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Role of donor macrophages after heart and lung transplantation

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Abstract

Since the 1960s, heart and lung transplantation has remained the optimal therapy for patients with end-stage disease, extending and improving quality of life for thousands of individuals annually. Expanding donor organ availability and immunologic compatibility is a priority to help meet the clinical demand for organ transplant. While effective, current immunosuppression is imperfect as it lacks specificity and imposes unintended adverse effects such as opportunistic infections and malignancy that limit the health and longevity of transplant recipients. In this review, we focus on donor macrophages as a new target to achieve allograft tolerance. Donor organ-directed therapies have the potential to improve allograft survival while minimizing patient harm related to global suppression of recipient immune responses. Topics highlighted include the role of ontogenically distinct donor macrophage populations in ischemia–reperfusion injury and rejection, including their interaction with allograft-infiltrating recipient immune cells and potential therapeutic approaches. Ultimately, a better understanding of how donor intrinsic immunity influences allograft acceptance and survival will provide new opportunities to improve the outcomes of transplant recipients.

Keywords

basic (laboratory) research/science; heart (allograft) function/dysfunction; heart disease: immune/inflammatory; heart transplantation/cardiology; immunosuppressant; immunosuppression/immune modulation; lung (allograft) function/dysfunction; lung disease: immune/inflammatory; translational research/science

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DATA AVAILABILITY STATEMENT

The data that support the findings of this study are available from the corresponding author upon reasonable request.

1 | INTRODUCTION

Organ transplant continues to represent the optimal therapy for individuals with end-stage organ failure since the first lung transplant in 1963 and heart transplant in 1967. Given increased demand for organ transplant and enhanced regulatory scrutiny, waitlist times have increased¹ out of proportion to donor availability.² Expanding the donor organ pool and enhancing immunologic compatibility have become priorities within the transplant community. In addition, limiting posttransplant complications, including ischemia–reperfusion injury (IRI), primary graft dysfunction, acute cellular rejection (ACR), antibody-mediated rejection (AMR), graft vasculopathy, and chronic airway rejection, remains an area of intense focus.

Current immunosuppression strategies globally suppress recipient immune responses and as a direct result lead to many complications (renal failure, infections, malignancy). Balancing allograft protection against risks of infection and malignancy is a problematic and challenging issue. These observations highlight the clinical need to explore alternative options to reduce rejection and improve allograft survival.

Recent paradigm shifting studies have uncovered that macrophages represent heterogeneous populations composed of distinct cell types with differing developmental origins, recruitment dynamics, and functions. Several timely studies have investigated the role of macrophages in solid organ transplant.^{3–5} Macrophages have dichotomous roles in inducing allograft injury and promoting allograft survival.⁵ These contrasting functions highlight the complex and dynamic functions of monocytes and macrophages. Macrophages within the heart and lung can be initially divided into tissue resident macrophages and infiltrating monocyte-derived macrophages.^{6–8} In the context of transplant, this subclassification is particularly informative as tissue resident macrophages and monocyte-derived macrophages originate from the donor and recipient, respectively. In the following sections, we provide a comprehensive discussion of tissue resident macrophage composition within the heart and lung, review the functions of tissue resident macrophages, and highlight known and presumptive roles for tissue resident macrophages in allograft rejection and tolerance.

2 | HEART AND LUNG MACROPHAGE POPULATIONS ARE DIVERSE WITH DIVERGENT FUNCTIONS

Macrophages are an essential component of the innate immune system. Although there are many approaches to categorize macrophage subtypes, division into tissue resident and infiltrating monocyte-derived populations is widely accepted and highly applicable to the transplant field. Tissue resident macrophages typically seed organs during embryonic or early postnatal development and exist throughout life largely independent of monocyte input.^{6,7,9} Many tissue resident macrophage populations (heart, lung, liver, kidney, brain) are derived from early hematopoietic progenitors located within the yolk sac and/or fetal liver.^{10–12} However, some tissue resident macrophages originate from peripheral monocytes (stomach, intestine, colon) and are later maintained through local proliferation.^{13,14} A general theme from studies of macrophage ontogeny is that tissue resident macrophages acquire unique functions based on the organ in which they reside.¹⁵ Thus, macrophage

phenotype is influenced by both ontogeny and environmental cues. Current work in this space is focused on delineating the mechanistic basis by which developmental and environmental information is integrated at the signaling, transcriptional, and epigenetic levels.

The heart contains distinct macrophage populations with divergent origins and functions.^{16–18} Cardiac tissue resident macrophages are readily distinguished based on the expression of C-C chemokine receptor 2 (CCR2).^{6,9} CCR2A and CCR2B are widely conserved among mammals^{17,19,20} and are expressed on the cell surface of monocytes, dendritic cells, and some T cells,²¹ and their primary ligands are CCL2 and CCL7.²² Under resting conditions, the adult heart contains CCR2⁺MHC-II^{hi} macrophages, CCR2⁻MHC-II^{lo} macrophages, and CCR2⁻MHC-II^{hi} macrophages. Monocytes are differentiated from cardiac macrophages because they are CCR2⁺MHC-II^{lo} and lack the expression of MertK.⁷

CCR2⁻ macrophages are derived from embryonic hematopoietic progenitors, seed the heart during fetal development, are long lived, and are maintained independent of monocyte input throughout life via local proliferation. CCR2⁻ macrophages promote coronary angiogenesis and cardiomyocyte proliferation and have anti-inflammatory effects potentially by secreting interleukin (IL)-10 and transforming growth factor (TGF)- β .^{7,18,23–25} CCR2⁻ macrophages suppress neutrophil and monocyte recruitment. CCR2⁺ macrophages are derived from circulating monocytes, seed the heart during postnatal life, and are maintained through a combination of gradual monocyte recruitment and local proliferation. CCR2⁺ macrophages are enriched in proinflammatory genes, and their activation represents a mechanism driving inflammation.^{18,24} CCR2⁺ macrophages orchestrate neutrophil and monocyte recruitment. Removal of CCR2⁺ macrophages is sufficient to reduce infarct area and adverse cardiac remodeling following myocardial infarction.²⁶ Recently, it was discovered that the human heart contains macrophage populations that are developmentally and functionally analogous to CCR2⁻ and CCR2⁺ macrophages found in the mouse heart (Table 1).¹⁶

The lung contains 2 distinct macrophage populations: alveolar and interstitial macrophages.^{27,28} They constitute 80% and 20% of the lung resident macrophage pool, respectively. Alveolar macrophages line the surface of alveoli and are long-lived lung-resident cells. Alveolar macrophages are derived from embryonic hematopoietic progenitors located predominantly within the fetal liver.²⁹ However, recent studies show that alveolar macrophages can also be derived from circulating bone marrow–derived monocytes, which can play a role in the development of lung fibrosis.³⁰ Interstitial macrophages are localized in the narrow space between the alveolar epithelium and vascular endothelium, largely originate from circulating blood monocytes, and are maintained through monocyte turnover.²⁷ Some investigators have suggested that a small population of interstitial macrophages may be derived from yolk sac hematopoietic progenitors (Table 2).³¹

Lung classical monocytes survey the parenchyma and are recruited to sites of inflammation by CCL2, CX3CL1, and CCR5 ligands.^{32,33} Classical monocytes can differentiate into interstitial macrophages, alveolar macrophages, and dendritic cells. Nonclassical monocytes patrol the endothelium in a crawling-type motility and are recruited to sites of inflammation by CX3CL1.^{32,34,35}

3 | ALLOGRAFT MACROPHAGE POPULATIONS FOLLOWING HEART AND LUNG TRANSPLANT

The transplanted heart contains a compilation of myeloid cells consisting of “donor-derived” CCR2⁺ and CCR2⁻ macrophages and infiltrating recipient neutrophils, monocytes, monocyte-derived macrophages, and monocyte-derived dendritic cells.^{16,36,37} Classical recipient monocytes derive from blood monocytes and differentiate into macrophages and dendritic cells and contribute to both rejection and tolerance.³⁸ In murine models, we can separate donor from recipient cells using cell-tracing strategies such as congenic (i.e., CD45.1/CD45.2) or reporter (i.e., GFP/RFP) mice. It is also possible to distinguish donor and recipient immune cells in humans by analyzing sex-mismatched transplant recipients or allelic HLA expression. For example, we analyzed endomyocardial biopsy samples (average 8.8 years after transplant, no active rejection, normal allograft function) involving a female donor and a male recipient. Donor-versus-recipient origin was based on the presence or absence of a “Y” chromosome.¹⁶ Combined in situ hybridization and immunostaining revealed the presence of a Y chromosome in a very small percentage of CCR2⁻ macrophages (<2%). In contrast, a Y chromosome was detected in 30% of CCR2⁺ macrophages, suggesting that cardiac CCR2⁺ macrophages represent a compilation of donor- and recipient-derived cells. Little is known regarding whether donor- and recipient-derived CCR2⁺ macrophages are functionally distinct in humans. However, mouse studies suggest that recipient-derived CCR2⁺ macrophages express higher levels of inflammatory chemokines (CXCL11, CXCL2, CCL2, CCL7, CCL19), cytokines (IL1- β , IL-10), and adverse cardiac remodeling genes (*AREG*, *EREG*, *GDF3*) compared with donor-derived CCR2⁺ macrophages. Donor-derived CCR2⁺ macrophages expressed higher levels of type I interferon responsive genes.²⁶

After lung transplant, the vast majority of alveolar macrophages in the allograft are donor derived.^{39,40} Donor-derived alveolar macrophages were positive for *Ki67*, suggesting they have the capacity to proliferate locally. Although donor-derived alveolar macrophages are the predominant macrophage for at least 2 to 3 years after transplant, the contribution of recipient monocyte recruitment is less well understood. Conflicting data exist regarding whether monocytes contribute to alveolar macrophages in the lung allograft over extended transplant durations.^{39,40} Collectively, these observations indicate that macrophages resident within the donor heart and lung exist within the graft for extended periods of time and bring to light the possibility that donor macrophages constitute unique and functionally important cell types that likely contribute to allograft health and longevity (Figure 1).

4 | ROLE OF DONOR MACROPHAGES IN IRI AND PRIMARY GRAFT DYSFUNCTION

A critical goal during organ transplant is to minimize ischemic time. Longer ischemic times are associated with poor clinical outcomes after heart^{41,42} and lung transplant.^{43,44} IRI is thought to be the predominant mechanism of primary graft dysfunction, a major cause of early allograft loss and mortality.^{45,46} Recently, the United Network for Organ Sharing modified the heart and lung allocation system to promote broader organ sharing and increase

allocation to sicker individuals. Although comprehensive data are not yet available, it is clear that the new allocation system prolongs allograft ischemic time due to increased distances between donor procurement sites and transplant centers. As such, there is growing interest to improve our understanding of the mechanisms that contribute to allograft IRI.

Macrophages have been shown to have dynamic roles on allograft function after IRI in kidney and liver transplant models.^{47–49} Distinct macrophage populations contribute to early proinflammatory responses⁵⁰ and later postinjury resolution.⁵¹ The precise identity of these macrophage subsets and mechanisms that orchestrate their activation and effector responses in the context of allograft IRI remain incompletely defined. Improved understanding of these topics could have profound clinical impact in this new era of more widespread organ sharing, as mitigating the effects of prolonged ischemic time may improve transplant outcomes and further increase the donor pool.

5 | IRI AFTER HEART TRANSPLANT

Incorporation of cold ischemic time into mouse heart transplant models allows investigators to dissect mechanisms that contribute to allograft IRI. These studies have observed evidence of cardiomyocyte cell death and infiltration of recipient neutrophils, monocytes, and monocyte-derived macrophages into the donor heart.²⁶ Intriguingly, macrophage populations resident within the donor heart differentially orchestrate graft inflammation following IRI. Depletion of tissue resident CCR2⁺ from the donor heart leads to reduced recipient neutrophil extravasation and monocyte recruitment into the donor heart following transplant. Within this context, donor CCR2⁺ macrophages governed recipient leukocyte recruitment through the generation of neutrophil and monocyte chemokines through a Toll-like receptor (TLR)9- and MYD88-dependent mechanism.^{26,52} Conversely, depletion of tissue resident CCR2⁻ macrophages from the donor heart resulted in a marked increase in recipient neutrophil, monocyte, and monocyte-derived macrophage recruitment. Intriguingly, single-cell mRNA sequencing uncovered remarkable diversity among infiltrating monocytes and monocyte-derived macrophages and revealed that donor macrophages play pivotal roles in monocyte fate decisions.²⁶ These findings implicate the importance of donor macrophages in allograft inflammation after IRI. In theory, pretreating donor macrophages before procurement may ameliorate IRI-associated allograft injury and reduce the incidence and severity of primary graft dysfunction. Consistent with this concept, studies assessing the feasibility of treating the procured donor heart have shown a significant role for macrophages on graft survival.^{53,54}

Release of damage-associated molecular patterns (DAMPs) and alarmins from injured and/or dying cells is thought to constitute the signal mechanistically linking IRI to immune cell activation. Necrosis, necroptosis, and ferroptosis are among the various forms of cell death implicated in DAMP and alarmin release. Although necroptosis has previously been implicated in allograft immune cell activation,⁵⁵ increasing evidence suggests that ferroptosis is responsible for the initial wave of cardiomyocyte and cardiac fibroblast cell death after IRI. The treatment of donor hearts with ferrostatin-1 (specific inhibitor of ferroptosis) is sufficient to dramatically reduce initial cardiomyocyte and fibroblast cell death and suppress early neutrophil graft infiltration. Inhibition of necroptosis had minimal

effect on cardiomyocyte cell death and neutrophil extravasation within 4 hours after transplant.⁵⁶ The exact identity of DAMPs/alarmins released by ferroptotic cardiomyocytes and fibroblasts remains to be elucidated. However, it is likely that these mediators engage TLRs. The deletion of TLR4 in donor endothelial cells was sufficient to prevent adhesion of neutrophils to venous endothelial cells, a critical upstream step in neutrophil extravasation. Mechanistically, TLR4 activation was transduced through a TRIF and type I interferon-dependent pathway.⁵⁶ Deletion of TLR9 and MYD88 in donor macrophages did not affect the initial adhesion of neutrophils to endothelial cells but instead resulted in impaired transendothelial migration.⁵² These findings highlight the possibility that DAMPs/alarmins released by dying myocardial cells initiate allograft inflammation via signaling to multiple donor resident cell types, including endothelial cells and macrophages. Future studies will be required to further delineate additional cell types and signaling mechanisms governing IRI-induced allograft inflammation and resultant allograft dysfunction.

6 | POSTTR ANSPL ANT ISCHEMIC LUNG INJURY

IRI causes severe graft dysfunction in up to 20% of lung transplant recipients.⁵⁷ IRI after lung transplant has been described as a biphasic process in which pulmonary macrophages play an important role. Donor macrophages were suggested to induce an initial response (independent of recipient neutrophils), and circulating leukocytes were thought to mediate downstream events.⁵⁸

In a rat lung transplant model, alveolar macrophages contribute to local inflammation via induction of cytokines (e.g., TNF- α).^{59–61} These findings were corroborated in a mouse model of pulmonary IRI.⁶¹ Consistent with a role for donor macrophages in generating inflammatory responses, the injection of gadolinium chloride (inhibitor of macrophage phagocytic and inflammatory responses) 24 hours before rabbit lung reperfusion resulted in improved oxygenation at 30 minutes.⁶²

We have shown that neutrophils play a critical role in mediating IRI and can augment alloimmunity.^{63,64} Interestingly, macrophage and monocyte populations in donors and recipients potentiate IRI by regulating trafficking of neutrophils into pulmonary grafts. For example, we have shown that lung-resident macrophages express the cell membrane-associated protein DAP12, which contributes to the local production of proinflammatory cytokines and neutrophil chemoattractants, in part by regulating the survival of macrophages.⁸ In a mouse lung transplant model, we demonstrated that deficiency of DAP12 in the donor reduces neutrophil extravasation and protects against IRI. Intravital 2-photon imaging experiments have revealed that circulating monocytes facilitate extravasation of neutrophils into reperfused lung grafts.⁶⁵ Subsequent work demonstrated that spleen-derived recipient classical CCR2⁺ monocytes promote neutrophil entry in injured lung tissue through MyD88-dependent production of IL-1 β .^{33,66} Recent evidence also suggests that intravascular nonclassical monocytes of donor origin (carried over during transplant despite flushing of the pulmonary vessels) also regulate neutrophil recruitment to the graft through MyD88/Trif-dependent production of the neutrophil chemokine CXCL2. Interestingly, donor nonclassical monocytes also regulate the recruitment of recipient

classical monocytes. Notably, depletion of nonclassical monocytes in donor lungs results in amelioration of lung transplant-mediated IRI (Figure 2).

7 | MACROPHAGES AND ALLOGRAFT REJECTION

Given their persistence after organ transplant, it is possible that donor macrophages contribute to allograft rejection. There is growing evidence that macrophages play a role in both ACR⁶⁷ and AMR.^{68–70} In the context of ACR, macrophages contribute to graft injury and myocardial fibrosis through cytokine and reactive oxygen species production.^{71–73} Within the rejecting human kidney, macrophages account for 38%–60% of infiltrating leukocytes.^{5,74,75} Examination of human heart transplant recipients (n = 25) with ACR revealed increased abundance of CD16⁺ monocytes/macrophages compared with healthy controls.⁷¹ CD16⁺ intermediate monocytes are considered a proinflammatory population with elevated expression of TNF- α and IL-1 β . Consistent with a contribution of infiltrating monocytes, the authors observed increased expression of HLA-DR and CD54 within circulating CD16⁺ intermediate monocytes, suggesting an activated state with increased migratory potential.^{76,77} Despite the recognition that macrophages accumulate within the allograft during ACR, the discrete roles of macrophage subpopulations and their origins are incompletely understood.⁷⁸

A hallmark of AMR is the perivascular and intravascular accumulation of neutrophils and macrophages. The clinical diagnosis is in part made by histopathologic evidence of intravascular CD68⁺ monocytes/macrophages.⁷⁹ Macrophages contribute to cardiac allograft injury in acute AMR.⁸⁰ Intravascular monocytes/macrophages displaying a proinflammatory phenotype contribute to AMR through antigen processing/presentation, cytokine production, and tissue remodeling.^{4,68,81–84} The signals responsible for recruiting intravascular monocytes/macrophages are not well established.

Donor macrophages contribute to allograft injury after lung transplant and may contribute to the development of chronic rejection. Because of their longevity in the lung allograft, donor alveolar macrophages serve as a long-term source of donor antigens. Donor alveolar macrophages secrete proinflammatory cytokines after stimulation with donor-specific antibodies, suggesting that donor alveolar macrophages contribute to antibody-mediated lung rejection.³⁹ Specifically, induction of zinc finger and BTB domain containing protein 7a, a transcription factor that helps regulate development of lymphocytes and tissue resident macrophages, in alveolar macrophages is a critical step in donor-specific antibody-induced chronic rejection.⁸⁵ Elimination of zinc finger and BTB domain containing protein 7a in alveolar macrophages was associated with decreased bronchiolar occlusion and chronic rejection.³⁹ Future studies will be required to define additional mechanisms by which donor macrophages participate in allograft rejection. Potential mechanisms include regulation of monocyte/neutrophil trafficking, monocyte fate specification, antigen presentation, and cytokine production.

8 | MACROPHAGES AND ALLOGRAFT TOLERANCE

The gold standard of transplant immunology is graft acceptance or graft tolerance. To date, tolerogenic protocols have targeted T cells, given their prominent role in graft rejection.^{86,87} Substantial evidence exists that macrophage subpopulations contribute to tolerance. Braza et al show that graft infiltrating macrophages expressed pattern recognition receptors dectin-1 and TLR4 (in response to DAMPs) and genetically deleting these proteins in recipient mice decreased recipient inflammatory Ly6c^{hi} macrophages and promoted graft tolerance with accumulation of Ly6c^{lo} macrophages.⁸⁷⁻⁸⁹ Interestingly, in this model, costimulatory blockade with CD40 inhibition (T cell receptor) was required to induce long-term tolerance, suggesting T cells may modulate macrophage phenotypes.

In addition, TIMD4⁺ and DC-SIGN⁺ macrophages suppress T cell activation, increase regulatory T cell abundance, and promote allograft tolerance.^{88,90,91} Although TIMD4⁺ macrophages represent long-lived tissue resident macrophages, DC-SIGN⁺ macrophages appear to be of monocytic origin. Blockade of the CD40L-CD40 costimulatory pathway promotes the differentiation of monocytes into DC-SIGN⁺CD169⁺ suppressive macrophages capable of secreting IL-10 and suppressing CD8 T cell activation.⁸⁸ Treatment of CD169-DTR recipient mice with diphtheria toxin was sufficient to deplete DC-SIGN⁺ macrophages and prevent allograft tolerance.⁸⁸ Conditional deletion of mammalian target of rapamycin (mTOR) in recipient myeloid cells provided further support for a role of monocyte-derived macrophages in tolerance. Deletion of mTOR in recipient monocytes and macrophages led to increased numbers of intragraft Foxp3⁺ T cells, long-term allograft survival, and reduced myocardial or vascular injury. Using high-density lipo-protein nanobiologic targeting the mTOR, Braza et al⁸⁷ were able to more selectively target myeloid cells and inhibit T cell proliferation and induce expansion of tolerogenic Foxp3 regulatory T cells suggesting an ability to promoting tolerogenic Ly6c^{lo} macrophages. Mechanistically, mTOR-deficient graft infiltrating macrophages upregulated programmed cell death 1 ligand and blockade of programmed cell death 1 ligand resulted in rapid graft rejection.⁹² Although these findings identify a role for monocytes and macrophages in establishing allograft tolerance, the exact role for donor macrophages remains to be clarified. An intriguing possibility is that donor macrophages may influence the ability of graft infiltrating monocytes to differentiate into macrophage subsets with regulatory activity.

9 | EFFECT OF IMMUNOSUPPRESSION ON MACROPHAGES

Current solid organ immunosuppression regimens consist of calcineurin inhibitors (tacrolimus, cyclosporine), antimetabolites (mycophenolate mofetil [MMF], azathioprine), mTOR inhibitors (sirolimus, everolimus), and glucocorticoids (prednisone). A fraction of patients additionally receive induction therapy with agents such as thymoglobulin and basiliximab. How these agents influence donor macrophage, recipient monocyte, and recipient monocyte-derived macrophage function is an area of interest.

Although the primary mechanism of calcineurin inhibitors is to inhibit T cell receptor activation through reductions in NFAT signaling and cytokine secretion (IL-2, interferon- γ), these agents also influence macrophage behavior.⁹³ Tacrolimus inhibits macrophage

calcineurin signaling but activates nuclear factor- κ B signaling and downstream production of IL-12 and TNF- α .^{93–95} Calcineurin inhibitors are also reported to inhibit TLR signaling and cytokine production (IL-1 β , TNF- α , IL-6, IL-10), bacterial phagocytosis, and regulation of macrophage polarization.^{96–98} The antimetabolite MMF inhibits the synthesis of guanosine nucleotides. In the context of rejection, circulating monocytes treated with MMF produced less IL-1 β , IL-10, and TNF- α .⁹⁹ Healthy human volunteer blood was exposed to tacrolimus or MMF in vitro.¹⁰⁰ There was mild inhibition of phosphorylation of CD14⁺ monocyte activation (p38MAPK with tacrolimus and AKT with MMF) but minimal effects on cytokine production and macrophage differentiation, suggesting that these agents do not dramatically influence macrophage function.¹⁰⁰ Glucocorticoids act through many pathways to control antigen presentation, cytokine production, and proliferation of lymphocytes.¹⁰¹ Glucocorticoids are associated with reduced CD14⁺CD16⁺⁺ monocytes, increased IL-10 expression, and reduced IL-1, IL-12, and TNF expression.^{102,103} These effects may be related to the ability of glucocorticoids to influence either macrophage signaling and/or monocyte differentiation.¹⁰⁴ Using a zebrafish amputation model for inflammation, Xie et al.¹⁰⁵ showed that glucocorticoids reduced neutrophil but not macrophage migration (no effect on chemoattractants Ccl2 or Cxcl11aa). They also show with RNA-seq that the glucocorticoid be-clometasone inhibits proinflammatory macrophage differentiation. mTOR inhibitors suppress macrophage CCL2, CCL3, CCL4, CCL5, IL-6, and IL-9 expression^{106,107} and impair antigen presentation through reduced CD80 expression.¹⁰⁸

The influence of induction therapy and biologic agents on donor and recipient macrophage function is less clear. Thymoglobulin induction therapy leads to complement-mediated T cell death and an increase in CD14⁺ monocytes.¹⁰⁹ Belatacept and abatacept (CTLA4-Ig recombinant proteins) prevent T cell activation through inhibition of costimulation: interactions between CD80/CD86 on macrophages and dendritic cells and CD28 molecule on T cells.¹¹⁰ These agents also reduce the production of IL-12 and TNF.¹¹¹ Whether immunosuppressive agents have differential effects on donor vs recipient macrophages remains unexplored.

10 | MACROPHAGES AS A THERAPEUTIC TARGET

There have been few publications suggesting that depleting macrophage populations, inhibiting their activation, or suppressing their effector mechanisms suppresses IRI, reduces rejection, and improves allograft survival.^{5,112–115} Although there are substantial gaps in knowledge delineating the exact contributions of donor- and recipient-derived macrophages, emerging data suggest that manipulating donor macrophage subsets could prove efficacious. If indeed the donor subpopulation must be targeted, the CCR2⁺ macrophages would be the presumptive target in heart transplant. These inflammatory populations could be targeted with nanoparticles or micelles that target short peptides,^{116,117} standard monoclonal antibodies (i.e., MLN1202), or newer constructs like bispecific or multispecific antibodies where multiple synergistic proteins could be targeted simultaneously or a putative protein with downstream effectors simultaneously such as the CCR2 and CCR5 dual receptor blocker cenicriviroc (TAK-652).^{118,119} Bispecific or multispecific antibodies can be used for several therapeutic purposes. These reagents can facilitate efficient cell removal by bringing

target cells in proximity to activated effector T or natural killer cells. In addition, they can be used to suppress cell signaling on specific cell types, such as donor CCR2⁺ macrophages.⁸⁸

Ultimately, there is tremendous potential for developing therapeutics that are applied directly to the donor organ before transplant (during cold storage or normothermic perfusion). Such agents would minimize risks associated with systemic immunosuppression including life-threatening infection and malignancy. Targeting upstream mechanisms that initiate allograft inflammation and rejection or promote the differentiation and survival of infiltrating monocytes and monocyte-derived macrophages with regulatory activity may be particularly advantageous over current approaches that block downstream inflammatory mechanisms.

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11 | CONCLUSIONS

Heart and lung transplant have transformed care for patients with end-stage organ failure. To prevent organ rejection, current standard of care requires the use of immunosuppression that lacks specificity and is wrought with untoward toxicities. Macrophages are implicated in many aspects of transplant pathology ranging from IRI to allograft rejection and tolerance. During the past several years, paradigm-shifting studies have provided captivating insights into the extent and functional importance of macrophage diversity. We are now breaking ground into the differential roles and interactions between donor macrophages and recipient monocytes and monocyte-derived macrophages. These studies have raised the possibility that targeting donor macrophages before transplant may be an avenue to improve transplant outcomes. Donor organ-based approaches provide distinct advantages including reductions in therapeutic toxicity. Targeting donor macrophages requires a comprehensive understanding of the respective roles of donor and recipient macrophages over the entire lifespan of the allograft. Large gaps in our knowledge base exist in this area. Key questions to be addressed include, What are the roles of distinct cardiac macrophage subsets in rejection and long-term allograft tolerance? What are the mechanisms that mediate cardiac macrophage activation and monocyte differentiation? How do donor and recruited monocytes and macrophages interact with the adaptive immune system to influence transplant outcomes? It is clear that answering these questions will yield new opportunities and promising targets for safe and effective immunosuppression.

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Abbreviations:

ACR	acute cellular rejection
AMR	antibody-mediated rejection
CCR2	C-C chemokine receptor 2
DAMP	damage-associated molecular pattern
IL	interleukin
IRI	ischemia–reperfusion injury
MMF	mycophenolate mofetil
mTOR	mammalian target of rapamycin
TGF	transforming growth factor
TNF	tumor necrosis factor
TLR	toll-like receptor

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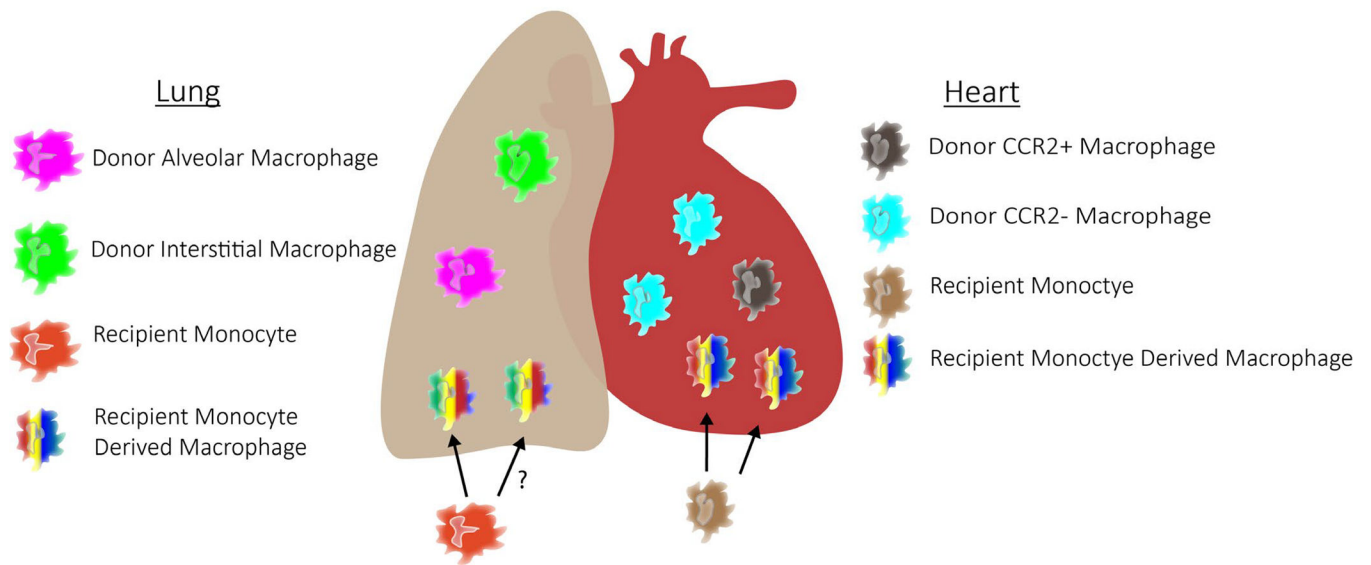
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Donor and Recipient Macrophage Populations after Transplantation

**FIGURE 1.**

Schematic of donor-vs recipient-derived macrophages after lung and heart transplant. Lung donor-derived macrophages consist of donor alveolar and donor interstitial macrophages. Circulating monocytes can infiltrate the lung and differentiate into macrophages, including recipient monocyte-derived alveolar macrophages and recipient monocyte-derived interstitial macrophages. Similarly, after cardiac transplant, the heart consists of donor CCR2⁺ macrophages, donor CCR2⁻ macrophages, in addition to recipient monocyte-derived macrophages. Some of these recipient monocyte-derived macrophages are CCR2⁺

Role of Donor Macrophages in Ischemic Reperfusion Injury

Heart

- Deletion of donor CCR2+ macrophages leads to reduced neutrophil extravasation and monocyte recruitment
- Deletion of donor CCR2- macrophages results in increase in recipient neutrophil, monocyte, and monocyte-derived macrophage recruitment
- Donor CCR2+ macrophages govern leukocyte recruitment through TLR9 and MYD88
- Ferroptosis is responsible for initial wave of cardiomyocyte cell death following transplantation and may supply DAMPs that activate donor macrophages
- Inhibition of ferroptosis was sufficient to reduce cardiomyocyte death and suppress neutrophil infiltration

Lung

- Donor macrophages induce initial response
- Recipient monocytes mediate downstream events
- Donor macrophages express DAP12 which enhances local production of neutrophil chemoattractants
- Deletion of DAP12 in donor macrophages was protective of IRI
- Nonclassical intravascular donor and classical spleen-derived recipient monocytes play important roles in promoting neutrophil recruitment to lung grafts

FIGURE 2.
Roles of donor macrophages after ischemia–reperfusion injury

TABLE 1

Surface phenotype, origin, and presumed function of donor and recipient macrophages and monocytes in the heart transplant

	Heart			
	Donor		Recipient	
	Donor CCR2⁺ macrophage	Donor CCR2⁻ macrophage	Classical recipient monocyte	Nonclassical recipient monocyte
Surface phenotype	CD11b ⁺ CD64 ⁺ CCR2 ⁺ MHC-II ^{hi}	CD11b ⁺ CD64 ⁺ CCR2 ⁻ MHC-II ^{hi or lo}	CD11b ⁺ Ly6G ⁻ CD64 ^{int} Ly6C ^{hi} CCR2 ⁺	CD11b ⁺ Ly6G ⁻ NK1.1 ⁻ SiglecF ⁻ CD64 ⁻ Ly6C ^{lo} MHCII ^{lo}
Origin	Blood monocytes	Extraembryonic progenitors	Blood monocytes	Blood monocytes
Function	Orchestrate recipient neutrophil and monocyte recruitment	Suppress recipient neutrophil and monocyte recruitment	Differentiate into macrophages and dendritic cells. Contribute to both rejection and tolerance	Unknown. It is thought that nonclassical recipient monocytes do not give rise to tissue macrophages

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TABLE 2

Surface phenotype, origin, and presumed function of donor and recipient macrophages and monocytes in the lung transplant

		Lung		
		Donor	Recipient	
		Donor alveolar macrophage	Donor interstitial macrophage	Donor nonclassical monocyte
Surface phenotype		CD11b ⁺	CD11b ⁺	CD11b ⁺
		CD64 ⁺ SiglecF ⁺ CD11c ⁺	CD64 ⁺ MHC-II ^{hi} CD11c ⁺	Ly6G ⁻ NK1.1 ⁻ SiglecF ⁻ CD64 ⁻ Ly6C ^{lo} MHCII ^{lo}
Origin		Embryonic progenitors or blood monocytes	Embryonic progenitors or blood monocytes	Blood monocytes
Function		Cytokine/chemokine production	Unknown	Orchestrate recruitment of recipient neutrophils and classical monocytes
				Differentiate into macrophages or dendritic cells Promote extravasation of neutrophils