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Article

# Infectious Diseases in the Caribbean and South America: Epidemiological Insights, Antibiotic Resistance, and Associated Infectious in Immunological Diseases

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Abstract: This paper explores various aspects of microbiology and immunology, focusing on the epidemiology and molecular characterisation of infectious diseases in the Caribbean and South America. Key areas of investigation include tuberculosis (TB), nosocomial infections, *Salmonella* infections, and bloodborne pathogens. A retrospective study in Jamaica underscores the high mortality rates in HIV-positive TB patients and the need for early HIV diagnosis and integrated care. Research in Trinidad and Tobago reveals the genetic diversity of *Mycobacterium tuberculosis* strains, highlighting high HIV-TB co-infection rates. The paper also examines the challenges posed by nosocomial infections, particularly those caused by antibiotic-resistant *Gram-negative* bacteria and methicillin-resistant *Staphylococcus aureus* (MRSA), and emphasises the importance of infection control. In addition, the study investigates *Salmonella* prevalence in poultry in Jamaica and the risks associated with foodborne infections. The paper further explores transfusion-transmitted infections and the development of IgM antibodies in egg whites for bacterial antigen detection. Lastly, it addresses immune responses in patients with immunodeficiency disorders, such as severe combined immunodeficiency (SCID) and chronic granulomatous disease (CGD), highlighting their susceptibility to a wide range of infections.

**Keywords:** Tuberculosis; HIV; Nosocomial infections; Antibiotic resistance; Salmonella; Blood transfusion safety; Immunology; West Indies; Mycobacterium tuberculosis; Gram-negative bacteria; IgM antibodies

#### 1. Introduction

Nosocomial infections, which are a major concern in healthcare settings, are also examined in this study. These infections, caused by drug-resistant pathogens like Gram-negative bacteria and methicillin-resistant *Staphylococcus aureus* (MRSA), are prevalent in Trinidad and Tobago. They lead to increased morbidity, mortality, and healthcare costs, underlining the importance of effective infection control measures and the prudent use of antibiotics [1,2].

Another critical issue addressed in this paper is foodborne illnesses, particularly Salmonella infections linked to poultry. The study evaluates the prevalence of Salmonella in Jamaica's poultry

industry, examining its implications for food safety and public health. Effective control of Salmonella requires stringent biosecurity measures and public education.

In addition to bacterial infections, the safety of blood transfusions is of paramount concern.

The paper explores novel approaches in antibody production and vaccine development, particularly using IgM antibodies in egg whites, highlighting their potential for immunological research and applications in the West Indies. *Peptide candidate vaccines to chicks induces specific anti-HIV gp120 and gp41 antibodies* that neutralize the original HIV antigens [3].

This multidisciplinary study provides insights into tackling infectious diseases through microbiological and immunological advancements [2]. Infectious diseases continue to pose significant public health challenges worldwide, particularly in resource-limited regions like the Caribbean. Microbiology and immunology play a crucial role in understanding the mechanisms of disease transmission, pathogenesis, and host immune responses. In the Caribbean, the dual burden of tuberculosis (TB) and HIV is especially concerning, with TB remaining one of the leading causes of morbidity and mortality in HIV-infected individuals. This paper explores the epidemiology of TB and HIV co-infection in Jamaica [4], among other themes including frequent infectious microorganisms associated to primary immunodeficiency disorders, autoimmune diseases and inflammatory immune disorders, highlighting the urgent need for early diagnosis and integrated care

# 2. Nosocomial Infections Globally

Nosocomial infections, particularly in intensive care, surgical, and medical wards, are a recurrent issue globally, with rates varying by region. In Trinidad and Tobago, limited data exist, but between 1992-1995, 7,158 infections were recorded, primarily urinary tract infections, postoperative wound infections, pneumonia, and bloodstream infections. Gram-negative bacilli, especially Pseudomonas aeruginosa, and methicillin-resistant Staphylococcus aureus were prevalent. Bloodstream infections significantly increased mortality, length of hospital stays, and costs. Skin and soft tissue infections (SSTIs) are also common, with management depending on severity, involving antibiotics like penicillin, cephalosporins, and aminoglycosides. Antibiotic resistance remains a concern [5–9].

The study by Amaral et al. (2005) examines the characteristics of a predominant variant within the Brazilian epidemic clonal complex of methicillin-resistant Staphylococcus aureus (MRSA). This research focuses on the enhanced abilities of this MRSA variant to produce biofilm and to adhere to and invade airway epithelial cells. The researchers characterised the variant by evaluating its biofilm production, which is a key factor in its pathogenicity. They found that this Brazilian MRSA variant exhibits a significantly increased capacity to form biofilms compared to other strains. Biofilms are dense clusters of bacteria that adhere to surfaces and are protected by a matrix, making infections harder to treat. Additionally, the study assessed the variant's ability to adhere to and invade airway epithelial cells, which are crucial in respiratory infections. The findings demonstrated that this MRSA variant has an enhanced capacity to attach to and penetrate these cells, potentially leading to more severe respiratory infections. The results highlight the variant's aggressive nature and its potential impact on patient outcomes, particularly in respiratory infections. The study underscores the need for targeted strategies to address infections caused by this MRSA variant, including novel therapeutic approaches and improved infection control measures [10].

Gram-negative bacterial infections pose a global health challenge, with beta-lactamase production being a key factor in beta-lactam resistance. Extended spectrum beta-lactamases (ESBLs), particularly in *Escherichia coli* and *Klebsiella pneumoniae*, are widespread, with over 200 ESBLs characterised worldwide [12,13]. ESBLs, such as TEM, SHV, and CTX-M, cause resistance to beta-lactam antibiotics, including cephalosporins and monobactams, but not cephamycins or carbapenems [14]. ESBL-producing organisms contribute to nosocomial infections, especially urinary tract infections, respiratory infections, and bloodstream infections [15,16]. Risk factors for ESBL-associated UTIs include antibiotic use, previous hospitalisations, and underlying conditions.

Preventive measures, such as hand hygiene and environmental cleaning, are crucial to controlling the spread of these resistant organisms [15].

#### 3. Salmonellosis

The Salmonella pandemic, first noted in the late 1980s, has been linked to contaminated eggs, with a rising proportion of *S.* serovar *Enteritidis* infections globally [18–20]. Control of *Salmonella* is challenging due to multiple contamination sources in poultry operations, such as chicks, feed, rodents, and farm environments [21]. Reducing *Salmonella* in poultry and livestock could significantly reduce human exposure [22], as seen in Denmark, where treated animal feed decreased Salmonella infections [23]. Early detection and understanding of cross-contamination points are critical to controlling salmonellosis outbreaks. Limited data exist on the prevalence of *Salmonella* in Jamaica's poultry industry, prompting this study, which involved 6,693 specimens collected from poultry farms, processing plants, and supermarkets across six parishes. The study revealed a low prevalence of *Salmonella* (1%), with environmental specimens, such as pig faeces, showing higher contamination rates [24,25]. Notably, *Salmonella Montevideo* and *Salmonella Kentucky* were isolated from abattoir specimens, marking their first identification in Jamaica [26,27].

Fly specimens also tested positive for *Salmonella*, raising concerns about their potential as vectors in food contamination. Despite these findings, no *Salmonella* was detected in poultry products from marketing outlets, offering some reassurance to consumers. Improved hygiene and surveillance in poultry farms, coupled with public education, are essential to controlling foodborne salmonellosis in Jamaica [28]. This study provides valuable insights into *Salmonella* prevalence and highlights the need for continued preventive measures in the poultry industry as a specific DNA into a cationic liposome to protect against salmonellosis in the Poultry industry, which show to be an effective as an attenuated vaccine [29].

Salmonella is a common cause of foodborne illness, particularly linked to poultry. This study investigated the effectiveness of three antibiotics—Trisulvitrim, Menorox, and Neochlore—in preventing Salmonella Typhimurium infection in growing chickens. Four groups of 60 chicks were studied, with three groups receiving one of the antibiotics and the fourth serving as a control. Results showed that chickens treated with antibiotics had a significantly lower prevalence of Salmonella infection (1.6%) compared to the control group (23.3%), demonstrating the effectiveness of the medications. Salmonella was predominantly found in the caeca and crops of infected birds. The study emphasises the need for effective antibiotic use in reducing Salmonella in poultry, although it acknowledges that medication may not provide 100% protection. It also highlights the increased risk of infection in organic farming practices [30]. Antibiotic resistance remains a concern, particularly due to the widespread use of antibiotics in both human and animal treatment [31].

#### 4. The Risk of Transfusion-Transmitted Infections

Blood transfusions are critical in medical treatments, saving lives by replenishing blood lost during surgery, injury, or illness. However, transfusion carries risks, including the transmission of bacterial, viral, and parasitic pathogens. Among these, the risk of transfusion-transmitted bacterial infections (TTBI) is notably higher than that of viral infections, particularly in platelet transfusions, which present the highest risk of bacterial contamination. This makes blood safety a major concern in healthcare systems worldwide [32–37].

Bacterial proteins such as staphylococcal protein A, peptostreptococcal protein L, and streptococcal protein G have been utilized in laboratory diagnostic studies to substitute diagnostic antibodies in tests such as ELISA, western blot, immunofluorescence, flow cytometry, and molecular biology assays. These proteins bind to immunoglobulins, enabling detection and identification of pathogens without the need for disease-specific antibodies. This application is especially useful in diagnosing infections, improving the accuracy and speed of results, and enhancing the overall effectiveness of diagnostic tools in transfusion safety [38].

Viral organisms transmitted through blood transfusions include HIV, hepatitis C (HCV), hepatitis B (HBV), hepatitis A (HAV), West Nile virus (WNV), cytomegalovirus (CMV), severe acute

respiratory syndrome coronavirus (SARS-CoV-1), human T-cell lymphotropic viruses (HTLV), Zika virus, and parvovirus B19. These viruses pose significant risks, particularly for immunocompromised patients, and may cause long-term health complications if transmitted through contaminated blood products [39–45].

Screening and pathogen reduction technologies have significantly reduced the incidence of viral infections transmitted through transfusions. For example, nucleic acid amplification tests (NAT) are now standard in blood screening processes to detect viral RNA or DNA in blood samples. However, despite advances in viral detection, bacterial infections continue to be a significant concern. There is association between some cancers and ABO groups. Bacteria can thrive in stored blood products, particularly platelets, which are stored at room temperature. This creates a breeding ground for bacterial growth, making the timely detection and prevention of TTBI essential for ensuring the safety of blood transfusions [38,46,47]

# 5. Egg Antibody Technology

The study focuses on investigating IgM antibodies in egg whites, a topic with limited prior research. Antibodies targeting bacterial antigens were developed, utilizing ELISA for detection and affinity chromatography for purification. By day 9 post-immunization, large amounts of anti-protein A antibodies were detected in egg whites, confirmed through affinity chromatography. Samples containing anti-SpA antibodies demonstrated agglutination inhibition, whereas negative samples exhibited agglutination. This indicates the successful production of IgM anti-protein A antibodies in egg whites, along with the inhibition of bacterial growth observed in vitro. Protein A-affinity chromatography played a crucial role in characterizing these antibodies, underscoring its value in antibody production and research [48–51].

The research also explores the complexities of experimental vaccine development, specifically focusing on immunogenicity outcomes. A key aspect highlighted is the innovative concept of idiotypic-antiidiotypic interactions, which can modulate antibody responses against bacterial and viral proteins. Three experimental vaccines targeting HIV, Salmonella, and Staphylococcus aureus were examined methodologically [52]. The author proposes that immunized chicken eggs could provide a valuable platform for producing IgY antibodies against these pathogens. This research holds significant relevance for epidemiology, particularly in regions such as the West Indies, where infectious diseases are a major public health concern. By outlining vaccine development methods and potential applications, the study offers the groundwork for future clinical trials aimed at combating important infectious microorganisms affecting human populations. Additionally, other researchers have reported the production of IgY antibodies with similar results [53–57].

# 6. Confronting Tuberculosis

This retrospective study examined 537 hospital records of patients with pulmonary tuberculosis at Jamaica's National Chest Hospital between 1995 and 2001. Of the 406 eligible patients, 11.6% were HIV-1 positive, primarily males aged 30-39. The mortality rate in HIV-positive patients (23.4%) was significantly higher than in HIV-negative patients (3.9%). All patients received standard quadruple drug therapy, with no cases of multiple drug resistance. Early HIV diagnosis and antiretroviral therapy are critical in reducing mortality [58].

This study aimed to determine the seroprevalence of HIV-1 infection among patients with pulmonary tuberculosis (TB) in Jamaica and assess the epidemiological links, characteristics, and mortality associated with coinfection. HIV infection is the strongest risk factor for TB, with individuals over 100 times more likely to develop TB than non-infected persons. Pulmonary TB remains a leading cause of HIV-related morbidity and mortality globally. By 2003, an estimated 22,000 people were living with HIV/AIDS in Jamaica. The study's findings are essential for developing strategies and policies to control the spread of HIV and TB in resource-limited settings [58].

This study examined 132 culture-positive Mycobacterium tuberculosis cases in Trinidad and Tobago over a year. The findings revealed a high male-to-female ratio of 4:1 and a significant 30%

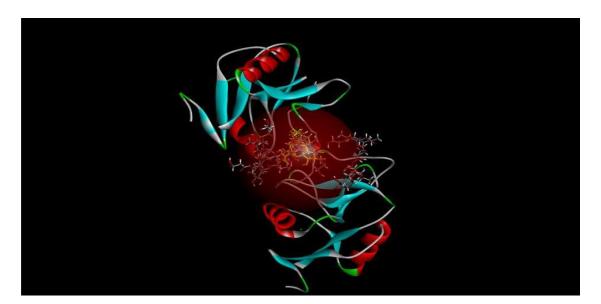
HIV-TB co-infection rate. African descendants, who make up 37.5% of the population, accounted for 69.7% of TB cases (P < 0.001). Spoligotyping identified 25 patterns and 12 clusters, with the SIT566 clone being the most prevalent [59].

A year-long investigation in Trinidad and Tobago characterized Mycobacterium tuberculosis strains using molecular and phylogenetic methods. The SIT566 clone, an "evolutionary modern" strain, represented over half of the cases, predominantly affecting younger populations, particularly in Port-of-Spain. Unique strain distributions, such as LAM-10CAM and EAI, set Trinidad and Tobago apart from other Caribbean countries. SIT566, also identified in Trinidadian patients in the United States, exhibited a distinct MIRU profile. Phylogenetic data suggest SIT566 evolved from X lineage strains, potentially linked to Anglo-Saxon ancestry. This study provides insights into the genetic diversity and origins of TB strains in the region [60].

Another study compared the QuantiFERON®-TB Gold (QFT-G) assay with the tuberculin skin test (TST) for latent tuberculosis infection (LTBI) screening among high-risk individuals in Trinidad & Tobago. Of 560 participants (including TB contacts, HIV patients, healthcare workers, inmates, and TB patients), QFT-G detected LTBI in 51% compared to TST's 39.4% (P = 0.001), except in the control group, where TST was preferred. Although QFT-G provided faster results, it was more expensive (\$18.60 vs. \$3.70 per subject). Importantly, QFT-G demonstrated lower sensitivity in immunocompromised individuals, raising concerns about its use in resource-limited settings with high HIV prevalence, such as Trinidad & Tobago [61,62].

In another investigation, spoligotyping and 15-loci MIRU-VNTR were used to characterize Mycobacterium tuberculosis strains in 74 Guyanese and 80 Surinamese patients. The average age in both countries was around 38 years, and male-to-female ratios were 2.25 in Guyana and 4.27 in Suriname. The techniques revealed 41 and 65 distinct patterns, with clustering rates of 83.8% and 68.8%, respectively. Combined analyses identified 18 clusters (2-41 isolates), with an overall clustering rate of 67.5% and a recent transmission rate of 55.8%. Clustering was significantly higher in Guyana compared to Suriname (79.7% vs. 56.3%; p=0.0019), and shared clusters between the two countries suggest potential transmission routes, highlighting the SIT53/15-MIT861 clone as a key target for intervention [63].

Mincle, a major C-type lectin receptor in macrophages, plays a vital role in the immune response against mycobacteria. Computational tools like Discovery Studio and Molegro Virtual Studios predict ligand-receptor binding positions, while ChemDraw assists in ligand preparation due to the time-consuming nature of protein preparation. Mincle's involvement in mycobacterial-induced inflammatory signaling, particularly in tuberculosis, is of growing interest. However, Molegro's inability to depict metallic atom interactions is a noted limitation. Mincle detects mycobacterial components like trehalose dimycolate (TDM) and trehalose dibenzenate (TDB), activating CARD9 signaling and prompting the release of pro-inflammatory cytokines and chemokines. This research shows potential for therapeutic applications, particularly in tuberculosis research in the West Indies [64]. Figure 1 presents a diagram of the five most feasible poses for TDB, a derivative of TDM, binding to mincle [64].



**Figure 1.** The diagram above shows the five most feasible poses in which TDB, the derivative analog of TDM, binds to mincle because of the low force field energy shown above in a reddish hemisphere, which is the area in which most hydrogen bond interactions occur. Taken from [64].

## 7. Infections in the Course of Immunological Disorders

## 7.1. Severe Combined Immunodeficiency Disorders (SCID)

Patients with severe combined immunodeficiency disorders (SCID) have profound defects in both their T and B cells, leaving them highly vulnerable to a broad spectrum of infections. These individuals are susceptible to bacterial, viral, fungal, and parasitic infections caused by both common and rare microorganisms. Among bacteria, patients often experience infections from *Staphylococcus aureus*, *Pseudomonas aeruginosa*, and *Streptococcus pneumoniae*. However, they are also vulnerable to rare organisms such as *Nocardia* and *Mycobacterium avium complex*. Viral infections are frequent, with cytomegalovirus (CMV), Epstein-Barr virus (EBV), and adenovirus being common culprits. Unusual viral pathogens, such as the measles virus and certain strains of enterovirus, can cause severe complications in these patients. Fungal infections are a significant threat, particularly from *Candida* species and *Aspergillus*. Less common fungal pathogens, such as *Cryptococcus neoformans* and *Pneumocystis jirovecii*, may also lead to life-threatening infections. In terms of parasitic infections, *Toxoplasma gondii* and *Cryptosporidium* are frequently seen. Additionally, these patients are susceptible to rare parasites like *Strongyloides stercoralis*. In SCID patients, even microorganisms that are relatively harmless to healthy individuals can cause severe, recurrent, or disseminated infections, making prompt diagnosis and treatment critical [66].

#### 7.2 Chronic Granulomatous Disease (CGD)

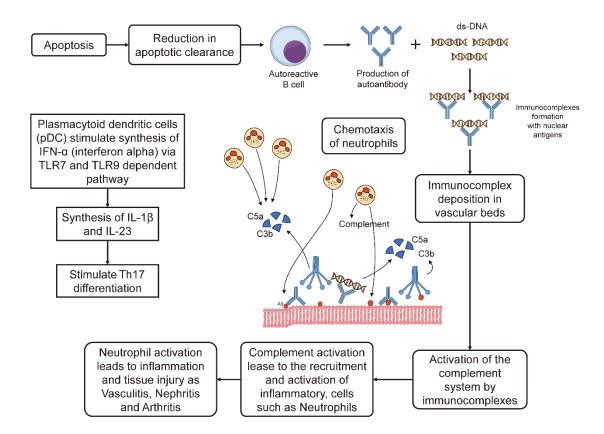
Chronic granulomatous disease (CGD) is a primary immunodeficiency disorder characterised by the inability of phagocytes, particularly neutrophils, to effectively kill certain bacteria and fungi. This defect results from mutations affecting the production of reactive oxygen species, essential for pathogen destruction. As a consequence, individuals with CGD are highly susceptible to recurrent and severe infections caused by specific pathogens. Common bacterial infections in CGD include those caused by *Staphylococcus aureus*, *Serratia marcescens*, and *Nocardia*. Fungal infections, particularly from *Aspergillus* and *Candida* species, are also frequent and often life-threatening. In addition, *Burkholderia cepacia* and *Salmonella* are known pathogens in these patients. Chronic infections and the formation of granulomas in organs such as the lungs, liver, and lymph nodes are typical features of CGD, requiring lifelong antimicrobial prophylaxis and, in some cases, stem cell transplantation for treatment [67].

# 7.3 Transient Hypogammaglobulinemia of Infancy (THI)

In transient hypogammaglobulinemia of infancy (THI), infants are prone to recurrent infections due to their low levels of immunoglobulin G (IgG), which plays a crucial role in the immune response. The most common infections in THI are upper and lower respiratory tract infections, including sinusitis, otitis media (ear infections), and bronchitis. These infections are often caused by bacteria such as *Haemophilus influenzae*, *Streptococcus pneumoniae*, and *Moraxella catarrhalis*. Children with THI may also experience gastrointestinal infections, although less frequently. Viral infections like respiratory syncytial virus (RSV) and adenovirus can be more severe in infants with THI, further complicating their clinical course. However, severe or life-threatening infections are uncommon in THI compared to other primary immunodeficiencies. Management of infections in THI usually involves the use of antibiotics to treat active infections or, in some cases, prophylactic antibiotics to prevent recurrent infections. Immunoglobulin replacement therapy is rarely needed, as most infants outgrow the condition by age 2 to 4 [68].

#### 7.4 Neuropsychiatric Systemic Lupus Erythematosus (NPSLE)

Neuropsychiatric systemic lupus erythematosus (NPSLE) refers to the involvement of the central and peripheral nervous systems in patients with systemic lupus erythematosus (SLE), leading to a range of neurological and psychiatric manifestations. These may include cognitive dysfunction, mood disorders, seizures, psychosis, and even stroke. Patients with NPSLE are particularly vulnerable to infections due to a combination of immunosuppressive therapy, disease-related immune dysfunction, and the involvement of the central nervous system. Infections in these patients can complicate the clinical picture, mimicking or exacerbating neuropsychiatric symptoms, which makes diagnosis challenging. Common pathogens include Staphylococcus aureus, Escherichia coli, and Pneumocystis jirovecii, as well as viral infections like herpes simplex virus (HSV) and cytomegalovirus (CMV). The dual burden of neuropsychiatric manifestations and infection increases morbidity in NPSLE patients, highlighting the need for careful monitoring and management to differentiate between lupus flares and infection-related complications [69,70]. Systemic Lupus Erythematosus (SLE) is an autoimmune disease characterized by the formation of immune complexes that trigger inflammation. It affects multiple organs, leading to conditions such as nephritis, arthritis, and vasculitis. Inflammatory cytokines and complement activation play key roles in tissue damage and disease progression [71–73]. Figure 2 shows the mechanism of SLE pathogenesis. Taken from [69].



**Figure 2.** In Systemic Lupus Erythematosus (SLE), a type III hypersensitivity reaction plays a crucial role. This process involves the formation of immune complexes, which subsequently activate the complement system. Complement components such as C3a and C5a act as potent chemotactic factors, attracting neutrophils to the sites where immune complexes are deposited. The activation of these neutrophils leads to local inflammation and tissue damage, contributing to conditions like vasculitis, nephritis, and arthritis. In addition to complement activation, cytokines such as TNF- $\alpha$ , IL-6, and IFN- $\gamma$  are also involved, further amplifying the inflammatory response and perpetuating tissue injury. Other mechanisms may also contribute to the pathology. Taken from [69].

## 7.5 Interleukin-2 Receptor Alpha (IL-2Rα) Deficiency

CD25 immunodeficiency, also known as interleukin-2 receptor alpha (IL- $2R\alpha$ ) deficiency, is a rare form of severe combined immunodeficiency (SCID). This condition results from mutations in the *IL2RA* gene, which impairs the function of regulatory T cells (Tregs) that play a critical role in controlling immune responses. Without functional CD25, Tregs cannot effectively modulate immune activity, leading to dysregulated immune responses, autoimmunity, and an increased susceptibility to infections. Individuals with CD25 immunodeficiency are prone to a wide range of infections caused by bacterial, viral, fungal, and parasitic pathogens. Common bacterial infections include those caused by *Staphylococcus aureus* and *Streptococcus pneumoniae*. Viral infections, such as those from cytomegalovirus (CMV), Epstein-Barr virus (EBV), and enteroviruses, are frequent and can lead to severe, life-threatening complications. Fungal infections, particularly by *Candida* and *Aspergillus* species, are also prevalent. Due to the severe immune dysregulation, infections in CD25 immunodeficiency are often persistent and recurrent, requiring aggressive antimicrobial treatment. Hematopoietic stem cell transplantation is the only curative option for restoring immune function in these patients, as it replaces the defective immune system [74].

## 7.6. Multiple Sclerosis (MS)

Multiple sclerosis (MS) is a chronic autoimmune disease in which the immune system attacks the myelin sheath surrounding nerve fibres in the central nervous system. This leads to a range of neurological symptoms, including motor and sensory impairments, vision problems, and cognitive dysfunction. While MS itself does not directly cause infections, the disease and its treatments can increase a person's susceptibility to infections. Patients with MS are often treated with immunomodulatory or immunosuppressive therapies, such as corticosteroids, disease-modifying drugs (DMDs), or biologics like natalizumab and ocrelizumab. These treatments can weaken the immune system, increasing the risk of infections. Respiratory tract infections, such as those caused by Streptococcus pneumoniae and Haemophilus influenzae, are common in MS patients. Urinary tract infections (UTIs) are also prevalent, particularly in those with bladder dysfunction, a common MS complication. Viral infections, such as those caused by herpes simplex virus (HSV) and varicellazoster virus (VZV), can be more frequent or severe in MS patients, particularly when on immunosuppressive therapies. Furthermore, certain infections, like Epstein-Barr virus (EBV), have been associated with an increased risk of developing MS, although this connection is still being studied. Careful infection monitoring and management are crucial in MS patients, especially during immunosuppressive treatment [75].

## 7.7. Ataxia-Telangiectasia (A-T)

Ataxia-telangiectasia (A-T) is a rare, inherited disorder that affects the nervous and immune systems, leading to progressive neurological problems, weakened immune function, and increased susceptibility to infections. A-T is caused by mutations in the *ATM* gene, which is responsible for DNA repair, immune regulation, and cellular responses to stress. Individuals with A-T typically present with ataxia (loss of coordination), telangiectasia (small dilated blood vessels), and immunodeficiency. The immunodeficiency in A-T mainly affects the production and function of immunoglobulins (particularly IgA and IgG), leading to recurrent respiratory infections. These infections are primarily caused by bacteria such as *Haemophilus influenzae*, *Streptococcus pneumoniae*, and *Staphylococcus aureus*, which result in frequent sinusitis, bronchitis, and pneumonia. Viral infections, particularly from respiratory syncytial virus (RSV) and adenovirus, can also occur more frequently and with greater severity. Due to the weakened immune system, infections in A-T patients tend to be persistent and can lead to long-term complications, such as chronic lung disease. Management of infections in A-T typically includes regular monitoring, antibiotic treatment, and sometimes immunoglobulin replacement therapy to prevent severe or recurrent infections [76,77].

## 7.8. Interleukin-12 Receptor Deficiency

Interleukin-12 receptor deficiency is a primary immunodeficiency that impairs the immune system's ability to effectively combat intracellular infections. This deficiency disrupts the signalling pathway of interleukin-12, a cytokine critical for activating natural killer (NK) cells and promoting the differentiation of T cells into Th1 cells. Without proper IL-12 receptor function, individuals are more susceptible to infections caused by intracellular pathogens, including mycobacteria, Salmonella, and certain viruses. Patients often experience recurrent or severe infections, as their immune system cannot mount an adequate cell-mediated immune response. Treatment typically involves antimicrobial therapy and, in some cases, cytokine replacement [78].

# 7.9. Allergic Bronchial Asthma

Allergic bronchial asthma is a chronic respiratory condition characterised by airway inflammation and narrowing, often triggered by allergens and microorganisms. Allergic asthma, the most common form, occurs when the immune system overreacts to allergens such as pollen, dust mites, and mould, leading to airway inflammation. Microorganisms, particularly respiratory viruses like rhinovirus and influenza, can worsen asthma by increasing inflammation and disrupting the balance of the airway microbiome. Some bacteria, such as *Haemophilus influenzae*, have also been linked to asthma exacerbations. Additionally, the hygiene hypothesis suggests that reduced exposure

to microorganisms early in life may contribute to the development of asthma by impairing immune system regulation [79,80].

# 7.10. Leukocyte Adhesion Deficiency (LAD)

Leukocyte Adhesion Deficiency (LAD) is a rare primary immunodeficiency characterised by defective leukocyte migration and impaired immune responses due to genetic mutations affecting cell adhesion molecules. These molecules are essential for leukocytes to adhere to and migrate across the endothelium toward sites of infection and inflammation. Persistent infections and impaired healing are hallmark features. Patients often experience severe bacterial infections, particularly in soft tissues, skin, respiratory tract, and mucosa. The associated microorganisms are bacteria including *Staphylococcus aureus* (skin and soft tissue infections), *E. coli* (urinary tract and bloodstream infections), *Pseudomonas aeruginosa* (opportunistic infections), *Klebsiella pneumoniae* respiratory and bloodstream infections), fungal pathogens such as *Candida albicans* (oral and systemic candidiasis), and *Aspergillus* species (less common but severe infections) [81–84].

# 7.11 Chediak-Higashi syndrome (CHS)

Chediak-Higashi Syndrome (CHS) is a rare autosomal recessive disorder caused by mutations in the LYST (lysosomal trafficking regulator) gene. This mutation impairs intracellular vesicle trafficking, resulting in defective lysosomal fusion and impaired immune responses. CHS primarily affects neutrophils, natural killer (NK) cells, cytotoxic T lymphocytes, and melanocytes. Impaired phagosome-lysosome fusion in neutrophils and macrophages leads to ineffective killing of engulfed pathogens. Giant granules accumulate in granulocytes, monocytes, and lymphocytes, reducing cell efficiency. NK cells exhibit impaired cytotoxic activity due to defective exocytosis of cytolytic granules, limiting the immune response against virus-infected cells and tumour cells. Cytotoxic T lymphocytes (CTLs) fail to release lytic granules, diminishing the immune response against intracellular pathogens. Melanin-containing cells are affected, causing partial albinism, often seen in CHS patients. Defective dense granules in platelets lead to prolonged bleeding times and increased risk of haemorrhage. Associated pathogens include bacteria such as Staphylococcus aureus causes recurrent skin abscesses, furuncles, and cellulitis, Streptococcus pneumoniae causes pneumonia, otitis media, and sepsis. Salmonella species are associated with gastrointestinal infections and Escherichia coli causes urinary tract infections and sepsis. Fungal infections caused by Candida albicans include oral thrush and systemic candidiasis due to compromised phagocyte function. Epstein-Barr Virus (EBV): Strongly linked to the development of the life-threatening accelerated phase of CHS, resembling haemophagocytic lymphohistiocytosis (HLH). Hematopoietic stem cell transplantation (HSCT) is the only curative option [85–90].

# 8. Conclusions

This comprehensive study delves into the critical challenges of infectious diseases in the Caribbean and South America, particularly tuberculosis and nosocomial infections, and their intersections with immunodeficiency and antibiotic resistance. The high HIV-TB co-infection rates in Jamaica and Trinidad and Tobago underscore the need for early detection and integrated treatment strategies. Furthermore, the spread of antibiotic-resistant pathogens, such as Gram-negative bacteria and MRSA, poses a significant threat in healthcare settings, necessitating stringent infection control measures and prudent antibiotic use. The findings related to *Salmonella* infections in poultry in Jamaica highlight the importance of biosecurity measures and public education to reduce foodborne disease risks.

The paper also brings attention to the potential of IgM antibodies in egg whites for immunological research, offering innovative approaches to vaccine development and bacterial antigen detection. Additionally, the study emphasises the vulnerability of immunodeficient patients, such as those with SCID, CGD, and other immunological disorders, to various infections. In conclusion, a multidisciplinary approach involving molecular characterisation, immunological

research, and effective public health strategies is essential for controlling infectious diseases in the region, particularly in resource-limited settings. Primary immunodeficiency disorders (PIDs) and autoimmune conditions in South America and the Caribbean exhibit diverse immunological characteristics due to genetic, environmental, and infectious factors. These immunological disorders pose significant health challenges in South America and the Caribbean, highlighting the need for tailored diagnostic and therapeutic approaches.

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