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Authors

Hua, Xiaoqin
Deuse, Tobias
Chen, Yi-Je
[et al.](#)

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The potassium channel KCa3.1 as new therapeutic target for the prevention of obliterative airway disease

Xiaoqin Hua¹, Tobias Deuse¹, Yi-Je Chen², Heike Wulff², Mandy Stubbendorff¹, Ralf Köhler³, Hiroto Miura⁴, Florian Länger⁵, Hermann Reichenspurner¹, Robert C. Robbins⁶, and Sonja Schrepfer^{1,6}

¹University of Hamburg, University Heart Center Hamburg, Transplant and Stem Cell Immunobiology (TSI) Lab, CVRC (Cardiovascular Research Center Hamburg) and DZHK (Deutsches Zentrum für Herz- und Kreislaufforschung, E.V., University Hamburg).

²University of California, Davis, Department of Pharmacology, California, USA.

³University of Southern Denmark, Cardiovascular and Renal Research, Institute of Molecular Medicine, Odense, DK; Aragon Institute of Health Sciences I+CS, Zaragoza, Spain.

⁴University of Nevada, Department of Physiology and Cell Biology, Reno, Nevada, USA.

⁵Institute of Pathology, Hannover Medical School, Hannover, Germany.

⁶Stanford University, School of Medicine, CT Surgery, Stanford, USA.

Abstract

Background—The calcium-activated potassium channel KCa3.1 is critically involved in T cell activation, as well as in the proliferation of smooth muscle cells and fibroblasts. We sought to investigate whether KCa3.1 contributes to the pathogenesis of obliterative airway disease (OAD) and whether knockout or pharmacological blockade would prevent the development of OAD.

Methods—Tracheas from CBA donors were heterotopically transplanted into the omentum of C57Bl/6J wild-type or KCa3.1^{-/-} mice. C57Bl/6J recipients were either left untreated or received the KCa3.1 blocker TRAM-34 (120mg/kg/d). Histopathology and immunological assays were performed on postoperative days (POD) 5 or 28.

Results—Subepithelial T cell and macrophage infiltration on POD 5, as seen in untreated allografts, was significantly reduced in the KCa3.1^{-/-} and TRAM-34 groups. Also, systemic Th1 activation was significantly, and Th2 mildly reduced by KCa3.1 knockout or blockade. After 28 days, luminal obliteration of tracheal allografts was reduced from 89±21% in untreated recipients

Address for Correspondence: Sonja Schrepfer, M.D., Ph.D., Department of Cardiothoracic Surgery, Stanford University School of Medicine, 300 Pasteur Dr., CVRB MC 5407, Stanford, CA 94305, USA, Tel: (650) 736-8592; Fax: (650) 725-3846, schrepfer@stanford.edu.

Authorship

T.D., H.W., R.C.R. and S.S. participated in research design

X.H., T.D., H.W., R.K., H.R., R.C.R., and S.S. wrote the paper

X.H., Y.C., M.S., F.L., H.M. and S.S. performed the research

X.H., T.D., Y.C., H.W., R.K. and S.S. participated in data analysis

H.W. and R.K. contributed the new reagent (TRAM-34) and the KCa3.1^{-/-} mouse strain

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none

to 53±26% (p=0.010) and 59±33% (p=0.032) in KCa3.1^{-/-} and TRAM-34-treated animals, respectively. The airway epithelium was mostly preserved in syngeneic grafts, mostly destroyed in the KCa3.1^{-/-} and TRAM-34 groups, and absent in untreated allografts. Allografts triggered an antibody response in untreated recipients, which was significantly reduced in KCa3.1^{-/-} animals. KCa3.1 was detected in T cells, airway epithelial cells and myofibroblasts. TRAM-34 dose-dependently suppressed proliferation of wild-type C57B/6J splenocytes, but did not show any effect on KCa3.1^{-/-} splenocytes.

Conclusions—Our findings suggest that KCa3.1 channels are involved in the pathogenesis of OAD and that KCa3.1 blockade holds promise to reduce OAD development.

Keywords

KCa3.1; Obliterative airway disease; TRAM-34; Chronic rejection; Heterotopic tracheal transplantation

Introduction

Obliterative airway disease (OAD) remains the most common chronic complication and major obstacle to the long-term graft survival in lung transplant recipients (1, 2). Alloimmunogenic T cell activation drives the development of fibroproliferative lesions, but the detailed pathogenesis of OAD remains incompletely understood (3–7). No specific or effective treatments have been developed yet.

Recent studies have shown that the intermediate-conductance Ca²⁺-activated potassium channel KCa3.1 plays an important role in Ca²⁺-signaling and T cell activation (8, 9). The KCa3.1 channel is composed of four α -subunits each containing 6 trans-membrane segments with calmodulin complexed to its C-terminus as calcium sensor (10). Opening of this channel due to elevated intracellular Ca²⁺ leads to K⁺ efflux, membrane hyperpolarization, and increases the driving force for store-operated Ca²⁺-entry through calcium-release activated calcium (CRAC) channels (11). The resulting increase in cytosolic Ca²⁺ turns on the calcineurin pathway and induces T cell activation. Interestingly, the expression of KCa3.1 increases from 5–35 channels per cell in resting T cells to 500 channels per cell in activated naïve and memory T cells (12), suggesting that activated T cells might be more sensitive to selective KCa3.1 blockade.

Pharmacological KCa3.1 blockade depolarizes T cells, reduces Ca²⁺-influx, and inhibits T cell proliferation and cytokine production *in vitro* (13–15), while *in vivo* studies have demonstrated that KCa3.1 blockers can prevent experimental autoimmune encephalomyelitis and anti-collagen antibody-induced arthritis in mice and contribute to the prevention of kidney graft rejection in rats (16, 17). Based on the additional involvement of KCa3.1 in smooth muscle cell and fibroblast proliferation and the efficacy of the KCa3.1 blocker TRAM-34 in models of restenosis (18, 19), atherosclerosis (20), and kidney fibrosis (21), KCa3.1 has also been proposed as a possible therapeutic target for cardiovascular diseases. However, whether the KCa3.1 channel could be considered as a novel therapeutic target for the prevention of OAD has not been investigated before.

Results

Tracheas from CBA donors were heterotopically transplanted into the greater omentum of C57Bl/6J mice. Recipients in the treatment group received TRAM-34 (120mg/kg/d, i.p.) for 5 days or 28 days. KCa3.1^{-/-} mice receiving grafts from CBA donors and C57Bl/6J receiving syngeneic grafts were used as control (see table 1).

5-day study

Inflammatory Cell Infiltration—In untreated heterotopic tracheal allografts harvested on POD5, massive F4/80+ macrophage and CD3+ T lymphocyte infiltration occurred in the subepithelial area (Fig. 1A). The degree of infiltration was two- to three-fold higher than in the KCa3.1^{-/-} and TRAM-34 groups ($p < 0.001$ for both CD3+ and F4/80+ cells, Fig. 1B). Only a few infiltrating cells were found in syngeneic grafts and the differences to the KCa3.1^{-/-} and TRAM-34 groups did not reach statistical significance.

Elispot—Elispot assays on POD 5 revealed that knockout (KCa3.1^{-/-}) or pharmacological blockade (TRAM-34) of the KCa3.1 channel resulted in decreased cellular immune activation. Spot frequencies for IFN- γ in the no medication group were significantly higher than those in the KCa3.1^{-/-} ($p = 0.007$) and TRAM-34 groups ($p = 0.015$; Fig. 1C). Spots were lowest in the syngeneic group ($p = 0.004$ vs. KCa3.1^{-/-}, $p = 0.002$ vs. TRAM-34, $p < 0.001$ vs. no medication). The IL-4 spot frequencies, representing the Th2 response, were significantly lower in the TRAM-34 than the no medication group ($p = 0.005$; Fig. 1D).

28-day study

Luminal obliteration—In the 28-day study, we observed myoproliferative tissue of high cellularity in the allogeneic no medication group causing luminal obliteration of $88.7 \pm 20.9\%$ (Fig. 2A). Tracheal grafts in the TRAM-34 and KCa3.1^{-/-} groups showed significantly reduced luminal obliteration ($p = 0.032$, $p = 0.010$; Fig. 2B). However, KCa3.1 blockade or knockout did not completely prevent obliteration ($p = 0.014$ and $p = 0.007$, respectively vs. the syngeneic group). Only syngeneic grafts presented without fibrotic tissue growth in the epithelial or subepithelial areas.

Donor-Specific Antibody (DSA) Assay—DSAs were evaluated on POD 28. The mean value of donor-reactive IgG in the no medication group was significantly higher than that of the syngeneic group ($p = 0.001$). DSAs of the KCa3.1^{-/-} group ($p = 0.018$) and the TRAM-34 group ($p = \text{ns}$) were lower than those of the no medication group (Fig. 2C).

Epithelial coverage—Tracheal allografts of the no medication group showed a complete loss of epithelial coverage. In contrast, the epithelium in the syngeneic group was well preserved for most of the luminal circumference, with 91.8% respiratory morphology ($p < 0.001$ vs. no medication). Syngeneic transplants using transgenic donors expressing firefly luciferase showed that the epithelium was donor-derived and not recipient-type ‘neo-epithelium’ (SDC, Fig. 1). In the KCa3.1^{-/-} and TRAM-34 groups, scattered areas of flattened cuboidal epithelia (7.8% and 10.1%, respectively) remained ($p = 0.453$ and $p = 0.356$, vs. no medication, respectively), but we did not observe any epithelium of respiratory type ($p < 0.001$ for both vs. no medication; Fig. 3).

mRNA-Expression of KCa3.1 in Tracheal Grafts—Semi-quantitative RT-PCR on POD 28 revealed similar amounts of KCa3.1 mRNA in whole grafts from the allogeneic no medication group and the syngeneic group (Fig. 4A), while KCa3.1 mRNA was hardly detectable in allografts of KCa3.1^{-/-} recipients ($p < 0.001$ vs. no medication; $p = 0.005$ vs. syngeneic). KCa3.1 mRNA amounts were significantly lower in the TRAM-34 group than in the no medication group ($p = 0.008$), which most likely reflected lower numbers of infiltrating KCa3.1-expressing mononuclear cells, as well as reduced luminal obliteration by KCa3.1-expressing fibroblasts.

KCa3.1 Protein Expression in Tracheal Grafts—In the no medication group, intense KCa3.1 staining was found in the subepithelial area, which was confined mostly to immune cell infiltrates. Within the luminal granulation tissue, we observed very intense staining of

fibroblast-like cells as well as T lymphocytes and macrophages. In the syngeneic group, KCa3.1-staining was most intense within the intact respiratory epithelium, which is in line with the reported physiological expression of the channel in this tissue (Fig. 4B). Significantly less KCa3.1 staining was observed in the KCa3.1^{-/-} and TRAM-34 groups, which showed mostly destroyed epithelium, little myoproliferation, and only a few infiltrating cells.

Side Effects—Mice of all groups recovered well from surgery and there were no significant differences in body weight over the study period (data not shown). The mice did not show any obvious signs of discomfort, or side effects arising from TRAM-34 treatment or KCa3.1 knockout. Complete blood counts and blood biochemistry (AST, ALT, creatinine, and BUN) were in the normal range in all groups (data not shown). To screen for epithelial toxicity of TRAM-34, native C57B/6J mice and C57B/6J recipients of syngeneic tracheal grafts were treated for 28 days with TRAM-34 (SDC, Fig. 2). We did not observe any epithelial damage in the native lung or GI tract, nor in syngeneic tracheal grafts demonstrating that TRAM-34 does not exhibit any epithelial toxicity despite KCa3.1 being expressed in epithelia

Proliferation Assay *in vitro*

In vitro proliferation of ConA-stimulated splenocytes from C57B/6J wild-type (WT) or KCa3.1^{-/-} mice under increasing concentrations of TRAM-34 is shown in Fig. 4C. In WT splenocytes, TRAM-34 dose-dependently suppressed proliferation ($p=0.007$ for 100nM, $p=0.006$ for 250nM, $p=0.0007$ for 1 μ M, and $p=0.0006$ for 5 μ M). However, the same TRAM-34 concentrations did not affect the proliferation of ConA-stimulated KCa3.1^{-/-} splenocytes, confirming that the TRAM-34 effect was mediated through inhibition of KCa3.1 and not through an unspecific off-target effect.

KCa3.1 in human OAD

To assess the relevance of the KCa3.1 channel in human disease, tissue specimens retrieved from lung transplant patients with OAD were studied (SDC, Fig. 3). KCa3.1 staining was abundant in human lung tissue, most prominent in the epithelium and myoproliferative areas.

Discussion

Based on previous studies showing that KCa3.1 is involved in the activation and proliferation of inflammatory cells (14, 22) the channel has been proposed as a novel target for immunomodulation (8). In this study, we demonstrate that KCa3.1 is also involved in the pathogenesis of OAD, and that KCa3.1 blockade or knockout slows disease progression.

KCa3.1 protein expression was observed in different cell populations of tracheal grafts, mainly infiltrating mononuclear cells, proliferating myofibroblasts, and respiratory epithelium. Semi-quantitative RT-PCR showed only negligible amounts of KCa3.1-mRNA in the KCa3.1^{-/-} recipients. In keeping with the previously reported increased expression of KCa3.1 in activated T cells (23), we observed high amounts of KCa3.1-mRNA and intensive KCa3.1 protein staining on inflammatory cells in the no medication group demonstrating marked KCa3.1 channel up-regulation in this allogeneic transplant setting. Compared to the no medication group, KCa3.1-mRNA expression was significantly lower in the TRAM-34 group, most likely due to lower numbers of KCa3.1-expressing inflammatory cells (14), destruction of the epithelium, and reduced myoproliferation.

OAD development has been shown to be mediated by alloimmune-activated T cells (24–26). Macrophages play a role in further recruiting inflammatory cells (27, 28) and producing pro-proliferative cytokines and growth factors (29, 30). We observed dense CD3⁺ T cell and F4/80⁺ macrophage infiltration in the subepithelial areas of untreated allografts, which were significantly reduced in both the KCa3.1^{-/-} and TRAM-34 groups. Also, IFN-γ Elispot frequencies, reflecting the degree of cellular immune activation, were significantly reduced in both groups. IFN-γ increases the expressions of MHC-I and -II, adhesion molecules, and co-stimulatory ligands on APCs after lung transplantation (31, 32), and is considered a central cytokine in cellular rejection. Pharmacologic blockade or knockout of KCa3.1 alone was sufficient to mitigate allo-immune Th1 activation. However, it was reported that the functions of Th17 and regulatory T cells in KCa3.1^{-/-} mice seemed unchanged (33). Furthermore, while Ca²⁺ influx and IL-2 production following TCR ligation was reduced in KCa3.1^{-/-} T cells, the absolute numbers of peripheral T cells and the CD4/CD8 ratio, as well as the macroscopic appearance of all lymphoid organs was normal. KCa3.1-deficiency may either be compensated for by the up-regulation of other channels or KCa3.1 may not be crucial for maintaining T cell numbers. The specificity of TRAM-34 for KCa3.1 was confirmed *in vitro*. Proliferation assays showed that TRAM-34 dose-dependently reduced cell proliferation in WT, but not KCa3.1^{-/-} splenocytes.

The DSA assays revealed reduced donor-specific IgG production in the KCa3.1^{-/-} group compared to untreated animals. Cumulating evidence suggests that allogeneic antibody responses play an important role in acute lung graft rejection as well as in the development of OAD. Lung transplant recipients with pre-existing anti-HLA antibodies show a significantly higher risk for early graft dysfunction and have a poor prognosis (26, 34). The reduced antibody response of the KCa3.1^{-/-} group may result from impaired T cell-mediated B cell activation.

The donor airway epithelium is considered to be the primary target for the allogeneic immune response (35), and epithelial injury plays a pivotal role in triggering rejection and OAD formation (36, 37). Preserved epithelial coverage of the airway lumen was found to slow the progression of OAD (7, 38). In our study, no airway epithelium was found in untreated animals after 28 days. In KCa3.1^{-/-} and TRAM-34-treated animals, there were some scattered, flattened cuboidal epithelial cells visible, which occupied less than 10% of the whole circumference, while the syngeneic group exhibited well preserved ciliated epithelium. It therefore seems that neither KCa3.1 channel knockout nor blockade protect the epithelium from being destroyed or facilitate epithelial recovery. Because this is a non-ventilated model, the airway epithelium is more prone to injury and destruction and our results may exaggerate the epithelial damage with KCa3.1 blockade. In the clinical setting and ventilated trachea transplant models, the airway epithelium is preserved despite OAD development (39).

TRAM-34 treatment induced no significant side-effects in keeping with previous studies (18–20, 40). We also did not observe any differences between the KCa3.1^{-/-} mice and the WT animals before or after transplantation and the KCa3.1^{-/-} mouse strain is viable and fertile (41). Senicapoc, another potent and selective KCa3.1 blocker, already passed through clinical phase I to III trials for sickle cell disease and proved efficacy in its biological endpoint of reducing hemolysis (42, 43). Although, senicapoc did not prevent clinically-relevant vaso-occlusive pain crises, these studies showed that KCa3.1 blockade was safe and well-tolerated in humans (44).

Human OAD is a form of chronic lung allograft dysfunction from allo- and innate immune injury, autoimmunity, environmental pathogens, and contributing conditions like acid reflux disease. Its aggressiveness and clinical course are likely related to the extent of immunologic

and non-immunologic insults (45). It is pathohistologically characterized by collagen-rich myoproliferative tissue that progressively obliterates the small airway lumen in the terminal and respiratory bronchioles (46, 47). Current treatment strategies involve switching to other calcineurin (48) or mTOR inhibitors (49), the addition of azithromycin (50) or statins (51), and extracorporeal photopheresis (52). Some patients with more active immune responses seem to be better amenable to the treatment of OAD, while the disease progresses steadily despite aggressive therapy in others. Although the heterotopic murine tracheal transplant model may not well represent human OAD in its whole complexity, it nicely mimics immune activation, graft infiltration, epithelial damage, and mice also mount a DSA response. The lack of environmental and viral challenges as well as the lack of airway clearance from secretions, however, somewhat restrict the transferability of results to the lung transplant setting.

To link our results to the human disease, we demonstrated similar KCa3.1 channel expression in mouse and human lung tissue. In accordance with the murine distribution, human KCa3.1 expression was most prominent in epithelial and myofibrotic cells. KCa3.1 blockade might therefore also benefit patients with OAD in the clinic. Because KCa3.1 channel blockers are mild immunosuppressants that double up as anti-proliferative agents, they are unlikely to replace current immunosuppressive agents, but may be useful additions for long-term maintenance therapy.

In conclusion, the present study demonstrates that KCa3.1 blockade or knockout significantly decreases T cell activation and reduces the development of OAD suggesting KCa3.1 blockade as an additional maintenance strategy for patients after lung transplantation.

Materials and Methods

Animals

Male CBA mice were used as allogeneic and C57Bl/6J mice as syngeneic trachea donors, and male C57Bl/6J mice, or KCa3.1^{-/-} mice on C57Bl/6J background, were used as recipients. Mice weighing 25 to 35 g were purchased from Charles River Laboratories (Sulzfeld, Germany) and KCa3.1^{-/-} mice derived from our own breeding colonies were genotyped as described previously (53). All animals received humane care in compliance with the guide for the principles of laboratory animals, prepared by the Institute of Laboratory Animal Resources, and published by the National Institutes of Health. The animals were maintained in the animal care facilities of the University Hospital Hamburg-Eppendorf.

Heterotopic Tracheal Transplantations, Graft Recovery, and Tissue Processing

The heterotopic tracheal transplant model in mice was chosen because of its reliability in presenting the characteristics of OAD pathogenesis and its high reproducibility (54, 55). Transplantations were performed as previously described (55). For analysis, tracheal grafts were recovered from the greater omentum and cut into two segments. One segment was fixed in 4% paraformaldehyde solution, dehydrated, and embedded in paraffin. The second segment was snap-frozen in liquid nitrogen and stored at -80°C.

Experimental Groups

Eight groups were involved in this study (see table 1). All animals were randomly assigned to one of the groups prior to tracheal transplantation and there were no animal deaths due to technical failures. Mice in groups 4 and 8 were treated with 120mg/kg TRAM-34 intra peritoneally (18, 56) once daily. TRAM-34 was freshly dissolved in Miglyol 812 (Pharmacy

of University Hospital Hamburg-Eppendorf) prior to use. Grafts were recovered 5 days after transplantation to investigate cellular rejection and immune activation (groups 1–4) or 28 days after transplantation for the study of OAD (groups 5–8).

Side Effect Screening

Blood was drawn during graft recovery for complete blood count, liver, and kidney toxicity. All recipients were examined daily for signs of discomfort or diarrhea.

Histology

General Histology—Sections of 5µm were cut and stained with hematoxylin and eosin (H&E) or Masson-Goldner trichrome. Histologic analyses were done using image-processing software (Leica, Bensheim, Germany). Luminal obliteration was quantified as previously described (57). Since the tissue inside the tracheal cartilage contains submucosal and epithelial tissue, the value for luminal obliteration in native tracheas is approximately 10% (58). The airway epithelium was classified as intact respiratory epithelium or flattened cuboidal epithelium. By computerized morphometry, the length of the respective epithelium along the whole luminal circumference was measured and calculated in percent.

Mononuclear infiltrating cells were identified using monoclonal antibodies against CD3 (DakoCytomation, Glostrup, Denmark) and the macrophage marker F4/80 (Clone BM8, BMA Biomedicals, Augst, Switzerland). Antigens were retrieved in epitope-retrieval solution (Dako, pH 9) by the heating method in a steamer, or by trypsin digestion (Dako), respectively. After overnight incubation with the primary antibody at 4°C, the biotinylated rabbit-anti-rat secondary antibody (Dako) was used if necessary. These antibodies were further bound with alkaline phosphatase conjugated enzyme polymer (AP Polymer Kit, Zytomed Systems, Berlin, Germany) and finally visualized by New Fuchsin Substrate System (Dako). Sections were counterstained with hematoxylin. Three high power fields were analyzed per animal to quantify cell infiltration (expressed as cells/mm²).

KCa3.1 channel expression—After antigen retrieval in sodium citrate (10mM, pH 6), KCa3.1 channel proteins were identified by an anti-KCNN4 antibody (Sigma-Aldrich, St. Louis, MO). A biotinylated secondary antibody was further labeled by horseradish peroxidase-conjugated avidin complex (Vector Laboratories, CA) and finally visualized by DAB (Vector Laboratories) as previously described (40).

Elispot Assays

The cellular immune response was investigated on POD5. Recipient spleens were harvested and splenocytes were isolated and freshly used. Elispot assays using 1×10⁵ mitomycin-inhibited donor splenocytes as stimulator cells and 1×10⁶ recipient splenocytes as responder cells were performed according to the manufacturer's protocol (BD Biosciences, CA) using IFN- and IL-4-coated plates. Duplicates or quadruplicates were done for each animal. Spots were counted automatically by an ELISPOT plate reader (CTL, St. Louis, OH) for scanning and analyzing.

DSA Assay

On POD28, sera from recipient mice were decompartmented by heating to 56°C for 30 min. Equal amounts of sera and donor splenocyte suspensions (5×10⁶/ml) were incubated for 30 min at 37°C. Bound allogeneic IgG antibodies were labelled with the FITC-conjugated goat anti-mouse IgG F(ab')₂ fragment (Sigma-Aldrich) and analyzed by flow cytometry (BD Bioscience).

KCa3.1 mRNA-Expression

mRNA was extracted and purified from tracheal transplants using the RNeasy Mini Kit (Qiagen). Semi-quantitative RT-PCR was conducted using iQ™ CYBR® Green Supermix (Bio-Rad) and a Stratagene MX3000p cyclor. Primer sequences were as follows: mKCa3.1: F, GTGGCCAAGCTGTACATGA; R, GCCACAGTGTGTCTGTGAGG; mGAPDH: F, CAATGAATACGGCTACAGCAAC; R, AGGGAGATGCTCAGTGTGG. Expression levels were normalized to the house keeping gene, glyceraldehyde-3-phosphate dehydrogenase (GAPDH) as reference gene by calculating the Ct values (Ct = Ct(target) – Ct(GAPDH)) and the ratios to GAPDH (ratio = 2^{-Ct}). Results were expressed as percentage of GAPDH (% GAPDH).

Proliferation Assay *in vitro*

Proliferation assays were performed by stimulating 1×10⁵ splenocytes from either C57Bl/6J WT or KCa3.1^{-/-} mice with 5µg/ml ConA (Sigma-Aldrich). Cells were incubated with different concentrations of TRAM-34 in flat-bottom 96 well plates for 48 hours. [³H]-TdR incorporation was measured, normalized to ConA-activated controls, and the background was subtracted.

Statistical analysis

Data are presented as mean ± standard deviation. Comparisons between groups were performed by analysis of variance (ANOVA) between groups with Fisher's Least Significant Difference (LSD) post hoc tests. Probability values (p) smaller than 0.05 were considered significant. Statistical analysis was performed using the SPSS statistical software package 17.0 for Windows (SPSS Inc., Chicago, IL).

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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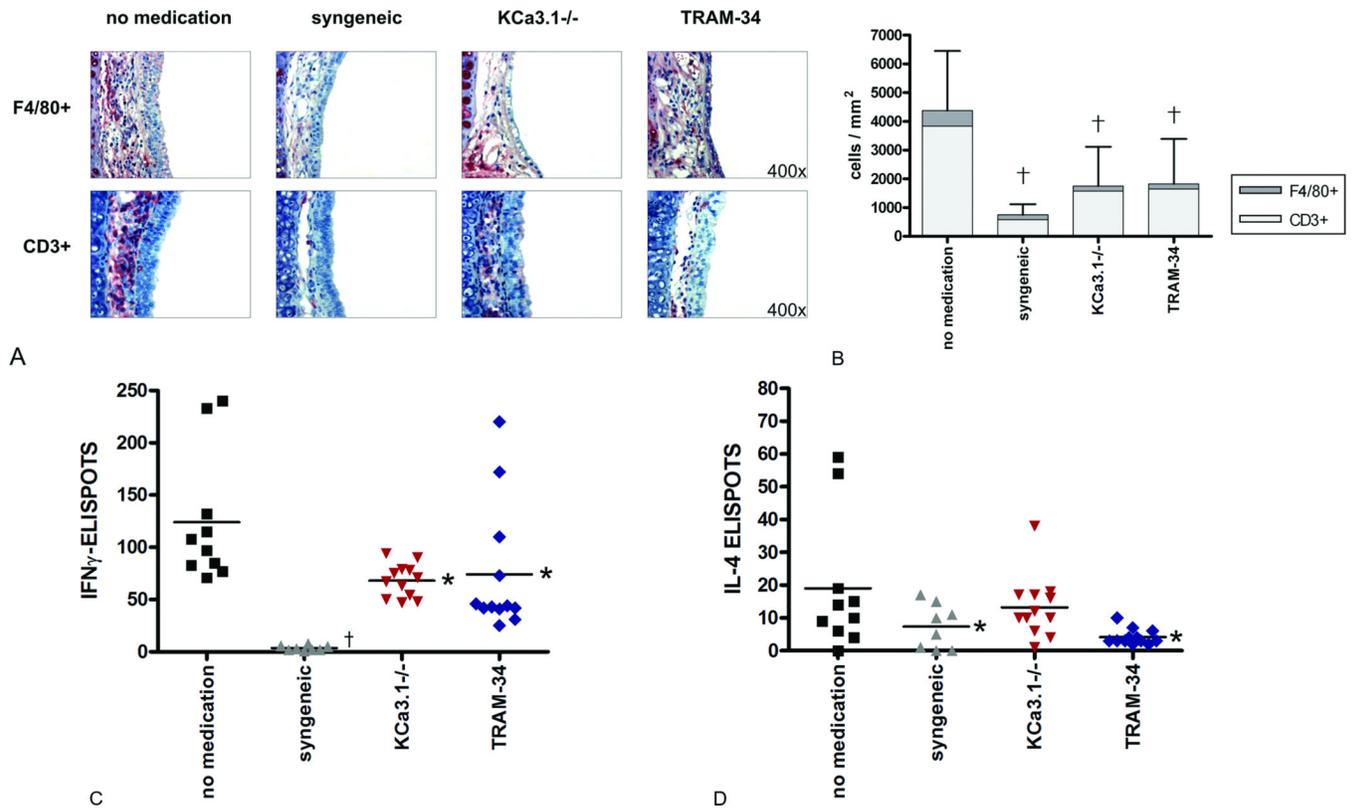


Figure 1. Graft infiltrating cells and systemic cellular immune response

Graft infiltration of F4/80+ macrophages and CD3+ lymphocytes within the subepithelial area on POD5 is shown by immunohistochemistry (A; magnification 400 \times). Mean numbers of F4/80+ macrophages (N=5 for no medication, N=6 for syngeneic, N=6 for KCa3.1^{-/-}, N=5 for TRAM-34) and CD3+ T cells (N=4 for all groups) are expressed as cells/mm² (B; †p<0.001 vs. no medication). On POD5, Elispot assays revealed attenuated systemic responses of IFN- γ producing Th1 cells (C) and IL-4 producing Th2 cells (D) in the KCa3.1^{-/-} (N=4) and TRAM-34 (N=5) groups (*p<0.05 vs. no medication; †p<0.001 vs. no medication (N=3), N=3 for syngeneic).

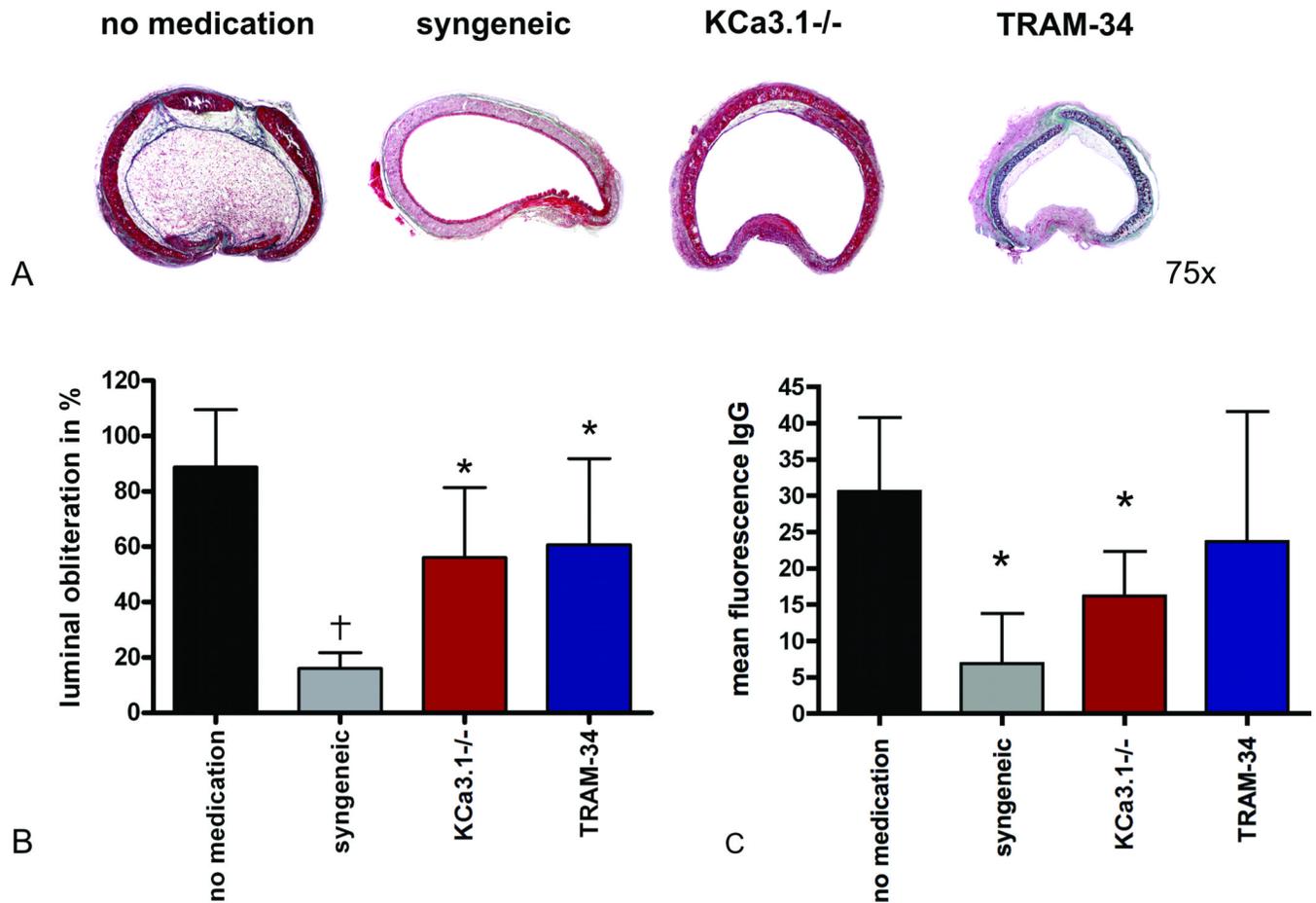


Figure 2. Histopathology, luminal obliteration, and donor-reactive antibodies

Representative cross sections of tracheal grafts on POD 28 stained with Masson-Goldner Trichrome at a magnification 75 \times are depicted (A). The average percent luminal obliteration is shown (B; * $p < 0.05$ vs. no medication; † $p < 0.001$ vs. no medication; N=7 for no medication and TRAM-34, N=5 for syngeneic, N=8 for KCa3.1 $^{-/-}$). Mean fluorescence of IgG demonstrates a significant reduction in DSAs in the syngeneic (N=5) and KCa3.1 $^{-/-}$ (N=9) groups (C; $p = 0.001$ and $p = 0.018$, respectively, vs. no medication (n=7); N=7 for TRAM-34).

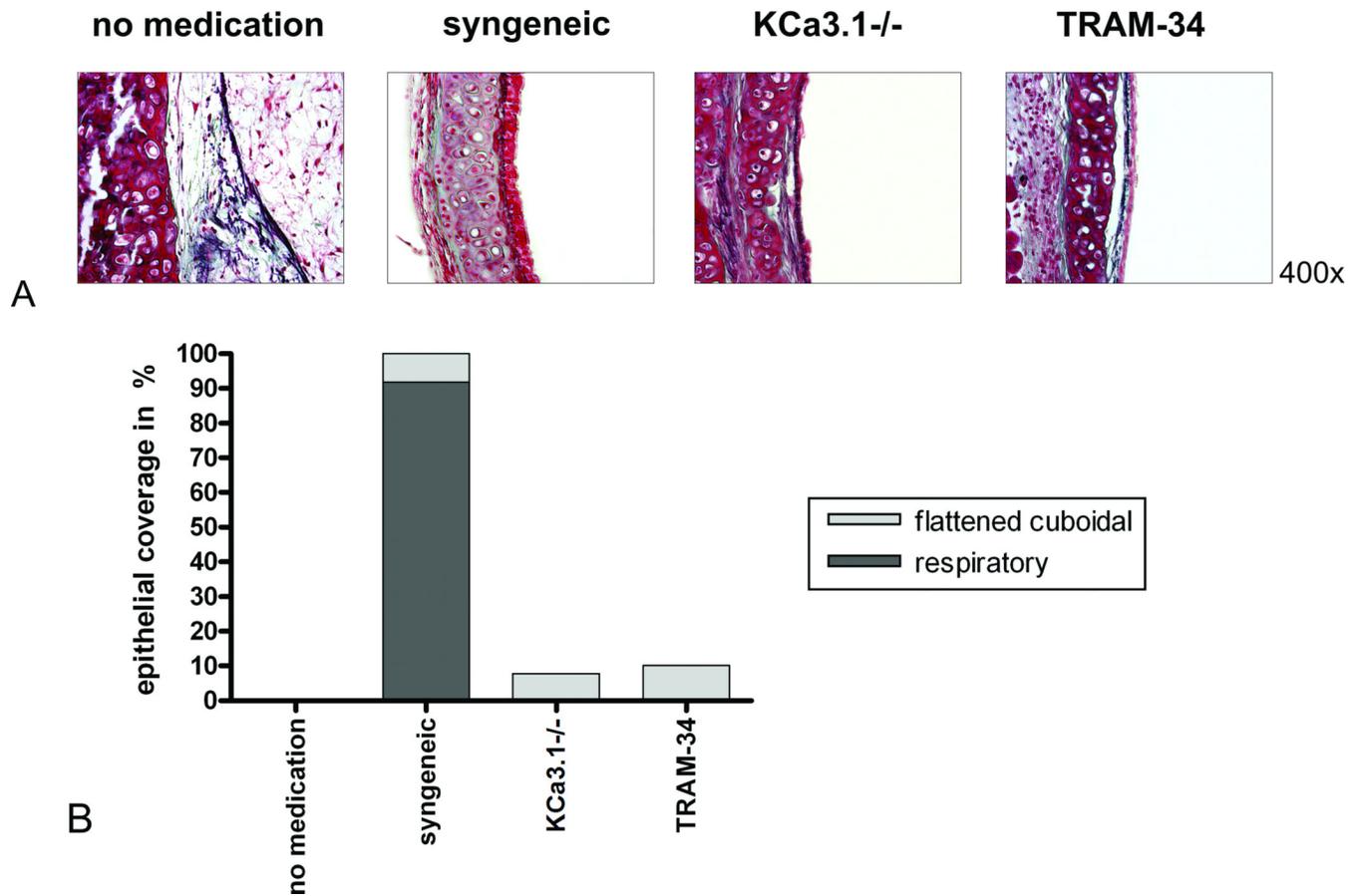


Figure 3. Airway epithelium

Representative sections of the tracheal epithelia on POD 28, stained with Masson-Goldner trichrome, are shown at a magnification of 400× (A). The respiratory epithelium of the syngeneic group (N=5) is widely preserved (B). The KCa3.1^{-/-} (N=10) and TRAM-34 (N=8) groups show cuboidal or flattened epithelia, whereas the epithelium in the no medication group (N=8) is totally destroyed.

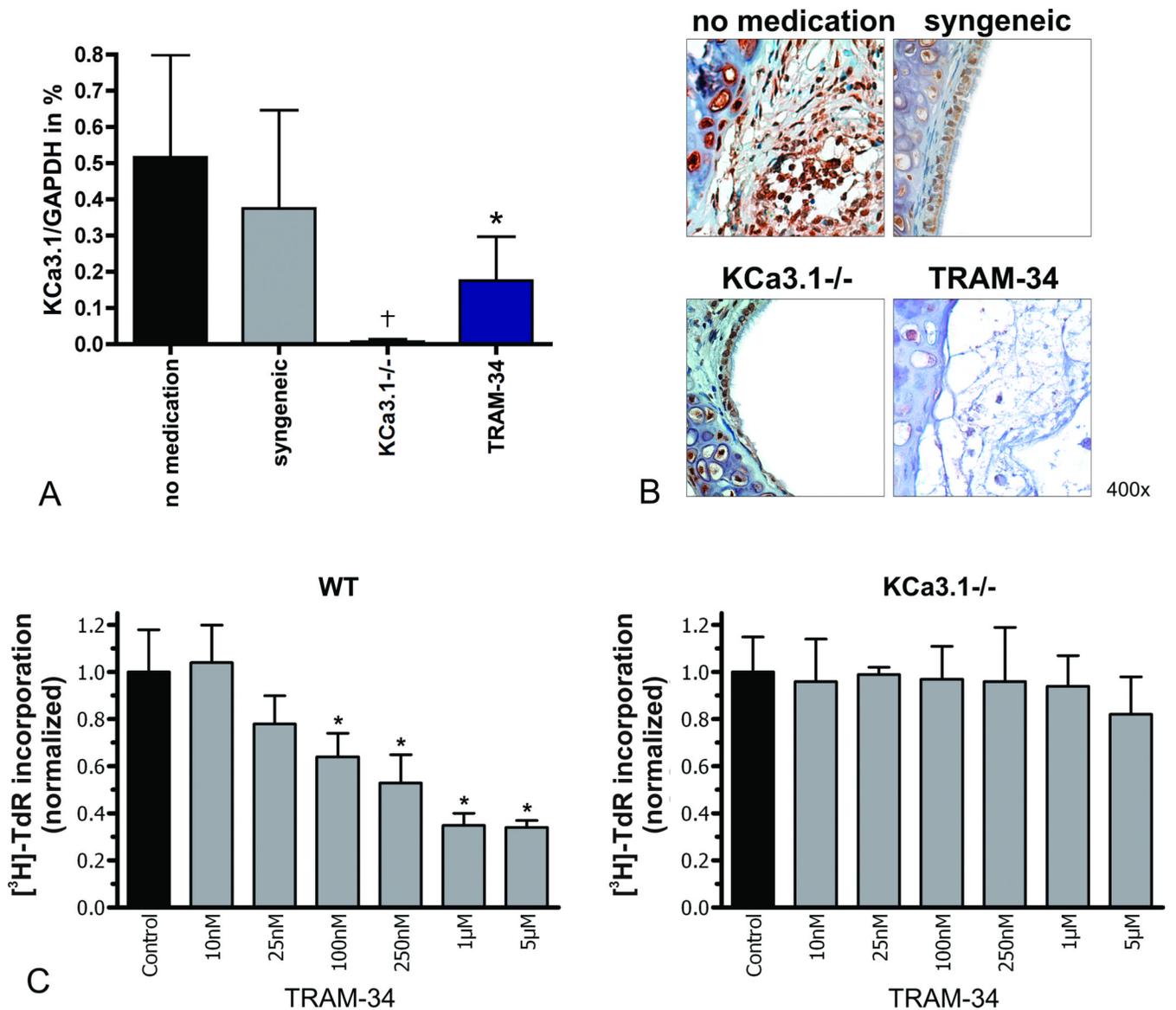


Figure 4. KCa3.1 expression and in vitro proliferation assay

KCa3.1 mRNA-expression in tracheal grafts was analyzed by semi-quantitative RT-PCR (A; * $p < 0.008$; † $p < 0.001$ vs. no medication (N=6), N=4 for syngeneic, N=8 for KCa3.1^{-/-}, N=5 for TRAM-34). Representative stainings for the KCa3.1 channel in tracheal graft are shown at a magnification of 400 \times (B). The results of the *in vitro* proliferation assay for WT or KCa3.1^{-/-} splenocytes are shown as [³H]-TdR incorporation normalized to the ConA-stimulated controls (C; * $p < 0.05$ vs. controls).

Table 1

Study groups

Grafts were recovered on POD5 in groups 1–4 to investigate acute rejection and immune activation, or after 28 days in groups 5–8 to assess OAD development

Study No.	Group	Donor	Recipients	Treatment
<i>Grafts recovered on POD5</i>				
1	no medication	CBA	C57Bl6	-
2	syngeneic	C57Bl6	C57Bl6	-
3	knockout	CBA	KCa3.1 –/–	-
4	TRAM-34	CBA	C57Bl6	TRAM-34
<i>Grafts recovered on POD28</i>				
5	no medication	CBA	C57Bl6	-
6	syngeneic	C57Bl6	C57Bl6	-
7	knockout	CBA	KCa3.1 –/–	-
8	TRAM-34	CBA	C57Bl6	TRAM-34