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Regulatory T cell expansion prevents retinal degeneration in type 2 diabetes

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Abstract

Background The global incidence of type 2 diabetes (T2D) is rapidly increasing, with retinopathy being its most common complication and a leading cause of preventable blindness. Although the precise mechanisms involved in the development of diabetic retinopathy (DR) are not fully understood, defective immunomodulation is a recognized key factor in its pathophysiology. Regulatory T cells (Treg) regulate inflammation and promote regeneration, and while they are known to have important anti-inflammatory and neuroprotective roles in other tissues, including central nervous system, their role in the diabetic retina remains largely unknown. The aim of the present study is to examine the effect of Treg expansion of retinal neurodegeneration, an early event in the pathogenesis of DR.

Methods Treg expansion was achieved by co-injecting recombinant mouse IL-2 with anti-IL-2 monoclonal antibody or its isotype in db/db mice as an established model of T2D. Treg expansion was confirmed via flow cytometry in blood, spleen, and retina. Fundus angiography was performed in the days prior to animal sacrifice at 18 weeks. To study the effect of Tregs on retinal neurons, glia and vascular permeability, immunohistochemistry against Cone-Arrestin, PKC α , synaptophysin, ChAT, TH, GFAP, Iba-1, calbindin, Brn3a, RBPMS, isolectin B4, and albumin was used. Retinal VEGF levels were measured with a magnetic bead-based immunoassay, and NLRP3, Casp1, p20 and IL-18 were analyzed by Western Blot in retinal homogenates.

Results There was a significant decrease in Treg in db/db mice blood. When this deficiency was corrected in db/db mice by systemic Treg expansion, there was an effective protection against retinal neurodegenerative, gliotic, inflammatory changes and vascular leakage associated with T2D. Importantly, Treg expansion did not impact the T2D phenotype in db/db mice as evaluated by blood glucose, HbA1c and circulating insulin.

Conclusion Treg modulation in T2D offers a promising therapeutic approach to prevent early stages of DR. This strategy focuses on reducing neuroinflammation and mitigating the associated neuronal, glial, and vascular degenerative changes characteristic of DR.

Keywords Regulatory T cells, Diabetic retinopathy, Immunomodulation, Retina, Neurodegeneration, Inflammation, Type 2 diabetes

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Introduction

T2D is one of the most common metabolic diseases in western societies and it is considered “the epidemic of the 21st century”: affecting 1 in 11 people in Europe [1]. DR is a common T2D complication and the leading cause of preventable visual impairment in the working-age population [2], accounting for 15–17% of total blindness in Europe [3]. Current DR treatments, such as laser photocoagulation, intravitreal injections of corticosteroids or anti-vascular endothelial growth factor (VEGF) agents, are focused on the late-stages of the disease. Besides, IV agents have shown limited efficacy, significant risk of side-effects and they are very expensive. Therefore, new and more effective therapeutic strategies that limit the escalation of DR from its early stages and thereby prevent blindness and its associated social and economic burden, are an urgent need for the global healthcare.

DR is currently considered a highly specific neurovascular disease rather than merely a microvascular disease [4]. From a pathophysiological point of view, special emphasis has been placed on the retinal Neurovascular Unit (NVU) [5], which consists of vascular elements, basement membrane, neurons, glial cells (Müller glia, astrocytes) and immune cells such as microglia and perivascular macrophages. It is widely accepted that chronic inflammation is the main common pathogenic driver in the natural course of T2D [6] and its complications, including DR [7–9]. Chronic low-grade inflammation leads to a maladaptive innate immune system, contributing to DR development via infiltration of circulating immune cells and serum proteins into the retina [8–10]. This immune infiltration drives localized retinal inflammation mediated by microglia, Müller cells and endothelial cells, which in turn leads to NVU damage and subsequent neurodegeneration [7, 8, 11]. Glial cell activation and the subsequent release of pro-inflammatory and apoptotic cytokines and angiogenic factors is one of the most important underlying mechanisms of this prolonged neuroinflammation, which exacerbates oxidative stress and retinal neurovascular damage [7, 12].

Treg are a key component of the adaptive immune system and play a vital role in immune tolerance due to their potent immunosuppressive properties [13–15]. Treg migrate into tissues and dampen inflammation, suppressing other adaptive or innate immune cells [14]. In recent years, new functions of Treg beyond their immune regulatory properties have been discovered, with a growing evidence for Treg regenerative and neuroprotective functions in various tissues, including the central nervous system (CNS). Treg promote remyelination in models of multiple sclerosis (MS) [16], while in Alzheimer’s disease (AD) and Parkinson’s disease (PD), Treg restore homeostasis and prevent cognitive decline (CD) by eliminating neuroimmunogens, polarising microglia, modulating

astrocytes and restoring normal neuronal function [13, 15]. Furthermore, the enhancement of astrocyte-mediated interleukin (IL)-2 production to locally expand Treg numbers in the CNS prevents inflammatory damage and facilitates repair following traumatic brain injury and age-associated CD [17, 18].

Given these neuroprotective, pro-regenerative and immunomodulatory roles, the beneficial effects of Treg expansion are currently under investigation in numerous clinical trials for AD, PD or MS [15, 19]. Since immunomodulation plays a significant role in the pathogenesis of DR it is plausible that Treg expansion may also be beneficial in this neurodegenerative disease [7–9].

Here, we provide experimental evidence that Treg expansion in early T2D rescues T2D-associated retinal pathology, preventing retinal neurodegeneration and reactive gliosis and reducing inflammasome activation. Our findings demonstrate that the well-established immunomodulatory role of Treg can be harnessed to prevent early-stage damage in DR and may help slow or prevent neurodegeneration, representing a new therapeutic strategy for treating this devastating complication of T2D.

Materials and methods

Animals

Diabetic male db/db (BKS.Cg-Dock7m $+/+$ Leprdb/J) mice and their control, non-diabetic mice (db/+; (BKS.Cg-Dock7m+Leprdb/+)) were acquired from Charles River Laboratories (Calco, Italy) at 6 weeks of age. The mutated leptin receptor carried by db/db mice gave rise to an obesity-induced type 2 diabetes phenotype. Mice were bred and maintained in the animal facilities of the Vall d’Hebrón Research Institute (VHIR). With the aim of minimizing variability, mice were randomly distributed (block randomization) into groups of two mice per cage in Tecniplast GM-500 cages (36 cm \times 19 cm \times 13.5 cm) under standard laboratory conditions at 22 ± 2 °C, with relative humidity of 50–60% and cycles of 12 h light/darkness. The cages were equipped with nesting material, absorbent bedding (BioFresh Performance Bedding 1/800 Pelleted Cellulose, Absorption Corp, Ferndale, WA, USA), ad libitum food (ENVIGO Global Diet Complete Feed for Rodents, Mucedola, Milan, Italy), and filtered water. Glycemia and glycosylated haemoglobin A1c (HbA1c) were measured through tail-vein blood sampling and detection with Bayer A1C Now+ (Cat. No. 08842610) and a blood glucose meter (71371-80, FreeStyle Optium Neo; Abbott).

All animal experiments were conducted in agreement with the European Community (86/609/CEE) and the guidelines of the Association for Research in Vision and Ophthalmology (ARVO) for the utilization of laboratory

animals. The Animal Care and Use Committee of VHIR (CEEA 14/21) authorized the present study.

Expansion

The expansion of endogenous Treg was performed as previously described [20] through the injection of IL-2/anti-IL-2 mAb (JES6-1) complexes. Briefly, recombinant mouse IL-2 (PeproTech, 1 µg per animal; Cat. No. 212-12) was mixed with either anti-IL-2 monoclonal antibody (mAb) (clone JES6-1; BE0043; Abyntek; 5 µg per animal, referred as Treg group) or its isotype control (rat IgG2a, referred to as “isotype” group from here on), and incubated at 37 °C for 30 min. 9-week-old mice were injected intraperitoneally (i.p.) in a final volume of 200 µL. Mice received a daily injection for 3 consecutive days on the first week and then a reminder injection once a week for 5-weeks (Summarised in Figure S1A). An additional group of db/db animals non-injected was added to the study to ascertain possible alterations driven by the isotype treatment in comparison to basal db/db mice. Animals were 18 weeks at the end point of the experiment, representing an established but not yet advanced stage of diabetes.

Nine-weeks after the first expansion, mice were terminally anaesthetised with i.p. Ketamine (75 mg/kg) and Xylazine (12 mg/kg) injection and transcardially perfused with ice-cold phosphate buffered saline (PBS) followed by 4% paraformaldehyde (PFA) (Sigma-Aldrich). Eyes were enucleated and immersed in 2% PFA for 2 h. Then, eyes reserved for cryosectioning were cryoprotected with 30% sucrose in PBS for 72 h, snap-frozen in OCT (Tissue-Tek, Sakura Finetek, Cat. No. 4583), cryosectioned at 15 µm thickness and immunostained as indicated below. Eyes that were used for flatmount staining were fixed overnight and transferred to PBS. Eyes for western blot were dissected and immediately frozen.

In vivo fundus imaging

All procedures were conducted under general anesthesia with a mixture of 2% isoflurane (Arrane®. Baxter laboratories, Victoria, Australia)/1% O₂ using an induction camera and a rodent mask. Body temperature was maintained with a heating pad. Pupils were dilated with Tropicamide eyedrops (Colorcusi Tropicamida® 10 mg/mL, Alcon Laboratories, El Masnou, Spain), and 2% Methocel gel (OmniVision, Puchheim, Germany) was administered to the cornea to favor contact with the lens. Animals were subjected to fundus fluorescence angiography (FA) on the week prior to the end point. Once the animals were anesthetized and pupils dilated, 10 µL/g sodium fluorescein (0.1% Fluorescein Oculous, Thea Laboratories, S.A., Spain) was administered intraperitoneally. Eye fundus images from the central retina were taken 1 min after

fluorescein administration using the Micron III platform (Phoenix Research Labs., Pleasanton, CA, USA).

Flow cytometry

Blood, spleen and retina

Treg levels were investigated in the blood, spleen and retina of all groups by flow cytometry at the time points specified in Figure S1A. In brief, spleens were mashed through a 70 µm strainer and then treated with red blood cell lysis buffer (STEMCELL Technologies) for 2 min at room temperature. Cells obtained from spleens were then washed with PBS and centrifuged at 300 g for 5 min at 4 °C. Cells were resuspended in 200 µL PBS and stained with a cell viability dye with eFluor 506 viability dye (1:500; Thermo Fisher Scientific, Cat. No. 65-0866-18) and cell surface stained with antibodies for CD45 (eBioscience, clone 104, Cat. No. 47-0454-82), CD4 (1:500; eBioscience, clone GK1.5, Cat. No. 17-0041-81), CD25 (1:500; eBioscience, clone PC61.5, Cat. No. 12-0251-82), CD3 (1:500; eBioscience, clone 17A2, Cat. No. 11-0032-82), all 1:500, for 15 min at RT. Cells were washed with flow cytometry wash buffer (PBS) and centrifuged at 300 g for 5 min at 4 °C. Cells were then fixed with Fix & Perm A (Thermo Fisher Scientific, Cat. No. GAS004) for 10 min at RT. Fixative was washed off with PBS and centrifuged at 300 g for 5 min at 4 °C. Then, cells were resuspended in 100 µL Fix & Perm B (Thermo Fisher Scientific; Cat. No. GAS004) with an anti-Foxp3 antibody (1:100; eBioscience, clone FJK-16 S, Cat. No. 48-5773-82) and T-bet antibodies (1:100, eBioscience, clone eBio4B10 (4B10), Cat. No. 45-5825-82) overnight at 4 °C. Cells were then washed with PBS and centrifuged at 300 g and 4 °C for 5 min. Final pellet was resuspended, data were acquired on a Fortessa (LSRFortessa™ High-Parameter Flow Cytometer, BD Biosciences) and analysed using FlowJo software version 10.0 (BD). To calculate Treg percentages, singlets were identified by FSC-H versus FSC-A and viable cells gated on CD4⁺CD3⁺ expression, and within the CD4⁺ population Treg markers Foxp3⁺ was delineated.

Blood

Blood obtained from the animals at the time of perfusion was collected in EDTA-tubes to avoid blood coagulation. 30 µL of the blood were incubated with surface antibodies in PBS as indicated per spleen for 15 min at RT. Cells were washed with flow PBS and centrifuged at 300 g for 5 min at 4 °C. Red blood cells were then lysed and cells were fixed with Optilys B (Beckmann and Coulter; Product No: IM1400) for 10 min at RT. Fixative was washed with ddH₂O to finish red blood lysis for 15 min and centrifuged at 300 g for 5 min at 4 °C. Then, cells were resuspended in Fix & Perm B (Thermo Fisher Scientific) with an anti-Foxp3 antibody (1:100; eBioscience,

clone FJK-16 S) and T-bet antibodies (1:100, eBioscience, clone eBio4B10 (4B10)) overnight at 4 °C. Cells were then washed with FCSB and centrifuged at 300 g and 4 °C for 5 min. The final pellet was resuspended, data were acquired on a Fortessa and analysed using FlowJo software version 10.0 (BD). Treg percentages were calculated as indicated above.

Retina

Retinas were dissected from enucleated eyes and separated from the RPE choroid. Retinas of two eyes from the same animal were put together, cut into small pieces and subjected to papain (Worthington) digestion for 30 min at 37 °C in hibernate A with DNase type I (Worthington). Upon digestion papain was washed-off with Hanks Buffered Salt Solution (HBSS) (Thermo Fisher Scientific) and cells were centrifuged at 300 g for 5 min at 4 °C. The pellet was resuspended in 4 ml trituration buffer (Hibernate A with 2% B27 (Gibco Life Technologies; Cat. No. 17504044) and 4mM Sodium Pyruvate (Thermo Fisher Scientific) and gently triturated with a 5mL serological pipette. Cell suspension was transferred to a 50mL tube through a 70 µm strainer and the remaining tissue resuspended in another 2 ml of trituration buffer and gently triturated using a fire-polished glass pipette. After trituration supernatant was again transferred to a 50mL tube through a 70 µm strainer. This step was repeated again twice, with a glass pipette of decreased diameter and a 1mL pipette tip. Cells were resuspended in 250 µl PBS and stained following the same protocol as the spleen. Data was acquired on a Fortessa and analysed using FlowJo software version 10.0. Considering the limited number of Treg in the retina, to adequately find the population, CD45 and CD4 gates were established and then CD4⁺ cells from each mouse were grouped (concatenated) based on the received treatment (db/+, db/db with Isotype and db/db with Treg). Within each treatment group the same number of CD4⁺ cells were selected (508 cells per group) using downsample plugin in FlowJo. This allowed to group a higher number of CD4⁺ T cells, facilitating the identification of the Foxp3⁺ Treg population within the retina despite its low number. Treg population was identified by positively selecting Foxp3⁺ cells within CD4⁺ population. Both, the percentage of Foxp3⁺ cells as well as the number of Foxp3⁺ events within CD4⁺ T cells were established per treatment group. The number of total Foxp3⁺ events were graphed in an histogram representing the counts for Foxp3⁺ events against the Foxp3-e450 intensity per group. Once the gate was established Treg count per group within the CD4⁺ gate was obtained and the percentage of Foxp3⁺CD4⁺ Treg was established per individual animal.

Immunostaining

Cryosection staining

Cryosection staining was performed as previously described [21]. Briefly, eye sections were dried for 30 min at RT and washed in PBS for 10 min. Antigen retrieval was performed using Citrate buffer pH 6.0 (Abcam, ab93678) at 95°C for 5 min in a water bath and after cooling, and additional 5 min with 10% SDS. Slides were then blocked in 10% Donkey serum (Sigma Aldrich, D9663) with 0.2% Triton-X in PBS for 1 h at room temperature. Primary antibodies (Table 1) were added and incubated overnight at 4°C in 0.5% donkey serum and 0.2% Triton-X followed by incubation with secondary antibodies (Table 1) for 1 h at room temperature, in PBS.

Flatmount staining

The retinal flatmount tissues were first washed in PBS, followed by permeabilization with 2% Triton-X 100 overnight at 4 °C. Afterward, the samples were blocked using 5% donkey serum in 0.2% Triton-X 100 with 1mM CaCl₂, incubating overnight at 4 °C. They were then treated with the appropriate primary antibody (Table 1) diluted in 0.2% Triton-X PBS for 96 h at 4 °C. Following a 6-hour wash, the tissues were incubated overnight at 4 °C with the relevant secondary antibodies (Table 1), washed again for 6 h, and finally mounted for analysis.

Image acquisition and analysis

Tissue sections were cover-slipped with Vectashield (Vector Labs, Cat. No. H-1000-10) and examined by Leica Thunder DMI8 or confocal microscope (Zeiss LSM980, Jena, Germany). Consistent cryosectioning patterns were applied across all animals using the same slides for each histological marker analysed and ensuring that all retinal sections were obtained from the middle-center eccentricities to minimize variability and potential positional errors. Images were captured from 3 to 4 regions across 2–4 different sections of the middle-center retina. Further image processing and analysis was performed using Fiji software and blindly manually counted.

Western blot

Samples were homogenized sonicating the frozen retinas in 70µL of RIPA buffer containing protease inhibitor cocktail (Sigma, Cat. No. 04693159001 and R0278, respectively) and PhosSTOP™ (Roche, Cat. No. 04906837001). Protein concentration was determined using a Bradford Assay (Thermo Fisher Scientific, Cat. No. 23246). The blot was performed using 60 µg of protein according to previously described methods [12] but with nitrocellulose membranes. Primary and secondary antibodies used are detailed in Table 1. Membranes were visualized with enhanced chemiluminescence WesternBright ECL HRP substrate kit

Table 1 Antibodies used in the study

Primary antibody	Cat. No.	Company	Host	Dilution
Brn3a Antibody (C-20)	sc-31,984	Santa Cruz Biotechnology	Goat	1:200
Ionized calcium binding adaptor molecule 1 (IBA-1) antibody	ab5076	Abcam	Goat	1:200
Recombinant Anti-PKC alpha antibody [133]	ab11723	Abcam	Mouse	1:500
Anti-Cone Arrestin Antibody	AB15282	Sigma	Rabbit	1:500
Guinea Pig Anti-Synaptophysin	AGP-144	Alomone	Guinea Pig	1:150
Anti-Glial Fibrillary Acidic Protein antibody (GFAP)	Z0334	Agilent	Rabbit	1:500
Biotinylated Isolectin B4	VEC.B-1205	Vector Labs	Biotin	1:50
Albumin	a90-134a	Bethyl	Goat	1:100
DAPI (4',6-diamidino-2-phenylindole)	D3571	ThermoFisher		1:1000
Tyrosine hydroxylase (TH) antibody	AB152	Merck	Rabbit	1:100
Choline acetyltransferase (ChAT)	Ab144P	Merck	Goat	1:100
Recombinant Anti-NLRP3 antibody	ab270449	Abcam	Rabbit	1:500
Vinculin antibody	sc-73,614	Santa Cruz Biotechnology	Mouse	1:1000
GAPDH antibody	ab8245	Abcam	Mouse	1:1000
IL-18 (H-173) antibody	sc-72,954	Santa Cruz Biotechnology	Rabbit	1:500
Caspase-1 Antibody and p20	#2225	Cell Signaling	Rabbit	1:1000
RBPM5	PA5-115831	ThermoFisher	Rabbit	1:100
CD68	ab53444	Abcam	Rat	1:100
Tmem119	ab209064	Abcam	Rabbit	1:100
Secondary antibody	Company	Cat. No.		
Donkey anti-Rabbit IgG (H+L) Highly Cross-Adsorbed Secondary Antibody, Alexa Fluor 568	ThermoFisher	A10042		
Donkey anti-Rabbit IgG (H+L) Highly Cross-Adsorbed Secondary Antibody, Alexa Fluor 647	ThermoFisher	A-31,573		
Donkey anti-Mouse IgG (H+L) Highly Cross-Adsorbed Secondary Antibody, Alexa Fluor 488	ThermoFisher	A21202		
Donkey anti-Mouse IgG (H+L) Highly Cross-Adsorbed Secondary Antibody, Alexa Fluor™ 647	ThermoFisher	A32787		
Donkey anti-Goat IgG (H+L) Cross-Adsorbed Secondary Antibody, Alexa Fluor™ 568	ThermoFisher	A11057		
Donkey Anti-Goat IgG H&L (Alexa Fluor® 488) (ab150129)	Abcam	ab150129		
Donkey Anti-Goat IgG H&L (Alexa Fluor® 647) (ab150131)	Abcam	ab150131		
Goat anti-Guinea Pig Alexa Fluor 568	Abcam	ab175714		
Goat Anti-Rabbit Immunoglobulins/HRP	Agilent DAKO	P0448		
Streptavidin, Alexa Fluor™ 647 conjugated	ThermoFisher	s21374		

(Advansta, K-12045-D50CA) and bands detected using LI-COR Licor Odyssey FC 2800 Western Blot Imaging System (LICORbio).

Immunoassays

HbA1 and insulin quantification

Plasma insulin levels were determined using an ultrasensitive mouse enzyme-linked immunosorbent assay (10-1251-01, Mercodia Inc., Winston-Salem, NC, USA).

Magnetic bead-based immunoassay for retinal VEGF detection

Protein from retinas were extracted by sonicating the isolated tissue in RIPA buffer containing protease inhibitor cocktail (Sigma) and PhosSTOP™ (Roche). Protein concentrations were determined by using Pierce BCA protein assay kit (Thermo Fisher Scientific, Cat. No. 23225).

VEGF concentration in 100 µg of retinal homogenates was determined using the ProcartaPlex Mouse and Rat Mix & Match kit (Thermo Fisher Scientific, PPX-10-MXT2CFM) following the manufacturer's instructions. Samples were processed in a 96-well flat bottom plate and analysed with a Luminex MAGPIX instrument (Luminex Corporation, Austin, TX, USA) utilizing xPONENT 4.2 software (Luminex Corporation). The MAGPIX system employs a sandwich ELISA methodology, where the capture antibody is attached to fluorescently dyed magnetic microspheres, facilitating bead-based quantitative protein detection. The protein concentration was derived from standard curves through five-parameter logistic (5 PL) curve fitting conducted by the software.

Statistical analysis

Statistical analysis was performed using Graph Pad Prism (GraphPad Software, Inc. version 10). First, normal

distribution was assessed using Kolmogorov-Smirnov tests. When the data was described as percentage, an *arc-sin* conversion was performed to analyse the data using parametric tests for normally distributed data. When only two groups were compared, e.g. Treg percentage changes upon Treg expansion, Student t-test was used. For comparisons involving more than two groups, 1-way ANOVA analysis was performed followed by Tukey's multiple comparisons test. For all statistical tests, differences were considered significant with p values below 0.05.

Results

db/db mice have a systemic and retinal decrease in Treg proportions which are recovered after systemic expansion

Due to the lack of consensus regarding the alteration in Treg cell numbers in T2D patients [22–24], our initial objective was to ascertain whether Tregs are altered in db/db T2D mice. In blood, db/db animals present almost half the percentage of Foxp3⁺CD4⁺ cells within their CD4⁺T lymphocytes than their control counterparts, the db/+ animals (Figure S1B). To overcome this decrease, we systemically expanded endogenous Treg following the protocol described by Webster et al. [20] with minor modifications (Figure S1A). We confirmed Treg expansion upon IL-2/mAb IL-2 treatment in blood 4–5 weeks after the initial injection (half-way the treatment) (Figure. S1C) and 9-weeks after the experiment (experimental endpoint) (Figure S1D).

Considering that Treg numbers in the healthy retina are very low [21], we next determined whether the impact of Treg expansion was detectable in the retina. As described for peripheral Treg, db/db mice showed a decrease proportion of Treg and Treg number within the CD4⁺ T cell population in the retina. Similarly to blood and spleen, despite still being found in small numbers, as described before [21], IL-2 mediated Treg systemic expansion was also effective in the retina [25] (Figure S1E–G).

Once local and systemic Treg expansion were confirmed, we next sought to determine if systemic Treg expansion in db/db mice had any consequence in metabolic control including hyperglycemia, HbA1c and blood insulin levels. Treg expansion did not significantly change these parameters, (Figure S2A–C, respectively).

Treg expansion prevents retinal neurodegeneration in db/db mice

We next aimed to determine whether Treg expansion rescued changes in retinal pathology and neurodegeneration that are characteristic of db/db mice. Histological analysis using hematoxylin-eosin staining revealed a significant decrease in total retinal thickness in both non-injected db/db mice and db/db animals receiving isotype treatment, in line with previous reports [26]. This

reduction of retinal thickness was not observed in the Treg-expanded group. The retinal thinning in db/db and isotype-treated mice primarily resulted from decreased thickness in the innermost layers, particularly the inner plexiform layer (IPL) and inner nuclear layer (INL). In contrast, the db/db mice undergoing Treg expansion exhibited no significant loss of IPL or INL thickness, and their retinal layers appeared similar to those of non-diabetic db/+ mice (Fig. 1A–B).

It is well-established that the altered vision associated with DR in db/db is linked to neurodegenerative pathology, including loss of photoreceptors, amacrine cells and retinal ganglion cells (RGC) [26, 27]. Thus, to determine the effect of Treg expansion in these defined retinal neuronal populations, we examined the changes in the total number of photoreceptors (DAPI⁺ cell rows) in the ONL and Cone Arrestin (CA) (Fig. 1C–E). We observed a significant decrease in the total number of photoreceptors in both diabetic non-injected and isotype injected animals, which was not present in animals that underwent Treg expansion. When looking at cone photoreceptors, we observed a slight but significant reduction in number in isotype-injected db/db, not present in Treg expanded. These results suggest that the reduction of photoreceptors seen in diabetic animals, whether they were non-injected (db/db) or isotype-injected, appears to be primarily linked to a reduction in rod photoreceptors. This loss was not observed in animals with Treg expansion.

To further explore the effect of Treg expansion on diabetic neurodegeneration, we also examined rod bipolar second-order neurons (Fig. 1F–H). In healthy retinas (db/+), rod bipolar cell bodies are positioned in the outer region of the INL, near the OPL. In non-injected diabetic mice (db/db), there was a mild but significant reduction in the total number of PKCα⁺ somas across the INL (Figure S4A). Degenerative changes, such as shorter and distorted processes and fewer cell bodies [21, 28, 29], were evident in both non-injected and isotype-treated diabetic retinas. Specifically, near the OPL, both groups had fewer PKCα⁺ somas, while the number of PKCα⁺ somas in Treg-expanded diabetic mice was comparable to that in non-diabetic db/+ mice (Fig. 1F, G). Beyond changes in cell number near the OPL, PKCα⁺ rod-bipolar cells showed altered laminarity, with a significant increase in ectopic PKCα⁺ somas located below the OPL, due to shorter axons, in both non-injected and isotype-treated db/db mice that was absent in non-diabetic db/+ and Treg expanded diabetic animals (Fig. 1F, H). We also observed bipolar dendrites and synaptophysin sprouts areas extending from the OPL towards the ONL, a recognised sign of neurodegeneration [28], in both non-injected control and isotype db/db mice, which were absent in diabetic animals that underwent Treg

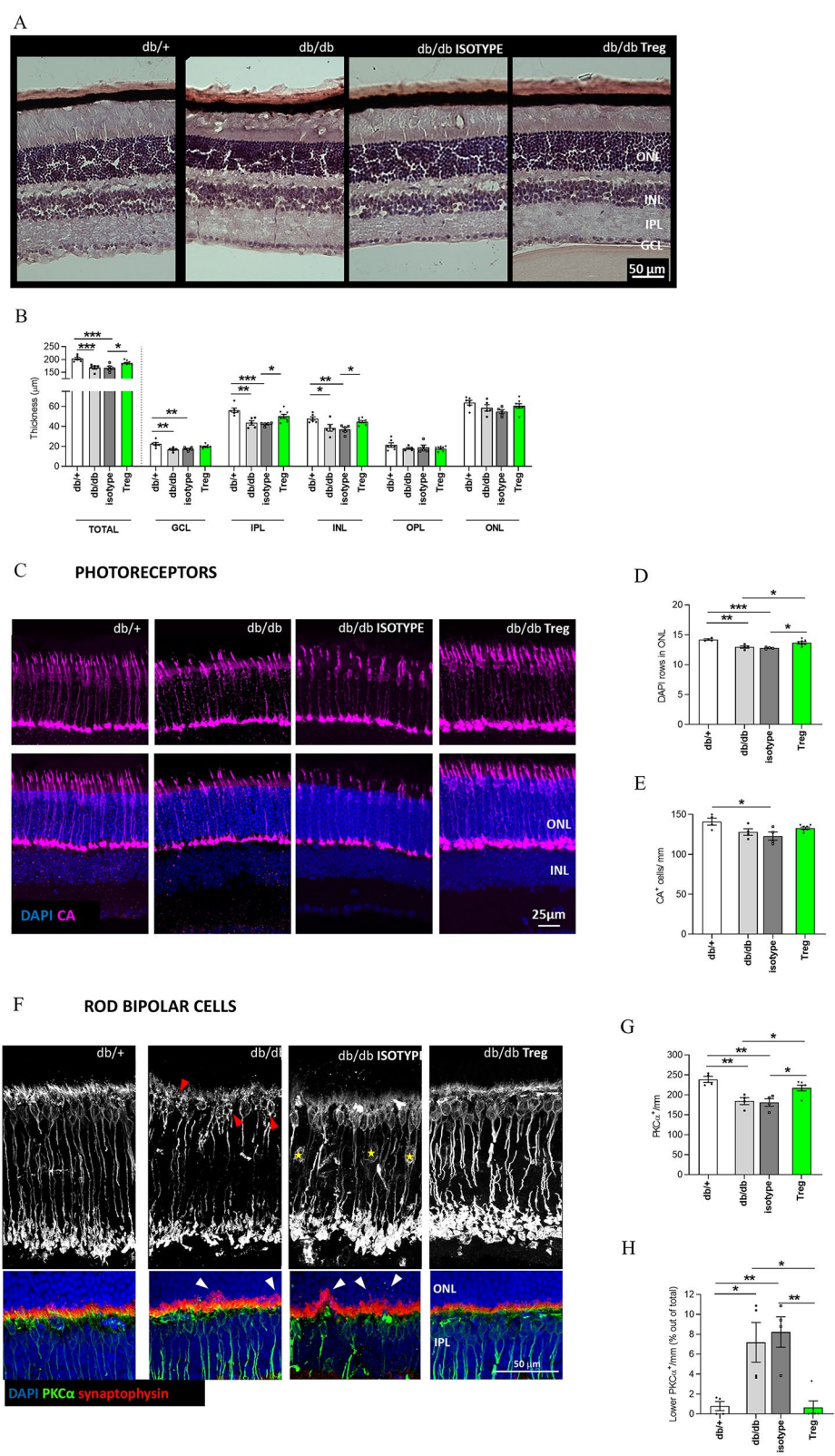


Fig. 1 (See legend on next page.)

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Fig. 1 Treg expansion limits DR neurodegeneration. **A–B**, Hematoxylin and eosin analysis of retinal thickness. Representative images showing hematoxylin and eosin staining of db/+, db/db non-injected and db/db injected with either IL-2/mAb IL or its control (**A**). Quantification of the total retinal layer thickness as well as the thickness of the different cell layers (**B**). Scale bar is 50 μm . $n=5–8$ mice. **C–E**, Immunostaining and quantitative analysis of photoreceptors. Representative images showing photoreceptors in the ONL (**C**) (Cones, CA⁺) in diabetic db/db animals non-injected, injected with the isotype, db/db animals receiving a Treg expansion and non-diabetic db/+ (scale bar = 25 μm). Quantitative analysis of the number of rows of DAPI⁺ nuclei in the ONL (**D**). Quantification of the number of cone photoreceptor cells (**E**) (CA⁺ cells) $n=4–5$ animals. **F–H**, Immunostaining and analysis of rod bipolar cells and synaptic vesicles. Representative image of PKC- α (**F**) (upper panel, grey. Lower panel, green) and synaptophysin (red, lower panels). Stars indicate abnormal location of some PKC- α cell soma lower within the INL. Arrowheads indicate abnormal rod bipolar dendrite and synaptic vesicles sprouts into the ONL (scale bar = 25 μm). Quantitative analysis of the number of PKC- α somas close to OPL (**G**) ($n=4–6$ mice) and (**H**), percentage of PKC- α somas found in a lower location out of the total of PKC- α cells ($n=4–6$ animals). Data expressed as mean \pm s.e.m. 1-way ANOVA followed by Tukey's multiple comparisons test. * $P < 0,05$; ** $P < 0,01$; *** $P < 0,005$; **** $P < 0,001$

expansion and in non-diabetic db/+ mice (Fig. 1F, white arrowheads).

To further address potential cellular changes linked to neurodegeneration in DR, we next examined retinal interneurons. Although no changes were observed in calbindin⁺ horizontal cells (Figure S4B), alterations were evident in amacrine cells of db/db animals (Fig. 2A–C). We observed a decrease in the number of ChAT⁺ (Fig. 2A, upper panel, and 2B) and TH⁺ (Fig. 2A, lower panel, and 2C) somas, representing cholinergic and dopaminergic amacrine cells, respectively. This finding aligns with previous reports from various experimental models of diabetic retinopathy [29–31]. Diabetic animals that underwent Treg expansion showed a significantly higher number of ChAT⁺ and TH⁺ amacrine cells, in line with the numbers observed in db/+ mice (Fig. 2A–C). Likewise, the decrease in displaced cholinergic amacrine cells observed in both db/db and isotype treated diabetic groups was absent in diabetic animals that underwent Treg expansion (Fig. 2A, D).

Lastly, we examined alterations in the retinal ganglion cell layer (GCL) due to the endogenous expansion of Treg in T2D animals (Fig. 2E–H). Treg-expanded diabetic animals exhibited a significant increase in the total number of DAPI⁺ cells in the GCL compared to their non-expanded diabetic controls. Notably, the total number of DAPI⁺ cells in the GCL of Treg-expanded diabetic animals was comparable to that of db/+ non-diabetic animals (Fig. 2E, F). As previously described, non-injected db/db mice presented fewer RGC cells (RBPMS⁺ cells) and Brn3a⁺ RGC when compared to non-diabetic db/+ mice (Fig. 2G and H, respectively). This decrease was also observed in the isotype treated db/db mice, while the expansion of Treg showed a significant increase in the number of total RGC cells (RBPMS⁺ cells) and Brn3a⁺ RGCs, bringing numbers up to those of non-diabetic db/+ mice (Fig. 2E, G and H).

Treg expansion reduce gliosis and retinal inflammation

We next studied the consequences of Treg expansion in the microglia and Müller cells that are responsible for initiating and perpetuating inflammation in DR [32, 33]. As expected, we observed an increased number of

Iba-1⁺ cells, the majority of which were Tmem119⁺ (Figure S4C, D), indicating that they were microglial cells, in both non-injected control and isotype-treated diabetic animals compared to non-diabetic controls. Treg expansion significantly decreased Iba-1⁺ microglial cells in the db/db retina, to levels that were comparable to non-diabetic controls (Fig. 3A–B). This finding is consistent with observations from retinal flatmounts (Fig. 3C). In line with our previous findings, the number of Iba-1⁺ microglial cells in the INL was elevated in isotype-treated diabetic animals, but this increase was significantly reduced following Treg expansion, bringing the levels closer to those observed in non-diabetic controls (Figure S4E).

Increased expression of glial fibrillary acidic protein (GFAP) as a response of reactive gliosis, is one of the earliest and more prominent signs of Müller cells activation in DR. db/db mice subjected to Treg expansion showed a significant reduction in the number of GFAP⁺ filaments that crossed the IPL when compared to either isotype or non-injected db/db animals (Fig. 3C–D).

Inflammasomes are central regulators of innate immunity, and their involvement in several neuroinflammatory pathologies has been widely described [34]. In the diabetic eye, the NLRP3 inflammasome has been shown to be activated within the retina, specifically in Müller cells, which are a primary source of NLRP3 [35, 36]. Thus, considering the decreased gliosis observed in DR upon Treg expansion, we next examined the expression of NLRP3 inflammasome in the retina (Fig. 3E, F). NLRP3 protein levels were significantly increased in isotype-treated diabetic animals, while animals exposed to Treg expansion present similar NLRP3 protein levels to those of non-diabetic db/+ mice. This was further confirmed by assessing the protein levels of two key downstream effectors of the NLRP3 inflammasome pathway: IL-18 and Caspase-1, including the cleaved form, Caspase-1 p20. All three—IL-18, Caspase-1, and Caspase-1 p20—were significantly reduced in db/db animals subjected to Treg expansion compared to isotype-treated db/db animals (Fig. 3G and H, J, respectively).

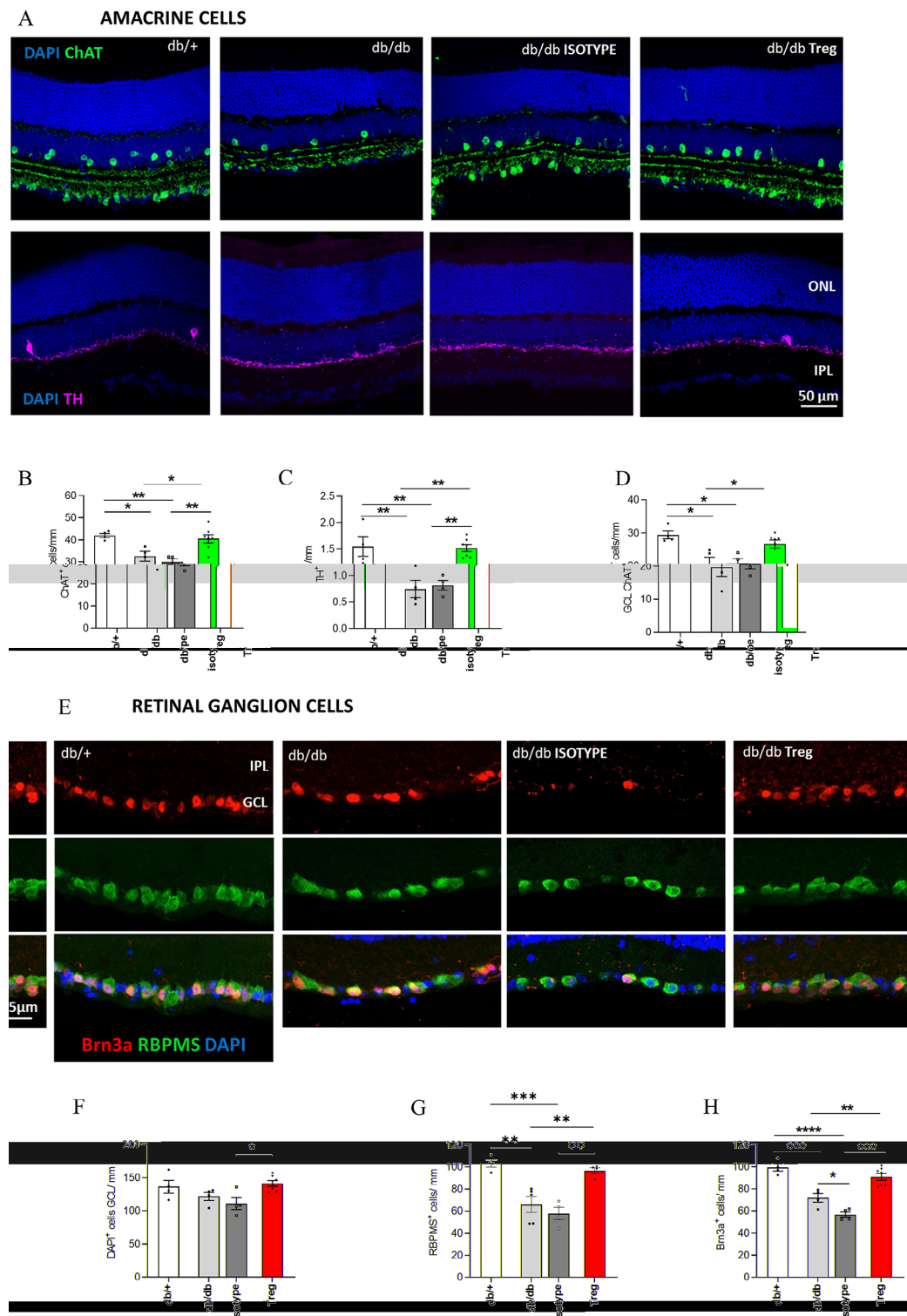


Fig. 2 Effect of Treg cell expansion on DR-related neuronal damage. **A–C**, Immunostaining and quantitative analysis of cholinergic (ChAT⁺) and dopaminergic (TH⁺) cells amacrine cells. **A**, Representative image of ChAT⁺ cells (green, upper panel) and TH⁺ amacrine cells (purple, lower panel) (Scale bar = 50 μ m). **B**, **C**, quantification of the number of ChAT⁺ (**B**) and TH⁺ (**C**) somas per mm of retina. **D**, quantification of the number of displaced ChAT⁺ amacrine cells into the GCL. $n=4-7$. **E–H**, Immunostaining and analysis of retinal ganglion cells (RGC). **E**, Representative image of RBPM and Brn3a⁺ RGC (Scale bar = 25 μ m). **F**, Quantification of the number of DAPI⁺ cells in the GCL; **G–H**, quantification of RBPMs and Brn3a⁺ RGC, respectively. Data presented as mean \pm s.e.m. * $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$; 1-way ANOVA followed by Tukey's multiple comparisons test

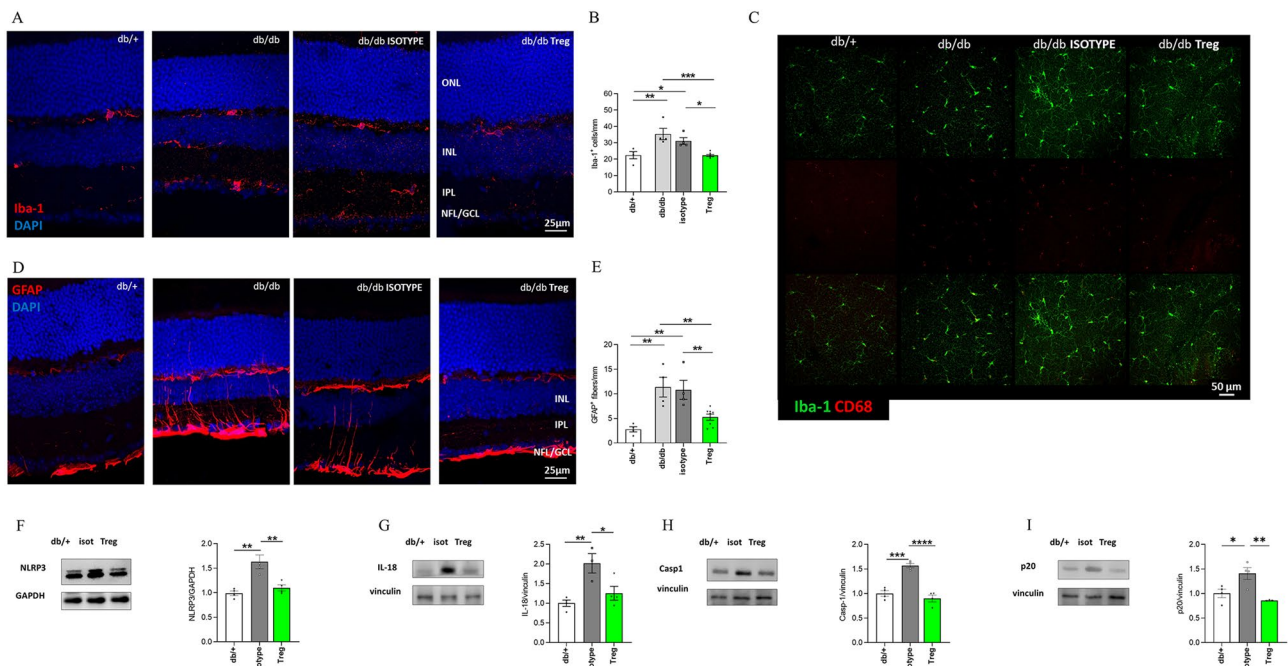


Fig. 3 Animals undergoing Treg expansion exhibited reduced DR-related inflammatory features. **A–E** Analysis of Iba-1⁺ microglial cells and GFAP⁺ Müller cells determined by immunofluorescence. Representative image of microglial cells in sections (**A**) (Iba-1⁺, red), in retinal flatmounts (**C**) (Iba-1 green, CD68 red) and **D**, gliotic Müller cells (GFAP⁺ fibers, red). Scale bars = 25 μ m in sections; 50 μ m in flatmount. **B**, Quantitative analysis of the total number of Iba-1⁺ cells in all layers of the retina. **E**, number of GFAP⁺ fibers crossing the IPL per mm. $n = 4–8$. **F–I**, Western blot analysis and representative blot of NLRP3 (**F**), mature form of IL-18 (**G**), Caspase-1 (**H**) and p20 Caspase-1 cleaved form (**I**) ($n = 4$). Data presented as mean \pm s.e.m. * $P < 0.05$; ** $P < 0.01$; *** $P < 0.005$; 1-way ANOVA followed by Tukey's multiple comparisons test

Peripheral Treg expansion improves diabetic retinopathy vasculopathy

Vascular damage is a hallmark of DR. To investigate this, we performed in-vivo fundus angiography with sodium fluorescein, which revealed markedly tortuous retinal vessels in db/db mice, both in non-injected and isotype-treated groups (arrows, Fig. 4A). This feature is reflective of the vasodegenerative pathology in DR [37] but was absent from db/db mice that underwent Treg expansion (Fig. 4A).

To determine if the inner blood retinal barrier was affected, as previously described in db/db mice [12, 38], we performed immunostaining with isolectin B4 and albumin with subsequent quantification of the number and size of the leakage spots, reflecting areas where blood-borne albumin had leaked into the neuropile (Fig. 4B–E). As shown in Fig. 4B–C, db/db non-injected animals or mice receiving the isotype exhibited several large, focal leakage areas (indicated with white arrowheads, Fig. 4B and Figure S4F–G in retinal flatmounts), while db/db mice with expanded Treg showed less and smaller leakage spots (Fig. 4B, yellow arrowhead; Fig. 4D and E, respectively and Figure S4F–G in retinal flatmounts). These findings indicate that Treg expansion ameliorates the diabetes-induced blood-retinal barrier breakdown. Since VEGF, also named vascular

permeability factor, plays an essential role in inducing vascular leakage in diabetic retina [39], we also investigated changes in this pivotal factor in our study. We found that control db/db mice showed an expected increase in the concentration of this growth factor that was absent in animals subjected to Treg expansion, with levels closer to those of db/+ mice (Fig. 4E).

Discussion

In this study we found a marked reduction of Tregs in db/db mice, a model of T2D that features hyperglycemia, insulin resistance [40], and early stages of DR [26]. Systemic Treg expansion led to increased retinal Tregs and prevented key DR hallmarks, such as neuron loss, glial activation, and vascular leakage. These effects could not be attributed to an improvement of blood glucose levels. This highlights Tregs as a potential new immunomodulatory approach to prevent retinal damage in DR.

Low-grade inflammation is a well-known underlying pathogenic factor of T2D, leading to several studies investigating Treg cells in the context of T2D. A systematic review and meta-analysis indicated that patients with T2D have a lower percentage of peripheral CD4⁺Foxp3⁺ Tregs compared to healthy controls, with even lower levels in those with complications [22]. However, other studies have reported not a decrease

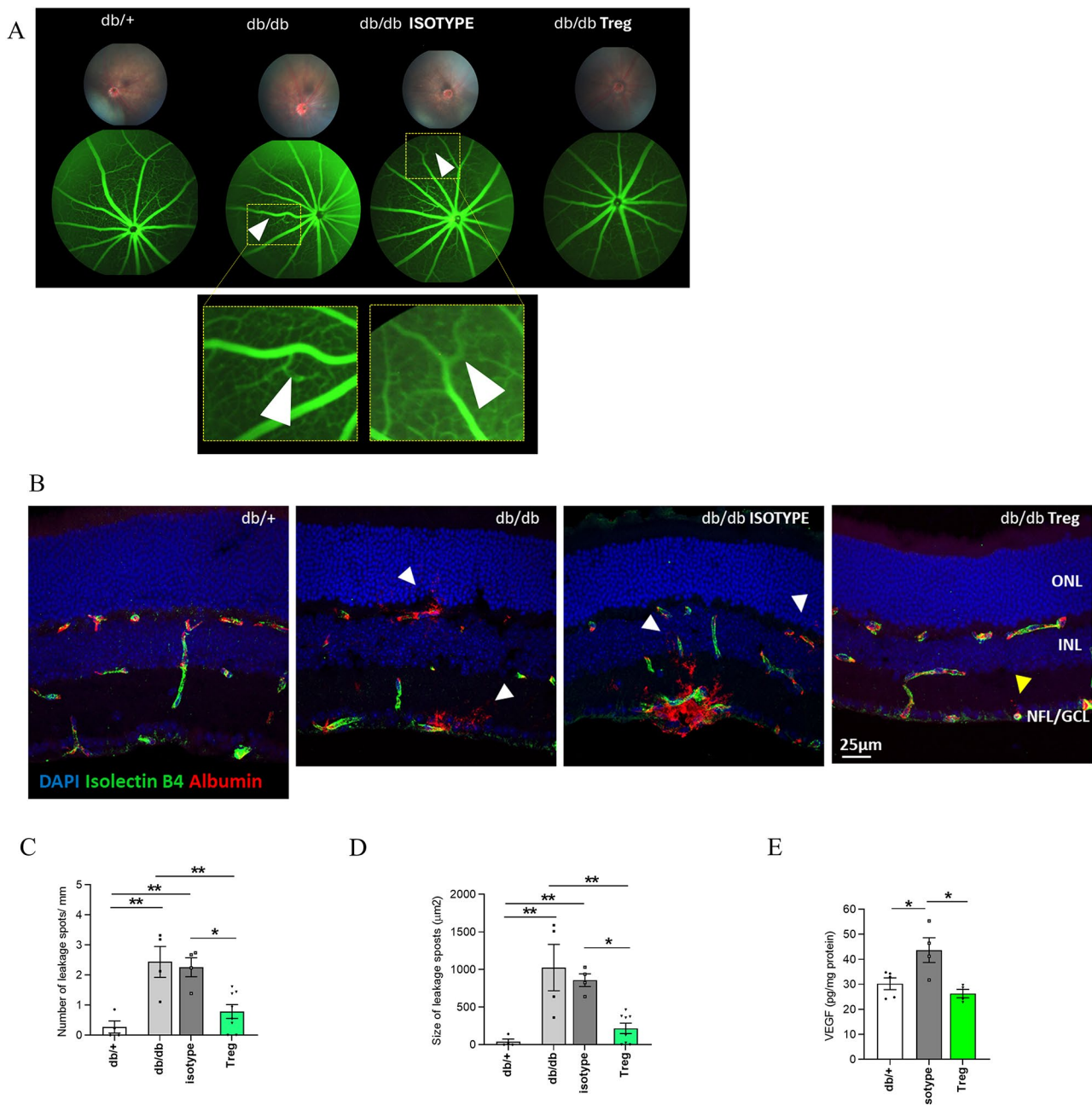


Fig. 4 Treg expansion reduces diabetes-associated vascular damage and alteration of thickness. **A**, Fundus fluorescein (fluorescein sodium) image of db/+, db/db and db/db receiving either IL-2/anti-IL-2 or its isotype. In yellow, magnified image of the area. The arrowhead indicates vascular tortuosity in db/db animals and db/db treated with IL-2/isotype. **B–D**, Representative images of retinal sections (**B**) showing retinal barrier leakage, visualized using isolectin B4 (green) and albumin (red) staining. The quantification includes the number of leakage spots per mm of retina (**C**) and the average size of the leakage spots (in μm²) (**D**). The leakage spots/areas were determined by counting the number of albumin⁺ and isolectin B4⁺ spots and measuring the area of albumin⁺ regions in retinal sections. White arrowheads indicate larger or more extensive areas of leakage, while yellow arrowheads mark smaller areas of leakage. Scale bar 25 μm. *n* = 4–8 (**C**) and *n* = 3–6 (**D**). **E**, Quantification of VEGF concentration in the retina, normalized to the total protein in the retina. *n* = 4–5. Data presented as mean ± s.e.m. **P* < 0.05; ***P* < 0.01; 1-way ANOVA followed by Tukey's multiple comparisons test

in the percentage of Tregs but rather an imbalance between Th17 and Treg cells [24], or a reduction in Treg functionality [41]. Despite the conflicting data, it seems plausible that alterations in Treg cells, either in reduced numbers or decreased functionality, could

contribute to the inflammatory environment characteristic of T2D. In experimental models of T2D, including studies with db/db mice, Treg levels have been sparsely studied, with results showing some variability that may depend on the specific mouse strain and T2D model

utilized [42, 43]. In the present study we have observed less CD4⁺CD25⁺Foxp3⁺ Treg in the blood of db/db mice compared with non-diabetic animals. This is particularly relevant considering the resemblance of this model with human T2D features. In an experimental model of T2D, Treg had been shown to reverse or improve insulin resistance [44]. However, we did not observe any significant changes in hyperglycemia, HbA1c, or circulating insulin. These findings strongly suggest that the observed effects in the retina are independent of any changes in the diabetic status of the animals. Instead, they can be attributed to the immunosuppressive and potentially to their suggested role in supporting neuronal health, as indicated in the literature and possibly relevant to the effects observed in this study [13–15]. Nonetheless, we cannot rule out that a different treatment regimen or a different Treg approach could influence the metabolic control or even the course of T2D and its associated complications.

Several investigations have demonstrated that Treg are integral to immune regulation, tissue regeneration and neuroprotection within the CNS, however, very few studies have specifically focused on their role in the retina [13, 21, 25]. While CNS has traditionally been considered as immune privileged, recent findings indicate the presence of Treg within the healthy CNS, including the retina, albeit in minimal quantities [21]. This suggests that Treg may possess mechanisms enabling them to traverse the blood-brain and blood-retinal barriers. In fact, Treg are actively recruited to the retina to repair pathological angiogenesis [25]. In addition, we have recently described a role of Treg in preserving retinal homeostasis and thereby restraining age-related degenerative changes [21], which share some common hallmarks with DR, such as neurodegeneration, inflammation and gliosis. On this basis it is tempting to speculate that the reduced systemic Treg reported in diabetes, and particularly its deficit in the retina, could be involved in the development and progression of DR. Our finding showing that Treg expansion was not only detected in blood and spleen but also in the retina of db/db mice, indicates that Treg have the capacity not only to migrate to the retina but to exert a myriad of beneficial actions.

One of the most evident results of Treg expansion has been the prevention of retinal thinning as a consequence of the preservation of retinal neurons (i.e. photoreceptors, bipolar cells, amacrine cells and RGCs). The other meaningful effect has been the abrogation of both macroglia and microglia activation and one of its main consequences: neuroinflammation. Although the vast majority of the detected Iba1⁺ cells were microglia, the role of the remaining 8% in the development of DR has yet to be determined. Similar effects have already been reported in other CNS pathologies [13, 16, 17]. It should be noted that in the setting of DR, inflammation

is as a key pathogenic factor that occurs before vascular changes can be detected [7–9] and as in many other neurodegenerative diseases, inflammasome activation is a well-established pathogenic mechanism [45] contributing to the heightened inflammatory response and the breakdown of the blood-retinal barrier (BRB) [35, 46, 47]. Our results have shown that Treg mitigate these effects by decreasing the retinal levels of NLRP3, IL-18 and Caspase-1, neutralizing at least one of the key innate-immune system driven inflammatory components and thus, facilitating the preservation of the neuroretina and the BRB.

Treg cells are able to mitigate vascular damage in chronic inflammatory conditions and in ocular diseases [25, 48]. In the retina, Treg have demonstrated the ability to repair vasculopathy in a model of oxygen-induced retinopathy [25]. Treg are recruited to the retina and, upon expansion, effectively reduced neovascularization, vascular leakage, and VEGF expression by altering the activation of microglia and inflammatory status [25]. We have broadened the understanding of Treg cells in mitigating vascular damage in retinal pathologies, as Treg expansion significantly decreased vascular tortuosity and leakage in the diabetic retina. This signifies a restoration of BRB function that was aligned to decrease in VEGF level, which were brought to levels observed in non-diabetic animals. Given that VEGF, besides a proinflammatory cytokine, is also a primary pathogenic mediator of vascular leakage [39], its normalization is an important mechanistic explanation supporting the vasculotropic effects of Treg expansion. The reason by which Treg prevented the diabetes-induced upregulation of VEGF within the retina could be associated with the concomitant decrease of inflammasome. In fact, proinflammatory cytokines are one of the most important triggers of VEGF production by the retina in situations in which the hypoxia, as is the case, is not predominant [49].

The role of Treg expansion in preventing retinal neurodegeneration in DR is evident in various retinal neural populations, including photoreceptors, bipolar cells, and retinal ganglion cells. We found significant retinal thinning in diabetic animals, which Treg expansion effectively prevented. Measurements of rod and cone density, amacrine cells, bipolar morphology, and retinal ganglion cell density were reduced in diabetic groups but restored by Treg expansion, highlighting its effect on DR-associated neurodegeneration. However, retinal neurodegeneration also involves inflammation, glial cell contributions and vascular damage that exacerbate the condition in db/db mice. Our study revealed significant changes in glial reactivity, inflammasome activation, and vascular leakage, all mitigated by Treg expansion. We cannot attribute the beneficial effects of Treg expansion to a single cell type or establish a precise sequence of events based

on our current data. Given the well-documented pleiotropic effects of Tregs across multiple cell types [13–15], the protective impact of Treg expansion in the DR retina likely results from their combined action across various retinal cells.

Current treatments for DR are only addressed to advanced stages of the disease when vision could already be compromised. These treatments include laser photocoagulation, intravitreal injections of corticosteroids or anti-angiogenic agents (i.e. VEGF) and vitreoretinal surgery. All of them present limited effectiveness and are associated with significant adverse effects [50]. At present, treatment of early stages of DR consists in controlling risk factors such hyperglycemia, hypertension and dyslipidemia, but there are no specific treatments to prevent or arrest the progression to advanced stages [51]. Topical (i.e. eye drops) drug formulations based on neuroprotective or anti-inflammatory agents capable of penetrating the retina have emerged as a new therapeutic approach for early stages of DR [52]. This route of administration could reduce systemic side effects and enable patients to self-administer treatment over extended periods. However, clinical evidence in terms of effectiveness and safety is scarce and specific trials are needed [53]. Therefore, the treatment of early stages of DR is an important unmet medical need. Our results have shown that endogenous Treg expansion prevents DR-associated neurodegeneration, inflammation and vascular leakage. In fact, Treg have demonstrated neuroprotective, promyelinating, and immunomodulatory roles in various CNS diseases with pathophysiological features shared with DR [13, 54], and also in aged retinas [21]. Taken together, Treg expansion seems a promising therapeutic avenue for treating DR.

The present study has several limitations. First, we cannot assure that the beneficial effects can be exclusively due to retinal Treg expansion. However, the lack of changes in metabolic profile and diabetic milieu suggests that Treg expansion is primarily responsible for directly reducing inflammation and preventing neuronal and vascular damage associated with DR, instead of changing the overall metabolic profile. The second limitation of this study is the reliance on morphological analysis to infer visual function, which carries the risk of over-extrapolation. While morphological changes provide valuable insights, they do not directly correlate with functional outcomes. To address this, further experiments, such as electroretinography (ERG) and optokinetic assessments, are needed to evaluate visual function more comprehensively and validate the implications of our findings. The last limitation is that the induction of peripheral immunosuppression can be associated risk of infection, especially in patients with T2D. This is a serious problem to translate our findings to the clinical arena. However, new

therapeutic strategies that increase Treg exclusively in damaged neural tissues, such as adeno-associated virus (AAV)-based delivery of IL-2, have proven effective in various experimental CNS-neuroinflammatory models [17, 18] and may open a new therapeutic avenue to build upon our findings.

Conclusions

Treg expansion is a promising new therapeutic strategy to prevent the most important pathological features of DR: neurodegeneration, vascular leakage and glial cell activation. Future work is needed to explore methods to exploit Treg expansion exclusively in the eye to avoid potential systemic side effects and also to delve in the understanding of the intricate molecular mechanisms involved in Treg- mediated retinal protection in T2D.

Abbreviations

AAV	Adeno-Associated Virus
AD	Alzheimer's Disease
CA	Cone Arrestin
CD	Cognitive Decline
ChAT	Choline Acetyltransferase
CNS	Central Nervous System
DAPI	4',6-diamidino-2-phenylindole
db/db	BKS.Cg-Dock7m +/- Leprdb/J
DR	Diabetic Retinopathy
GCL	Ganglion Cell Layer
GFAP	Glial Fibrillary Acidic Protein
HbA1c	glycosylated haemoglobin A1
HBSS	Hanks Buffered Salt Solution
IBA-1	Ionized calcium binding adaptor molecule 1
IL-2	Interleukin-2
INL	Inner-Nuclear Layer
i.p.	Intraperitoneal
IPL	Inner Plexiform Layer
mAb	Monoclonal Antibody
MS	Multiple Sclerosis
NVU	Retinal Neurovascular Unit
ONL	Outer-Nuclear Layer
PBS	Phosphate-Buffered Saline
PD	Parkinson's Disease
RBPMS	RNA-Binding Protein with Multiple Splicing
RGC	Retinal Ganglion Cells
T2D	Type 2 Diabetes mellitus
TH	Tyrosin Hidroxylase
Treg	Regulatory T cells
VEGF	Vascular endothelial growth factor

Supplementary Information

The online version contains supplementary material available at <https://doi.org/10.1186/s12974-024-03323-0>.

Supplementary Material 1
Supplementary Material 2
Supplementary Material 3
Supplementary Material 4
Supplementary Material 5
Supplementary Material 6
Supplementary Material 7
Supplementary Material 8

Supplementary Material 9

Supplementary Material 10

Supplementary Material 11

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Author contributions

Experiments were designed by M.L.S., M.T., A.G.F., A.W.S., C.H. and R.S. Experiments were performed by M.L.S., A.G.F., J.G.S., D.P.M., A.D. M.G.R., A.D., A.A., P.B. and L.R. Experiments were analysed by M.L.S., M.T., A.G.F. and D.P.M. Manuscript was written by M.L.S., A.G.F., C.H. and R.S. and with contributions from all authors. M.L.S. and R.S. oversaw the study.

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Data availability

No datasets were generated or analysed during the current study.

Declarations

Ethics approval and consent to participate

All animal experiments were directed in agreement with the European Community (86/609/CEE) and the guidelines of the Association for Research in Vision and Ophthalmology (ARVO) for the utilization of laboratory animals. The Animal Care and Use Committee of VHIR (CEEA 14/21) authorized the present study.

Consent for publication

N/A.

Competing interests

The authors declare no competing interests.

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