

# Transfusion Transmission of Pathogens from Asymptomatic Donors: A Review

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## ABSTRACT

**Background:** Ensuring the safety of blood transfusion has improved significantly over the years due to better donor screening and advanced testing methods. Techniques such as serological testing and nucleic acid amplification testing (NAT) have greatly reduced the risk of transmitting infections like HIV, hepatitis B, and hepatitis C. However, the risk has not been completely eliminated. One of the major concerns is the presence of infections during the "window period," when the pathogen is present in the blood but not yet detectable. Additionally, some donors may carry infections without showing any symptoms, making detection even more challenging. The emergence of new and re-emerging infectious diseases further complicates blood safety, especially in developing regions.

**Methods:** This review is based on a systematic analysis of research articles published over the last 20 years, sourced from databases such as PubMed, Scopus, and Web of Science. Relevant studies focusing on transfusion-transmitted infections, asymptomatic carriers, and advancements in screening technologies were selected. Information was carefully analyzed to understand patterns of infection, diagnostic challenges, and the effectiveness of current preventive strategies.

**Results:** The findings show that a wide range of pathogens can be transmitted through blood from donors who do not show symptoms. These include well-known viruses like HIV, HBV, and HCV, as well as emerging infections such as dengue and Zika virus. Although modern testing methods have improved early detection, they still have limitations, especially during the early stages of infection. Moreover, many emerging pathogens are not routinely screened in blood banks. Factors like globalization, climate change, and increased travel contribute to the spread of new infections. New approaches such as pathogen reduction technologies and advanced genetic testing methods offer promising solutions but are not yet widely used.

**Conclusion:** The risk of infection from asymptomatic donors remains a critical challenge in transfusion medicine. Continuous improvement in screening technologies, better surveillance of emerging diseases, and global cooperation are essential to ensure blood safety. Adopting innovative technologies and flexible strategies will play a key role in reducing future risks.

**Keywords:** Transfusion-transmitted infections, asymptomatic donors, blood safety, emerging infections, nucleic acid testing, pathogen reduction, window period.

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## Introduction

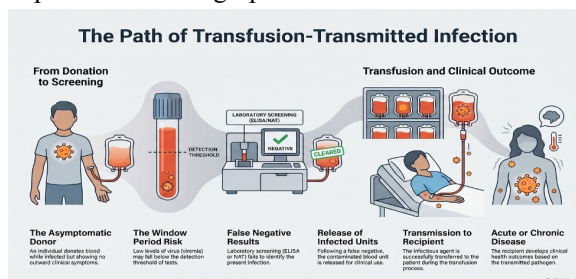
Blood transfusions, critical medical interventions, transmit infectious agents from donors to recipients, termed transfusion-transmitted infections (TTIs) (1). This constitutes an inherent risk. Asymptomatic blood

donors harboring pathogens persistently challenge transfusion safety. This necessitates continuous vigilance and enhanced detection techniques despite stringent screening measures (2). Despite advanced donor screening and blood processing, novel

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pathogens and detection limitations threaten the global blood supply. This threat is particularly acute during the infection phase prior to seroconversion or symptom manifestation (2). This review examines TTI mechanisms, obstacles, and countermeasures originating from asymptomatic donors. It covers both traditional and novel infectious agents. Specifically, it explores pathogen identification challenges during asymptomatic or pre-symptomatic stages. This includes the critical window period, during which standard tests fail to detect infected donors (3). The review also investigates implicated pathogens, their asymptomatic dissemination processes, and advancing screening and mitigation approaches. These include pathogen inactivation methods (2). Additionally, it highlights specific infection risks: SARS-CoV-2, dengue, Zika, and mpox. Asymptomatic viremia precedes symptoms for these agents, thereby heightening transfusion transmission potential (4). The paper further considers global and Indian viewpoints on TTI monitoring and risk control. This highlights regional differences and efforts to bolster blood safety. It then discusses prospective strategies to reduce residual TTI risks. These include cutting-edge diagnostics, refined donor criteria, and unified 'One Health' surveillance models. This involves tackling emerging infections lacking routine screening, which can spread undetected and enable accidental transmission (4). Blood-borne pathogen adaptability and novel agent emergence necessitate persistent surveillance and screening enhancements to safeguard the blood supply (5). Transfusion medicine evolution integrates emerging science and technologies to protect public health (6). Understanding donor TTI seroprevalence is vital for effective screening and risk reduction. This is critical for asymptomatic carriers and during infection window periods (7). Nucleic acid amplification technology (NAT) has lowered risks for established agents—HIV, HCV, and HBV—to  $<1:10^6$  units in high-resource nations. Conversely, vector-borne threats (e.g., West Nile Virus, Chikungunya virus) require swift screening or deferral adjustments during outbreaks (8,9). Globalization accelerates novel pathogen spread, necessitating coordinated global surveillance and responses. This shift mandates a forward-looking, cross-disciplinary 'One Health' framework. The framework links human, animal, and environmental monitoring to preempt blood supply risks (10). It recognizes interconnectedness as a driver of disease emergence, requiring interdisciplinary collaboration for transfusion safety (11). Combining serological virus

antibody/antigen tests with NAT for HIV, HCV, and HBV has substantially improved global blood product viral safety. This has slashed classic pathogen transmission risks (12). Conversely, resource-limited areas face higher TTI burdens. This highlights the need for flexible, robust screening (13). For instance, Ghanaian research reports a 21.0% overall TTI rate. This rate varies by pathogen (e.g., HBV, HCV, HIV, syphilis), stressing the critical need for blood safety improvements in high-prevalence contexts (14).



**Figure 1:** Flowchart illustrating the mechanism of transfusion-transmitted infections from asymptomatic donors. Despite routine screening, infections may escape detection during the window period or due to low-level viremia, leading to transmission in recipients.

### Methods

This section delineates the systematic methodology underpinning this comprehensive review, detailing the search strategy, selection criteria for relevant literature, and the analytical framework used to synthesize the findings. The methodology encompassed a comprehensive search of peer-reviewed articles and relevant guidelines from reputable scientific databases, focusing on publications from 2018 to 2024 to ensure currency and relevance. Specific databases interrogated encompassed PubMed, Scopus, Web of Science, and Google Scholar, using a combination of keywords such as "transfusion-transmitted infection (TTI)," "asymptomatic donors," "blood safety," "emerging infectious diseases," "screening strategies," "window period," and specific pathogen names (e.g., "SARS-CoV-2," "Zika," "Dengue," "Hepatitis E," "Mpox"). The initial search yielded a substantial number of articles, whereupon subsequent systematic screening was conducted based on their titles and abstracts to identify studies directly pertinent to the transfusion transmission of pathogens from asymptomatic donors. Full-text articles, including original research, review articles, and systematic reviews, which met the defined inclusion criteria and focused on human blood transfusion safety and pathogen transmission, were

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subsequently retrieved and subjected to critical appraisal for methodological rigor and relevance to the review's objectives. Data extraction focused on identifying common and emerging transfusion-transmissible infections, screening methodologies, challenges posed by the window period, risk mitigation strategies, and global epidemiological trends, with a particular emphasis on recent outbreaks and their impact on blood safety. This synthesis of extracted information facilitated a comparative analysis of screening efficacy, the identification of persistent gaps in blood safety protocols, and the delineation of future research directions. This structured approach ensured a thorough and evidence-based examination of the complex interplay between asymptomatic donors and TTI (TTI) risks, providing a robust foundation for the subsequent sections of this review. A medical librarian experienced in systematic review retrieval provided search strategy assistance (15).

### Types of Pathogens Transmitted Through Transfusion

Blood transfusion-transmissible pathogens include established viral, bacterial, and parasitic agents. Emerging zoonotic infectious diseases increasingly contribute to this pathogen spectrum. These transfusion-transmissible infections cause substantial morbidity and mortality in recipients, necessitating rigorous donor screening and blood product testing protocols (16). Historically, pathogens such as HBV, HCV, HIV, and *Treponema pallidum* (syphilis) have been major concerns, prompting development of sensitive serological and nucleic acid amplification tests (NATs) (17,18). Despite these advancements, numerous bloodborne agents beyond routinely screened pathogens persist as concerns (18,19). The evolving infectious disease landscape, coupled with global travel and climate change, introduces new or re-emerging pathogens into the blood supply. This poses ongoing challenges for blood safety agencies worldwide (18,20). These agents include bacterial, viral, parasitic, and prion entities. Each presents unique detection and prevention challenges for transfusion transmission (21),(19),(17,22). Prioritized pathogens include HIV, HBV, HCV, and *Treponema pallidum* (syphilis). These pathogens, despite advanced screening methodologies, present a substantial, albeit attenuated, risk. This risk is attributable to nascent infection's asymptomatic presentation and dynamic pathogen evolution (23),(24),(25).

### Bacterial Pathogens

Routine screening for *Treponema pallidum* is standard. Other bacterial pathogens lack routine screening in blood donations. These agents clearly present a substantial risk, particularly considering evolving demographics and increased global mobility (26). Bacterial contamination, though infrequent, may constitute a primary etiology of transfusion-associated sepsis, often originating from donor bacteremia or suboptimal venipuncture site preparation (27),(28),(29). This can result in significant morbidity and mortality, especially in immunocompromised recipients (27,30). Principal bacterial contaminants frequently implicated in septic transfusion reactions encompass *Staphylococcus aureus*, coagulase-negative staphylococci, *Bacillus cereus*, and *Yersinia enterocolitica*, with cold-storing pathogens like *Yersinia* clearly present specific challenges for refrigerated blood products (31),(32). The implementation of bacterial culture and rapid detection assays, alongside enhanced donor screening questionnaires and improved collection procedures, has been crucial in mitigating this risk (33,34). The emergence of multidrug-resistant bacteria poses an additional threat to transfusion safety. This necessitates continuous surveillance and the development of rapid detection technologies (35).

### Viral Pathogens

Viral pathogens comprise a diverse, continuously evolving agent class. This class significantly impacts transfusion safety, encompassing characterized viruses and novel strains (36). Historically, the most critical viral agents appear to encompass human immunodeficiency virus (HIV), hepatitis C virus (HCV), and hepatitis B virus (HBV). These three viruses drove substantial advancements in donor screening and blood product testing protocols (37,38). Nonetheless, despite stringent testing protocols, residual risk of transfusion-transmitted viral infections (TTVIs) persists. This risk demonstrably stems from: the "window period" phenomenon, false-negative test results, and novel viral threats unaddressed by routine screening panels (39,40). Additionally, certain viral pathogens may not cause severe acute symptoms in donors. However, they establish persistent infections or significant disease in immunocompromised recipients. This complicates risk assessment and mitigation strategies. (30,41).

### Parasitic Pathogens

Transfusion-transmitted parasitic infections (TTPIs) represent a critical concern in endemic regions. Global travel and migration patterns increase this risk, despite

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lower incidence in developed countries (42), (43). TTPIs exhibit prolonged asymptomatic carriage in donors. This presents unique diagnostic challenges, stemming from parasitic life cycle diversity and the absence of universal screening assays for all relevant parasites (42). Malaria, Chagas disease (*Trypanosoma cruzi*), and babesiosis (*Babesia microti*) constitute significant parasitic threats. These necessitate specific donor deferral criteria or targeted testing strategies in high-prevalence areas (42), (43). Toxoplasmosis and leishmaniasis also pose transfusion risks. This is due to varying geographical pathogen distribution and current broad-spectrum screening methodology limitations (44), (45). Protozoan parasites (*Plasmodium* species, *Trypanosoma cruzi*) present specific challenges. Their long-term persistence in asymptomatic carriers facilitates inadvertent transmission via blood transfusions (42). These cryptic infections frequently circumvent conventional donor screening. This is particularly true in non-endemic regions with low clinical suspicion (42). The low incidence of TTPIs compared to bacterial and viral contamination masks their severe illness potential. These infections significantly impact immunocompromised recipients (43).

### Prion Diseases

Prion diseases, including variant Creutzfeldt-Jakob disease (vCJD), present a distinct challenge to transfusion safety. This is primarily due to prolonged asymptomatic incubation periods, absence of reliable ante-mortem diagnostic tests, and resistance to conventional pathogen inactivation methods (46). The potential for transfusion-transmitted vCJD, despite its low documented incidence, prompted extensive precautionary measures. These include donor deferral policies and leukoreduction to minimize theoretical risks (47), (48).

### Mechanisms of Asymptomatic Transmission

The asymptomatic nature of certain infections in blood donors constitutes a central tenet for transfusion transmission risk assessment: infected individuals carry pathogens without overt clinical symptoms, thus eluding standard health assessments (49). This phenomenon spans a wide pathogen spectrum: viruses, bacteria, and parasites. Each employs distinct silent carriage mechanisms (50). These mechanisms include pathogen immune evasion, host genetic factors influencing disease presentation, or dynamic equilibrium between pathogen replication and immune control, preventing overt disease expression in donors (51), (52). Consequently, infected asymptomatic donors present an insidious challenge to blood safety. This

necessitates advanced detection strategies beyond symptomatic presentation (53), (49). Pathogen establishment of latent or chronic infections, coupled with host immune response variability, contributes significantly to the asymptomatic carrier state, complicating detection (51). Moreover, the "window period" following initial infection further complicates early detection. During this period, a donor is infectious, but standard serological tests remain negative, expanding the asymptomatic carrier pool (54).

### Window Period Infections

The window period (WP) represents the interval between infection and detectability of pathogen-specific markers (antibodies, antigens, or nucleic acids) by routine screening assays (25). This WP constitutes a critical vulnerability within the blood supply chain (55). During this phase, infected donors can transmit pathogens through transfusion even when conventional serological or nucleic acid tests (NAT) yield negative results. This phenomenon presents significant challenges for emerging infections and those with highly variable incubation periods. Seroconversion dynamics frequently do not align with established testing protocols, facilitating the ingress of infectious donations into the blood supply. WP duration and characteristics vary significantly among disparate pathogens. These factors directly influence the residual risk of transfusion transmission (25). For example, dengue virus-infected individuals present asymptomatic states for extended periods. Their blood may test negative via conventional methods, yet transmission occurs (56). Similarly, HIV and HCV infections exhibit high viral loads during the pre-seroconversion phase. This necessitates NAT implementation for transmission risk mitigation during this critical interval (57), (58).

### Latent Infections

Latent infections involve pathogen persistence in a quiescent state within the host, without active replication or overt disease. This poses a significant challenge to transfusion safety.

### Classical Transfusion-Transmitted Infections (TTIs)

Hepatitis B virus (HBV), hepatitis C virus (HCV), and human immunodeficiency virus (HIV) are historically prevalent pathogens. They necessitate rigorous monitoring due to established asymptomatic carriage and profound clinical transmission implications (59). Despite nucleic acid testing (NAT) implementation and stringent donor selection, residual risks persist. These risks originate particularly from donations made during infection window periods (60), (61). These

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classical transfusion-transmitted infections (TTIs) drive diagnostic innovation in sensitivity and specificity. Robust surveillance and adaptable screening protocols are thus required to maintain blood product safety (61). Historically, serological testing for TTIs served as a blood safety cornerstone. However, it often fails to detect early window period infections; only NAT identifies low-level viremia or early infection stages (60). However, novel and re-emerging infectious agents necessitate continuous re-evaluation of existing screening paradigms. This requires the development of assays with high sensitivity and specificity for broader pathogen detection. This clearly demonstrates the continuous challenge of achieving zero risk. False negatives and transmissions, even if rare, demonstrably occur with advanced technologies like nucleic acid amplification tests (NATs) (62).

### Human Immunodeficiency Virus (HIV)

The introduction of highly sensitive nucleic acid testing for human immunodeficiency virus (HIV) significantly curtailed the window period, reducing the risk of transmission to approximately 11 days post-exposure, thereby, the substantial enhancement of blood safety(63). Consequently, this advancement has clearly decreased the residual risk of transfusion-associated HIV, thereby transforming it from a significant concern to an extremely rare event in regions with advanced screening infrastructure. Model-based estimates clearly establish the residual risk of HIV transmission via transfusion at approximately one in 1,800,000 to one in 2,000,000 donations (64),(63). This reduction results from antigen and antibody testing combined with NAT. This combination significantly shortens the diagnostic window period (65). Despite these improvements, ongoing surveillance and the development of even more sensitive assays remain crucial. This necessity stems from viral strain evolution and variations in diagnostic performance across diverse donor populations.

### Hepatitis B Virus (HBV)

Hepatitis B virus (HBV) exhibits persistent global prevalence. This necessitates stringent, evolving screening protocols due to its chronic infection capacity and its long, often asymptomatic incubation period. The implementation of highly sensitive HBsAg assays and HBV DNA NAT has been pivotal. These assays reduce HBV transfusion-transmission risk by detecting occult and acute infections prior to seroconversion (66). Despite these advances, challenges persist: HBV's occult infection phase (HBsAg undetectable, HBV DNA present) and mutations in the

'a' determinant region impacting HBsAg detection. These necessitate vigilant monitoring and exploration of new diagnostic strategies. Moreover, the potential for immune escape mutants complicate diagnostic efforts. This underscores the necessity for multi-target NAT and improved serological assays capable of detecting diverse viral variants. The residual risk of HBV transmission, even with advanced testing, persists due to the prolonged, often asymptomatic window period, estimated at approximately 11.6 days when tested individually(67).

### Hepatitis C Virus (HCV)

Rigorous screening for Hepatitis C Virus (HCV) has also demonstrably enhanced transfusion safety. This was achieved via anti-HCV antibody testing and Nucleic Acid Testing (NAT) for HCV RNA, which critically reduced both the window period and the residual risk of transfusion-transmitted HCV (68). These implemented methods effectively reduced the residual HCV transmission risk to 1 in 0.83 million donations (69). Challenges persist. These include detection of low viral loads in immunocompromised donors or during early seroconversion. Furthermore, novel HCV genotypes or variants may evade current detection methods. Historically, post-transfusion hepatitis, specifically HBV and HCV infections, presented substantial challenges in blood safety. HBV exhibited a higher residual transmission risk than HCV or HIV due to its prolonged serologically negative window period (70).

### Syphilis

Although less common than viral transfusion-transmissible infections (TTIs), bacterial infections like syphilis constitute a persistent risk, particularly due to the challenges inherent in the identification of asymptomatic donors with latent infections. Current syphilis screening utilizes serological assays for treponemal and non-treponemal antibodies. These methods exhibit detection limitations for early-stage or latent infections, where spirochetemia levels may be intermittent or below assay detection thresholds. Molecular diagnostics, such as PCR-based assays targeting specific treponemal DNA sequences, demonstrate augmented sensitivity for direct pathogen detection in early and latent syphilis. This clearly mitigates limitations of traditional serological screening (71). Integration of advanced molecular methods into routine screening reduces residual syphilis risk. These methods detect window-period and asymptomatic infections(72). Rapid multiplex assays for diverse bacterial pathogens—and adaptive screening, novel biomarkers, and pathogen-agnostic

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detection—are essential for counteracting evolving strains and resistance. The development of broad-spectrum pathogen reduction technologies presents a promising avenue. These technologies inactivate a broad spectrum of bacterial contaminants, including syphilis-implicated pathogens, within blood products prior to transfusion. This approach provides an additional safety stratum beyond donor screening and testing. It addresses inherent limitations in identifying all asymptomatic bacterial carriers<sup>(73)</sup>.

### **Malaria**

Malaria, caused by Plasmodium, constitutes a notable transfusion-transmissible infection, particularly in endemic regions, thereby necessitating stringent donor screening attributable to its frequently asymptomatic parasitemia in semi-immune individuals. Malaria screening strategies in blood donations exhibit geographical variance. Non-endemic areas employ serological and molecular testing; endemic regions rely on donor deferral questionnaires and microscopy<sup>(74),(75)</sup>. Antigen and antibody detection tests are common. However, low parasitemia levels and Plasmodium species genetic diversity limit their sensitivity. Robust diagnostic modalities are imperative<sup>(76),(77)</sup>. Molecular diagnostic techniques, such as PCR, offer enhanced sensitivity for detecting low-level parasitemia and identifying different Plasmodium, thereby providing a more reliable method for mitigating transfusion-transmitted malaria<sup>(78)</sup>.

### **Emerging Transfusion-Transmitted Infections (TTIs)**

The landscape of transfusion-transmitted infections is continuously reshaped by the emergence of novel pathogens and the re-emergence of established infectious agents. Factors include globalization, climate change, and evolving human-animal interfaces<sup>(35)</sup>. This dynamic environment necessitates continuous vigilance and adaptable screening strategies to safeguard the blood supply against emerging threats<sup>(79)</sup>. The absence of established testing or mitigation interventions for certain diseases currently in widespread use poses a complex challenge for transfusion medicine. The implications of screening for these emerging transfusion-transmissible infections extend beyond direct pathogen detection. It encompasses resource allocation, test limitations, and donor deferral ethical dimensions<sup>(80)</sup>. New pathogen reduction technologies (PRTs) further mitigate these risks. PRTs inactivate a broad spectrum of infectious agents within blood products, thus enhancing overall transfusion safety<sup>(81),(82),(83)</sup>.

### **Dengue Virus**

Dengue virus constitutes a significant risk to blood safety, particularly in endemic regions, owing to its high prevalence and the frequently asymptomatic nature of infection during the viremic phase<sup>(41)</sup>. The challenge is further compounded by several factors: the absence of widespread licensed screening assays, the potential for occult viremia in afebrile donors, and the possibility of transfusion-associated dengue<sup>(84)</sup>. Nucleic acid testing (NAT) for dengue virus RNA functions as a crucial screening tool. NAT detects the virus during pre-symptomatic and early symptomatic phases. This effectively reduces transfusion-transmitted dengue (TTD) risk in high-incidence regions<sup>(85)</sup>.

### **Zika Virus**

Zika virus (ZIKV) emergence constitutes a significant public health concern. Its association with severe neurological complications necessitated rapid implementation of effective blood donation screening strategies<sup>(86)</sup>. Primary concern: prolonged viremic window in asymptomatic individuals and potential for sexual and congenital transmission. This necessitates comprehensive donor deferral policies and NAT screening in affected regions<sup>(87)</sup>.

### **Chikungunya Virus**

Chikungunya virus (CHIKV), a mosquito-transmitted arbovirus, poses a blood safety risk. This risk, similar to dengue and ZIKV, stems from asymptomatic viremia in infected donors, especially during outbreaks<sup>(88)</sup>. Given the rapid global spread of arboviruses and the high proportion of asymptomatic infections, A universal blood safety approach is imperative. This approach encompasses proactive surveillance and adaptable screening methodologies. Such methods mitigate the persistent threat posed by emergent infectious diseases to the blood supply<sup>(84),(89)</sup>.

### **West Nile Virus**

West Nile Virus (WNV) constitutes a significant transfusion-transmitted infection (TTI) in North America. Seasonal outbreaks mandate robust nucleic acid amplification test (NAT) screening of blood donations in affected regions<sup>(90)</sup>. Emergent and re-emergent arboviruses pose a persistent challenge. This necessitates continuous global surveillance and the development of highly sensitive, specific diagnostic platforms. These measures ensure transfusion safety<sup>(91)</sup>.

### **SARS-CoV-2**

The SARS-CoV-2 pandemic clearly demonstrated potential risks from respiratory viruses to blood safety. This instigated research into transfusion

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transmissibility and donor screening protocols. Studies consistently indicate minimal transfusion risk, attributable to low, transient viremia. This redirects primary focus toward nosocomial prevention and supply continuity amidst prevailing restrictions. Future pandemics require ongoing variant monitoring, adaptable protocols, targeted screening, and pathogen inactivation. These efforts must mirror efficacious rapid response protocols for West Nile Virus (WNV), particularly regarding Nucleic Acid Testing (NAT) <sup>(92)</sup>.

### Hepatitis E Virus (HEV)

Hepatitis E Virus (HEV) demonstrates escalating recognition as a transfusion-transmissible pathogen. This is particularly true in highly endemic regions, where asymptomatic viremia contributes to chronic infections in immunocompromised recipients. Chronic HEV infection potential, particularly in vulnerable populations, necessitates robust screening strategies. Nucleic acid testing (NAT) detects viremic donations, especially in high-endemicity areas <sup>(93)</sup>.

### Mpox Virus

Mpox virus (MPXV), formerly monkeypox, emerged as a pathogen of concern for transfusion safety. Its potential for asymptomatic or oligosymptomatic presentation during prodromal and early eruptive phases could coincide with blood donation <sup>(94)</sup>.

### Trypanosoma cruzi (Chagas Disease)

Protozoan parasites cause babesiosis and Chagas disease. These organisms present significant, regionally specific transfusion transmission risks. Risk factors include prolonged asymptomatic parasitemia in infected donors and limitations in current routine blood center screening protocols<sup>(42)</sup>. The persistent asymptomatic carrier state in endemic areas necessitates the development and implementation of highly sensitive, specific diagnostic assays. These assays detect low-level parasitemia, preventing transfusion-associated infections. The varying geographical distribution of these parasitic infections necessitates a dynamic risk assessment framework. This framework enables the development of targeted screening strategies. These strategies account for donor travel history and epidemiological data.

Table 1: Comparison of Classical vs Emerging Transfusion-Transmitted Infections (TTIs)

Category	Pathogen	Type	Primary Mode of Transmission	Risk from Asymptomatic Donors	Key Challenges
Classical	HIV	Virus	Blood, sexual contact	High (long window period)	Early detection during window period
Classical	HBV	Virus	Blood, body fluids	Very high (occult infection possible)	Occult HBV infection, low viral load
Classical	HCV	Virus	Blood	Moderate to high	Asymptomatic chronic infection
Classical	<i>Treponema pallidum</i> (Syphilis)	Bacteria	Blood	Low to moderate	Reduced viability in stored blood
Emerging	Dengue virus	Virus	Vector-borne, transfusion	High (during viremic phase)	Seasonal outbreaks, asymptomatic donors
Emerging	Zika virus	Virus	Vector-borne, transfusion	High	Often asymptomatic, congenital risk
Emerging	Chikungunya virus	Virus	Vector-borne, transfusion	Moderate	Limited routine screening
Emerging	West Nile virus	Virus	Vector-borne, transfusion	High	Silent infections in donors

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Category	Pathogen	Type	Primary Mode of Transmission	Risk from Asymptomatic Donors	Key Challenges
Emerging	Hepatitis E virus (HEV)	Virus	Feco-oral, transfusion	Moderate to high	Chronic infection in immunocompromised
Emerging	SARS-CoV-2	Virus	Respiratory (possible blood)	Low (uncertain)	Limited evidence, evolving data
Emerging	Mpox virus	Virus	Contact (potential blood)	Emerging risk	Insufficient transfusion data

A comparative overview of classical and emerging transfusion-transmitted infections is presented in Table 1.

### Screening Strategies for Asymptomatic Donors

Many transfusion-transmissible infections exhibit silent incubation or chronic phases. Therefore, effective screening is essential for blood safety [\(95\)](#). Comprehensive strategies implement donor questionnaires, serological tests, and nucleic acid testing (NAT) to detect asymptomatic cases. Deferrals based on geography, travel, and lifestyle risks confer a preliminary protective effect. However, questionnaires exhibit limitations. These limitations result from potential unawareness or non-disclosure of infectious status by asymptomatic carriers. Laboratory assays primarily afford objective detection.

Documentation of transfusion-associated Mpox virus (MPXV) cases is unrecorded. Asymptomatic viremia and outbreaks necessitate heightened vigilance [\(96\)](#). The MPXV pandemic initiated significant deliberation concerning universal PCR testing, despite deferral guidelines for cases and contacts [\(97\)](#). Implementation of sensitive assays is paramount. MPXV DNA is detectable in the bloodstream during viremia; however, elucidation of associated donor risks warrants further investigation [\(98\)](#).

Table 2: Screening Methods in Blood Transfusion – Advantages and Limitations

Screening Method	Principle	Advantages	Limitations
ELISA (Enzyme-Linked Immunosorbent Assay)	Detects antigen/antibody reaction	Cost-effective, widely used, high throughput	Cannot detect infections during window period
NAT (Nucleic Acid Testing)	Detects viral RNA/DNA directly	High sensitivity, early detection, reduces window period	Expensive, requires technical expertise
Rapid Diagnostic Tests (RDTs)	Immunochromatographic assays	Quick results, easy to perform, useful in remote areas	Lower sensitivity and specificity
Donor Health Questionnaire (DHQ)	Behavioral and medical risk assessment	Identifies high-risk donors, inexpensive	Depends on donor honesty, subjective
Chemiluminescence Immunoassay (CLIA)	Detects antigen/antibody using luminescence	Higher sensitivity than ELISA, automated	Costlier than ELISA

Table 2 summarizes commonly used screening methods along with their advantages and limitations.

### Donor History Questionnaire

The questionnaire-based approach, while foundational, possesses inherent limitations. These include reliance on donor candor and exposure awareness, which is challenging for emerging or unrecognized infections. Furthermore, asymptomatic individuals cannot effectively self-defer due to absent overt clinical symptoms. This necessitates

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complementary objective screening methodologies. Consequently, deployment of advanced laboratory testing, including Nucleic Acid Amplification Tests (NAATs), becomes indispensable. This enables infection identification during the pre-symptomatic or asymptomatic window period when conventional methods are insufficient<sup>(99)</sup>.

### Serological Testing

Serological testing detects infection-induced antibodies or specific antigens. This identifies past or ongoing infections, critical for pathogens with prolonged seroconversion windows or persistent humoral responses. Serological assays detect established infections. However, they may not detect acute infections during the pre-seroconversion window period. Molecular diagnostics integration is necessary for comprehensive donor screening<sup>(99)</sup>. Nucleic Acid Testing (NAT), a nucleic acid amplification technology, constitutes a cornerstone of modern blood safety protocols. NAT enables direct pathogen genetic material detection, significantly reducing the window period for numerous transfusion-transmissible infections<sup>(100)</sup>. This approach facilitates viremic donation identification prior to antibody development. It mitigates transmission risk for infections like HIV, HCV, and HBV during early, highly infectious stages<sup>(100)</sup>. This technological advancement demonstrates enhanced the safety of the blood supply, particularly for pathogens like West Nile Virus; NAT proves instrumental in transmission prevention<sup>(101)</sup>.

### Nucleic Acid Testing (NAT)

Nucleic Acid Testing (NAT) represents a pivotal advancement in transfusion safety. It offers the most sensitive, direct methodology for viral genetic material detection. This yields a significantly abbreviated infection-detectability window period compared to serological methods<sup>(102)</sup>. This mechanism facilitates infection identification prior to seroconversion. It addresses a critical vulnerability of traditional antibody-based screening<sup>(103)</sup>. The implementation of NAT demonstrates high efficacy in reducing the residual risk of transfusion-transmitted infections such as HIV, HCV, and HBV, resulting in per-unit risks less than 1:1,000,000 in high-income countries<sup>(104)</sup>.

### Pathogen Reduction Technologies

Pathogen reduction technologies constitute an additional stratum of safety through the inactivation of pathogens within donated blood components. This contributes to further residual risk reduction for transfusion-transmitted infections<sup>(105),(106)</sup>.

### Challenges in Detecting Pathogens During the Window Period

The "window period" defines the post-infection interval. Pathogens persist in the bloodstream before detectable antibody levels develop, despite negative diagnostic outcomes<sup>(107)</sup>. This interval impedes blood safety. Donors may unknowingly transmit pathogens, circumventing rigorous screening protocols. This demonstrates the persistent threat emergent infectious agents pose to blood supply integrity<sup>(107),(105)</sup>. This interval varies based on pathogen type and diagnostic assay sensitivity. The range typically spans several days to multiple weeks. For HIV, Nucleic Acid Testing (NAT) yields a window period of 7–11 days. This duration is significantly shorter than the 22-day interval for antibody-based tests<sup>(102),(104)</sup>. Despite this reduction, NAT presents a considerable challenge to blood product safety.

### Limitations of Current Assays

Despite technological advancements, inherent limitations persist. These manifest in infections with low viral loads or in emergent pathogens where established diagnostic assays lack sufficient sensitivity or specificity<sup>(108)</sup>. Furthermore, pathogen genotype evolution facilitates detection evasion by assays calibrated for antecedent strains. This necessitates perpetual assay refinement and heightened surveillance protocols<sup>(5)</sup>. Consequently, the diagnostic window period remains a significant impediment in transfusion medicine. This mandates innovative strategy development for early pathogen detection in asymptomatic donors<sup>(109),(110)</sup>. For instance, heterogeneous immune responses influence antigen expression and antibody production rates. This complicates early detection and augments the residual risk of transfusion-transmitted infections<sup>(58)</sup>.

### Cost-Effectiveness of Advanced Screening

Highly sensitive, broad-spectrum screening methodologies present economic considerations. These include infrastructure investment, operational expenses, and increased donor deferral rates. A rigorous cost-benefit analysis is required. This analysis must balance enhanced safety with blood product sustainability and accessibility across diverse healthcare systems. Advanced screening methods, such as nucleic acid amplification testing (NAAT), significantly reduce the window period. However, they do not eliminate transmission risk from asymptomatic donors, especially for novel or evolving pathogens<sup>(108),(111)</sup>.

### Global and Indian Scenario of Transfusion-Transmitted Infections

Globally, transfusion-transmitted infection (TTI) prevalence and incidence demonstrate substantial

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variation. This variation is due to disparate epidemiological landscapes, socioeconomic determinants, and blood screening protocol sophistication. Resource constraints in developing nations preclude advanced screening technology adoption, thus elevating TTI prevalence among blood donors (112). Developed countries achieve substantial TTI incidence rate reduction via comprehensive screening. This includes nucleic acid testing (NAT), robust donor selection criteria, and stringent quality control measures. However, transmission risk persists during the window period or from emerging pathogens (113). Specific challenges include hepatitis E virus genotype 3 in European regions and Babesia in USA regions; these highlight localized threats. Globally, continuous emergence of new infectious agents necessitates dynamic monitoring and adaptive safety measures (6). In India, the existing scenario complicates TTI risk profiles (16). Elevated TTI prevalence in India, primarily due to window-period donations, mandates comprehensive strategies. This includes mandatory NAT for enhanced blood safety within resource-constrained environments (114). Furthermore, dependence on replacement donors in Indian blood banks introduces additional risks. These donors often do not undergo the rigorous screening protocols applied to voluntary, professional donors (7).

### Future Perspectives and Research Directions

Further advancements in pathogen detection technologies, including multiplex PCR assays and next-generation sequencing, demonstrate capability in reducing the diagnostic window period and rapid identification of novel infectious agents (115). Additionally, the development of pan-pathogen detection systems capable of identifying a broad spectrum of known and unknown infectious agents proves crucial for the proactive response to emerging threats (116). Moreover, the integration of artificial intelligence and machine learning algorithms enhances predictive modeling for outbreak detection and risk assessment. This optimizes donor screening protocols (117,118). Ongoing research into pathogen reduction technologies also offers a pathway the neutralization of a diverse array of viruses, bacteria, and parasites in donated blood components. This provision provides an additional safety layer beyond donor screening (18).

### Conclusion

Transfusion transmission of pathogens from asymptomatic donors continues to represent a significant challenge in ensuring the safety of the blood supply. Despite considerable advancements in

donor screening and laboratory testing, the risk cannot be completely eliminated due to factors such as the diagnostic window period, occult infections, and the emergence of new pathogens. Both classical transfusion-transmitted infections and recently emerging infectious agents highlight the need for continuous vigilance and improvement in transfusion practices.

The integration of stringent donor selection criteria with advanced screening technologies, including nucleic acid testing and highly sensitive immunoassays, has substantially reduced the residual risk of infection. However, disparities in infrastructure and resource availability, particularly in developing regions, limit the universal implementation of these strategies. Therefore, strengthening hemovigilance systems, enhancing public awareness, and promoting voluntary blood donation remain essential components of a safe transfusion system.

Future efforts should focus on the adoption of innovative approaches such as pathogen reduction technologies and the incorporation of advanced diagnostic tools to address both known and emerging threats. A comprehensive and adaptable blood safety framework, supported by global collaboration and equitable resource distribution, is crucial to minimizing transfusion-related risks and ensuring safe and effective blood transfusion practices worldwide.

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