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Essay

# A Mathematical Model of Disease

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**Abstract:** The identification of fetal cells in the blood and tissues of healthy mothers raised questions about the potential alloreactive origins of various diseases, such as autoimmunity, diabetes, arteriosclerosis, and cancer. However, the extent to which alloreactivity contributes to these diseases remains unverified. This study presents a basic mathematical disease model based on the genes' origin and immunological interactions. This model, grounded in set theory, classifies these immunological interactions based on their intrinsic, extrinsic, and allogeneic nature. Therefore, this model offers an unexpected common theoretical foundation for understanding alloreactivity, genetic diseases, and infections. Furthermore, this model provides a valuable framework for scrutinizing the role of alloreactivity in health and disease.

**Keywords:** alloreactivity; fetal microchimerism; relativity; set theory; non-communicable diseases; autoimmunity; diabetes; arteriosclerosis; cancer

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## Introduction

The identification of genes and pathogens has paradigmatically changed our understanding of diseases. Traditionally, we attributed diseases to two primary categories: infectious diseases caused by pathogens and genetic diseases governed by Mendel's law of heredity, arising from genetic mutations. However, a significant group defies these classifications within the realm of diseases. This group encompasses disorders such as arteriosclerosis, type 2 diabetes, autoimmunity, chronic respiratory diseases, neurological diseases, and cancer. These diseases are classified as non-communicable diseases and constitute approximately 70% of global mortality, as estimated by the World Health Organization [1]. The etiology of these diseases is presumed to be multifactorial, arising from a complex interplay of lifestyle, behavior, environmental influences, and genetic factors.

Moreover, research has focused on additional factors, such as the gut microbiome [2] and epigenetics [3]. Studies on pregnancy and transplantation medicine have also suggested the potential involvement of alloreactivity in the development of these diseases. This idea is mainly supported by the fact that allogeneic cells transferred during pregnancies persist in small numbers and remain viable for decades after childbirth. These cells are referred to as microchimerism.

## Microchimerism

The existence of chimerism contradicts the inherent idea that cells exchanged between individuals cannot persist in a foreign immune system environment and should be rejected. Nevertheless, there is increasing evidence that most, if not all, individuals carry a small fraction of cells with an allogeneic origin [4–6]. Microchimerism has been detected in the blood [4] and multiple tissues, including the brain [7], liver [8], and pancreas [9]. Furthermore, chimerism has been observed in cattle, pigs, and other animals, suggesting it is not a rare evolutionary occurrence [10,11]. The discovery of microchimerism fundamentally challenges our concept of alloreactivity, as it indicates that allogeneic interactions are not confined to pregnancy or childbirth but may persist long after these events. Considering the lifelong persistence of these cells, it has been hypothesized that alloreactivity may be involved in non-communicable diseases [12,13].

## Autoimmune Diseases

Notably, a strong association exists between auto- and alloimmunity [12,14]. Chronic graft-versus-host disease (cGvHD) following allogeneic stem cell transplantation has similar clinical components as autoimmune diseases, such as dry mouth and eyes, joint stiffness, skin lesions, and fibrosis. Moreover, autoantibodies, which indicate autoimmune diseases, are frequently observed in patients with cGvHD. Antinuclear antibodies have been reported in approximately 65% of patients with cGvHD [15]. Detecting anti-topoisomerase I antibodies in cGvHD [16] is noteworthy as these antibodies are highly specific for systemic sclerosis [17]. Moreover, anti-platelet-derived growth factor (anti-PDGF) receptor antibodies have been identified in nearly all patients with systemic sclerosis. These antibodies are suspected to be significant in the disease's pathogenesis due to their capacity to activate collagen gene expression [18]. Similarly, anti-PDGF-receptor antibodies have been detected in approximately all patients with extensive cGvHD, whereas they are absent in controls without cGvHD [19].

Additional direct evidence linking auto- and alloreactivity stems from discovering allogeneic DNA in the tissues of patients with autoimmune diseases. Fetal DNA has been found in the salivary glands of women with Sjögren's syndrome [20], the skin lesions of women with systemic sclerosis [21], and the thyroid tissue of women with autoimmune thyroiditis [22,23]. Conversely, myocardial cells of maternal origin have been found in newborns with lupus congenital heart block [24]. However, the evidence of an association with autoimmune diseases is complicated by the fact that microchimeric cells can also be detected in healthy individuals [20–23].

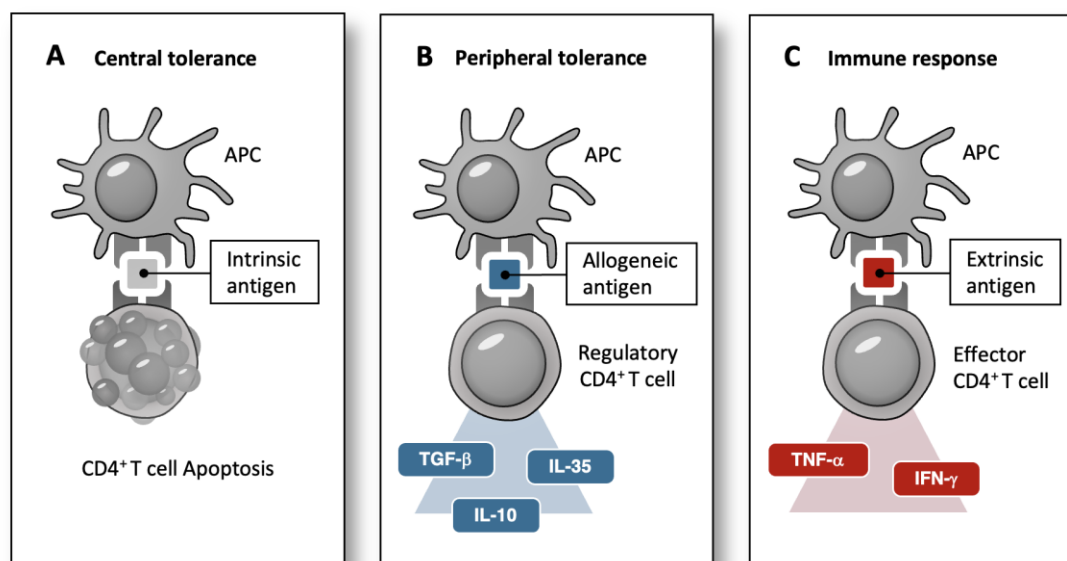
Nevertheless, studies with inbred mouse models provide compelling evidence supporting alloreactivity's involvement in autoimmune diseases. The injection of immune cells from one inbred mouse strain into another frequently induces autoimmunity, characterized by typical clinical and histopathological patterns [25].

## Type 2 Diabetes

Approximately 60% of women with gestational diabetes develop type 2 diabetes later in life [26]. Consequently, gestational diabetes represents one of the most significant risk factors for type 2 diabetes. Furthermore, gestational diabetes closely resembles type 2 diabetes in two clinical components: impaired glucose tolerance and insulin resistance. The development of gestational diabetes is attributed to the hormonal and metabolic changes that occur during pregnancy. However, pregnancy presents a unique immunological condition characterized by direct allogeneic cell contact at the fetomaternal interface and bidirectional exchange of antigens, antibodies, and cells. Fetal cells enter the maternal circulation during pregnancy and are not entirely removed after that [27,28]. It has been established that allogeneic encounters during pregnancy primarily result in immunological tolerance [29,30]; however, there is also evidence indicating that tolerance may be incomplete, with excessive alloreactivity being managed by peripheral tolerance [31].

In contrast to central tolerance, which involves the thymic depletion of autoreactive immune cells (Figure 1A), peripheral tolerance is closely associated with the secretion of cytokines, such as the transforming growth factor- $\beta$  (TGF- $\beta$ ) in response to antigens (Figure 1B). These cytokines can further affect the body, as TGF- $\beta$  has been shown to regulate glucose tolerance and pancreatic islet  $\beta$ -cell function in mice [32,33]. Furthermore, increased TGF- $\beta$  levels have been correlated with obesity in humans [33]. Therefore, an alternative explanation for gestational diabetes could be that fetal allogeneic cells in the maternal body stimulate maternal regulatory T cells to produce cytokines, such as TGF- $\beta$ , subsequently leading to gestational diabetes. Recently, a correlation between increased fetal cells in the maternal circulation and poor glucose control has been demonstrated in diabetic pregnancies [34].

Moreover, eclampsia, a pregnancy-associated disorder characterized by hypertension, proteinuria, and thrombotic microangiopathy, leads to endovascular damage and organ dysfunction. In eclampsia, placental damage increases fetal cell translocation into the maternal blood and tissues [35]. Pre-eclampsia is associated with gestational diabetes and a twofold increase in diabetes in later life [36].



**Figure 1.** Predominant immune reactions to intrinsic, allogeneic, and extrinsic antigens. **(A)** During T-cell maturation in the thymus, CD4<sup>+</sup>T-cells undergo apoptosis when they recognize self-antigens presented by thymic epithelial cells. **(B)** The contact of the immune system with allogeneic antigens primarily leads to peripheral tolerance. Regulatory CD4<sup>+</sup>T-cells secrete cytokines such as TGF- $\beta$ , IL-10, and IL-35 following antigen recognition. **(C)** Effector-CD4<sup>+</sup>-T-cells secrete cytokines like IFN- $\gamma$  and TNF- $\alpha$  in response to antigens of extrinsic origin, such as viral or bacterial antigens. APC, Antigen presenting cell; TGF- $\beta$  Transforming growth factor  $\beta$ ; IL-10, Interleukin-10; IL-35, Interleukin-35; IFN- $\gamma$  Interferon- $\gamma$  TNF- $\alpha$  Tumor necrosis factor- $\alpha$ .

## Arteriosclerosis

Human arteriosclerosis is a chronic inflammatory disorder with significant clinical manifestations, such as myocardial infarction and stroke [37]. Similarly, graft arteriosclerosis, which often occurs following organ transplantation, is the primary clinical factor limiting the long-term survival of allogeneic organ grafts [38]. The formation of arteriosclerotic plaques by allogeneic triggers is noteworthy. Perivascular inflammation and consecutive graft arteriosclerosis have been documented in heart, kidney, lung, and liver transplantations [38,39]. Histologically, there are similarities and differences between spontaneous and transplant arteriosclerosis [40–42]. The accumulation of lymphocytes, macrophages in the artery wall, and lipids in atherosclerotic plaques are common features in spontaneous and transplant arteriosclerosis. Transplant arteriosclerosis exhibits more diffuse concentric intimal thickening compared with spontaneous arteriosclerosis. Furthermore, spontaneous arteriosclerosis typically develops over several decades, whereas transplant arteriosclerosis may develop within 1 or 2 weeks. However, calcification occurs late in transplant arteriosclerosis.

Arteriosclerosis has also been observed during pregnancy, where acute atherosclerosis of the spiral arteries at the fetomaternal interface closely resembles early-stage atherosclerosis [43]. Acute atherosclerosis is rare in healthy pregnancies but is frequently observed in abnormal pregnancies, such as those complicated by pre-eclampsia [43,44]. The role of alloreactivity in acute atherosclerosis has also been previously discussed [44]; alloreactivity leads to transplant vasculopathy and is implicated in the etiology of spontaneous arteriosclerosis.

## Cancer

Chronic infections caused by human papillomavirus, hepatitis B and C, and *Helicobacter pylori* contribute significantly to cancer development [45]. In addition to viral oncogenes and cytokines, chronic inflammation is involved in cancer development and increases the risk of malignant transformation [46]. Allogeneic cells have been found in various cancer tissues, including melanoma,

breast, lung, thyroid, and brain tumors [13,47]. A malignant transformation caused by allogeneic cells is yet to be directly proven; however, their presence in tumor tissues has been extensively documented. Their active role in malignant transformation has been discussed in addition to the potential benefits of microchimerism, such as anti-tumor effects and tissue repair [13,47,48].

### **Alloreactivity in Non-Communicable Diseases**

In 1893, Georg Schmorl described fetal cells in the lungs of women who died of eclampsia [49]. He was the first to postulate a connection between fetal cells in the maternal circulation and a disease. Over a century later, in 1996, J. Lee Nelson advanced the exploration of this idea by introducing the question, "Is some autoimmune disease auto-alloimmune or allo-autoimmune?" [12]. Unfortunately, the low frequency of microchimeric cells prevents comprehensive tracking of their local and systemic effects. Even today, the theory that cells exchanged between individuals subsequently cause alloreactive effects in the host's body remains speculative. None of the previously discussed correlations offer definitive evidence, leaving the hypothesis neither validated nor disproven.

### **Mathematical Aspects of Alloreactivity**

Questioning whether microchimerism leads to disease goes beyond pinpointing a cause; it raises the intriguing question of whether diseases themselves might follow mathematical patterns, suggesting a deeper, rule-based structure to their occurrence and progression.

In contrast to microchimerism, alloreactivity and its role in health and disease have been more clearly defined. Focusing on alloreactivity, three fundamental mathematical concepts may help unravel its role in health and disease: relativity, set theory, and asymmetry. Importantly, these mathematical concepts of alloreactivity originate from concrete properties of the immune system.

### **The Immune System Works in a Relativistic Mode**

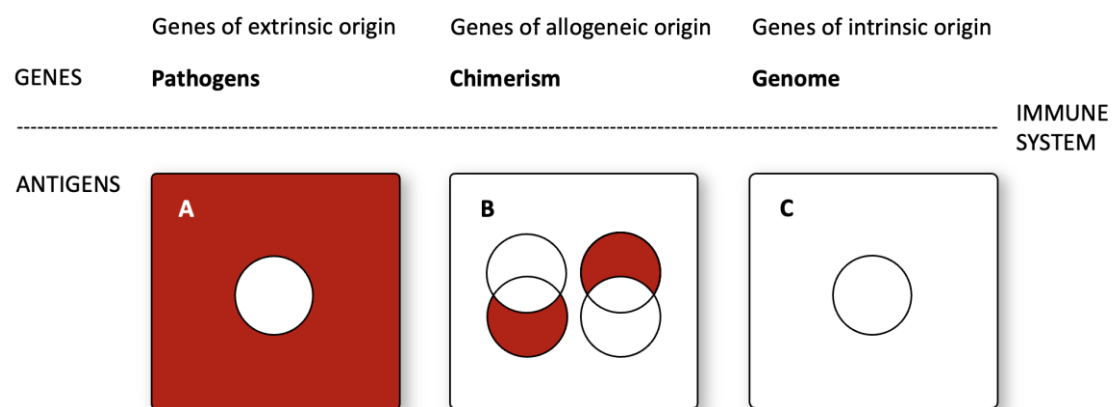
Relativity, a fundamental concept in physics, elucidates the intricate relationships between space, time, mass, energy, and gravity. Notably, the principles of relativity extend beyond the realm of physics to biological systems [50]. Based on the following findings, relativity is pivotal in the interaction between the immune system and genes: The immune system establishes a natural barrier against invasion, safeguarding the body against pathogens. Moreover, the immune system adapts to the body's genome, and during T-cell maturation, central immune tolerance eliminates T-cells in the thymus that react with self-antigens. This process, however, leaves an intriguing gap in immune surveillance (Figure 2A). Consequently, the immune system contains an individual's immunological information, characterized by a complete set of specific immune cells devoid of a depleted set of self-reactive cells. Therefore, each person possesses a distinct set of self-antigens, representing their immunological frame of reference. Since there is an individual and no universal or privileged frame of reference, the immune system operates in a relativistic mode [50].

### **A Relativistic Perspective on Alloreactivity**

Across different species, a substantial genetic commonality exists, with most genes being shared. However, alloantigens are encoded outside the overlapping genetic pool. Alloreactivity is the general term for immune responses to alloantigens not found within an individual's genome but instead encoded in the genomes of others within the same species. Alloantigens commonly arise due to genetic mutations. Notably, many blood group systems are founded on genetic variants, such as the Rhesus D blood group alloantigen resulting from a gene deletion and the H-Y alloantigen exclusively encoded on the Y chromosome. A central feature of alloantigens is their presence alongside genes shared by other individuals. Importantly, the information that triggers an alloreactive immune response is not contained within the host or donor genomes but depends on the exclusive antigen expression in one but not the other individual. Each immune system represents a distinct frame of reference encoded by an individual genome. Following transmission into a foreign body, the donor immune cells can detect host antigens not encoded by the donor's genome. Therefore, the term

“relativistic” implies that the ensuing effect defies explanation or prediction through the lens of a single genome. The personal genome provides only an individual frame of reference, and immunological incompatibility emerges in response to genomic differences between individuals.

In addition to minor alloantigens, variations in the major histocompatibility complex (MHC) determine alloreactivity; T-cells recognize a compound of the MHC and its presented antigen. T-cells may directly target MHC variants, but alterations in the MHC molecules lead to a shift in antigen selection. Therefore, transplantation of MHC-mismatched allogeneic organs or stem cells may lead to complex and unpredictable immunological interactions. This complexity can be markedly reduced if the immune system is considered a binary system that distinguishes self- and non-self-antigens, regardless of their MHC and minor antigen components. Since the term “non-self” pertains to a single individual other than oneself, the binary system, featuring two conditions, “self” and “non-self,” reflects the relativity of genes.



**Figure 2.** A mathematical model of disease: Immunological interaction of genes with extrinsic, allogeneic, and intrinsic origin.

The genes are classified based on extrinsic, allogeneic, or intrinsic origins. The immune system then converts the genes into antigens.

(A) Within the universe of all existing antigens (red rectangle), central and peripheral tolerance leads to a gap in immune surveillance (white circle). Extrinsic antigens are defined as the absolute complement of self-antigens (white circles) within all antigens (red rectangles). Pathogens are extrinsic antigens.

(B) Following the transfer of cells into another individual, donor immune cells detect host antigens by recognizing host antigens (left red circle) in a graft-versus-host-like reaction. Therefore, allogeneic antigens can be defined as a set of host antigens (left red circle) that do not overlap the self-antigens of the donor (left white circle): allogeneic antigens of the host are the relative complement of host antigens (left red circle) without donor antigens (left white circle). Host immune cells can recognize donor antigens through the antigen recognition of donor antigens (right red circle). Therefore, allogeneic antigens can be defined as a set of donor antigens (right red circle) without the self-antigens of the host (right white circle): allogeneic antigens of the donor are the relative complement of the set of donor antigens (right red circle) without the host antigens (right white circle).

(C) Individual genomes encode intrinsic antigens (self-antigens). Central and peripheral tolerance deletes or silences self-reactive immune cells. Therefore, antigens of intrinsic origin do not induce an immune response (white circle).

## A Mathematical Model of Disease

The interaction of T cells, antigen-presenting cells, and costimulatory signals commonly determines immune reactions. For this analysis, we will set aside the complex relationships of these cells and explore the situation from the antigen's perspective.

Alloantigens are defined by their absence in an individual's genome but their presence in the genomes of others within the same species. This definition implies that alloantigens, and indeed antigens more broadly, can be precisely characterized using the rules of set theory and their ability to describe the inclusion and exclusion of elements within sets. In addition to a mathematical description, antigens can be categorized by set theory. Self-antigens encoded within an individual's genome are of intrinsic origin (Figure 2C). Pathogenic antigens, such as bacterial and viral antigens, are extrinsic (Figure 2A). Antigens encoded by some, but not all, members of a species, such as the Rhesus-D-protein, are allogeneic (Figure 2B). Therefore, antigens can be categorized based on their intrinsic, extrinsic, or allogeneic origins.

Moreover, this classification serves as a mathematical model of disease integrating genetic, infectious, and allogeneic diseases (Figure 2). This model offers a mathematical approach by quantifying the relationships among these disease categories. It provides a mathematical framework for differentiating immune responses to alloantigens from those against extrinsic antigens. Furthermore, it accurately predicts that, unlike these antigens, the genome and genetic disorders are unlikely to trigger an immune reaction. Starting from intrinsic self-antigens that are genomically defined, extrinsic and allogeneic antigens can be mathematically calculated using two different axioms of set theory: the absolute and relative complements.

#### **Definition of Non-Self-Antigens as Absolute Complement by Set Theory**

B- and T-cells commonly recognize non-self-antigens, such as bacterial or viral antigens. The set theory provides a useful framework for classifying non-self-antigens. Extrinsic antigens, such as those from pathogens, represent the absolute complement of self-antigens (Figure 2A). Mathematically, a set of extrinsic antigens encompasses the entire spectrum without a set of self-antigens. These antigens are not encoded in an individual's genome; therefore, this set includes all viral and bacterial antigens and allogeneic antigens, such as the Rhesus protein. Allogeneic antigens are categorized as extrinsic; however, they can be further defined as follows:

#### **Definition of Allogeneic Antigens as Relative Complement by Set Theory**

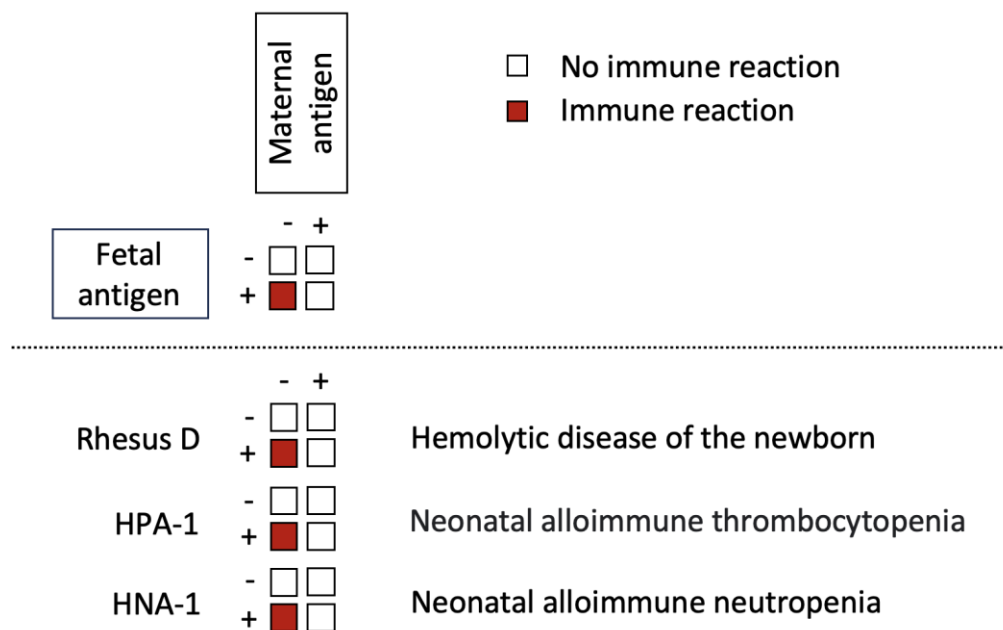
Allogeneic antigens represent a specific subset of non-self-antigens. During the exchange of cells between individuals, both immune and non-immune cells encounter foreign immune systems and antigens. When fetal immune cells enter the maternal circulation, the immunogenic targets can be more precisely characterized using another principle of set theory: the set of allogeneic minor antigens is defined as a relative complement of the host's (maternal) antigens without the self-antigens of the (fetal) donor (Figure 2B, left) and vice versa (Figure 2B, right).

Alloreactivity is a bidirectional process that manifests in response to maternal and fetal antigens. Maternal genes not present in the fetal genome encode proteins that can serve as allogeneic antigens. Consequently, when fetal immune cells enter the maternal circulation, they may recognize maternal antigens that are not contained in the set of self-antigens in a graft-versus-host-like reaction. Conversely, maternal immune cells or antibodies entering the fetal circulation may recognize fetal non-self-antigens in a host-versus-graft response, considering the fetus as a graft within the female's body. These interactions have clinical implications. For instance, following immunization with the Rhesus-D-protein during a first pregnancy, a Rhesus-negative woman's Rhesus-D-specific antibodies can destroy the blood cells of Rhesus-positive babies in subsequent pregnancies. This potentially fatal condition can be prevented by administering Rhesus prophylaxis to mothers who are Rhesus-negative.

#### **Asymmetry as a Marker of Alloreactivity**

Our diverse genetic backgrounds often result in unpredictable immunological interactions when cells are transferred from one individual to another. Asymmetry can reveal alloreactivity in specific scenarios. It arises due to the direction of the immune system; antigen-negative individuals respond to antigens that are not encoded in their genomes. For instance, when Rhesus D-negative mothers are immunized during pregnancy with a Rhesus D-positive child, it can lead to hemolytic disease in the newborn in subsequent pregnancies. Notably, other examples of alloreactive diseases include neonatal alloimmune thrombocytopenia, primarily caused by the human platelet antigen-1 (HPA-1), and neonatal alloimmune neutropenia, which depends on the human neutrophil antigen-1 (HNA-1) expression in the fetus but not in the mother (Figure 3).

A sex-dependent immunological imbalance is another example of asymmetry resulting from alloreactivity. The male genome carries genes on the Y chromosome that are absent in the female genome. This genomic distinction results in a sex-dependent immunological imbalance and has clinical implications for donor selection in human stem cell transplantation. It is widely acknowledged that using a female donor for a male recipient should be avoided, as this combination significantly increases the risk of graft-versus-host disease [51]. The HY antigen encoded in the Y chromosome has been identified as a relevant alloantigen in transplantations. In inbred mouse strains, such as C57BL/6, all individuals are genetically identical except males, who possess the Y chromosome. The HY antigen prompts female mice to reject skin grafts from male donors, whereas they tolerate female skin grafts, and male mice tolerate both female and male skin grafts [52]. This asymmetric pattern may help uncover the contribution of alloreactivity to human diseases.



**Figure 3.** The asymmetrical pattern of alloreactivity.

The fetus can be considered as a graft for the mother. Allogeneic immune responses occur when the lymphocytes of an antigen-negative mother recognize the antigen of an antigen-positive child (red squares). No immune reactions occur if the mother and child are antigen negative, both are antigen positive, or if the mother is antigen positive and the child is antigen negative (white squares). Examples include hemolytic disease of the newborn, in which lymphocytes of a Rhesus D-negative mother recognize the Rhesus D-antigen on the child's red blood cells; neonatal alloimmune neutropenia, in which lymphocytes of an HPA-1-negative mother recognize the HPA-1-antigen on the child's platelets; and neonatal alloimmune thrombocytopenia, in which lymphocytes of an HNA-1-negative mother recognize the HNA-1-antigen on the child's neutrophils.

HNA-1, Human neutrophil antigen 1; HPA-1, Human platelet antigen 1; Rhesus D, Rhesus blood group antigen D.

## Discussion

The genome encodes linear information stored within the solid molecular structure of the DNA. However, spontaneous mutations introduce dynamism into this otherwise static condition, maintaining an ongoing process where the DNA is subject to continuous modifications throughout evolution. Consequently, the genetic diversity has increased over time. During pregnancy, genetic disparities between mothers and children gain relevance and are primarily shaped by alloreactivity. Nevertheless, the continuous maternal effector immune cell response to fetal alloantigens would lead to severe fetal damage during pregnancy. This underscores the predominance of alloantigen tolerance within the immune system before birth [29]. However, it is crucial to consider three important aspects that suggest a limit of allogeneic tolerance: First, pregnancy directly proves that genomic variability, such as blood group incompatibility, can cause immunological incompatibility and subsequent pregnancy loss. Second, maternal microchimerism fosters peripheral tolerance to non-inherited maternal antigens and positively affects offspring survival in the next generation [6]. Conversely, this strongly implies a detrimental effect on the offspring of mothers lacking this acquired tolerance. Third, peripheral tolerance is, to some extent, incomplete, as it involves the secretion of cytokines, such as TGF- $\beta$  (Figure 1B).

Immunosuppression can be discontinued three months after allogeneic stem cell transplantation. This is widely accepted as strong evidence that stem cell transplantation results in tolerance. However, cGvHD is a common adverse effect and acute, chronic, and transfusion-associated GvHD have demonstrated that transferred immune cells can cause lethal damage to the host [53]. Remarkably, the extent of cGvHD following allogeneic stem cell transplantation increases with the sex, relationship, and ethnic backgrounds of the donor and host. Therefore, cGvHD is a gradual process, reflecting the accumulation of immunological incompatibility with increasing genetic distance [54,55]. Notably, relativity operates with modest effects at lower volumes and markedly amplifies effects at higher volumes. When tolerance is incomplete, accumulating immunological incompatibility may limit genomic variability within a species [56]. Therefore, alloreactivity, like an evolutionary clock, may be an immunological counterbalance to genomic variability.

Allogeneic immune responses during pregnancy, transfusion, and transplantation have been extensively studied. Naturally occurring alloreactive diseases include various conditions, such as hemolytic diseases in newborns, neonatal alloimmune neutropenia, and neonatal alloimmune thrombocytopenia. Notably, most red blood cells, neutrophils, and thrombocyte alloantigens are based on single nucleotide polymorphisms, underscoring the exceptional sensitivity of the immune system in detecting the smallest differences between the two genomes. The substantial and increasing number of alloantigens (> 300) in red blood cells, neutrophils, and platelets suggests the existence of additional undiscovered alloantigens in blood and tissues.

However, detecting the effects caused by alloreactivity remains difficult. Notably, most studies on microchimerism are based on quantifying allogeneic cells in blood or tissues; however, this approach does not provide insights into their function. In addition, the frequency of microchimeric T-cells is very low, suggesting that research on antibodies could be a more viable alternative. Antibodies are compelling evidence of a B-cell effector activity. As such, identifying antibodies from microchimeric B-cells or those targeting microchimeric antigens could offer a new perspective. The source of these antibodies can be identified through their distinct (non-self) allotype. Therefore, detecting antibodies with a foreign allotype indicates the immunological activity of microchimeric cells. For instance, patients with rheumatoid arthritis were found to produce antibodies targeting foreign immunoglobulin allotypes [57,58]. Moreover, the corresponding genes of these extrinsic immunoglobulins were detected in synovial tissues by polymerase chain reaction, suggesting a microchimerism involvement [57]. Unfortunately, there are also limitations on antibody studies: antibodies targeting foreign antigens are continuously cleared from the bloodstream through their binding to their respective tissues. Notably, multiple other strategies can be used to explore the impact of alloreactivity on diseases, such as animal models for type 2 diabetes and arteriosclerosis, in which antigens can be actively defined.

## Conclusion

The mathematical disease model presented here effectively integrates infectious, genetic, and alloreactive diseases. However, while this model does not prove that alloreactivity is involved in non-communicable diseases, it provides a valuable mathematical framework. This framework emphasizes that alloreactivity adheres to both predictable and quantifiable principles. Therefore, it can serve as a tool for evaluating the role of alloreactivity in health and disease. In this framework, diseases' primary origins are no longer attributed to misbehavior, dysfunction, or dysregulation. Instead, we understand that the immune system adapts and responds to the inherent relativity of genes.

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