Cutting Edge: Immunological Consequences and Trafficking of Human Regulatory Macrophages Administered to Renal Transplant Recipients


*J Immunol* 2011; 187:2072-2078; Prepublished online 29 July 2011; doi: 10.4049/jimmunol.1100762

http://www.jimmunol.org/content/187/5/2072

Supplementary Material

http://www.jimmunol.org/content/suppl/2011/07/29/jimmunol.1100762.DC1

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Copyright © 2011 by The American Association of Immunologists, Inc. All rights reserved.
Print ISSN: 0022-1767 Online ISSN: 1550-6606.
Cutting Edge: Immunological Consequences and Trafficking of Human Regulatory Macrophages Administered to Renal Transplant Recipients


Regulatory macrophages (Mregs) were administered to two living-donor renal transplant recipients. Both patients were minimized to low-dose tacrolimus monotherapy within 24 wk of transplantation and subsequently maintained excellent graft function. After central venous administration, most Mregs remained viable and were seen to traffic from the pulmonary vasculature via the blood to liver, spleen, and bone marrow. By 1 y posttransplantation, both patients displayed patterns of peripheral blood gene expression converging upon the IOT-RISET signature. Furthermore, both patients maintained levels of peripheral blood FOXP3 and TOAG-1 mRNA expression within the range consistent with nonrejection. It is concluded that Mregs warrant further study as a potential immune-conditioning therapy for use in solid-organ transplantation. The results of this work are being used to inform the design of The ONE Study, a multinational clinical trial of immunomodulatory cell therapy in renal transplantation. The Journal of Immunology, 2011, 187: 2072–2078.

Conditioning the response of organ transplant recipients to donor alloantigens using cell-based therapies is now becoming a clinically feasible strategy, and, as the potential risks are better understood and minimized, such approaches are gaining credibility. Compared to longer established techniques, such as donor-specific blood transfusion and bone marrow transplantation, newer methods involving the ex vivo induction, expansion, and purification of tolerance-promoting cells offer the substantial advantages that the quality and dose of cell products can be tightly controlled. Moreover, by expansion or induction of cells in culture and by specifically purifying tolerogenic cell types, the number of viable suppressor cells administered to the patient can be maximized, and the dangers of sensitization and graft-versus-host reactions may be reduced.

Efforts in our laboratory to develop a cell product for promoting transplant tolerance in the clinical setting have focused on a type of suppressor macrophage, the human regulatory macrophage (Mreg) (1–7). Mregs exhibit a number of properties that might make them particularly suitable for clinical purposes, in particular, the cells are fully differenti- ated and potently T cell suppressive (8). Mregs derive from CD14+ peripheral blood monocytes in the absence of other cell types when monocytes are cultured for 6 d in medium supplemented with human AB serum before stimulation with IFN-γ for a further 24 h. Cell populations generated in this manner are homogeneously CD14+/low HLA-DR+CD80−/lowCD86+CD16−/lowCD64−/lowTLR2−/lowTLR4− and CD163+/low (8).

This work was supported in part by The ONE Study (European Union Seventh Framework Programme Grant 260687), the Reprogramming the Immune System for Establishment of Tolerance network (http://www.risetfp6.org), and the Deutsche Forschungsgemeinschaft (Grant GE-1188/1-1).

The sequences presented in this article have been submitted to the National Center for Biotechnology Information/Gene Expression Omnibus (http://www.ncbi.nlm.nih.gov/geo/) under accession number GSE24172.

Address correspondence and reprint requests to Dr. James A. Hutchinson, Department of Surgery, University Hospital Regensburg, Franz-Josef-Strauß-Allee-11, Regensburg 93053, Germany. E-mail address: james.hutchinson@klinik.uni-regensburg.de

The online version of this article contains supplemental material.

Abbreviations used in this article: 1-D/L-MT, 1-methyl-D/L-tryptophan; IOT, Indices of Tolerance; Mreg, regulatory macrophage; RISET, Reprogramming the Immune System for Establishment of Tolerance; SPECT, single photon emission computed tomography.

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*Laboratory for Transplantation Research, Department of Surgery, University Hospital Regensburg, Regensburg 93053, Germany; †The ONE Study Consortium of the European Union Seventh Framework Programme; ‡Institute for Medical Immunology, Berlin Charité University Hospital, Berlin 10117, Germany; ‡‡Miltenyi Biotec GmbH, Bergisch Gladbach 51429, Germany; ‡‡‡Tel-Express, Nantes 44200, France; †Department of Nuclear Medicine, University Hospital Schleswig-Holstein, Campus Kiel, Kiel 24105, Germany; ††Institute for Immunology, University Hospital Schleswig-Holstein, Campus Kiel, Kiel 24105, Germany; ‡Department of Visceral and Transplantation Surgery, Charité University Medicine, Campus Virchow, Berlin 13353, Germany; ‡‡Division of Nephrology, Department of Medicine, University Hospital Schleswig-Holstein, Campus Kiel, Kiel 24105, Germany; ‡‡‡Division of Nephrology, Department of Medicine, University Hospital Regensburg, Regensburg 93053, Germany; †Division of Nephrology, Department of Medicine, University Hospital Hamburg-Eppendorf, Hamburg 20246, Germany; ‡‡Division of Nephrology, Department of Medicine, University Hospital Frankfurt am Main, Frankfurt am Main 60590, Germany; ‡Laboratory of Immunobiology, Rene Descartes University, Pariss 75743, France; ‡‡Medical Research Council Center for Transplantation, King’s College London, London WC2R 2LS, United Kingdom; ‡‡‡Nuffield Department of Surgery, University of Oxford, Oxford OX3 9DU, United Kingdom; and ***Clinic for Applied Cell Therapy, University Hospital Schleswig-Holstein, Campus Kiel, Kiel 24105, Germany

Received for publication March 24, 2011. Accepted for publication July 5, 2011.

www.jimmunol.org/cgi/doi/10.4049/jimmunol.1100762
This report describes the treatment of two living-donor renal transplant recipients with preoperative infusions of donor-derived M regs. Both patients were successfully weaned to low-dose tacrolimus monotherapy within 24 wk of transplantation and remain in excellent clinical condition. Throughout their follow-up, the patients were intensively monitored using cutting-edge immunomonitoring assays provided by the European Union Framework Program 6 Reprogramming the Immune System for the Establishment of Tolerance (RISET) consortium, including flow cytometry, serial gene expression profiling, and analysis of TCR VB usage by peripheral blood T cells. The two patients are now 3 y posttransplantation with stable graft function and receive minimal maintenance immunosuppression.

Materials and Methods

Ethics

In a modification of the TAIC-II trial protocol (http://www.clinicaltrials.gov, NCT00223067), two patients were treated with M reg therapy at the discretion of two senior consultants as “individueller heilversuch” (individual healing attempts). The patients and their donors gave full, informed, written consent to the procedure and follow-up investigations. Human leukocytes for experimental use were obtained with approval of the local ethics committee (ethics vote 09/100) and informed consent of the donors.

In vitro characterization of human M regs

M regs were prepared according to published methods (1). For flow cytometry, harvested M regs were resuspended in ice-cold staining buffer (Dulbecco’s modified PBS, 1% BSA, and 0.1% NaN₃) and blocked for 30 min with 10% FCS block (Miltenyi Biotec) before staining with fluorochrome-conjugated primary Abs for 1 h. 7-aminominoacetylcysteine D exclusion was used for live/dead discrimination. To assess the T cell-suppressive capacity of M regs, CFSE-labeled CD3⁺ T cells and M regs were set in direct coculture for 5 d. Subsequently, T cell proliferation and absolute numbers were assessed by flow cytometry, as described elsewhere (1). The mechanism of M reg-mediated T cell suppression was investigated in direct 1:1 M reg/T cell cocultures. 1-methyl-D/L-tryptophan (1-D/L-MT; Sigma-Aldrich), an inhibitor of IDO, was completely dissolved in 5N HCl with gentle heating and agitation, before application.

Production of M regs for infusion into patients

M regs for administration to patients were prepared under strict GMP conditions according to an adaptation of a previously published method (1). Briefly, donor PBMC were obtained by leucapheresis and Ficoll density gradient separation. Plastic-adherent PBMC were plated at a density of 35 × 10⁶ monocytes/175 cm² culture flask (Cell+ T175 flask; Sarstedt) in 30 ml RPMI 1640-based medium without phenol red (Lonza) supplemented with 10% human AB serum (Lonza), 2 mM L-glutamine (Lonza), 100 U/ml penicillin, 10 µg/ml streptomycin (Lonza), and recombinant human M-CSF (R&D Systems) at a final concentration of 5 ng/ml carried on 0.1% human serum albumin (Aventis). The cells were cultured for 6 d with complete medium exchanges on days 1, 2, and 4. On day 6, cultures were stimulated with 25 ng/ml recombinant human IFN-γ (Inukin; Boehringer Ingelheim). On day 7, the adherent cell fraction was recovered by trypsin-FcR block (Miltenyi Biotec) before staining with fluorochrome-conjugated Abs and analysis by flow cytometry.

Clinical management of patients CA and MM

M regs were administered to patients CA and MM in the context of an immunosuppressive protocol comprising tacrolimus, steroids, and azathioprine. To reduce the risk of preoperative infusion of M regs sensitizing recipients, infusions were given under cover of 2 mg/kg/d azathioprine, commencing 3 d prior to administration of M regs and continuing for 8 wk postoperatively (9). M regs were administered 6 (MM) or 7 (CA) d prior to transplantation by central venous infusion. From the day of transplantation onwards, the patients were immunosuppressed with tacrolimus and steroids; steroids were weaned over the first 10 wk postoperatively, leaving the patients with maintenance tacrolimus monotherapy, aiming for trough serum levels between 4 and 8 ng/ml.

Radiological investigations

Radiological investigations were performed by the Department of Nuclear Medicine at Universitätsklinikum Schleswig-Holstein according to standard clinical protocols. For short-term cell tracking studies, patient MM received a total of 5 × 10⁶ M regs labeled with 45 MBq [⁹⁹mTc]-oxine (Coviden). Single photon emission computed tomography (SPECT) images were interpreted by an expert radiologist.

Gene expression profiling

Gene expression profiling data have been deposited in the National Center for Biotechnology Information/Gene Expression Omnibus (http://www.ncbi.nlm.nih.gov/geo/) with accession numbers GSE14655 and GSE24172.

Statistics

Reported values are mean ± SD in all cases. All n-values signify the number of independent biological replicates using cells from separate donors. Error bars shown in Fig. 1 represent SEM. A paired t test was used for all significance testing. Statistical treatment of microarray data is described in the text and figure legends.

Results

Production and quality control of clinical M reg preparations

A basic scientific understanding of the development, function, and possible immunological role of naturally occurring counterparts of the human M reg has lead to an optimized method of cell production for clinical purposes. The quality of M reg preparations produced by this method can be tightly specified in terms of cell surface phenotype, cell morphology, and potency in suppressing mitogen-driven T cell responses. Specifically, M regs are consistently CD14⁻/low-HLA-DR⁺/CD80⁻/high-CD86⁺/CD16⁻/low-HLA-DR+CD80⁺/low-CD86⁻/CD16⁺/low (Fig. 1A).

M regs suppress T cell responses in vitro

To assess the T cell-suppressive capacity of M regs, a flow cytometric assay was used to quantify mitogen-driven CD4⁺ and CD8⁺ T cell proliferation and, in parallel, to make absolute counts of T cells in direct coculture with M regs. T cells cultured alone did not proliferate and, by the fifth day of culture, 17 ± 6.5% of CD4⁺ and 22.1 ± 11.1% of CD8⁺ T cells had died spontaneously (Fig. 1B, 1C). Direct coculture of M regs with allogeneic T cells did not stimulate an MLR reaction, nor did M regs rescue cocultured lymphocytes. The strong proliferation of both CD4⁺ and CD8⁺ T cells observed after PHA stimulation was profoundly suppressed by coculture with allogeneic M regs (Fig. 1B). Strikingly, fewer CD4⁺ and CD8⁺ T cells remained in coculture with M regs when stimulated with PHA (p < 0.001 and p = 0.009, respectively), which points to an M reg-mediated elimination of activated T cells (Fig. 1C). This disappearance of T cells during M reg cocultures is most likely due to phagocytosis of dying T cells by M regs because examination of Mregs by transmission electron microscopy after coculture with T cells revealed the presence of numerous phagolysosomes (Supplemental Fig. 1). Suppression of T cell proliferation by M regs was observed at M reg/T cell ratios >1:8 (Fig. 1D). Inhibition of IDO activity with 1 mM 1-D/L-MT restored the ability of T cells cocultured with the presence of allogeneic M regs to proliferate in response to PHA (Fig. 1E).

Administration of M regs to patient MM

Patient MM, a 23-y-old female with renal failure owing to IgA nephropathy, received a kidney transplant from her 58-y-old mother, with whom she had single HLA-A and HLA-DR...
mismatches. CMV and EBV serology was negative in donor and recipient. Six days prior to transplantation, MM was given 4.3 \times 10^8 viable donor-derived Mregs (an equivalent of 8.0 \times 10^6 cells/kg) by central venous infusion under cover of 1 mg/kg/d azathioprine (Fig. 2A, Supplemental Fig. 2). Transplantation was without complications and initial graft function was satisfactory, in that serum creatinine levels had fallen to 1.4 mg/dl within 1 wk. Azathioprine was discontinued from week 8 postoperatively without adverse effect, and steroid therapy was discontinued by week 14, after which MM was maintained on tacrolimus monotherapy. Protocol biopsies at 8 and 24 wk posttransplantation showed no signs of rejection. Currently, MM is 3 y posttransplantation and in a stable clinical condition on tacrolimus 2 mg twice daily with trough levels of 4 to 5 ng/ml.

Profiling of peripheral blood TCR V\beta usage showed no mono- or oligoclonal T cell expansion, which is indicative of T cell nonreactivity to the transplanted kidney (Supplemental Fig. 3A) (10). Peripheral blood TOAG-1 mRNA expression levels are usually reduced during acute rejection of kidney allografts, but MM consistently maintained levels of TOAG-1 expression similar to healthy controls (Supplemental Fig. 3B) (11). No anti-HLA Abs, which were assayed on a monthly basis, were detected following the administration of Mregs.

### FIGURE 1.
Phenotypic and functional attributes of Mregs in vitro.

A. Mregs are uniquely identified by the CD14- HLA-DR+CD80-CD86+CD16+TLR2- and CD163-low phenotype. Expression ranges of these cell-surface Ags by Mregs were determined in cell preparations from six normal healthy donors (mean \pm SD). Unfilled trace, specific staining; filled trace, isotype control.

B. Over 5 d in direct 1:1 coculture, Mregs did not stimulate allogeneic T cell proliferation and were suppressive of PHA-stimulated T cell responses (n = 12).

C. Absolute quantification of CD4+ and CD8+ T cell numbers in 5-d 1:1 cocultures with allogeneic Mreg suggest that Mregs eliminated activated T cells.

D. Suppression of T cell proliferation by Mregs was observed at Mreg/T cell greater ratios >1:8.

E. 1-D/L-MT, an inhibitor of IDO, disinhibited the mitogen-stimulated proliferation of T cells in 1:1 cocultures with Mregs (n = 3). *p < 0.05.
trapped in the lungs, but after 2.5 h, cells were detected in circulation and had begun to accumulate in the liver and spleen. Twenty-two hours after M reg administration, the signal from the lung fields had much diminished, the cells having accumulated in the liver, spleen, and hematopoietically active bone marrow. The absence of signal from the patient’s urinary tract throughout the investigation indicates that the majority of infused M regs remained alive (12).

Administration of M regs to patient CA

Patient CA was an athletic 47- y-old male patient with severe chronic renal failure owing to nephrosclerosis as a consequence of long-standing arterial hypertension. CA was transplanted with a kidney from an unrelated 40- y-old living donor (Fig. 3A). Donor and recipient were fully HLA mismatched (donor: HLA-A3, 2; HLA-B41, 55; HLA-DR11, 13; recipient: HLA-A2, 2; HLA-B7, 35; HLA-DR4, 14), and both were CMV positive. One week prior to transplantation, a total of \(7.5 \times 10^8\) viable donor-derived M regs (equivalent to \(7.1 \times 10^6\) cells/kg) were administered to CA by slow central venous infusion (Supplemental Fig. 2). No evidence of impaired pulmonary perfusion caused by M reg administration was found (Fig. 3B). Seven days later, the patient underwent transplantation without complications. Serum creatinine levels gradually declined over the subsequent 10 wk to <2 mg/dl. Azathioprine treatment was stopped after the 8th week postoperatively, and steroids were discontinued by the 10th week, leaving the patient with tacrolimus as his sole maintenance therapy. Protocol biopsies taken at weeks 8, 24, and 52 showed no signs of acute cellular rejection, although occasional clusters of CD20+ B cells of unknown pathological significance were observed (data not shown). At 36 mo posttransplantation, CA had stable renal function with a creatinine of 1.43 mg/dl. Patient CA is being maintained on once-daily 5 mg sustained-release tacrolimus (Advagraf; Astellas Pharma) monotherapy; at 36 mo, his trough tacrolimus level was 2.7 ng/ml.

After administration of M regs, but prior to transplantation, the expression of TOAG-1 mRNA in the peripheral blood of CA increased 5.5-fold (Supplemental Fig. 3B). A similar increase in TOAG-1 was observed in patient MM after M reg treatment. As observed in MM, patient CA’s expression of FOXP3 mRNA began to rise from the fourth week postoperatively (Supplemental Fig. 3B), which corresponded...
to a small increase in the number of circulating Tregs observed by flow cytometry (Supplemental Fig. 4).

Discussion

Several alternative cell types are now approaching the point of preclinical development that might allow them to be properly investigated as adjunct immunosuppressive therapies in early-stage clinical trials, including certain tolerogenic DC subsets, regulatory T cells, mesenchymal stem cells, and Mregs. Yet clinical studies using cellular therapies in transplantation remain controversial because many in the field doubt the clinical practicality of such therapies and their safety. It is certainly true that administration of cell preparations to patients is not without potential complications, but these risks can be minimized by refining cell production and clinical monitoring of the recipient: in both these respects, we have learned much from the cases of MM and CA.

Should we be surprised by the clinical outcomes of MM and CA? Although a high proportion of renal transplant recipients with stable graft function several years after transplantation may be weaned to tacrolimus monotherapy, quite a different situation pertains to our patients. Both MM and CA underwent an early, rapid reduction in immunosuppression, such that both received tacrolimus monotherapy by 14 and 10 wk, respectively, and were further minimized to 4–6 ng/ml trough tacrolimus levels by 24 wk. At 3 y posttransplantation, patient CA is clinically stable with once-daily 5 mg sustained-release tacrolimus, and patient MM is being stably maintained with tacrolimus 2 mg twice daily. Despite their early minimization of immunosuppressive therapy, neither patient MM nor CA underwent an acute rejection episode during the 3-y follow-up period. Shapiro et al.’s study (14) remains a benchmark trial of minimized tacrolimus monotherapy in renal transplant recipients: In this study, 150 patients were treated with 5 mg/kg antithymocyte globulin with bolus prednisone as induction and were subsequently treated with tacrolimus monotherapy. Under this regimen, 37% of patients underwent acute rejection by 4 mo. After the fourth month, 113 patients underwent a stepwise minimization of tacrolimus dosing; these patients were followed up for a mean of 11 ± 5.4 mo, during which time 23% of patients underwent acute rejection. Other studies with the aim of establishing renal transplant patients on tacrolimus monotherapy after antithymocyte globulin induction (15) or alemtuzumab induction (16) achieved similar outcomes in terms of acute rejection rates and the proportion of patients tolerating monotherapy. Despite receiving no conventional induction therapy, neither MM nor CA underwent acute rejection or showed any signs of subclinical rejection at their last protocol biopsy. From clinical experience, for the patients to have tolerated such an abrupt weaning of

FIGURE 4. Using the RISET 2.0 Agilent custom microarray platform, serial analyses of gene expression in the peripheral blood of patients MM and CA were performed. The resulting dataset was then compared with that obtained from the IOT-RISET patients. Expression profiles of the 10 most discriminatory biomarkers of tolerance identified by the IOT study were extracted from the complete quantile normalized, log2-transformed dataset. From this data, pairwise correlation coefficients were calculated comparing the medians of the IOT-tolerant patient group with MM and CA at each separate time point, and heatmaps of reporter-wise median-centered log2 data were generated.

Throughout her postoperative course, the profile of gene expression displayed by patient MM was very similar to that of the tolerant patient group. Patient CA followed a different course, upregulating SH2D1B, H535T1, TCL1A, FCRL1, FCRL2, and CD79b and downregulating SLC8A1 and TLR5 only later in the follow-up period. In consequence, CA initially most resembled the group of chronically rejecting and stably immunosuppressed patients, but after week 32 closely resembled the clusters of patients classified as tolerant or healthy controls.

FIGURE 5. Heatmaps of reporter-wise median-centered log2 expression levels from the RISET 2.0 pilot study. Gene expression data were normalized and log-transformed (x-axis). The y-axis represents the 20 most discriminatory biomarkers of tolerance identified by the IOT-RISET study. Gray diagonal lines denote equal expression ratios. For visualization purposes, 10 clusters were considered; each cluster represents a group of patients with similar expression patterns.

Serial analysis of Indices of Tolerance marker gene expression in the peripheral blood of patients MM and CA

A pattern of gene expression in peripheral blood that correlates with a drug-free, tolerant state in kidney transplant recipients has been defined by the Indices of Tolerance (IOT) research network (13). Using the RISET 2.0 Agilent custom microarray platform (Agilent Technologies), serial analyses of gene expression in the peripheral blood of patients MM and CA were performed (Fig. 4). The dataset from MM and CA was then compared with the dataset obtained from the cohort of kidney transplant recipients studied by IOT. Expression profiles of the 10 most discriminatory biomarkers of tolerance identified by the IOT study were extracted from the complete quantile normalized, log2-transformed dataset. From this data, pairwise correlation coefficients were calculated comparing the medians of the IOT-tolerant patient group with MM and CA at each separate time point, and heatmaps of reporter-wise median-centered log2 data were generated.

The convergence over time of patient MM and CA’s marker gene expression upon the tolerance-associated IOT-RISET gene signature is best appreciated by considering pairwise Pearson correlation coefficients comparing the medians of the IOT-RISET drug-free, tolerant patient group with MM and CA at each sampling time point (bottom right panel).
immunosuppression is a surprising result and suggestive of attenuated antidonor reactivity.

Serial analyses of the immunological status of the two patients, especially by gene expression profiling, support the contention that the reactivity of MM and CA against their grafts was attenuated. As time progressed, the expression pattern of the IOT-RISET gene set in MM and CA tended toward that of a group of tolerant kidney transplant recipients or healthy untransplanted individuals (13). The frequency of individuals displaying the IOT-RISET gene signature among the cohort of stably immunosuppressed patients on calcineurin inhibitor monotherapy studied by IOT-RISET was only 14.7%. Considering that this group of patients were all 6 to 9 years after transplantation and had been purposefully selected for their stable clinical state, we can reasonably expect the frequency of patients displaying the IOT-RISET gene signature within the general pool of renal transplant recipients to be lower.

Could selection of MM and CA have biased our interpretation of their clinical outcomes? Patient CA was fully mismatched against his unrelated donor. Patient MM, who received a kidney from her 58-y-old mother, was at risk for being presensitized against noninherited maternal Ags (17). Therefore, neither donor-recipient pair could be said to be especially more likely to have a better outcome than the general pool of living-donor transplant recipients.

After systemic administration, human M regs could influence recipient responses against donor alloantigen by multiple mechanisms: M regs might have direct effects on alloreactive and regulatory T cells or may act indirectly through a systemic release of cytokines or alloantigen. At the cellular level, though, it has been shown that M regs profoundly suppress mitogen-stimulated T cell proliferation in vitro and that this activity is partly mediated by the tryptophan-catabolizing enzyme IDO. In mice, IDO is indispensable for maintaining maternal tolerance of allogeneic conceptuses (18) and participates in the establishment of allograft tolerance by adoptive transfer of T regs (19). Therefore, it is at least plausible that human M regs might influence responses to allografts through the action of IDO.

Central to any understanding of the possible mechanisms by which M regs might act is the question of whether they transiently or stably engraft in the recipient or die shortly after administration. Tracking studies with \(^{111}\)In-labeled cells in transiently or stably engraft in the recipient or die shortly after which M regs might act is the question of whether they influence responses to allografts through the action of M regs (19). Therefore, it is at least plausible that human M regs establish allograft tolerance by adoptive transfer of T erance of allogeneic conceptuses (18) and participates in the partly mediated by the tryptophan-catabolizing enzyme IDO.

Acknowledgments

The authors have no financial conflicts of interest.

References


