Immunomodulatory Strategies Directed Towards Tolerance of Vascularized Composite Allografts

Maria Lucia L. Madariaga#1, Kumaran Shanmugarajah#2, Sebastian G. Michel3, Vincenzo Villani4, Glenn M. La Muraglia Il5, Radbeh Torabi6, David A. Leonard7, Mark A. Randolph8, Robert B. Colvin9, Kazuhiko Yamada10, Joren C. Madsen11, Curtis L. Cetrulo Jr12, and David H. Sachs13

Transplantation Biology Research Center, Department of Surgery, Massachusetts General Hospital, Boston, MA, USA

# These authors contributed equally to this work.

Abstract

Background—Achieving tolerance of vascularized composite allografts (VCAs) would improve the risk-to-benefit ratio in patients who undergo this life-enhancing, though not life-saving, transplant. Kidney co-transplantation along with a short course of high-dose immunosuppression enables tolerance of heart allografts across a full MHC mismatch. In this study, we investigated whether tolerance of VCA across full MHC disparities could be achieved in animals already tolerant of heart and kidney allografts.

Methods—Miniature swine that were tolerant of heart and/or kidney allografts long-term underwent transplantation of myocutaneous VCA across the same MHC barrier. Prior to VCA transplant, Group 1 (n=3) underwent Class I-mismatched kidney transplantation; Group 2 (n=3) underwent two sequential Class I-mismatched kidney transplantations; Group 3 (n=2) underwent two sequential Class I-mismatched kidney transplantations; Group 3 (n=2) underwent

Address for Correspondence: David H. Sachs, M.D. Transplantation Biology Research Center Building 149, 13th Street Charlestown, MA, USA. Telephone: 617-726-4065 Fax: 617-726-4067 David.Sachs@tbrc.mgh.harvard.edu.

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Kumaran Shanmugarajah, M.D. participated in research design, performance of the research, data analysis, and writing of the paper. kumaran.shanmugarajah@tbrc.mgh.harvard.edu

Sebastian G. Michel, M.D. participated in research design, performance of the research, data analysis, and writing of the paper. sebastian.michel@med.unimuenchen.de

Vincenzo Villani, M.D. participated in writing of the paper. vincenzo.villani@tbrc.mgh.harvard.edu

Glenn M. La Muraglia Il, B.S. participated in performance of the research. gnlamuraglia@partners.org

Radbeh Torabi, M.D. participated in writing of the paper. rtorabi@yahoo.com

David A. Leonard, M.D. participated in writing of the paper and is supported by the 2012 ASTS-Genentech Basic Scientist Scholarship and 2013 ASTS-Novartis Scientist Scholarship Grant. dleonard5@partners.org

Mark A. Randolph participated in writing of the paper. marandolph@mgh.harvard.edu

Robert B. Colvin, M.D. participated in writing of the paper. colvin@helix.mgh.harvard.edu

Kazuhiko Yamada, M.D. Ph.D. participated in performance of the research and writing of the paper. kaz.yamada@tbrc.mgh.harvard.edu

Joren C. Madsen, M.D., D.Phil. participated in research design and writing of the paper. jcmadsen@mgh.harvard.edu

Curtis L. Cetrulo, Jr. M.D. participated in research design and writing of the paper. ccetrulo@partners.org

David H. Sachs, M.D. participated in research design, performance of the research, data analysis, and writing of the paper. david.sachs@tbrc.mgh.harvard.edu

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haploidentical MHC-mismatched heart/kidney transplantation; and Group 4 (n=2) underwent full MHC-mismatched heart/kidney transplantation.

Results—All three animals in Group 1 and two of three animals in Group 2 showed skin rejection ≤85 days; one animal in Group 2 showed prolonged skin survival >200 days. Animals in Groups 3 and 4 showed skin rejection ≤80 days and regained in vitro evidence of donor responsiveness.

Conclusion—This is the first pre-clinical study in which hearts, kidneys, and VCAs have been transplanted into the same recipient. Despite VCA rejection, tolerance of heart and kidney allografts was maintained. These results suggest that regulatory tolerance of skin is possible but not generally achieved by the same level of immunomodulation that is capable of inducing tolerance of heart and kidney allografts. Achieving tolerance of skin may require additional immunomodulatory therapies.

INTRODUCTION

Vascularized composite allograft (VCA) transplantation is an emerging field that provides patients significant functional and psychological benefits over conventional reconstructive techniques (1,2). To date, close to 30 face transplants and 100 hand transplants have been performed world-wide (3-6). However, because VCA transplantation is a life-enhancing rather than life-saving procedure, these benefits are mitigated by the risks of chronic immunosuppression. Recipients of VCAs have suffered from infection, malignancy, metabolic complications and drug toxicity as a result of immunosuppressive therapy, and despite medication compliance, as many as 85% of VCA recipients experience acute rejection episodes within the first year alone (5-7).

One strategy to offset these risks is to apply tolerance induction protocols already successful in clinical kidney transplantation to VCAs (8,9). Establishing immunological tolerance would maximize long-term, rejection-free survival and abrogate the need for chronic immunosuppression. Indeed, our laboratory has recently demonstrated the ability to induce tolerance of VCAs across a haploidentical (single haplotype, Class I and Class II) major histocompatibility (MHC) mismatch in a pre-clinical large animal model using a nonmyeloablative preconditioning regimen to generate durable multilineage mixed chimerism (10,11).

However, in contrast to kidney transplantation, tolerance of VCAs presents a particular set of challenges. First, as deceased donors remain the source of VCAs, tolerance protocols cannot include extensive recipient conditioning prior to transplant. Second, VCAs are composed of tissues that have varying degrees of antigenicity, with skin being the most antigenic (12,13). Third, the morbidity of conditioning protocols, such as the risk of graft-versus-host disease, should be minimized in the context of quality-of-life VCAs (8,9,14-16).

Demonstrating that immunomodulatory mechanisms alone could induce tolerance of VCAs would be a significant step forward in applying a tolerance strategy to clinical VCA transplantation. We previously demonstrated that transplantation of kidney allografts followed by 12 days of high-dose immunosuppression uniformly induces long-term
tolerance across Class I alone or full MHC barriers (17,18). We have also demonstrated that kidney co-transplantation allows tolerance of cardiac allografts across a full MHC barrier (19). The mechanism underlying long-term acceptance involves systemic immunomodulation, as evidenced by in vitro studies identifying the necessary presence of a regulatory cell population (20,21) and the finding that long-term tolerant recipients of Class I mismatched renal allografts accepted subsequent donor MHC-matched kidney transplants without further immunosuppression (22).

VCAs placed in kidney recipients who had already achieved tolerance across a Class I alone MHC disparity rejected their skin component in five of the six animals tested (23). In this study, we investigated whether tolerance of VCA could be achieved in recipients already tolerant of kidney and heart allografts, hypothesizing that kidney-induced tolerance of a heart could be extended to a VCA. We compared VCAs that had been transplanted into recipients already tolerant of heart and/or kidney allografts across a (1) Class I alone MHC mismatch; (2) haploidentical MHC mismatch; and (3) full MHC mismatch (two haplotype, Class I and Class II).

**MATERIALS AND METHODS**

**Animals**

Transplant donors and recipients were selected from our herd of partially inbred miniature swine (age, 3-12 months; weight, 15-60kg). The immunogenetic characteristics of this herd have been described previously (24). In Group 1, to generate an MHC disparity across Class I but not Class II, SLA\(^{dd}\) (class I\(^{dd}/\)II\(^{dd}\)) animals received a kidney transplant from an SLA\(^{gg}\) (class I\(^{cc}/\)II\(^{dd}\)) donor with 12 days of cyclosporine, followed within 70 days by a myocutaneous VCA transplant from an SLA\(^{gg}\) (class I\(^{cc}/\)II\(^{dd}\)) donor without further immunosuppression (Table I, previously published in (23)). In Group 2, to generate an MHC disparity across Class I but not Class II, SLA\(^{dd}\) (class I\(^{dd}/\)II\(^{dd}\)) animals received a kidney transplant from an SLA\(^{gg}\) (class I\(^{cc}/\)II\(^{dd}\)) donor with 12 days of cyclosporine; after 100 days, Group 2 animals underwent nephrectomy of the primary kidney graft and a second donor-matched SLA\(^{gg}\) (class I\(^{cc}/\)II\(^{dd}\)) kidney transplant without further immunosuppression, followed by a donor-matched SLA\(^{gg}\) (class I\(^{cc}/\)II\(^{dd}\)) myocutaneous VCA transplant without further immunosuppression more than 100 days after the second kidney transplant (Table I, previously published in (23)). In Group 3, to generate an MHC disparity across a single haplotype with Class I and Class II, SLA\(^{ac}\) (class I\(^{ac}/\)II\(^{ac}\)) animals received heart and kidney transplants from an SLA\(^{ad}\) (class I\(^{ad}/\)II\(^{ad}\)) donor with 12 days of FK506, followed after 100 days with a donor-matched SLA\(^{ad}\) (class I\(^{ad}/\)II\(^{ad}\)) myocutaneous VCA transplant without further immunosuppression (Table I). In Group 4, to generate an MHC disparity across two haplotypes with Class I and Class II, SLA\(^{cc}\) (class I\(^{cc}/\)II\(^{cc}\)) animals received heart and kidney transplants from an SLA\(^{dd}\) (class I\(^{dd}/\)II\(^{dd}\)) donor with 12 days of FK506, followed after 100 days with a donor-matched SLA\(^{dd}\) (class I\(^{dd}/\)II\(^{dd}\)) myocutaneous VCA transplant without further immunosuppression (Table I). All recipients demonstrated significant in vitro anti-donor cytotoxic activity by CML and/or MLR before organ transplantation. All animal care and procedures were approved by the Massachusetts General Hospital Institutional Animal Care and Use Committee and conducted in compliance with the “Guide
Surgical procedures

The surgical procedures used for heart and kidney transplantation have been described in detail previously (25–27). Briefly, the recipients underwent bilateral nephrectomy. The aorta and inferior vena cava were used for end-to-side arterial and venous anastomoses for both the heart and kidney, with the heart placed at least 1 cm caudad to the kidney. The kidney transplantation was completed by performing a vesicoureteral anastomosis. Two indwelling silastic central venous catheters were placed surgically into the external or internal jugular veins. The catheters facilitated immunosuppression administration and frequent blood sampling for *in vitro* assays and for monitoring of renal function and whole blood tacrolimus or cyclosporine levels.

The surgical procedure used for gracilis myocutaneous VCA transplantation has been described previously (28). The VCA, composed of skin, subcutaneous tissue, muscle and its vascular pedicle (femoral artery and femoral vein) was anastomosed to the recipient's internal jugular vein and internal carotid artery. The animal did not receive immunosuppression post-operatively.

Rejection monitoring

Kidney function was monitored by serial serum creatinine levels. Renal allograft rejection was defined as sustained rise in serum creatinine to >10 mg/dL and/or uremia. Heart function was monitored by daily palpation and electrocardiogram (ECG) using the AliveCor Veterinary Heart Monitor (AliveCor, Inc., San Francisco, CA). Cardiac allograft rejection was defined by either loss of a ventricular impulse on palpation, and/or QRS-wave amplitude of less than 0.3mV, and/or the lack of ventricular contraction on echocardiography (29). The VCA was monitored for viability by checking capillary refill and monitored for rejection by visual inspection and serial biopsies. VCA rejection was defined as the point at which the skin became necrotic and was confirmed by biopsy.

Routine biopsies were performed on all transplant recipients at predetermined time intervals (POD 20-30, 50-60, 90-100) or after a rise in creatinine, a decrease in donor heart palpation/QRS-wave amplitude or a change in VCA appearance. Allograft rejection was confirmed histologically in all cases.

Immunosuppression

Cyclosporine (Novartis Pharmaceutical Corporation, Hanover, NJ) was mixed and administered as an intravenous suspension according to the specifications of the manufacturer. Cyclosporine was given as a daily intravenous infusion over 1 hour (13 to 16 mg/kg/day with target levels 400 to 800 ng/mL) for 12 consecutive days, starting on the day of primary kidney transplantation (day 0) for animals in Groups 1 and 2.

Tacrolimus (Haorui Pharma-Chem Inc., Irvine, CA) was mixed and administered as an intravenous suspension according to the specifications of the manufacturer. Tacrolimus was
given as a continuous infusion at a dose of 0.08-0.20 mg/kg (adjusted to maintain a whole blood level of 30-50 ng/ml) for 12 consecutive days, starting on the day of heart and kidney transplantation (day 0) for animals in Groups 3 and 4.

Pathology studies
Core needle biopsies were performed on cardiac allografts. Wedge biopsies were performed on kidney allografts. Kidney biopsies were taken at the same time as the heart samples. 6mm punch biopsies and wedge biopsies were performed on the VCA graft. Tissue was fixed in formalin and embedded in paraffin for routine light microscopy (H&E, PAS). Scoring of rejection was performed without knowledge of the functional status of the graft based on the International Society for Heart and Lung Transplantation System for hearts (30) and the current Banff consensus criteria for kidney (31) and vascularized cutaneous allografts (32). Complete necropsies were done upon completion of the experiments and tissue was similarly processed for pathological examination.

Preparation of peripheral blood leukocytes (PBLs)
Freshly heparinized whole blood was diluted approximately 1:2 with HBSS (Gibco BRL, Grand Island, NY), and the mononuclear cells were obtained by means of gradient centrifugation with Histopaque (Sigma, St. Louis, MO). The mononuclear cells were washed once with HBSS, and contaminating red cells were lysed with ammonium chloride potassium lysing buffer (Bio Whittaker, Inc, Walkersville, MD). Cells were then washed with HBSS and resuspended in tissue culture medium. All cell suspensions were kept at 4°C until used in cellular assays.

Cell-mediated lymphocytotoxicity (CML) assay
Cell-mediated lymphocytotoxicity (CML) assays with porcine cells have been described previously (33). The tissue culture media used for the CML assays consisted of RPMI-1640 (Gibco BRL) supplemented with 6% fetal bovine serum (Sigma Chemical Co, St Louis, MO), 100 U/mL penicillin, 135 mg/mL streptomycin (Gibco BRL), 50 mg/mL gentamicin (Gibco BRL), 10 mmol/L N–2-hydroxyethylpiperazine-N–2-ethanesulfonic acid (HEPES; Fisher Scientific, Pittsburgh, PA), 2 mmol/L L-glutamine (Gibco BRL), 1 mmol/L sodium pyruvate (Bio Whittaker, Inc), nonessential amino acids (Bio Whittaker, Inc), and 5 × 10^{-5} mol/L β2-mercaptoethanol (Sigma Chemical). The effector phase of the CML assay was performed with Basal Medium Eagle (Gibco BRL) supplemented with 6% controlled processed serum replacement 3 (Sigma Chemical) and 10 mmol/L HEPES. Briefly, lymphocyte cultures containing 4 × 10^6/mL responder and 4 × 10^6/mL stimulator PBLs (irradiated with 2500 cGy) were incubated for 6 days at 37°C in 5% carbon dioxide and 100% humidity in CML medium. Bulk cultures were harvested, and effectors were tested for cytotoxic activity on chromium 51–labeled (Amersham, Arlington Heights, IL) lymphoblast targets generated from phytohemagglutinin (M-form; Life Technologies, Gaithersburg, MD) stimulation. Effector cells were incubated for 5.5 hours with target cells at effector/target ratios of 100:1, 50:1, 25:1, and 12.5:1. Two target cells were tested in each assay: (1) PBLs SLA matched to the donor and (2) third-party PBLs. Supernatants were then harvested by using the Skatron collection system (Skatron, Sterling, VA), and ^{51}Cr release was
determined on a gamma counter (Micromedics, Huntsville, AL). The results were expressed as a percentage of specific lysis and calculated as follows:

$$\text{Percentage of specific lysis} = \left( \frac{\text{Experimental release} \ [\text{cpm}] - \text{Spontaneous release} \ [\text{cpm}]}{\text{Maximum release} \ [\text{cpm}]} \right) \times 100\%$$

**Mixed-lymphocyte reaction (MLR) assay**

Mixed lymphocyte reaction (MLR) responses to self, donor and third-party were determined in a single assay for each animal. MLR media consisted of RPMI 1640 (Life Technologies) supplemented with 6% fetal pig serum (Sigma; St. Louis, MO, USA), 100 U/mL penicillin (GIBCO-Invitrogen Corporation; Carlsbad, CA, USA), 135 μg/mL streptomycin (GIBCO-Invitrogen Corporation), 50 μg/mL gentamicin (GIBCO-Invitrogen Corporation), 10 mM HEPES (Cellgro Mediatech, Inc.; Manassas, VA, USA), 2 mM l-glutamine (Life Technologies), 1 mM sodium pyruvate (BioWhittaker–Cambrex; East Rutherford, NJ, USA), nonessential amino acids (BioWhittaker–Cambrex) and 5 × 10^{-5} M 2-beta-mercaptoethanol (Sigma). Cultures containing 4 × 10^6 responder and 4 × 10^6 irradiated (2500 cGy) stimulator PBMCs were incubated in 200 μL of media in 96-well flat-bottomed plates (Costar Corning; Lowell, MA, USA) for 5 days at 37°C in 5% CO2 and 100% humidity. After the 5-day incubation, 1 uCi of [3H]-thymidine was added to each well, followed by an additional 5-hr incubation under the same conditions. [3H]-thymidine incorporation was determined in triplicate samples by beta-scintillation counting. Absolute counts were compensated for background and then expressed as stimulation indices (SI), calculated as SI = average counts per minute for a responder–stimulator pair per c.p.m. of the same responder stimulated by an autologous stimulator.

**Assessment of alloantibody**

The presence of anti-donor immunoglobulin (IgM and IgG) in the serum of experimental swine was examined by indirect flow cytometry using a Becton Dickinson FACScalibur (Sunnyvale, CA) to determine the SLA-binding specificity of the antibody. FITC-labeled goat anti-swine IgM or IgG polyclonal antibodies were used as secondary reagents (Kirkegaard & Perry Laboratories Inc, Gaithersburg, MD). For staining, 1 × 10^6 cells per tube of donor-type PBLs (SLA^{dd} or SLA^{ad}) were resuspended in 100 μL HBSS containing 0.1% bovine serum albumin and 0.05% NaN₃ and incubated for 30 minutes at 4°C with 10 μL decomplemented test sera (neat). After two washes, a saturating concentration of FITC-labeled goat anti-swine IgM or IgG was added and incubated for 30 minutes at 4°C. After a final wash, cells were analyzed by means of flow cytometry with propidium iodide gating to exclude dead cells. Both normal pig serum and pretransplant sera from each experimental animal were used as controls for specific binding.

**RESULTS**

**Early epidermis and muscle loss in recipients of haploidentical MHC-mismatched VCAs**

To determine whether tolerance achieved across a single-haplotype Class I and Class II mismatch could confer tolerance to a VCA, two animals (SLA^{ad}) who had been long-term tolerant of heart and kidney allografts (heart and kidney allografts from the same SLA^{ad}
Early epidermis and muscle loss in recipients of full MHC-mismatched VCAs

To determine whether tolerance achieved across a two-haplotype Class I and Class II mismatch could confer tolerance to a VCA, two animals (SLA<sup>cc</sup>) who had been long-term tolerant of heart and kidney allografts (heart and kidney allografts from the same SLA<sup>dd</sup> donor) for >100 days underwent VCA transplantation (VCA allograft from a different SLA<sup>dd</sup> donor) (Group 4, Table I). The VCA on Group 4 animal #21740 showed visual and histological signs of epidermis rejection by POD14; VCA dermis and muscle were rejected by POD35. The VCA on Group 4 animal #22025 showed visual and histological signs of epidermis, dermis, and muscle rejection by POD30 (Figure 1).

Donor-MHC-specific responsiveness is regained after VCA transplant

To assess immune competence in recipients before and after VCA transplant, MLR and CML assays were performed. Group 3 animal #21517, which had been donor-specific unresponsive by CML and MLR prior to VCA transplant, regained donor-responsiveness after VCA transplant (Figures 3 and 4). Group 3 animal #21270 already showed positive donor-specific response by CML and MLR at the time of VCA transplant (data not shown). Both Group 4 animals showed donor-specific unresponsiveness by CML and MLR prior to VCA transplant. After VCA transplant, Group 4 animal #22025 regained donor-responsiveness by CML and MLR (Figures 3 and 4); Group 4 animal #21740 also regained donor-responsiveness by MLR, but remained unresponsive to donor by CML (data not shown). Animals in Groups 1 and 2 displayed donor-specific hyporesponsiveness or unresponsiveness by CML before and after VCA epidermis rejection (23).

Lack of circulating alloantibody following VCA rejection

To determine whether rejection of VCA led to alloantibody formation, flow cytometry analysis of anti-donor antibodies was performed. Animals in Group 3 and Group 4 did not develop any detectable circulating levels of anti-donor IgM or IgG antibody before or after VCA transplant (Figure 5). Animals in Groups 1 and 2 also did not develop alloantibody after VCA transplant (23).

Heart and kidney allograft tolerance maintained despite VCA rejection

Animals in Groups 3 and 4 were monitored by serial biopsies to determine whether VCA rejection affected tolerance of heart or kidney allografts. Heart and kidney allografts were biopsied up to 40 days after VCA transplant in Group 3 and up to 65 days after VCA transplant in Group 4. Animals in both groups showed no clinical or histological signs of
heart rejection (Table II). Animals in Group 3 showed no signs of kidney rejection, though animal #21270 developed pyelonephritis with a corresponding increase in creatinine (Table II). Group 4 animal #22025 showed no signs of kidney rejection; however, Group 4 animal #21740 had chronic rejection changes present prior to VCA transplantation (Table II). Nodular lymphocytic infiltrates associated with the vasculature of the kidney defined as “Treg-rich organized lymphoid structures” were found in all animals on kidney biopsies before VCA transplantation and after VCA rejection (Figure 6).

DISCUSSION

Because VCA transplantation is a life-enhancing rather than life-saving procedure, any risk to the patient must be minimized. Inducing tolerance of VCAs would alleviate the burden of chronic immunosuppression (5-7). The use of deceased donors and a small donor pool preclude MHC antigen matching, implying that tolerance of VCAs needs to be applicable across full Class I and Class II MHC disparities. However, current clinical and pre-clinical protocols for achieving tolerance across full MHC mismatches contain risks that are unacceptable for the VCA recipient. For example, tolerance of MHC-mismatched kidneys was achieved in humans through conditioning regimens that generate mixed chimerism, which carry up to a 34% risk of GvHD (8,9,34-36). In a pre-clinical large animal model, a mixed chimerism-based protocol resulted in tolerance of haploidentical VCAs but incurred the morbidity of GvHD in 12% of animals (11). Therefore, achieving tolerance of VCA using immunomodulatory pathways alone would be an important building block from which alternative strategies that do not require stable chimerism or intense conditioning can be developed.

Here we investigated whether immunomodulatory mechanisms alone are sufficient to induce tolerance of VCAs. Immunomodulation occurring during the maintenance phase of kidney-induced cardiac allograft tolerance across a full MHC mismatch is known to allow indefinite survival of heart allografts without the presence of kidney grafts and prolong donor skin graft survival (19). In this study, we found that animals who received hearts/kidneys and a delayed VCA across a haploidentical or full MHC barrier showed rejection of the skin component, rejection of the muscle and return of donor responsiveness by CML and MLR (Groups 3 and 4, Table I, Figures 3 and 4). In comparison, animals in Groups 1 and 2, who received kidneys and a delayed VCA across a Class I MHC barrier, showed rejection of the skin component, long-term acceptance of the muscle and maintenance of donor-specific unresponsiveness in vitro (23).

These findings demonstrate that (1) immunomodulatory mechanisms that maintain tolerance of heart and kidney allografts are usually, but not always, insufficient to induce tolerance of VCAs and (2) inducing tolerance across a full MHC barrier is more difficult than inducing tolerance across a lesser MHC disparity (e.g. prolonged skin survival in Group 1 and 2 versus Group 3 and 4 recipients). Indeed, in 1 of 3 animals who achieved tolerance across a Class I alone MHC mismatch (#18954), VCA epidermis survival was prolonged for over 200 days (Group 2, (23)). This finding indicates that tolerance of VCA via immunomodulatory mechanisms alone, although rare in these studies, is possible and can be
potentially due, in the case of animal #18954, to augmented T regulatory cell activity (23),
the reason for which may be worthy of further investigation.

The ability of systemic versus local factors to induce tolerance depends on the tolerogenicity
of the organ implanted. For example, with a tolerogenic organ such as the kidney, regulatory
mechanisms that maintain long-term tolerance of kidneys (18,20,37-39) are sufficient to induce tolerance. Long-term tolerant recipients of Class I mismatched renal allografts accepted second transplants from donors MHC matched to the donors of the first renal grafts without additional immunosuppression (22). Here we attempted to exploit the robust tolerance achieved across a full MHC mismatch for heart and kidney allografts to achieve tolerance of VCA. Tolerance across a full MHC mismatch, despite being more difficult to induce, appears harder to abrogate once gained. This hypothesis is based on recent work showing that recipients of Class I alone MHC mismatched heart and kidney allografts who undergo kidney graftectomy and subsequent skin grafting demonstrate heart rejection with severe cardiac allograft vasculopathy whereas recipients of full MHC mismatched heart and kidney allografts who are subject to the same immunologic challenge remain tolerant of their heart allografts indefinitely (Michel et al, manuscript in preparation). However, in the present study, despite robust tolerance across highly disparate MHC barriers, VCAs placed in a delayed fashion showed en-bloc rejection. One possibility is that the state of tolerance was not robust enough to induce tolerance of VCA. Another more likely possibility is that the VCA itself, due to tissue-specific or minor antigens, is not as tolerogenic as a kidney and requires additional conditioning for acceptance. Indeed, a 28-day course of high-dose FK506, which is sufficient to induce kidney allograft tolerance (18), could not induce tolerance of VCA (2 animals, Torabi et al, unpublished data).

Interestingly, despite en-bloc rejection of the VCA and return of donor-specific responsiveness in vitro (Figures 3 and 4), animals in Groups 3 and 4 remained tolerant of their heart and kidney allografts. This finding suggests the presence of intra-graft suppressive phenomena that maintain tolerance of hearts and kidneys despite sensitization to donor MHC (22,38,40,41). Indeed, nodular lymphocytic infiltrates associated with the vasculature of the kidney, termed “Treg-rich organized lymphoid structures” (42) were found in kidney samples taken before and after VCA rejection (Figure 6). A key difference between the tolerant heart/kidney grafts and rejected VCA grafts is timing of transplantation. Future work will investigate whether kidney transplantation at the time of VCA transplant (day 0 protocol) would generate sufficient systemic and local immunomodulatory mechanisms to enable tolerance of VCA.

Long-term acceptance of the skin component of VCAs without incurring risk of comorbidity remains an elusive goal. Several groups have attempted to offset risk by minimizing maintenance immunosuppression or by the addition of costimulatory blockade (43-47). Current mixed chimerism protocols, on the other hand, contain the risk of GvHD (8,11,35). Demonstrating that immunomodulatory mechanisms alone could induce tolerance of VCAs would be a significant step forward. To our knowledge, this is the first pre-clinical study in which hearts, kidneys, and VCAs have been transplanted into the same recipients. We demonstrate that a robust immunomodulatory milieu alone, which is able to induce tolerance of kidneys, is insufficient to achieve tolerance of VCA in most cases. In this regard,
tolerance of VCA, especially skin, is more difficult to achieve than tolerance of kidneys or hearts co-transplanted with kidneys, and will require additional strategies that address both the need for tolerance across highly disparate MHC and for minimizing the risks of a life-enhancing procedure.

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Abbreviations

ACR acute cellular rejection  
CML cell-mediated lympholysis  
HKtx heart kidney transplant  
HSC hematopoietic stem cell  
MHC major histocompatibility complex  
MLR mixed-lymphocyte reaction  
PAA pig allelic antigen  
PBL peripheral blood leukocytes  
POD postoperative day  
PSL percent specific lysis  
SLA swine lymphocyte antigen  
VCA vascularized composite allograft.

Reference List


Figure 1.
Gross appearance of VCA. Representative clinical images of VCAs from Group 3 animal #21517 (*top row*) and Group 4 animal #22026 (*bottom row*) by postoperative day. Animals in both groups showed patchy areas of necrosis starting on postoperative day 7 that progressed to epidermal sloughing.
Figure 2.
Histology from representative VCA biopsies taken on postoperative day 30-31. VCA biopsy from Group 3 animal #21517 (top row) shows grade 4 rejection of the epidermis and acute rejection of the muscle with endarteritis. VCA biopsy from Group 4 animal #22025 (bottom row) shows grade 4 rejection of the epidermis and acute rejection of the muscle with endarteritis.
Figure 3.
MLR assays from VCA recipients. Stimulation indices to self, donor-type (SLA^ad or SLA^dd), and third-party (YO) peripheral blood mononuclear cells before VCA transplant and 30-31 days after VCA transplant. MLR, mixed-lymphocyte reaction. YO, York.
Figure 4.
CML assays from VCA recipients. Percent specific lysis is plotted as a function of effector:target ratio. Response against donor-type (SLA^{ad} or SLA^{dd}) targets before VCA transplant and 30-31 days after VCA transplant. CML, cell-mediated lympholysis.
Figure 5.
Alloantibody response. Levels of circulating IgM and IgG alloantibody were measured by flow cytometry in recipients in Groups 3 and 4. Data were normalized to the mean fluorescence intensity of negative control values to plot normalized mean fluorescence intensity as a function of postoperative day (POD). HK tx, heart kidney transplant.
Figure 6.
Histology from representative kidney samples taken prior to VCA transplantation and after VCA rejection. Group 3 animal #21517 (top row) and Group 4 animal #22025 (bottom row) demonstrate the presence of organized lymphoid structures before and after VCA rejection.
### Table I

#### Survival of vascularized composite allografts

<table>
<thead>
<tr>
<th>Group</th>
<th>VCA MHC disparity</th>
<th>Animal #</th>
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<td>19842</td>
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<td>GG</td>
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<td>Single haplotype, Class I and II</td>
<td>21270</td>
<td>AD</td>
<td>AD</td>
</tr>
<tr>
<td></td>
<td></td>
<td>21517</td>
<td>AD</td>
<td>AD</td>
</tr>
<tr>
<td>4</td>
<td>Two haplotype, Class I and II</td>
<td>21740</td>
<td>DD</td>
<td>DD</td>
</tr>
<tr>
<td></td>
<td></td>
<td>22025</td>
<td>DD</td>
<td>DD</td>
</tr>
</tbody>
</table>

1 Group 1 animals previously published in Cetrulo et al, Transplantation 2013 (23). Group 1 animals received a Class I mismatched VCA transplant less than 70 days after primary kidney transplant.

2 Group 2 animals previously published in Cetrulo et al, Transplantation 2013 (23). Group 2 animals received a Class I mismatched VCA transplant at least 100 days after kidney re-transplantation, without further immunosuppression (i.e., >200 days after primary transplantation).
Table II

Histology and function of heart and kidney allografts before and after VCA transplantation

<table>
<thead>
<tr>
<th>Group</th>
<th>VCA MHC disparity</th>
<th>Animal #</th>
<th>Heart Histology<em>1/Function</em>2</th>
<th>Kidney Histology<em>1/Function</em>2</th>
</tr>
</thead>
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<td></td>
<td></td>
<td>Before VCA</td>
<td>After VCA</td>
</tr>
<tr>
<td>3</td>
<td>Single haplotype, Class I and II</td>
<td>21270</td>
<td>0/sinus</td>
<td>0/sinus</td>
</tr>
<tr>
<td></td>
<td></td>
<td>21517</td>
<td>0/sinus</td>
<td>0/sinus</td>
</tr>
<tr>
<td>4</td>
<td>Two haplotype, Class I and II</td>
<td>21740</td>
<td>0/sinus</td>
<td>1/sinus</td>
</tr>
<tr>
<td></td>
<td></td>
<td>22025</td>
<td>0/sinus</td>
<td>0/sinus</td>
</tr>
</tbody>
</table>

*1 Grading of heart acute rejection from 0 (no rejection) to 3 (severe rejection) based on ISHLT scoring system (28); grading of kidney acute rejection from 0 (no rejection) to 3 (severe rejection) based on Banff classification (29).

*2 Function of heart allografts assessed by ECG and palpation; function of kidney allografts assessed by creatinine measurements (mg/dL)

*3 Animal #21270 developed pyelonephritis.

*4 Animal #21740 had chronic rejection of the kidney allograft before and after VCA transplantation.