diseases present in (Hathaway et al., 2008:). For example, alpha toxin functions as a phospholipase, rupturing cell membranes and causing necrosis in tissue as in (Banal et al., 2019; Ural et al., 2018; Moore et al., 2016; Fernandez-Miyakawa et al., 2007).

The various toxin profiles of the various strains show how complex C. perfringens' pathophysiology is (Berryman et al., 1988; Rood and Abraham, 1985; Abraham and Rood, 1985). Three crucial stages of the life cycle of Candida perfringens are adhesion, colonization, and immune response evasion. conversations with its host of (Canaled et al., 2012; Cubes & Kolacz, 2013; Boettcher & Man, 2014; Borregaard & Nausea, 2014). Understanding these dynamics provides insightful viewpoints on the spread of infections and potential avenues for corrective action in (McLane et al., 2023; McLane et al., 2023; Anderson et al., 2012). C. perfringens foodborne infections are common and often associated with eating tainted meat or poultry, as in the case of (Fernley et al., 2016; Mahon et al., 2013). The aim of epidemiology is to comprehend the factors that facilitate the bacterium's spread, from food processing to environmental reservoirs. When C. perfringens causes necrotic enteritis in veterinary settings, it has an impact on both the animals' health and the poultry industry's bottom line (Kirk et al., 2015; Freedman et al., 2014; Greco et al., 2004). The intricate relationship that exists between the bacterium, gut health, and the chicken industry highlights the need for suitable preventive measures. View of C. perfringens in (Ritter et al., 2017; Buck et al., 2004). Infections pose serious challenges to public health, necessitating proactive surveillance, efficient outbreak response, and preventive measures (Hoekstra et al., 2011). Because the bacteria can cause acute gastroenteritis, food safety procedures are essential to preventing outbreaks of (Hoekstra et al., 2011; Hall et al., 2013; Hoekstra et al., 2011). Questions concerning animal welfare and the ethics of intensive farming practices are raised by animal necrotizing enteritis. (Broom et al., 2012; Sanded et al., 2006). Lessening the impact of C. perfringens animal experiments should be discussed in greater detail when discussing ethical and sustainable farming practices (Join et al., 2019; Field et al., 2020). The one health perspective in studies of the interconnectedness between human. animal. and environmental health (Sharp et al., 2011; Worobey et al., 2014). Collaboration with the environmental sciences is essential to fully comprehend the issues raised by C. perfringens in human and veterinary medicine.

## Conclusion

Gram-positive, anaerobic *Clostridium perfringens* is a complex bacterium with considerable pathogenic potential. It makes both humans and animals sick in a variety of ways. Its ability to produce a variety of toxins and its complex interactions with the host are closely related to its pathogenicity. The factors of virulence related to *C. perfringens*, specifically toxins such as beta, epsilon, and alpha toxins, seem to have a significant effect upon the acuteness and clinical presentation of illnesses associated to this organism. Adhesion, colonization, and immune response evasion seem to be good instances of host-pathogen interplay. In addition to inducing inflammation, damaging the tissue, and systemic effects, *C. perfringens* makes toxins that affect the growth of pathogens. To make up more focused therapeutic

interventions as well as preventive measures, it could be vital to grasp up such interactions. Enterotoxigenic strains of Clostridium perfringens are the cause of gastroenteritis, which is characterized by abdominal cramps and diarrhea, according to epidemiology. C. perfringens is a common cause of foodborne infections. Gas gangrene-related wound infections carry greater risks, especially in surgical or traumatized wounds. Poultry necrotizing enteritis is another serious issue that affects both agricultural productivity and animal welfare. The cons of C. perfringens infections on public health focus on the need for food safety regulations, good wound care, and the creation of vaccines and therapies. Moreover, the One Health method admits the relationship among human and animal health and stress the value of group-work in addressing C. perfringens-related problems. In brief, the pathogenicity of Clostridium perfringens focuses on the significance of meticulous assay, close monitoring, and public health campaigns. Via total comprehension of the virulence, epidemiology, and effects on human and animal health, we might work to make effectual disease prevention, rapid diagnosis, and improved treatment methods for ailments associated to C. perfringens

#### **Conflict of Interest**

The author hereby declares no conflict of interest.

#### **Consent for publication**

The author declares that the work has consent for publication.

#### **Funding support**

The author declares that they have no funding support for this study.

#### References

Abraham, L. J., & Rood, J. I. (1985). Cloning and analysis of the *Clostridium* perfringens tetracycline resistance plasmid, pCW3. *Plasmid*, 13(3), 155– 162.

https://doi.org/10.1016/0147-619x(85)90038-1

- Abraham, L. J., & Rood, J. I. (1985). Molecular analysis of transferable tetracycline resistance plasmids from *Clostridium perfringens*. *Journal of Bacteriology*, 161(2), 636–640. <u>https://doi.org/10.1128/jb.161.2.636-640.1985</u>
- Abraham, L. J., & Rood, J. I. (1987). Identification of Tn4451 and Tn4452, chloramphenicol resistance transposons from *Clostridium perfringens*. *Journal of Bacteriology*, 169(4), 1579–1584. https://doi.org/10.1128/jb.169.4.1579-1584.1987
- Abraham, L. J., & Rood, J. I. (1988). The *Clostridium perfringens* chloramphenicol resistance transposon Tn4451 excises precisely in Escherichia coli. *Plasmid*, 19(2), 164–168. https://doi.org/10.1016/0147-619x(88)90055-8
- Abraham, L. J., Berryman, D. I., & Rood, J. I. (1988). Hybridization analysis of the class P tetracycline resistance determinant from the *Clostridium perfringens* R-plasmid, pCW3. *Plasmid*, 19(2), 113–120. https://doi.org/10.1016/0147-619x(88)90050-9
- Alemayehu, T., Ali, M., Mitiku, E., & Hailemariam, M. (2019). The burden of antimicrobial resistance at tertiary care hospital, southern Ethiopia: a three years' retrospective study. *BMC Infectious Diseases*, 19(1). <u>https://doi.org/10.1186/s12879-019-4210-1</u>

- Alemayehu, T., Ali, M., Mitiku, E., & Hailemariam, M. (2019). The burden of antimicrobial resistance at tertiary care hospital, southern Ethiopia: a three years' retrospective study. *BMC Infectious Diseases*, 19(1). <u>https://doi.org/10.1186/s12879-019-4210-1</u>
- Alonso, D., McKane, A. J., & Pascual, M. (2006). Stochastic amplification in epidemics. *Journal of the Royal Society Interface*, 4(14), 575–582. <u>https://doi.org/10.1098/rsif.2006.0192</u>
- Alsayed, S. S. R., & Gunosewoyo, H. (2023). Tuberculosis: Pathogenesis, Current Treatment Regimens and New Drug Targets. International Journal of Molecular Sciences, 24(6), 5202. https://doi.org/10.3390/ijms24065202
- Aly, S. S., Anderson, R. J., Whitlock, R. H., Fyock, T. L., McAdams, S. C., Byrem, T. M., . . . Gardner, I. A. (2012). Cost-effectiveness of diagnostic strategies to identify Mycobacterium avium subspecies paratuberculosis super-shedder cows in a large dairy herd using antibody enzyme-linked immunosorbent assays, quantitative real-time polymerase chain reaction, and bacterial culture. *Journal of Veterinary Diagnostic Investigation*, 24(5), 821–832. https://doi.org/10.1177/1040638712452107
- Amon, J. D., Artzi, L., & Rudner, D. Z. (2022). Genetic Evidence for Signal Transduction within the Bacillus subtilis GerA Germinant Receptor. *Journal of Bacteriology*, 204(2). <u>https://doi.org/10.1128/jb.00470-21</u>
- Amulic, B., Cazalet, C., Hayes, G. L., Metzler, K. D., & Zychlinsky, A. (2012). Neutrophil Function: From Mechanisms to Disease. *Annual Review of Immunology*, 30(1), 459–489. https://doi.org/10.1146/annurev-immunol-020711-074942
- BAŞ, B. (2022). WHAT TELL US THIS BIONUMBERS IN PLANT DEFENSE PROTEIN PHOSPHORYLATION. Eskişehir Teknik Üniversitesi Bilim Ve Teknoloji Dergisi - C Yaşam Bilimleri Ve Biyoteknoloji, 11(1), 31–38. https://doi.org/10.18036/estubtdc.907029
- Benz, R., & Popoff, M. (2018). Clostridium perfringens Enterotoxin: The Toxin Forms Highly Cation-Selective Channels in Lipid Bilayers. Toxins, 10(9), 341.

https://doi.org/10.3390/toxins10090341

- Camiade, E., Peltier, J., Bourgeois, I., Couture-Tosi, E., Courtin, P., Antunes, A., . . . Pons, J. L. (2010). Characterization of Acp, a Peptidoglycan Hydrolase of *Clostridium perfringens* with N -Acetylglucosaminidase Activity That Is Implicated in Cell Separation and Stress-Induced Autolysis. *Journal of Bacteriology*, 192(9), 2373–2384. https://doi.org/10.1128/jb.01546-09
- Chauhan, A. (2023). Pearls in the Eye. Clinical Research and Clinical Reports, 02(03), 01–04.

https://doi.org/10.31579/2835-8325/017

Cheung, J. K., Adams, V., D'Souza, D., James, M., Day, C. J., Jennings, M. P., ... Rood, J. I. (2020). The EngCP endo α-N-acetylgalactosaminidase is a virulence factor involved in *Clostridium perfringens* gas gangrene infections. *International Journal of Medical Microbiology*, 310(2), 151398.

https://doi.org/10.1016/j.ijmm.2020.151398

- Edwards, R. T. (1905). Bacillus Mycogenes (Bacterium Mucogenum), Nov. Spec. An Organism Belonging To The Bacillus Mucosus Capsulatus Group. *Journal of Infectious Diseases*, 2(3), 431–435. https://doi.org/10.1093/infdis/2.3.431
- Fernandez-Miyakawa, M. E., Marcellino, R., & Uzal, F. A. (2007). Clostridium perfringenstype A Toxin Production in 3 Commonly Used Culture Media. Journal of Veterinary Diagnostic Investigation, 19(2), 184–186.

https://doi.org/10.1177/104063870701900208

- Fjeld, J., Achten, N., Hilligoss, H., Nagy, A., & Srikumar, M. (2020). Principled Artificial Intelligence: Mapping Consensus in Ethical and Rights-Based Approaches to Principles for AI. SSRN Electronic Journal. <u>https://doi.org/10.2139/ssm.3518482</u>
- Food Microbiology: Fundamentals and Frontiers. Fourth Edition. Edited by Michael P. Doyle and Robert L. Buchanan. Washington (DC): ASM Press. \$199.95. xix + 1118 p.; ill.; index. ISBN: 978-1-55581-626-1 (hc); 978-1-55581-846-3 (eb). 2013. (2013). *The Quarterly Review of Biology*, 88(2), 144–144. https://doi.org/10.1086/670570
- Fragoulis, A., Tohidnezhad, M., Kubo, Y., Wruck, C. J., Craveiro, R. B., Bock, A., . . . Suhr, F. (2023). The Contribution of the Nrf2/ARE System to Mechanotransduction in Musculoskeletal and Periodontal Tissues.

International Journal of Molecular Sciences, 24(9), 7722. https://doi.org/10.3390/ijms24097722

- Garcia, J. P., Adams, V., Beingesser, J., Hughes, M. L., Poon, R., Lyras, D., . . Uzal, F. A. (2013). Epsilon Toxin Is Essential for the Virulence of *Clostridium perfringens* Type D Infection in Sheep, Goats, and Mice. *Infection and Immunity*, 81(7), 2405–2414. https://doi.org/10.1128/iai.00238-13
- GHONEIM, N., & HAMZA, D. (2017). Epidemiological studies on Clostridium perfringens food poisoning in retail foods. Revue Scientifique Et Technique De L'OIE, 36(3), 1025–1032. https://doi.org/10.20506/rst.36.3.2734
- Grass, J. E., Gould, L. H., & Mahon, B. E. (2013). Epidemiology of Foodborne Disease Outbreaks Caused by *Clostridium perfringens*, United States, 1998–2010. *Foodborne Pathogens and Disease*, 10(2), 131–136. <u>https://doi.org/10.1089/fpd.2012.1316</u>
- Guidance on the risk assessment of food and feed from genetically modified animals and on animal health and welfare aspects. (2012). *EFSA Journal*, *10*(1), 2501.

https://doi.org/10.2903/j.efsa.2012.2501

- Ha, E., Chun, J., Kim, M., & Ryu, S. (2019). Capsular Polysaccharide Is a Receptor of a *Clostridium perfringens* Bacteriophage CPS1. *Viruses*, 11(11), 1002. https://doi.org/10.3390/v11111002
- Hall, A. J., Wikswo, M. E., Manikonda, K., Roberts, V. A., Yoder, J. S., & Gould, L. H. (2013). Acute Gastroenteritis Surveillance through the National Outbreak Reporting System, United States. *Emerging Infectious Diseases*, 19(8), 1305–1309. https://doi.org/10.3201/eid1908.130482

Harwood, V. J., Staley, C., Badgley, B. D., Borges, K., & Korajkic, A. (2014). Microbial source tracking markers for detection of fecal contamination in environmental waters: relationships between pathogens and human health outcomes. *FEMS Microbiology Reviews*, 38(1), 1–40. https://doi.org/10.1111/1574-6976.12031

Hassan, K. A., Elbourne, L. D., Tetu, S. G., Melville, S. B., Rood, J. I., & Paulsen, I. T. (2015). Genomic analyses of *Clostridium perfringens* isolates from five toxinotypes. *Research in Microbiology*, 166(4), 255– 263.

https://doi.org/10.1016/j.resmic.2014.10.003

Hatheway, C. L. (1990). Toxigenic clostridia. Clinical Microbiology Reviews, 3(1), 66–98.

https://doi.org/10.1128/cmr.3.1.66

- Havelaar, A. H., Kirk, M. D., Torgerson, P. R., Gibb, H. J., Hald, T., Lake, R. J., . . . Devleesschauwer, B. (2015). World Health Organization Global Estimates and Regional Comparisons of the Burden of Foodborne Disease in 2010. *PLOS Medicine*, *12*(12), e1001923. https://doi.org/10.1371/journal.pmed.1001923
- Huang, X. Y., Sun, W. Y., Yan, Z. Q., Shi, H. R., Yang, Q. L., Wang, P. F., . . Gun, S. B. (2019). Novel Insights reveal Anti-microbial Gene Regulation of Piglet Intestine Immune in response to *Clostridium perfringens* Infection. *Scientific Reports*, 9(1). https://doi.org/10.1038/s41598-018-37898-5

Hunter, S. E., Brown, J. E., Oyston, P. C., Sakurai, J., & Titball, R. W. (1993). Molecular genetic analysis of beta-toxin of *Clostridium perfringens* reveals sequence homology with alpha-toxin, gamma-toxin, and leukocidin of Staphylococcus aureus. *Infection and Immunity*, 61(9), 3958–3965.

https://doi.org/10.1128/iai.61.9.3958-3965.1993

- Immerseel, F. V., Buck, J. D., Pasmans, F., Huyghebaert, G., Haesebrouck, F., & Ducatelle, R. (2004). *Clostridium perfringens*in poultry: an emerging threat for animal and public health. *Avian Pathology*, 33(6), 537–549. https://doi.org/10.1080/03079450400013162
- Irikura, D., Monma, C., Suzuki, Y., Nakama, A., Kai, A., Fukui-Miyazaki, A., . . . Kamata, Y. (2015). Identification and Characterization of a New Enterotoxin Produced by *Clostridium perfringens* Isolated from Food Poisoning Outbreaks. *PLOS ONE*, 10(11), e0138183. <u>https://doi.org/10.1371/journal.pone.0138183</u>
- Jobin, A., Ienca, M., & Vayena, E. (2019). The global landscape of AI ethics guidelines. *Nature Machine Intelligence*, 1(9), 389–399. <u>https://doi.org/10.1038/s42256-019-0088-2</u>
- Johansson, A., Greko, C., Engström, B., & Karlsson, M. (2004). Antimicrobial susceptibility of Swedish, Norwegian and Danish isolates of *Clostridium perfringens* from poultry, and distribution of tetracycline

resistance genes. *Veterinary Microbiology*, *99*(3–4), 251–257. https://doi.org/10.1016/j.vetmic.2004.01.009

Kiu, R., & Hall, L. J. (2018). An update on the human and animal enteric pathogen *Clostridium perfringens*. *Emerging Microbes & Infections*, 7(1), 1–15.

https://doi.org/10.1038/s41426-018-0144-8

Kiu, R., & Hall, L. J. (2018). An update on the human and animal enteric pathogen *Clostridium perfringens*. *Emerging Microbes & Infections*, 7(1), 1–15.

https://doi.org/10.1038/s41426-018-0144-8

Kiu, R., & Hall, L. J. (2018). An update on the human and animal enteric pathogen *Clostridium perfringens*. *Emerging Microbes & Infections*, 7(1), 1–15.

https://doi.org/10.1038/s41426-018-0144-8

- Kolaczkowska, E., & Kubes, P. (2013). Neutrophil recruitment and function in health and inflammation. *Nature Reviews Immunology*, 13(3), 159–175. <u>https://doi.org/10.1038/nri3399</u>
- Lassen, J., Gjerris, M., & Sandøe, P. (2006). After Dolly—Ethical limits to the use of biotechnology on farm animals. *Theriogenology*, 65(5), 992– 1004.

https://doi.org/10.1016/j.theriogenology.2005.09.012

Li, C., Lillehoj, H. S., Gadde, U. D., Ritter, D., & Oh, S. (2017). Characterization of *Clostridium perfringens*Strains Isolated from Healthy and Necrotic Enteritis-Afflicted Broiler Chickens. *Avian Diseases*, 61(2), 178–185.

https://doi.org/10.1637/11507-093016-reg.1

- Li, J., Chen, J., Vidal, J. E., & McClane, B. A. (2011). The Agr-Like Quorum-Sensing System Regulates Sporulation and Production of Enterotoxin and Beta2 Toxin by *Clostridium perfringens* Type A Non-Food-Borne Human Gastrointestinal Disease Strain F5603. *Infection and Immunity*, 79(6), 2451–2459. https://doi.org/10.1128/iai.00169-11
- Li, J., Uzal, F., & McClane, B. (2016). *Clostridium perfringens* Sialidases: Potential Contributors to Intestinal Pathogenesis and Therapeutic Targets. *Toxins*, 8(11), 341.

https://doi.org/10.3390/toxins8110341

- Llanco, L. A., Nakano, V., Moraes, C. T. D., Piazza, R. M., & Avila-Campos, M. J. (2017). Adhesion and invasion of *Clostridium perfringens* type A into epithelial cells. *Brazilian Journal of Microbiology*, 48(4), 764–768. <u>https://doi.org/10.1016/j.bjm.2017.06.002</u>
- M'ikanatha, N. M. (2007). Book Review Food Safety: Old Habits, New Perspectives By Phyllis Entis. 400 pp. Washington, DC, ASM Press, 2007. \$49.95. 978-1-55581-417-5 Food Microbiology: Fundamentals and Frontiers Third edition. Edited by Michael P. Doyle and Larry R. Beuchat. 1038 pp., illustrated. Washington, DC, ASM Press, 2007. \$169.95. 978-1-55581-407-6. New England Journal of Medicine, 357(12), 1266–1267. https://doi.org/10.1056/nejmbkrev58331
- Ma, M., Gurjar, A., Theoret, J. R., Garcia, J. P., Beingesser, J., Freedman, J. C., . . . Uzal, F. A. (2014). Synergistic Effects of *Clostridium perfringens* Enterotoxin and Beta Toxin in Rabbit Small Intestinal Loops. *Infection* and Immunity, 82(7), 2958–2970. <u>https://doi.org/10.1128/iai.01848-14</u>
- Manz, M. G., & Boettcher, S. (2014). Emergency granulopoiesis. *Nature Reviews Immunology*, 14(5), 302–314. https://doi.org/10.1038/nri3660
- Mehdizadeh Gohari, I., A. Navarro, M., Li, J., Shrestha, A., Uzal, F., & A. McClane, B. (2021). Pathogenicity and virulence of *Clostridium perfringens*. *Virulence*, 12(1), 723–753. https://doi.org/10.1080/21505594.2021.1886777
- Mehdizadeh Gohari, Li, Navarro, Uzal, & McClane. (2019). Effects of Claudin-1 on the Action of *Clostridium perfringens* Enterotoxin in Caco-2 Cells. *Toxins*, 11(10), 582. <u>https://doi.org/10.3390/toxins11100582</u>

Melville, S., & Craig, L. (2013). Type IV Pili in Gram-Positive Bacteria. Microbiology and Molecular Biology Reviews, 77(3), 323–341. https://doi.org/10.1128/mmbr.00063-12

Miyamoto, K., Li, J., Sayeed, S., Akimoto, S., & McClane, B. A. (2008). Sequencing and Diversity Analyses Reveal Extensive Similarities between Some Epsilon-Toxin-Encoding Plasmids and the pCPF5603*Clostridium perfringens*Enterotoxin Plasmid. *Journal of Bacteriology*, 190(21), 7178– 7188.

https://doi.org/10.1128/jb.00939-08

- Miyamoto, K., Yumine, N., Mimura, K., Nagahama, M., Li, J., McClane, B. A., & Akimoto, S. (2011). Identification of Novel *Clostridium perfringens* Type E Strains That Carry an Iota Toxin Plasmid with a Functional Enterotoxin Gene. *PLoS ONE*, 6(5), e20376. <u>https://doi.org/10.1371/journal.pone.0020376</u>
- Moffatt, C. R., Musto, J., Pingault, N., Miller, M., Stafford, R., Gregory, J., . . . Kirk, M. D. (2016). Salmonella Typhimurium and Outbreaks of Egg-Associated Disease in Australia, 2001 to 2011. *Foodborne Pathogens and Disease*, *13*(7), 379–385. https://doi.org/10.1089/fpd.2015.2110

Morris, W. E., Dunleavy, M. V., Diodati, J., Berra, G., & Fernandez-Miyakawa, M. E. (2012). Effects of *Clostridium perfringens* alpha and epsilon toxins in the bovine gut. *Anaerobe*, 18(1), 143–147. https://doi.org/10.1016/j.anaerobe.2011.12.003

Nagahama, M., Ochi, S., Oda, M., Miyamoto, K., Takehara, M., & Kobayashi, K. (2015). Recent Insights into *Clostridium perfringens* Beta-Toxin. *Toxins*, 7(2), 396–406.

https://doi.org/10.3390/toxins7020396

Nauseef, W. M., & Borregaard, N. (2014). Neutrophils at work. *Nature Immunology*, 15(7), 602–611. https://doi.org/10.1038/ni.2921

Navarro, M., McClane, B., & Uzal, F. (2018). Mechanisms of Action and Cell Death Associated with *Clostridium perfringens* Toxins. *Toxins*, 10(5), 212. https://doi.org/10.3390/toxins10050212

Oda, M., Kabura, M., Takagishi, T., Suzue, A., Tominaga, K., Urano, S., . . . Sakurai, J. (2012). *Clostridium perfringens* Alpha-toxin Recognizes the GM1a-TrkA Complex. *Journal of Biological Chemistry*, 287(39), 33070– 33079.

https://doi.org/10.1074/jbc.m112.393801

- Patel, M. M., Hall, A. J., Vinjé, J., & Parashar, U. D. (2009). Noroviruses: A comprehensive review. *Journal of Clinical Virology*, 44(1), 1–8. <u>https://doi.org/10.1016/j.jcv.2008.10.009</u>
- Petit, L., Gibert, M., & Popoff, M. R. (1999). Clostridium perfringens: toxinotype and genotype. Trends in Microbiology, 7(3), 104–110. <u>https://doi.org/10.1016/s0966-842x(98)01430-9</u>
- Powell, D. A. (1992). Report of the committee on infectious diseases, 21st ed. Committee on Infectious Diseases. Elk Grove Village, IL: American Academy of Pediatrics, 1991, 670 pp. *Pediatric Pulmonology*, *13*(4), 269– 269.

https://doi.org/10.1002/ppul.1950130418

of

Journal

Revitt-Mills, S. A., Rood, J. I., & Adams, V. (2015). Clostridium perfringens extracellular toxins and enzymes: 20 and counting. Microbiology Australia, 36(3), 114. https://doi.org/10.1071/ma15039

Comero-García, N., Huete-Acevedo, J., Mas-Bargues, C., Sanz-Ros, J., Dromant, M., Badenes, R., & Borrás, C. (2023). Extracellular Vesicles: The Future of Diagnosis in Solid Organ Transplantation? *International* 

Molecular

https://doi.org/10.3390/ijms24065102
Rood, J. I., Adams, V., Lacey, J., Lyras, D., McClane, B. A., Melville, S. B., .
Van Immerseel, F. (2018). Expansion of the *Clostridium perfringens* toxin-based typing scheme. *Anaerobe*, 53, 5–10. https://doi.org/10.1016/j.anaerobe.2018.04.011

Sciences.

5102.

24(6).

- Rood, J. I., Adams, V., Lacey, J., Lyras, D., McClane, B. A., Melville, S. B., . . Van Immerseel, F. (2018). Expansion of the *Clostridium perfringens* toxin-based typing scheme. *Anaerobe*, 53, 5–10. <u>https://doi.org/10.1016/j.anaerobe.2018.04.011</u>
- Rood, J. I., Adams, V., Lacey, J., Lyras, D., McClane, B. A., Melville, S. B., . . Van Immerseel, F. (2018). Expansion of the *Clostridium perfringens* toxin-based typing scheme. *Anaerobe*, 53, 5–10. <u>https://doi.org/10.1016/j.anaerobe.2018.04.011</u>
- Rood, J. I., Adams, V., Lacey, J., Lyras, D., McClane, B. A., Melville, S. B., . . Van Immerseel, F. (2018). Expansion of the *Clostridium perfringens* toxin-based typing scheme. *Anaerobe*, 53, 5–10. <u>https://doi.org/10.1016/j.anaerobe.2018.04.011</u>
- Rood, J. I., Keyburn, A. L., & Moore, R. J. (2016). NetB and necrotic enteritis: the hole movable story. *Avian Pathology*, 45(3), 295–301. <u>https://doi.org/10.1080/03079457.2016.1158781</u>
- Sakaue, M., Ota, K., Nakamura, E., Nitta, M., Oka, M., Oishi, Y., . . . Takasu, A. (2019). Type A fulminant *Clostridium perfringens* sepsis indicated RBC/Hb discrepancy; a case report. *BMC Infectious Diseases*, 19(1). https://doi.org/10.1186/s12879-019-4350-3

#### Microbial Science Archives (2024) Vol. 4(2), 6-12

- Sakurai, J., & Duncan, C. L. (1978). Some properties of beta-toxin produced by *Clostridium perfringens* type C. *Infection and Immunity*, 21(2), 678– 680. https://doi.org/10.1128/iai.21.2.678-680.1978
- Scallan, E., Griffin, P. M., Angulo, F. J., Tauxe, R. V., & Hoekstra, R. M. (2011). Foodborne Illness Acquired in the United States—Unspecified Agents. *Emerging Infectious Diseases*, 17(1), 16–22. https://doi.org/10.3201/eid1701.p21101
- Scallan, E., Griffin, P. M., Angulo, F. J., Tauxe, R. V., & Hoekstra, R. M. (2011). Foodborne Illness Acquired in the United States—Unspecified Agents. *Emerging Infectious Diseases*, 17(1), 16–22. https://doi.org/10.3201/eid1701.p21101
- Scallan, E., Hoekstra, R. M., Angulo, F. J., Tauxe, R. V., Widdowson, M. A., Roy, S. L., . . Griffin, P. M. (2011). Foodborne Illness Acquired in the United States—Major Pathogens. *Emerging Infectious Diseases*, 17(1), 7– 15.

https://doi.org/10.3201/eid1701.09-1101p1

- Sharp, P. M., & Hahn, B. H. (2011). Origins of HIV and the AIDS Pandemic. Cold Spring Harbor Perspectives in Medicine, 1(1), a006841–a006841. https://doi.org/10.1101/cshperspect.a006841
- Shrestha, A., Uzal, F. A., & McClane, B. A. (2016). The interaction of *Clostridium perfringens* enterotoxin with receptor claudins. *Anaerobe*, 41, 18–26.

https://doi.org/10.1016/j.anaerobe.2016.04.011

- Uzal, F. A., Freedman, J. C., Shrestha, A., Theoret, J. R., Garcia, J., Awad, M. M., . . . McClane, B. A. (2014). Towards an understanding of the role of *Clostridium perfringens* toxins in human and animal disease. *Future Microbiology*, 9(3), 361–377. <u>https://doi.org/10.2217/fmb.13.168</u>
- Wade, B., Keyburn, A. L., Haring, V., Ford, M., Rood, J. I., & Moore, R. J. (2020). Two putative zinc metalloproteases contribute to the virulence of

#### How to cite this article

Khorsheed, S. A. (2024). Virulence and Pathogenicity of *Clostridium perfringens*. *Microbial Science Archives*, Vol. 4(2), 6-12. https://doi.org/10.47587/MSA.2024.4201

This work is licensed under a <u>Creative Commons Attribution 4.0 International License</u>

Publisher's Note: The Journal stays neutral about jurisdictional claims in published maps and institutional affiliations.

*Clostridium perfringens* strains that cause avian necrotic enteritis. *Journal* of Veterinary Diagnostic Investigation, 32(2), 259–267. https://doi.org/10.1177/1040638719898689

- Wagley, S., Bokori-Brown, M., Morcrette, H., Malaspina, A., D'Arcy, C., Gnanapavan, S., . . Titball, R. W. (2018). Evidence of *Clostridium perfringens* epsilon toxin associated with multiple sclerosis. *Multiple Sclerosis Journal*, 25(5), 653–660. <u>https://doi.org/10.1177/1352458518767327</u>
- Wambui, J., Cernela, N., Stevens, M. J. A., & Stephan, R. (2021). Whole Genome Sequence-Based Identification of Clostridium estertheticum Complex Strains Supports the Need for Taxonomic Reclassification Within the Species Clostridium estertheticum. *Frontiers in Microbiology*, 12.

https://doi.org/10.3389/fmicb.2021.727022

Wang, H., Latorre, J. D., Bansal, M., Abraha, M., Al-Rubaye, B., Tellez-Isaias, G., . . . Sun, X. (2019). Microbial metabolite deoxycholic acid controls *Clostridium perfringens*-induced chicken necrotic enteritis through attenuating inflammatory cyclooxygenase signaling. *Scientific Reports*, 9(1).

https://doi.org/10.1038/s41598-019-51104-0

Worobey, M., Han, G. Z., & Rambaut, A. (2014). A synchronized global sweep of the internal genes of modern avian influenza virus. *Nature*, 508(7495), 254–257.

https://doi.org/10.1038/nature13016

Yonogi, S., Matsuda, S., Kawai, T., Yoda, T., Harada, T., Kumeda, Y., Iida, T. (2014). BEC, a Novel Enterotoxin of *Clostridium perfringens* Found in Human Clinical Isolates from Acute Gastroenteritis Outbreaks. *Infection* and Immunity, 82(6), 2390–2399. https://doi.org/10.1128/iai.01759-14

 $\odot$   $\odot$