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The potential of costimulation blockade to serve as a novel transplant immunosuppression strategy has been explored for over 20 years, culminating in the recent clinical approval of belatacept for renal transplant patients. Despite improving long-term graft function and survival compared with calcineurin inhibitors, clinical acceptance of belatacept has been hindered by elevated rates of acute rejection. We examined the signaling pathways required to activate costimulation blockade–resistant alloreactive T cells and identified the OX40/OX40L secondary costimulatory pathway as a promising target. We next sought to improve the clinical efficacy of traditional costimulation blockade using belatacept by coupling it with anti-OX40L. Using a murine transplant model, we demonstrate that combined blockade enhances the suppression of alloreactive T cell proliferation and effector functions including both cytokine release and cytotoxic degranulation. We also show that anti-OX40L may be particularly useful in targeting alloreactive memory T cell responses that are relatively unaffected by traditional costimulation blockade regimens. Finally, we translated this therapy to a clinically relevant nonhuman primate renal transplant model, validating the efficacy of this regimen in a potentially novel steroid- and calcineurin inhibitor–free immunosuppression regimen.

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Interruption of OX40L signaling prevents costimulation blockade-resistant allograft rejection

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The potential of costimulation blockade to serve as a novel transplant immunosuppression strategy has been explored for over 20 years, culminating in the recent clinical approval of belatacept for renal transplant patients. Despite improving long-term graft function and survival compared with calcineurin inhibitors, clinical acceptance of belatacept has been hindered by elevated rates of acute rejection. We examined the signaling pathways required to activate costimulation blockade-resistant alloreactive T cells and identified the OX40/OX40L secondary costimulatory pathway as a promising target. We next sought to improve the clinical efficacy of traditional costimulation blockade using belatacept by coupling it with anti-OX40L. Using a murine transplant model, we demonstrate that combined blockade enhances the suppression of alloreactive T cell proliferation and effector functions including both cytokine release and cytotoxic degranulation. We also show that anti-OX40L may be particularly useful in targeting alloreactive memory T cell responses that are relatively unaffected by traditional costimulation blockade regimens. Finally, we translated this therapy to a clinically relevant nonhuman primate renal transplant model, validating the efficacy of this regimen in a potentially novel steroid- and calcineurin inhibitor-free immunosuppression regimen.

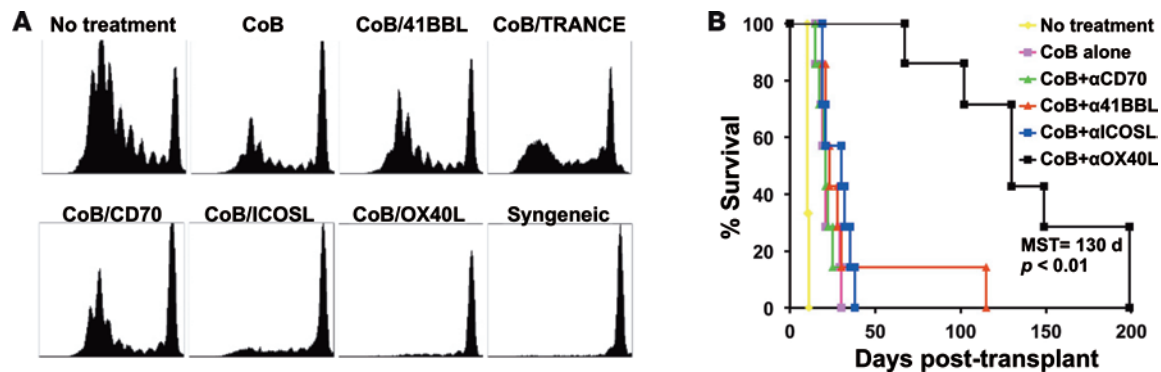


Figure 1. Combined costimulation blockade (CoB) and OX40L blockade inhibits naive alloreactive T cell expansion in vivo and prolongs skin graft survival. (A) In vivo mixed lymphocyte reaction was performed by adoptively transferring CFSE-labeled C57BL/6 splenocytes into sublethally irradiated BALB/c mice treated with CoB plus a secondary costimulatory receptor antagonist. Splenocytes were harvested after 72 hours and were assessed for CFSE-labeled cell division. Treatment with either CoB + anti-ICOSL or CoB + anti-OX40L suppressed alloproliferation ($n = 5$ per group, representative of

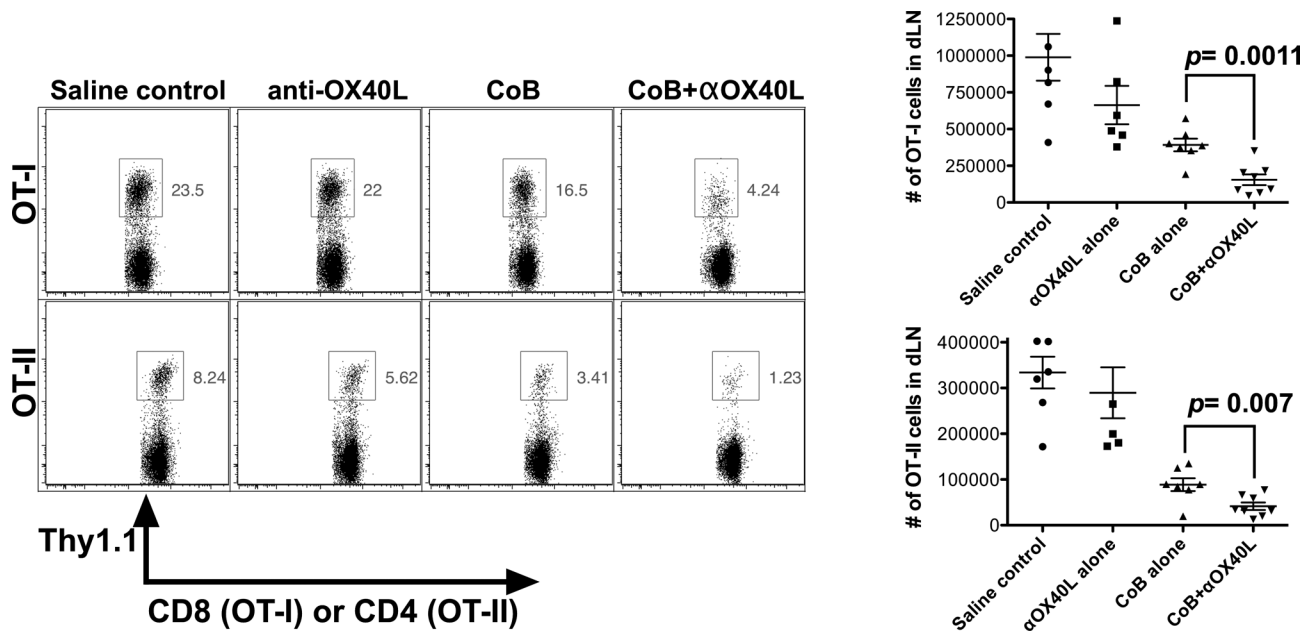
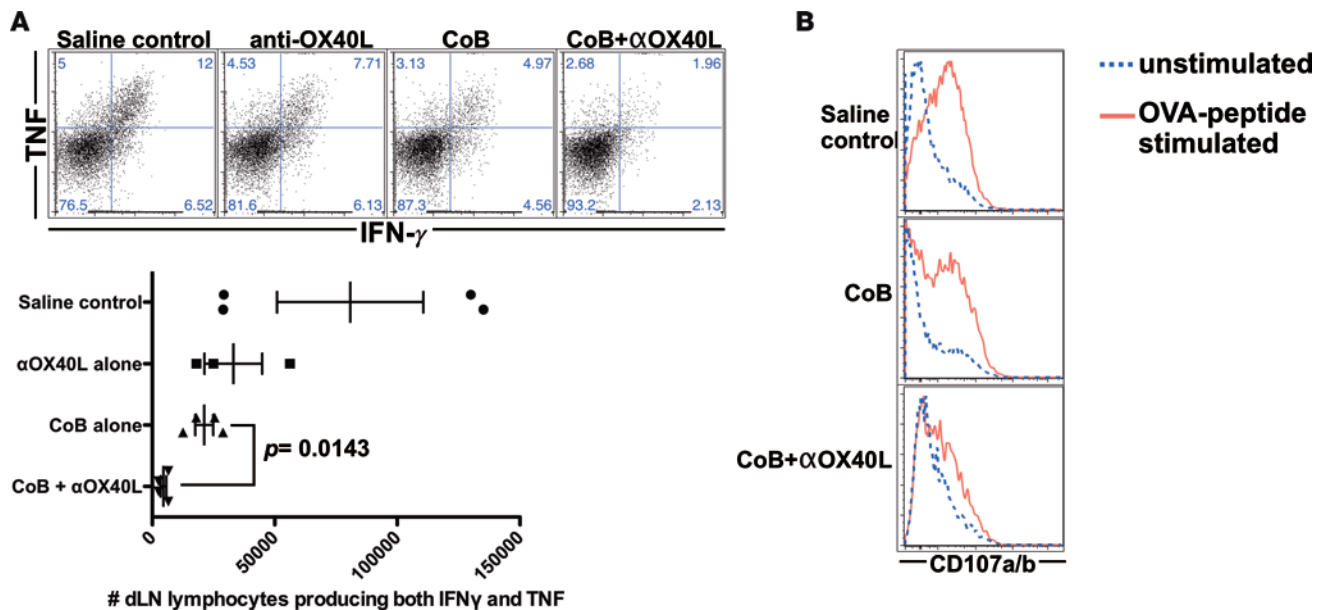


Figure 2. Combined costimulatory blockade (CoB) and OX40L blockade inhibits naive donor-specific T cell proliferation in vivo. C57BL/6 recipients were adoptively transferred with ovalbumin-specific OT-I and OT-II T cells and then transplanted with skin grafts from transgenic mouse donors ubiquitously expressing membrane-bound ovalbumin in all tissues (mOVA). Draining lymph nodes (dLNs) were harvested on posttransplant day 10 and analyzed ($n = 6$ –8 mice/group, combined data over 2 independent experiments). Expansion of Thy1.1⁺ OT-I and OT-II T cells was measured by analyzing the percentage of the total CD8⁺ and CD4⁺ T cell population in the dLNs that was also Thy1.



regulatory T cell (Treg) suppression of alloresponses, as Li and colleagues have demonstrated that OX40L signaling can inhibit Treg suppression (22, 23). We are actively exploring the impact of combined blockade therapy on Treg numbers and function to assess this hypothesis.

Previous groups have also examined the OX40/OX40L pathway in alloimmune responses. Monotherapy with anti-OX40L did not prolong rat heart or skin graft survival (24), but it did suppress development of cardiac allograft vasculopathy in another murine cardiac transplant system (25). Similar to our findings, the efficacy of OX40L blockade was enhanced when coupled with CD28/CD154 blockade (18). We have substantially extended these earlier findings, examining the molecular mechanisms at play in combined CoB and OX40L blockade and demonstrating the efficacy of this regimen in a rigorous preclinical primate transplant system that spared both calcineurin inhibitors and steroids.

This work highlights the redundancy of costimulatory pathways involved in alloreactive T cell activation, demonstrating that secondary pathways such as OX40/OX40L can assume a more vital role once the primary costimulatory pathways (such as CD28/B7) are blocked. Thus, successful suppression of alloreactivity may require the simultaneous blockade of multiple costimulatory pathways. Relatively high rates of CoB-resistant acute rejection have limited the widespread adoption of belatacept clinically, and strategies that may enhance the efficacy of CoB could offer real clinical benefits. Importantly, the humanized anti-OX40L monoclonal antibody employed in our experiments was utilized in human clinical trials for asthma as a monotherapy treatment and no safety concerns were revealed (NCT00983658). These experiments suggest that humanized anti-OX40L may also merit investigation as a transplant immunosuppressant for use in conjunction with belatacept.

Methods

Mice. Adult male 6- to 8-week-old C57BL/6 mice (Jackson Laboratories), TCR transgenic OT-I mice (Taconic Farms), μ MT mice (Jackson Laboratories), and Act-mOVA mice (gifted by Marc Jenkins, University of Minnesota, Minneapolis, Minnesota, USA) (26) were obtained. To generate mice with memory OT-I cells, 10^4 OT-I cells were adoptively transferred into naive C57BL/6 mice via tail vein injection, and then 2 days later the recipients were infected with 10^4 CFU of LM-OVA (27) by i.p. injection, followed by a 30-day waiting period prior to experimental use.

In vivo mixed lymphocyte reaction. C57BL/6 splenocytes were labeled for 5 minutes with 10 μ M CFSE, and 2×10^7 to 3×10^7 of these labeled responders were adoptively transferred i.v. into irradiated BALB/c mice (700 rads), which were treated with the different immunosuppression regimens as described below. Splenocytes were harvested after 72 hours and analyzed by flow cytometry to assess the CFSE dilution and thus proliferation of H-2K^d-negative (responder) T cells.

Skin grafting. Full-thickness tail skin grafts (~ 1 cm²) were transplanted onto the dorsal thorax of recipient mice. Where indicated, recipients of skin grafts received treatment with CoB (500 μ g each of hamster anti-mouse-CD154 mAb [Bio X Cell, clone MR-1] and human CTLA-4-Ig [Bristol-Meyers Squibb] and/or 250 μ g of rat anti-mouse OX40L mAb [Bio X Cell, clone RM134L]). All mAbs were administered i.p. on posttransplant days 0, 2, 4, and 6.

Flow cytometric analyses for frequency and absolute number. Splenocytes, blood, and/or cells obtained from axillary dLNs in skin graft recipients treated with CTLA-4-Ig and/or anti-OX40L were stained with Thy1.1-PerCP (clone OX-7), CD8A-APC (clone 53-6.7), CD11A-FITC (clone M17/4) and/or CD49D-PE (clone R1-2) (all BD Biosciences) for analysis on an LSRII flow cytometer (BD Biosciences). Absolute numbers of OT-I T cells were determined by TruCount Bead analysis according to the manufacturer's instructions. Cell sorting for CD8⁺Thy1.1⁺ cells was performed using MACS kits from Miltenyi Biotec. Data were analyzed using FlowJo software (Tree Star).

Intracellular cytokine staining. Splenocyte suspensions were incubated with 10 nM OVA₂₅₇₋₂₆₄ (SIINF-EKL) (Emory University Core Facility) and 10 μ g/ml Brefeldin A (BD Biosciences). Replicates without peptide were also performed. After 5 hours in culture, cells were processed using an intracellular staining kit (BD Biosciences) according to the manufacturer's instructions and stained with anti-TNF-PE (clone MP6-XT22) and anti-IFN- γ -FITC (clone XMG1.2) (both BD Biosciences). The adjusted percentage of dual producers of TNF and IFN- γ for each sample was calculated by subtracting the percentage of dual producers in the nonstimulated samples from the matched SIINF-EKL-stimulated samples.

CD107A/B degranulation assay. As previously described (28), splenocyte suspensions were incubated in R10 media at 37°C in a 96-well plate (4×10^6 cells/well) for 5 hours with monensin and anti-CD107A/B-FITC in the presence or absence of 10 nM SIINF-EKL peptide. After incubation, surface staining with

anti-Thy1.1-PerCP and anti-CD8A-Pacific Blue was performed. Degranulation was measured as the adjusted median fluorescence intensity (MFI) of CD107A/B (peptide-stimulated – unstimulated).

Immunohistochemistry. Explanted skin grafts were fixed in OTC and frozen. Sections were stained with anti-Thy1.1 mAb and developed with horseradish peroxidase to visualize infiltrating OT-I cells. Representative images are shown at an original magnification of $\times 40$.

RNA sequencing. C57BL/6 mice with memory OT-I cells were generated as above, and these received an mOVA skin graft as well as CoB $\pm$ anti-OX40L therapy. Recipients were sacrificed on POD 5, dLNs were harvested and processed into a single-cell suspension, and magnetically sorted with both CD8⁺ and Thy1.1⁺ MACS beads (Miltenyi Biotec) to enrich OT-I cells. Fifty thousand cells per mouse were then taken for RNA sequencing at the Yerkes Genomics Core. Data were analyzed using R. A heatmap was constructed containing a subset of genes of interest from whole-genome RNASeq. RNAseq transcriptome data were expressed in reads per million kilobases (RPKM) for normalization, and represented as low to high RPKM by color depth (white = low RPKM, dark blue = high RPKM).

Rhesus kidney transplants. Rhesus macaque renal transplants were performed in the sterile operating theater at the National Yerkes Primate Research Center using standard techniques. All rhesus subjects were MHC haplotyped at the University of Wisconsin to construct maximally mismatched donor-recipient pairings. Kidney recipients were treated with belatacept (20 mg/kg on POD 0, 3, 7, and 14, and every 2 weeks thereafter until POD 84), humanized anti-OX40L (Roche, 20 mg/kg on POD 0, 10 mg/kg on POD 3, 7, 14, 21, 28, 35, and 42), or combined therapy. Ratios of blood urea nitrogen to creatinine (BUN/Cr) were measured on POD 1, 3, and 7, and weekly thereafter, and weekly peripheral blood samples were analyzed by flow cytometry.

Donor-specific antibody development. The development of donor-specific antibody was monitored by flow cross-match. Donor peripheral blood mononuclear cells (3×10^5) were isolated and incubated with titrated recipient serum. Cells were subsequently stained with FITC-labeled goat anti-rhesus IgG (KPL, catalog 072-11-021), PE-labeled anti-CD20 (BD Biosciences, catalog 556633), and PerCP-Cy5.5-labeled anti-CD3 (BD Biosciences, catalog 552852). The positive control was determined by utilizing known rhesus-reactive rhesus IgG1 antibody. MFI was normalized against pretransplant MFI and changes in MFI were measured over the course of transplantation.

Study approval. Both murine and nonhuman primate experimental subjects received humane care and treatment in accordance with Emory University IACUC guidelines, and all experimental protocols utilizing animals were conducted with approval by this institutional review board.

Statistics. Skin graft experiments and primate renal transplants are presented on Kaplan-Meier survival curves and were compared with log-rank test. All other assays were compared with the 1-tailed Mann-Whitney nonparametric test. Statistical analyses were conducted using GraphPad Prism. A *P* value less than 0.05 was considered significant.

Author contributions

WK designed many of the experiments, helped conduct many of the murine skin grafts and primate kidney transplants, analyzed the data, and wrote the manuscript. YD helped perform murine skin grafts and conduct some flow cytometric analyses. DM performed the gene expression analysis experiments. CB performed many of the flow cytometric analyses of primate data and the donor-specific antibody experiments. ES was the supervising veterinarian for all primate experiments. MF and CPL provided help with experimental design and manuscript revision. AA conceived of many of the experiments, assisted with analysis, helped perform primate kidney transplants, and assisted with manuscript editing. MEF provided the anti-OX40L monoclonal antibody and contributed to experiment design.

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