REVIEW

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Complement networks in gene-edited pig xenotransplantation: enhancing transplant success and addressing organ shortage

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Abstract

The shortage of organs for transplantation emphasizes the urgent need for alternative solutions. Xenotransplantation has emerged as a promising option due to the greater availability of donor organs. However, significant hurdles such as hyperacute rejection and organ ischemia–reperfusion injury pose major challenges, largely orchestrated by the complement system, and activated immune responses. The complement system, a pivotal component of innate immunity, acts as a natural barrier for xenotransplantation. To address the challenges of immune responses and enhance the overall success of xenotransplantation. This comprehensive review aims to illuminate strategies for regulating complement networks to optimize the efficacy of gene-edited pig xenotransplantation. We begin by exploring the impact of the complement system on the effectiveness of xenotransplantation. Subsequently, we delve into the evaluation of key complement regulators specific to gene-edited pigs as a viable source of organs. These investigations provide valuable insights into the feasibility and potential success of xenotransplantation, offering a bridge between scientific advancements and clinical application.

Keywords Xenotransplantation, Complement systems, Genetically modified pigs, Clinical trials, α -1,3-galactosyltransferase gene-knockout

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Introduction

Transplantation is a crucial strategy for addressing endstage organ failure. However, the persistently low supply of donated human organs has resulted in a growing demand that far exceeds the available supply. In the U.S. alone, with 103,327 individuals awaiting organ transplants, only 42,000 transplants were conducted in 2022, leading to a tragic daily toll of 17 lives lost while awaiting organs [1]. Recognizing the striking similarities in size and various biological aspects between porcine and human organs, pigs have emerged as prime candidates for xenotransplantation. Their potential is further amplified through genetic engineering, enabling pigs to serve as optimized sources for cells, tissues, and organs



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in transplantation [2]. Recent strides in genome editing have significantly propelled advancements in this field, but graft rejection remains a pressing problem.

A critical factor contributing to xenograft failure is the activation of the complement system, resulting in hyperacute rejection, ischemia-reperfusion injury, coagulation disorders, and related inflammatory responses. Consequently, comprehending and mitigating immune rejection triggered by the complement system is paramount. In this article, we summarize the evolving landscape of complement in the context of xenotransplantation, explore preclinical applications involving gene-edited pigs related to complement, and outline strategies for regulating complement networks to enhance the efficacy of xenotransplantation. By navigating the intricate interplay between genetic engineering and complement biology, this review aims to contribute to the ongoing dialogue regarding xenotransplantation's potential to address the growing disparity between organ supply and demand.

Three paths of complement activation

The complement system, comprising over 50 diverse proteins and cleavage molecules such as proenzymes, proteases, anaphylatoxins, receptors, regulators, opsonins, multimolecular complexes, and pattern recognition molecules that provide host defense against foreign microbes or allografts, mediate inflammatory responses and maintain normal tissue homeostasis [3, 4]. The activation of complement relies on three precisely regulated activation systems: classical pathway (CL), alternative pathway (AP), and mannose-binding lectin (MBL) pathway (Fig. 1).

Classical complement activation pathway

The classical pathway of complement activation begins with the C1 complex comprising C1q, C1r, and C1s subcomponent proteins. Initially, the spherical head of C1q recognizes IgG/IgM antigen-antibody complexes, causing the rearrangement and activation of C1r, followed by the activation of C1s within the C1r-C1s tetramer. Active C1s cleaves C4 into C4a and C4b, and subsequently, C4b binds to C2, which is cleaved by C1s into C2a and C2b. This forms the C4b2b complex, known as the classical pathway C3 convertase [5]. This convertase catalyzes the conversion of C3 to C3b and C3a. C3b molecules are deposited together, and the substrate specificity is switched to form the classical pathway C5 convertase (C3bBb3b and C3b4b2b). C5 convertase (C3bBb3b and C3b4b2b) has a high level of complement C5 affinity, which allows C5 to be dissolved and activated quickly [6, 7]. C5, the first complement factor formed by the membrane attack complex (MAC). After being lysed, C5 initiates the complex assembly that can be inserted into the target cell membrane. The assembly is formed around C5b and includes four complements—C6, C7, C8 and C9 [8].

The classical complement activation pathway is a crucial arm of the immune response, providing a rapid and potent means of neutralizing pathogens marked by antibodies.

Alternative complement pathway

The alternative complement pathway operates independently of antibodies and involves only two main components: factors B and D [9]. Factor D, an active serine protease, binds to and activates Factor B [10]. The activated Factor B then joins with C3b(H2O) to create the C3 convertase C3(H2O)Bb [11]. This convertase triggers a positive feedback loop by cleaving natural C3 molecules into C3a and C3b. Additionally, C3b binding to C3 convertase forms the C5 convertase, which cleaves C5 into biologically active fragments, C5a and C5b [12]. C5b then recruits complement components C6, C7, C8, and C9 to form the membrane attack complex (MAC). The MAC inserts into the pathogen membrane, leading to cell lysis and destruction [13].

The alternative complement pathway offers rapid and innate protection against pathogens by identifying and flagging foreign surfaces for removal. Its ongoing, low-level activity plays a crucial role in immune surveillance, greatly enhancing the effectiveness of the complement system.

Lectin pathway

The lectin pathway of the innate immune system is activated by recognizing specific carbohydrate patterns on pathogens. This activation involves pattern recognition molecules such as MBL, ficolins, or collectins binding to the pathogen's surface, initiating a cascade that includes serine protease zymogens like MASP-1, MASP-2, MASP-3, and the nonenzymatic protein MAp19, where MASP-1 and MASP-2 serving as key enzymes [14]. Upon binding to the carbohydrate ligand, the MBL-MASP complex converts MASP from a zymogen to its activated form [15]. Subsequently, MASP generates C3 convertase (C4b2a), resulting in a reaction akin to the classical pathway.

The lectin pathway aids in opsonizing pathogens for recognition and phagocytosis by immune cells. Moreover, it initiates the complement cascade, resulting in the formation of the MAC for the destruction of targeted pathogens. It serves as a vital frontline defense in the immune response against infections.



Fig. 1 Simplified overview of the active complement cascade. Foreign surface-bound antigen–antibody (Ag/Ab) complexes initiate the classic pathway, while polysaccharides, lipopolysaccharides, and/or IgA activate the alternative complement pathway. Damage-associated molecular patterns (*DAMPs*) like mannose-binding lectin (*MBL*), ficolins (*Fcns*), and certain collections (*CLs*) can directly trigger the classic pathway or initiate the lectin pathway. The three complement activation pathways collectively cleave C3 into C3b and C3a, triggering terminal pathway activation, mainly involving C5-C9, which assembles to form the membrane attack complex (MAC). Key complement regulatory factors include C1 inhibitor (C1-INH), factor H (FH), factor I (FI), CD46, C4BP, and CD59. C1-INH inhibits C1r activation of C1s, preventing C4 and C2 cleavage. Simultaneously, C1-INH can inhibit the binding of MBL to MASP-1 or MASP-2. Factor I and factor H, aided by C4BP and CD46, can phagocytize C3 from the alternative pathway, inhibiting its activation. CD59 prevents C9 from binding to C5b678 to form MAC

Complement regulatory proteins

The complement regulatory system, comprising soluble and membrane-bound factors, plays a crucial role in all three complement activation pathways. These regulators engage with specific complement components to intricately balance activation and inhibition, ensuring precise homeostasis. This balance safeguards our tissues, effectively preventing damage from foreign microorganisms.

C1 inhibitor (C1-INH)

C1-INH is a key inhibitor that regulates both the complement system and coagulation cascade. It plays a key role in controlling the classical complement pathway by inhibiting activated C1s and C1r, as well as targeting factors XIIa, kallikrein, and factor XIa in the coagulation contact system [16]. By forming a covalent complex with C1s, it prevents its activation [17]. Additionally, it hinders the activation of MASP-1 and MASP-2, inhibiting the lectin pathway [18], and interferes with the interaction between C3b and factor B, preventing alternative pathway activation. This multifaceted activity underscores the critical role of C1-INH in maintaining immune balance and preventing harmful responses [19].

FI

Factor I (FI) is a crucial serine protease in the complement system, crucial for regulating complement activation. Through enzymatic cleavage, FI indirectly inhibits the complement system by hindering the activation of complement factors C4b and C3b [20]. Working in coordination with other complement regulatory proteins, FI cleaves C3b and C4b into their inactive forms, preventing excessive complement cascade amplification and maintaining a balance in immune responses [21]. This regulatory function is essential for protecting host cells and tissues from damage caused by excessive or inappropriate complement activation.

FH

Factor H (FH) is a vital complement regulatory protein that inhibits the alternative pathway of the complement system. FH accelerates the decay of C3 convertases (C3b, Bb) of the alternative pathway and acts as a cofactor for factor I-mediated C3b cleavage and inactivation [22]. Additionally, FH directly inhibits the formation of the C3bBC3bC3b complex by interacting with C3b, properdin (factor P), and FB in the presence of FI [23]. The regulatory actions of FH are crucial for preventing excessive and inappropriate complement activation, safeguarding host cells and tissues.

C4BP

C4b-binding protein (C4BP) is a crucial regulator of the complement system, controlling the activation cascade by interacting with C4b. It serves as a cofactor for FI-mediated cleavage of C4b, leading to the formation of inactive C4b fragments [24]. By facilitating C4b degradation, C4BP inhibits the classical and lectin pathways, preventing excessive complement activation. Moreover, it disrupts the assembly of the C3 convertase (C4b2a) in the classical pathway, further regulating the complement cascade [25]. This pivotal regulatory role maintains immune balance and protects host cells and tissues from damage.

CD46

CD46, also known as membrane cofactor protein (MCP), is pivotal in regulating complement activation by promoting the proteolysis and activation of FI. This activity results in the degradation of C3b and C4b, preventing the formation and amplification of C3 and C5 convertases, and ultimately inhibiting the downstream steps of the complement cascade [26]. The regulatory function of CD46 is crucial for maintaining immune homeostasis and preventing autologous cell lysis. By controlling complement activation on host cell surfaces, CD46 helps prevent inappropriate immune responses and protects cells from damage caused by uncontrolled complement activation [27, 28].

CD59

CD59, also known as protection, plays a pivotal role in immune regulation by inhibiting the terminal complement pathway. It prevents MAC assembly by preventing C8 and C9 from joining the MAC complex [29]. This protective action ensures the preservation of host cells, preventing cell lysis and maintaining cellular integrity [30]. CD59's function as a terminal pathway inhibitor is indispensable for averting unintended cell damage and upholding immune homeostasis.

Complement and transplantation

In transplantation, the complement system serves a dual role—acting as a protective mechanism against foreign tissues while also posing a potential risk for transplant-related complications [31]. Its importance is seen in graft rejection and incompatibility. When foreign tissues are transplanted, the complement system can activate through foreign tissue pathways. Studies indicate that xenotransplant rejection is primarily mediated by the classical and alternative pathways of complement, with no significant role played by the lectin (MBL) pathway [32]. Complement proteins C3a and C5a, along with the MAC activated by these pathways, play a crucial role in lysing xenografts.

Complement activation, especially when the recipient's blood reacts against the transplanted organ, plays a key role in rejection [33]. Poor outcomes in clinical islet transplantation may be attributed to the occurrence of a destructive instant blood-mediated inflammatory response (IBMIR) [34]. Complement activation is a vital component of IBMIR, triggered after a thrombotic reaction. During this phase, pancreatic islets exposed to blood in the portal vein undergo a direct assault by the complement system [35, 36], primarily due to the extensive binding of antibodies against C4 and C3 on the surface of transplanted pancreatic islet cells [37].

Hyperacute allograft rejection (HAR) and ischemia– reperfusion injury (IRI) in xenografts are central factors contributing to heterogeneous transplant failures, and the complement system plays a pivotal role in both HAR and IRI. HAR, characterized by rapid rejection within 48 h post-transplantation, results from preexisting cytotoxic antibodies in the recipient binding to graft antigens, leading to severe complement-dependent rejection [38, 39]. IRI is frequently encountered in kidney transplantation, often attributed to blood flow disorders. The complement component C3 plays a pivotal role in inflammatory processes, with its elevation worsening IRI-induced acute kidney injury and stimulating the production of secondary epithelial cell chemokines, thereby contributing to local inflammation [40, 41].

Understanding and managing complement activation in transplantation is vital for enhancing success rates and the long-term functionality of transplanted organs. Therapeutic complement inhibitors effectively protect organs from inflammatory damages [42], and ongoing research is exploring genetic engineering strategies in donor pigs with human complement regulatory proteins (hCRP) to minimize the impact of complement system activation on xenograft survival [43]. These studies aim to develop new complement inhibition and immunomodulation strategies, enhancing transplantation outcomes.

Gene-edited pigs

Pigs are considered excellent candidates for xenotransplantation due to their genetic, physiological, metabolic, and anatomical similarities to humans. They can be easily bred and raised in controlled environments, providing organs of suitable size for human transplantation [44, 45]. However, molecular incompatibility between pig donors and human recipients often leads to immune complications and xenotransplant rejection [46]. Technologies such as zinc finger nucleases [47], transcription activator-like effector (TALE) nucleases [48], and CRISPR/Cas [49–51] have enabled the efficient editing of pig genomes.

Deleting xenoreactivity antigens

In pig-to-primate xenotransplantation, a major challenge is hyperacute rejection (HAR) occurring shortly after transplantation [52]. HAR is primarily due to preexisting antibodies in human plasma targeting the Gala (1,3)-Gal antigen on porcine endothelial cells (Fig. 2). This antigen is absent in humans and higher primates. Over 80% of complement-fixing xenoreactivity antibodies in human serum recognize Gala epitopes [53]. When xenoreactivity



Fig. 2 Xenograft activates the complement system. The binding of IgG/IgM to the protein α -Gal on the graft surface activates the classical pathway of complement, while the interaction of IgA with α -Gal activates the alternative pathway. The combined activation of classical and alternative pathways leads to the generation of C5 convertase, ultimately resulting in the formation of the membrane attack complex (MAC) composed of C5b, C6, C7, C8, and C9. This MAC complex functions to attack the xenograft

antibodies recognize and bind to Gala (1,3)-Gal antigen, the classical complement pathway is activated, leading to HAR induction. Therefore, preventing HAR requires strategies to neutralize the impact of anti-Gal antibodies or complement.

To make pig organs suitable for xenotransplantation, it is essential to eliminate the Gala (1,3) Gal antigen from xenograft cell surfaces. One approach involves deactivating the GGTA1 gene, responsible for forming the Gala (1,3) Gal epitope. In a study by Lai et al. [54], nuclear transfer successfully knocked out α -1,3-galactosyltransferase, producing pigs with heterozygous GGTA1 inactivation. Co-expressing human α 1,2-fucosyltransferase and α -galactosidase significantly reduced Gal antigen levels on cell surfaces, thereby decreasing xenotransplant immunogenicity [55] (Fig. 2). Heterotopic heart transplantation from α -1,3-galactosyltransferase knockout pigs into baboons eliminated the galactose- α -1,3-galactose epitope, preventing HAR and extending porcine heart survival in baboons for 2-6 months. Additionally, homozygous α-1,3-galactosyltransferase knockout pigs were produced through breeding and somatic cell nuclear transfer (SCNT) [56].

Expression of human complement regulatory proteins

To mitigate complement-mediated graft injury in xenotransplantation, genetically modifying pigs to express human complement regulatory proteins (hCRPs) like CD46 (membrane cofactor protein, MCP), CD55 (decay accelerating factor, DAF), and CD59 (membrane inhibitor of reactive lysis, MIRL) is an effective strategy [57, 58]. These hCRPs play a crucial role in maintaining the balance between complement activation and inhibition, acting as inhibitors at all stages of complement activation. The creation of multitransgenic pigs expressing CD46, CD55, and/or CD59 indicates that simultaneous expression of multiple hCRPs enhances protection [59]. Successful kidney transplantation from pigs expressing both hCD55 and hCD59 into nonimmunosuppressed baboons demonstrated their protective effect against hyperacute rejection (HAR) without immunosuppression [<mark>60</mark>].

Transplantation GTKO porcine hearts and kidneys significantly extend transplant survival [61, 62]. Compared to GTKO or CRP alone, incorporating hCRPs into GTKO pigs further reduces antibody-mediated rejections [63, 64]. Human CD55 expression effectively blocks HAR and limits local complement activation in GTKO heart transplantation [64]. Additionally, GTKO combined with human CD46 transgenic (GKO/CD46) islets enhances xenograft survival by mitigating early platelet deposition and neutrophil infiltration [65]. These findings collectively suggest that GGTA1 knockout pigs with one or two hCRPs are more suitable donors for organ xenotransplantation.

Gene-edited pig organ transplantation model

Ongoing advancements in gene editing technology, paired with continual enhancements in immunosuppressive regimens, have significantly propelled the evolution of pig-to-non-human primate (NHP) organ transplant models. Pigs now serve as vital organ donors in the field of xenotransplantation (Fig. 3).

Heart transplant

Most porcine heart transplant studies involve heterotopic transplantation [66, 67]. Until 2005, the longest median survival time for porcine hearts transplanted heterotopically into baboons was 96 days [68]. With an immunosuppressive regimen comprising aTG, anti-CD154, and MMF, effective B cell depletion with the anti-CD20 antibody extended the survival time of heterotopic cardiac transplants to 236 days. However, delayed rejection eventually led to graft failure [69]. In 2016, the same researchers utilized GTKO/hCD46/hTBM donor porcine hearts for transplantation into baboons. Implementing an aCD40 antibody-based immunomodulatory regimen (2C10R4), the longest survival time for heterotopic heart transplantation was extended to an impressive 945 days, with a median survival time of 298 days [70]. Non-human primate studies show that both ectopic and orthotopic heart transplants maintain function and prolong survival. This is attributed to sustained cardioplegic solutions during hypothermic ischemia and an effective immunosuppressive regimen [71]. These preclinical findings pave the way for successful xenotransplantation of genetically modified porcine hearts into patients with end-stage heart failure.

In recent years, several international teams have initiated preclinical studies on gene-edited pig heart transplantation. On January 7, 2022, surgeons at the University of Maryland successfully performed the world's first transgenic pig heart transplant on a 57-year-old man. Despite the patient's death 60 days after surgery, this procedure marked a groundbreaking moment in xenotransplantation history [72]. Notably, it overcame potential postoperative obstacles such as hyperacute immune rejection, achieved favorable short-term outcomes, and emphasized the necessity for further clinical research. However, viral safety was overlooked during this xenotransplantation process, leading to the transmission of pig viruses (PCMV/PRV) to human recipients, resulting in patient fatalities [73]. In July 2023, Nader Moazami et al. transplanted hearts from 10 geneedited pigs into two brain-dead human recipients and



Fig. 3 Gene-edited pig & xenotransplantation. Organs cultivated from genetically modified cloned pigs, such as heart, liver, lung, kidney, etc., can be transplanted into patients

meticulously monitored transplant function, pre-existing xenoreactivity antibody injury, hemodynamics, and systemic reactions within 66 h. They found no safety risks in the deceased recipients [74]. While progress has been made, it's clear we're not fully prepared for human xenotransplantation.

Kidney transplant

The utilization of kidneys in pig-to-NHP models has progressed more slowly compared to heart transplants. Wild-type pig kidneys typically fail in NHPs within a few minutes [75]. In 2004, the transplantation of CD55 pig kidneys into a cynomolgus monkey model revealed that the average survival of pig renal xenotransplants was limited to several weeks, with 90 days being the longest reported survival in a pig-to-NHP model [76].

Until Higginbotham et al.'s breakthrough in 2015, progress in pig xenotransplantation had been limited. In their study, GGTA1 KO/hCD55 pig kidneys were transplanted into rhesus monkeys. After T cell exhaustion, the treatment was sustained by stimuli blocking in addition to the maintenance of mycophenolate mofetil and steroids. Remarkably, the graft survived for an impressive 310 days [77]. Subsequent experiments with gene-edited pigs have shown promising results. Iwase et al. conducted kidney transplants from GTKO/CD46/CD55/TBM/EPCR/CD39 pigs to baboons treated with an anti-CD40mAb-based regimen, with kidney function enduring for 136 days [78]. Kim et al. achieved the longest reported life-sustaining xenotransplantation, lasting 499 days, by transplanting GGTA1KO/hCD55 pig kidneys into rhesus monkeys with low anti-pig antibody titers and selective depletion of CD4+ or CD8+T cells, emphasizing the significance of CD4+T cell depletion [79].

Before the first global porcine heart transplantation to humans, three instances of genetically modified pig kidney transplants into humans occurred. Two recipients received genetically modified GTKO porcine kidneys, maintaining normal renal functions for 54 h without signs of HAR or antibody-mediated rejection [80]. In the third case, the donor kidneys (bilateral transplantation) had ten genetic modifications, and three pig carbohydrate antigens and the pig growth hormone receptor gene were deleted. These gene-modified kidneys functioned for 74 h without rejection or antibody or complement protein deposition [81].

In summary, the added expression of several hCRPs in GTKO transgenic pigs can further prevent rejection

development [82]. With continuous scientific efforts, lifesustaining kidney xenotransplantation is much closer to clinical reality than previously thought.

Liver transplant

Liver xenotransplantation from pigs to humans faces considerable hurdles compared to heart and kidney transplants, including immunological complexities, clotting abnormalities, and rejection risks. Ekser et al. pioneered the transplantation of GTKO or GTKO/hCD46 pig livers into baboons, resulting in severe thrombocytopenia with a survival of 7 days [83, 84]. In 2012, Kim transplanted GTKO livers into baboons and achieved up to 9 days of porcine liver xenotransplantation under the same immunosuppression regimen as xenotransplantation of heart and kidney [85]. The Shah team explored the effects of exogenous administration of human coagulation factors after pig-to-baboon liver xenotransplantation (LXT) using GTKO pig donors. The addition of costimulatory blockade to this regimen increased individual recipients' LXT survival from 9 to 25 days [86, 87]. In their modified experimental protocol, costimulation blockade (belatacept or anti-CD40mAb) extended the 25 day survival period to 29 days [88]. The outlook for pig-to-NHP liver transplantation depends on ongoing advancements in genetic engineering, immunosuppressive protocols, and a deeper understanding of the immunological and physiological factors involved in xenotransplantation.

Lung transplant

Xenotransplantation faces heightened challenges in lung transplantation due to the lung's sensitivity to injury and multiple immune rejection mechanisms. During transplantation, the pig lung is the organ most severely damaged due to rapid coagulation dysfunction [89]. While progress has been made in heart and kidney transplantation with GTKO and hCD46, lung xenografts still face challenges. In 2007, Nguyen et al. transplanted GTKO left lungs into baboons, but these lungs could only sustain life for 3.5 h due to severe coagulation disorders [90]. Laird et al. discovered that transgenic expression of human leukocyte antigen-E attenuated GTKO/hCD46 pig lung xenograft injury, prolonging survival ex vivo [91]. Watanabe et al. proposed that transgene expression of hCD47 on porcine blood vessels could alleviate acute vascular rejection in baboons, particularly in GTKO pig lungs that are highly susceptible [92]. Utilizing multigene donor pigs, combined with targeted complement activation (hCD46, hCD55), coagulation (hEPCR, hVWF, hTBM, hTFPI, hCD39), and anti-inflammatory pathway regulatory genes (HO-1, HLA-E), significantly improved the survival of xenogeneic swine lungs in both ex vivo human blood perfusion and in life-supporting in vivo models [93]. However, the survival rate of lung xenotransplantation remains measured in days rather than weeks or months [94].

Islet transplantation

The primary challenges in the early stages of porcine islet transplantation are HAR and IBMIR [95]. When the NHP immune system detects Gala (1,3) in porcine tissue, it triggers the classical complement pathway, resulting in the formation of membrane attack complexes and cell lysis. Thus, knockdown of the islet surface α -Gal epitope is therefore a logical choice [96]. In addition to removing the α -Gal antigen, reducing IBMIR also requires the expression of human complement regulatory factors (hCD46, hCD55, hCD59) [97]. When porcine pancreatic islets are transplanted into non-human primates (NHPs) with diabetes, a significant portion of the graft is typically lost early due to IBMIR and intense immunosuppressive therapy [98]. Windt et al. addressed complement activation by expressing hCD46 on pig islets, effectively limiting antibody-mediated rejection and preserving islet quality. While this approach didn't prevent the immediate loss of most transplanted islets, it significantly improved the outcomes of islet xenotransplantation in diabetic cynomolgus monkeys, maintaining normoglycemia for over 12 months [99, 100]. Hawthorne et al. achieved minimal IBMIR when transplanting α -Gal-deficient pigs with hCD55 and hCD59 transgenes onto neonatal islet cell clusters, combined with a clinically relevant immunosuppressive protocol [101]. Achieving clinical long-term survival of pancreatic islets may require more effective immunosuppression or further modification of donor genes.

Corneal transplantation

Porcine corneal xenotransplantation is considered feasible due to the cornea's immune privilege and avascular nature [102]. Recent nonhuman primate studies have shown promising results, indicating that porcine xenografts can endure for an extended period during corneal transplantation [103]. Dong et al. discovered that corneas from GTKO/CD46 pigs did not significantly improve graft survival compared to those from wild-type pigs. Prolonging the survival of corneal xenografts in pig-tomonkey corneal xenotransplantation encounters challenges in preventing anterior synechiae and retrocorneal membrane formation [104]. The use of decellularized corneas from wild-type pigs for anterior lamellar keratoplasty has shown graft transparency for over a year [105]. In treating corneal fungal ulcers and other clinical diseases, Zhang et al. reported that implantation of acellular porcine corneal stromata (APCS) resulted in no recurrence of infection during a 6 month follow-up



Fig. 4 Preclinical and clinical trials of complement therapy. Complement therapy progresses through different stages, spanning from preclinical work to market authorization. These stages include laboratory research, animal models, clinical phases I, II, and III, and final clinical implementation. The therapeutic goals are categorized into four quadrants, representing major complement categories: anaphylatoxins, active pathways, amplification and terminal pathways, and effectors. Each arrow denotes a specific agent and its development stage. Drugs targeting C1r/s and MASP include Cinryze (Shire), Berinert (CSL Behring), Cetor (Sanquin), and Ruconest (Pharming), which are already being used in clinics; drugs targeting C1q include ANX005 (Annexon); drugs targeting C1s include TNT003 (True North), TNT009 (True North), and BIVV020 (Sanofi); TP10 (CDX-1135; Celldex Therapeutics) targets the soluble form of complement receptor type 1 (CR1); OMS721 (Narsoplimab, Omeros) targets the MASP-2 target; Drugs targeting C3 include OMS906 (Omeros); drugs targeting Properdin include CLG561 (Novartis) and NM9401 (Novelmed); drugs targeting C3 include AMY-101 (Amyndas), APL-1 (Apellis), APL-2 (Apellis), CB2782 (Catalyst), Cp40 (Amyndas); drugs targeting C3b and convertases include AMY-201 (Amyndas), and Mirococept (MRC); drugs targeting FB include Bikaciomab (Novelmed); drugs targeting FD include Lampalizumab (Genentech), ACH-4471 (Achillion), and "Compound 6" (Novartis); drugs targeting C5 include Eculizumab (Soliris, Alexion), ALXN1210 (Alexion), ALXN5500 (Alexion), LFG316 (Novartis), Coversin (Akari), RA101495 (Ra Pharma), ALN-CC5 (Alnylam), RA101348 (RaPharma), ARC 1905 (Zimura; Ophthotech), and the affibody SOBI002 (Swedish, Orphan Biovitrum) targets C5 (programme recently terminated); drugs targeting C5a include IFX-1 (InflaRx), ALXN-1007 (Alexion), NOX-D21 (Noxxon Pharma); drugs targeting C5a Rinclude CCX168 (Chemocentryx); C6 target drug includes Regenemab (Regenesance); CR2–FH target drug includes TT30 (ALXN 1102; Alexi

period, and all corneal ulcers healed with neovascularization. APCS grafts proved safe and efficacious in lamellar keratoplasty for various clinical conditions [106]. As genetically engineered pigs become more available, pig corneas have the potential to address the global shortage of corneas soon.

Research and application of complement-related drugs

Advances in comprehending the complement system's composition, structure, and interactions have paved the way for developing drugs with both stimulating and inhibitory effects on complement activity. These medications show potential as therapies for a wide range of conditions, including infectious, inflammatory, traumatic,

cancerous, autoimmune, or age-related diseases, as well as for preventing transplant rejection.

Four C1-INH drugs have been approved [107]: Cinryze, Berinert, Ruconest, and Cetor. C1-INH inhibits the kinin B1 receptor, reduces the release of chemotactic microvesicles from damaged donor tissues, and effectively prevents late antibody-mediated rejection. C1-INH has a longstanding history of treating hereditary angioedema (HAE) with high safety and effectiveness [108]. While data on its use in organ transplantation are limited, experimental evidence suggests potential benefits in alleviating acute antibody-mediated rejection (ABMR) observed in baboons during transplantation [109]. A case report on Cinryze's use in treating acute ABMR postkidney transplantation indicates promising outcomes [110]. Berinert, according to data from a phase 1/2 study involving 20 patients, may enhance allograft function in kidney recipients with unresponsive acute ABMR [111]. However, there is still limited research on Ruconest and Cetor in organ transplantation.

Inhibiting the complement system at the C3 level effectively prevents unregulated activation, safeguarding host cells from damage. Cobra venom factor (CVF), an analog of complement component C3, exhibits specific anticomplement C3 biological activity. CVF induces lysis of C3 and C5, depleting complement C3 (and C5), inhibiting humoral immunity, and has found widespread use in xenotransplantation due to its long-term complement removal effects [112, 113]. Cp40, an analog of compstatin, is a potent inhibitor of complement C3. It effectively prevents C3 activation and mitigates complement-mediated injury triggered by endothelial antibody binding and extracorporeal circulation [114]. Cp40 demonstrates the ability to inhibit complement activation, promote antiinflammatory and anticoagulant effects in septic animals [115]. Notably, Cp40 can prevent the adhesion of leukocytes, specifically neutrophils, to porcine endothelium [116]. Considering these findings, Cp40 stands out as a promising adjunct for preclinical and future clinical cardiac xenotransplantation [117].

Several interventions are currently being explored to prevent xenograft injury and improve its survival rate. Eculizumab, a recombinant antibody targeting complement C5, holds the potential for reducing antibodymediated rejection [118]. As the first anti-complement drug, eculizumab offers a novel therapeutic approach for various human diseases, reshaping treatment strategies for conditions like PNH and significantly impacting their clinical outcomes [119, 120].

Identifying suitable complement inhibitors and defining therapeutic strategies is crucial for future studies. Genetically engineering pigs with appropriate human complement modulators emerges as a promising strategy in xenotransplantation [121]. Figure 4 outlines development programs focusing on inhibitors against various complement targets, with some undergoing clinical studies in both healthy individuals and patients [122–125].

Conclusions and perspective

The global shortage of organs for transplantation has led to the investigation of xenotransplantation as a potential solution. This method, which involves transplanting organs from genetically modified pigs into humans, offers hope in alleviating the scarcity of human organ donors. Research indicates that graft failure often stems from the activation of the complement system, affecting critical aspects of xenografts, including galactosidase binding, antibody interactions, and complex responses involving coagulation, inflammation, and adaptive immune reactions during transplantation.

The recent breakthroughs in gene-edited porcine heart transplantation represent significant progress and offer valuable insights for refining this approach [73]. However, the presence of pig viruses in gene-edited pig xenotransplantation cannot be overlooked [126]. Despite this, these successes not only shed light on the long-term viability of cardiac xenotransplants but also lay a foundation for transplanting other organs into humans. The lessons learned from porcine heart transplantation serve as a crucial reference for optimizing xenotransplantation, which involves strategically using genetically modified organs to evade the human immune response. The implementation of advanced gene-editing techniques, including CRISPR/Cas9, TALEN, and SCNT, in modifying potential pig donors has led to substantial advancements in xenotransplantation. Genetically engineered pig organs, combined with novel immunosuppressive therapies, have extended the survival rates in NHP xenotransplants.

The ongoing development of complement-related clinical drug candidates provides a diverse array of options for selective inhibition, targeting, and drug delivery, contributing to the progress of xenotransplantation. Despite facing immunobiological challenges, the increasing variety of genetically modified pigs and the expansion of immunosuppressant and anti-inflammatory drugs offer optimism. Clinical trials for pig kidney, heart, liver, lung, pancreatic islet, and corneal transplantation are anticipated, bringing animal organ transplantation into reality for human recipients shortly.

Abbreviations

ABMR	Antibody-mediated rejection
AP	Alternative pathway
APCS	Acellular porcine corneal stromata
C1-INH	C1 inhibitor
C4BP	C4b-binding protein
CL	Classical pathway

CVF	Cobra venom factor
EPCR	Endothelial protein C receptor
FI	Factor I
FH	Factor H
GHR	Growth hormone receptor
HAE	Hereditary angioedema
HAR	Hyperacute allograft rejection
hCRP	Human complement regulatory protein
hHO-1	Heme oxygenase-1
HLA	Human leukocyte antigen
IBMIR	Instant blood-mediated inflammatory response
IRI	lschemia–reperfusion injury
LXT	Liver xenotransplantation
MAC	Membrane attack complex
MAP	Membrane attack protein
MBL	Mannose binding lectin
MCP	Membrane cofactor protein
MIRL	Membrane inhibitor of reactive cleavage
NHP	Non-human primate
PERVs	Porcine endogenous retroviruses
SCNT	Somatic cell nuclear transfer
TALE	Transcription activator-like effector

Acknowledgements

Not applicable.

Author contributions

All the authors contributed to the manuscript and reviewed and approved it as presented here. YLY, GPZ established the review idea and the article structure. YLY, YYC, and DYZ drafted the article and designed the figure. YY, YSZ, DNL, and XMJ aided in revising the manuscript. GPZ improved and supervised the submitted and revised manuscript. All authors read and approved the final manuscript.

Funding

This work was supported by funding from the National Natural Science Foundation of China (8200062054, 81771723), the 68th general funded project of the China Postdoctoral Science Foundation (2020M683295), medical and industrial integration funded project of the University of Electronic Science and Technology of China (ZYGX2021YGLH024), the Department of Science and Technology of Sichuan Province (2022YFS0157), and the Department of Science and Technology of Sichuan Province (2021YFS0375).

Availability of data and materials

Not applicable.

Declarations

Ethics approval and consent to participate Not applicable.

Consent for publication

Not applicable.

Competing interests

All authors declare no competing interests related to this review.

Received: 26 December 2023 Accepted: 27 March 2024 Published online: 02 April 2024

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