

Interphotoreceptor retinoid-binding protein (IRBP) is downregulated at early stages of diabetic retinopathy

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Abstract

Aims/hypothesis Interphotoreceptor retinoid-binding protein (IRBP) plays a major role in the visual cycle and is essential to the maintenance of photoreceptors. The aim of this study was to determine whether a decrease in IRBP production exists in the early stages of diabetic retinopathy. **Methods** Vitreous samples from diabetic patients with proliferative and non-proliferative diabetic retinopathy

(PDR, NPDR), and from non-diabetic patients with macular hole (control group) were selected for IRBP quantitative assessment by proteomic analysis (fluorescence-based difference gel electrophoresis) and western blot. Human post mortem eyes ($n=16$) from diabetic donors without clinically detectable retinopathy and from non-diabetic donors ($n=16$) were used to determine *IRBP* (also known as *RBP3*) mRNA levels (RT-PCR) and protein content (western blot and confocal microscopy). Retinal neurodegeneration was assessed by measuring glial fibrillar acidic protein (GFAP) and the apoptotic rate. Y79 human retinoblastoma cells were used to test the effects of glucose, TNF- α and IL-1 β on *IRBP* expression and IRBP levels. **Results** Intravitreal IRBP concentration was significantly lower in PDR < NPDR < control in proteomic and western blot analysis. *IRBP* mRNA levels and IRBP protein content were significantly lower in the retinas from diabetic donors than in those from non-diabetic donors. Increased GFAP and a higher degree of apoptosis were observed in diabetic retinas compared with non-diabetic retinas. A dose-dependent downregulation of *IRBP* mRNA expression and IRBP content was detected with glucose, TNF- α and IL-1 β in cultures of Y79 human retinoblastoma cells. **Conclusions/interpretation** Underproduction of IRBP is an early event in the human diabetic retina and is associated with retinal neurodegeneration. The mechanisms leading to this deficit deserve further investigation.

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Keywords Diabetic retinopathy · Human study · Interphotoreceptor retinoid-binding protein · Neurodegeneration · Retina · Vitreous fluid

Abbreviations

DIGE Fluorescence-based difference gel electrophoresis
GFAP Glial fibrillar acidic protein

IRBP	Interphotoreceptor retinoid-binding protein
NPDR	Non-proliferative diabetic retinopathy
PDR	Proliferative diabetic retinopathy
RPE	Retinal pigment epithelium
TdT	Terminal deoxynucleotidyl transferase

Introduction

Diabetic retinopathy has been classically considered to be a microcirculatory disease of the retina. However, before any microcirculatory abnormalities can be detected, retinal neurodegeneration is already present [1–4]. In fact, we have detected apoptosis and glial activation, two characteristic features of retinal neurodegeneration, in retinas from diabetic donors, which were free of microcirculatory abnormalities in ophthalmoscopic examinations performed in the 2 years preceding death [5, 6]. Therefore, the study of the mechanisms that lead to neurodegeneration and of their effects on essential proteins involved in the visual cycle could be crucial for identifying new therapeutic targets in the early stages of diabetic retinopathy.

In the vertebrate retina, vision is initiated and maintained by the photolysis and regeneration, respectively, of light-sensitive pigments in the disk membranes of the photoreceptor outer segments. This cyclical process depends on an exchange of retinoids between the photoreceptors and the retinal pigment epithelium (RPE). There is considerable evidence that the transport of retinoids between these cellular compartments is mediated by the interphotoreceptor retinoid-binding protein (IRBP), a large glycoprotein synthesised in the photoreceptors and extruded into the interphotoreceptor matrix that fills the subretinal space [7, 8]. Apart from participating in the visual cycle, IRBP is important in fatty acid transport and is essential to the maintenance of photoreceptors [9, 10]. In this regard, a reduction of IRBP may precede the loss of photoreceptors seen in some animal models of hereditary retinal degeneration [11, 12]. In addition, knockout (*Irbp*^{-/-}) mice revealed a loss of photoreceptors and profound changes in the structural integrity of the receptor outer segments [10, 13]. Recently, a homozygous missense mutation in the *IRBP* gene (also known as *RBP3*) has been associated with autosomal recessive retinitis pigmentosa in children [14].

We recently compared the protein profiles of human vitreous from diabetic patients with proliferative diabetic retinopathy (PDR) with the vitreous fluid from non-diabetic patients with macular holes using the fluorescence-based difference gel electrophoresis (DIGE) strategy [15]. Our results provided evidence that IRBP is among the lowest produced proteins in the vitreous fluid of patients with PDR. However, it is unknown whether this finding is also

present in the early stages of diabetic retinopathy. This is important because a deficit of IRBP could be involved in the impairment of vision that occurs in diabetic patients even before clinical diabetic retinopathy can be detected. In addition, the lower IRBP content could lead to photoreceptor damage, which is one of the components of retinal neurodegeneration.

On this basis, the aim of the present study was to determine whether decreased IRBP production also exists in the early stages of diabetic retinopathy. For this purpose we used vitreous fluid from diabetic patients with non-proliferative diabetic retinopathy (NPDR) and retinas from diabetic donors without clinically detectable diabetic retinopathy.

Methods

Participants and sample selection

Vitreous fluid Samples of vitreous from four type 2 diabetic patients with PDR, four type 2 diabetic patients with NPDR and from eight non-diabetic patients with idiopathic macular hole (control group), all closely matched by age, were selected from our vitreous bank for proteomic analysis.

An additional 30 samples (ten PDR, ten NPDR and ten macular hole, matched by age) were used for western blot analysis. Exclusion criteria were: (1) previous vitreoretinal surgery; (2) photocoagulation in the preceding 6 months; (3) recent vitreous haemorrhage (less than 3 months before vitrectomy) or intravitreal haemoglobin higher than 5 mg/ml; (4) renal failure (creatinine ≥ 120 $\mu\text{mol/l}$); and (5) other chronic diseases apart from diabetes. For diabetic retinopathy classification we used the International Clinical Diabetic Retinopathy Severity Scale [16]. In accordance with this classification, NPDR samples were from patients with mild ($n=8$) and moderate ($n=2$) NPDR. Diabetic nephropathy was defined according to the criteria of the American Diabetes Association [17]. The main characteristics of controls, NPDR and PDR patients are summarised in Table 1.

Details of vitrectomy and sample collection have been previously described [18].

Retinas

We obtained 16 human post mortem eyes from eight diabetic donors (age 66.4 ± 5.9 years). Eyes were free of fundoscopic abnormalities in ophthalmological examinations performed during the preceding 2 years. Eye cups ($n=16$) obtained from eight non-diabetic donors matched by age (age 67.6 ± 6.9 years) were used as the control group.

Table 1 Main clinical features of participants included in the study

Variables	Participant groups			<i>p</i> value
	With PDR	With NPDR	Control group	
<i>n</i>	10	10	10	
Age (years)	60.3±15.3	61.3±16.2	59.6±10.7	0.85 ^a
Sex (men/women)	5/5	5/5	4/6	0.65 ^a
Hypertension (%)	70	60	50	0.36 ^a
Duration of diabetes (years)	15.9±6.2	6.9±3.1	–	<0.001 ^b
HbA _{1c} (%) ^c	8.1±2.9	7.9±3.5	–	0.47 ^b
Nephropathy (%) ^d	60	20	–	0.08 ^b
Neuropathy (%)	40	10	–	0.15 ^b
Macroangiopathy (%)	50	30	–	0.32 ^b

Unless otherwise indicated, data are expressed as the mean ± SD

^a For comparison among the three groups

^b For comparison between PDR and NPDR participants

^c Mean of the values obtained in the preceding year

^d Diabetic nephropathy was defined according to the criteria of the American Diabetes Association [17], but patients with renal failure (creatinine ≥120 μmol/l) were excluded

The main clinical features and causes of death of diabetic and non-diabetic donors are summarised in Table 2.

The time elapsing from death to eye enucleation was less than 4 h. After enucleation, one eye of each donor was snap-frozen at –80°C and stored until assayed for mRNA and western blot analyses. The other was fixed in 4% (wt/vol.) paraformaldehyde and embedded in paraffin for the immunohistochemical study.

All ocular tissues were obtained and processed following the recommendations of the Declaration of Helsinki for research involving human tissue and those of the Directive of the European Parliament and the Council of the European Union (Directive 2004/23/EC). In addition this study was approved by the Ethics Committee of our hospital.

Sample preparation, DIGE proteomic analysis and protein identification Sample preparation along with details of DIGE proteomic analysis and protein identification by mass spectrometry have been described elsewhere [15].

IRBP assessment

RNA extraction and IRBP mRNA quantification Neuroretinas and RPE were removed under microscopic dissection of isolated eye cups from donors. Total RNA was obtained from isolated human neuroretinas of eye donors using a kit (RNeasy Mini; Quiagen Distributor IZASA, Barcelona, Spain) according to the manufacturer's protocol.

We used 1 μg total RNA for reverse transcription. Quantitative real-time PCR was performed with 40 ng of cDNA and TaqMan Universal Mastermix (Applied Biosystems, Madrid, Spain) in an Applied Biosystems device (7900 model). TaqMan pre-made gene expression assays (Applied Biosystems) were used for specific gene amplifi-

Table 2 Clinical features of diabetic and non-diabetic donors included in the study

Variables	Diabetic donors ^a	Non-diabetic donors	<i>p</i> value
<i>n</i>	8	8	
Age (years)	66.4±5.9	67.6±6.9	0.47
Sex (men/women)	5/3	6/2	0.50
Diabetes duration (years)	5.8±3.1	–	
HbA _{1c} (%) ^b	7.4±0.8	–	
Diabetes treatment			
Diet only	3		
Oral agents	5		
Insulin	0		
Cause of death			
Cardiovascular disease	5	3	
Malignant neoplasm	3	4	
Traffic accident	0	1	

Data are expressed as *n* or as the mean ± SD

^a All diabetic donors had type 2 diabetes

^b Mean of the values obtained in the preceding 2 years

cation (human *IRBP*: Hs00161253_m1). Each sample was assayed in duplicate and negative controls were included in each experiment. Automatic relative quantification was obtained with a software package (ABI Prism 7900 SDS; Applied Biosystems) using β -actin as the endogenous gene expression reference (Hs9999903_m1; Applied Biosystems).

Western blot analysis IRBP was assessed in the vitreous fluid of diabetic and control patients, and in the neuroretina from donors. Total protein extracts from the neuroretina were quantified using an assay (Bradford Protein Assay; Bio-Rad, Barcelona, Spain). Equal amounts of protein (5 μ g) were loaded on a 7.5% (vol./vol.) SDS-polyacrylamide gel and electroblotted on to nitrocellulose membranes (Immobilon; Millipore, Barcelona, Spain). Membranes were probed with IRBP antibodies (kindly provided by J. M. Nickerson, Emory University, Atlanta, GA, USA) at a dilution of 1:2,000. Anti human β -actin (Calbiochem, Bionova, Madrid, Spain) was used for normalisation. Blots were developed by chemoluminescence (Supersignal West Dura Extended Duration Substrate; Pierce, Madrid, Spain). Densitometry of the developed bands was obtained in a calibrated GS-800 device (Bio-Rad).

Immunohistochemistry for IRBP Neuroretina sections (6 μ m) were dewaxed following standardised methods. Sections were permeabilised and blocked for 1 h in PBS buffer with 0.05% (vol./vol.) Tween and 2% (wt/vol.) BSA. Primary antibodies (anti-IRBP and anti-glia fibrillar acidic protein [GFAP] [Sigma, Madrid, Spain]) were incubated overnight at 4°C in the same blocking buffer. Afterwards, sections were extensively washed, then incubated with Alexa Fluor 488 or Alexa Fluor 568 anti-rabbit (1:200; Molecular Probes, Invitrogen, Madrid, Spain) for 1 h. A negative control section was performed by omitting the corresponding antiserum from the primary antibody solution. Double staining was performed including two antibodies in the same section. After three washes in PBS buffer, sections were mounted on fluorescence mounting medium with DAPI (H-1200; Vector, CA, USA) and observed under a spectral confocal microscope (FV1000; Olympus, Madrid, Spain). Quantification of immunofluorescence was performed directly on confocal images of 1.024 pixels with an Olympus Fluoview imaging system (Olympus, Barcelona, Spain). To quantify IRBP in the neuroretina, the total fluorescence intensity values corresponding to ten field frame images (40 \times numerical aperture 0.9) of each retina sample were measured. These results were then normalised taking into account the area analysed. All these calculations were made using specific software (Fluoview ASW 1.4; Olympus).

Measurements of retinal neurodegeneration

Apoptosis: TUNEL Neuroretina sections were processed using an in situ apoptosis detection kit (ApoTaq Plus Fluorescein; Millipore, Madrid, Spain) according to the manufacturer's protocol. Briefly, sections were deparaffinised with xylene and ethanol, and the tissue pretreated with proteinase K (20 μ g/ml) for 15 min before digoxigenin-labelled dUTP was applied in the presence of the terminal deoxynucleotidyl transferase (TdT) enzyme (TdT concentration was changed to half the recommended dilution). Nuclear staining in apoptotic cells was detected by FITC staining with anti-digoxigenin conjugate. To counterstain, propidium iodide antifade solution (Millipore) was used as a mounting media. A non-specific control was included by omitting the TdT enzyme in the incubation step. Positive controls were incubated with DNase I prior to incubation with TdT. Quantification was performed by counting individually fluorescence-labelled cells in three sections per sample. The apoptotic index was calculated by dividing the total number of apoptotic cells by the total number of cells in each layer and in a 0.5 cm² area. To avoid false positive results, only TUNEL-positive cells in which nuclear fragmentation and/or condensation was present were rated as apoptotic. For this purpose we used inter-differential contrast (phase-contrast microscopy) and propidium-iodide immunofluorescence.

Glial activation: GFAP immunofluorescence Tissue sections were incubated overnight at 4°C with the primary antibody anti-human GFAP (1:200; Sigma). After washing, sections were incubated with Alexa Fluor 488 (Molecular Probes, Eugene, OR, USA) secondary antibody for 1 h. GFAP immunofluorescence in the neuroretina was quantified using a laser confocal scanning microscope. The procedure was the same as mentioned above for IRBP.

Cell culture and treatments

Y79 human retinoblastoma cells (ATCC HTB 18) were maintained in suspension in RPMI 1640 under standard conditions with 10% (vol./vol.) fetal bovine serum, penicillin (100 U/ml) and streptomycin (100 μ g/ml) (Cultek, Barcelona, Spain) at 37°C and 5% (vol./vol.) CO₂. For glucose treatments, cells (2 \times 10⁴/ml) were serum-starved and cultured for 72 h in increasing glucose concentrations (5.5, 10, 15, 25 mmol D-glucose; Sigma) in RPMI 1640 with 0.1% (vol./vol.) fetal bovine serum. To rule out potential bias due to osmotic effect, D-mannitol (Sigma) was used at appropriate concentrations to equilibrate the osmosis. In another set of experiments, Y79 cells were plated on poly-D-

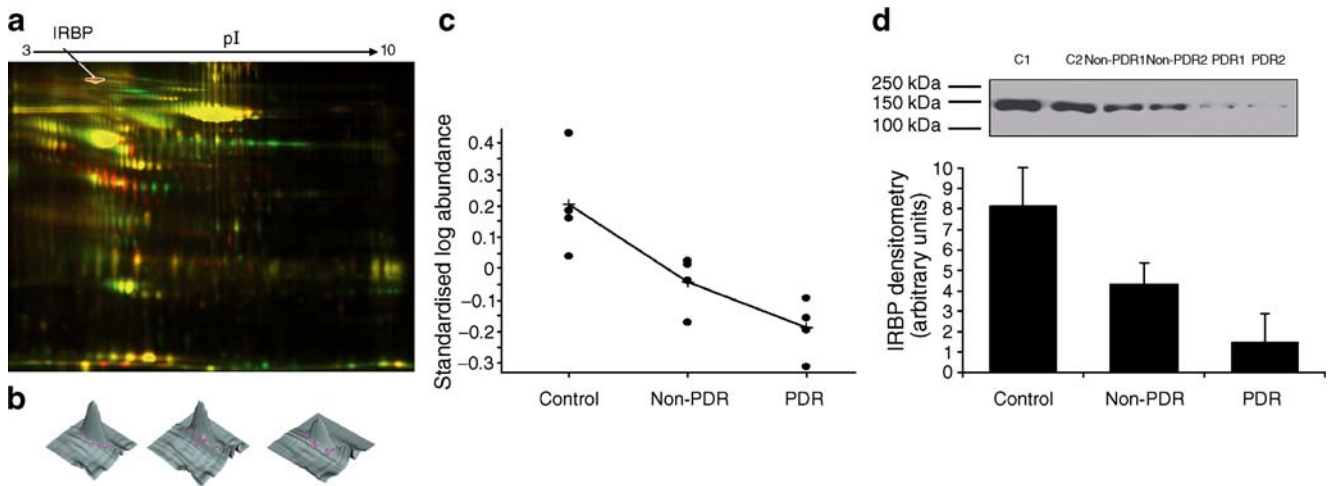


Fig. 1 **a** Superimposed images in pseudocolour from samples labelled with Cy3 (green, PDR vitreous proteins) and Cy5 (red, Control vitreous proteins), and run on a two-dimensional DIGE gel. The range of the horizontal dimension is in isoelectric points (pI). The vertical dimension ranges from 15 kDa (bottom) to 200 kDa (top). The position of the spot corresponding to IRBP is marked. **b** Three-dimensional images of the IRBP spot corresponding to the image of a Control, Non-PDR and PDR sample (left to right). **c** Standardised abundance plot for IRBP displaying the log of abundance observed for

the spot in each of the four gel images corresponding to control, non-PDR and PDR samples, as indicated, after standardising the values using the internal standard pool images (Cy2) of each of the four gels. The line links the average abundance values for each group of samples (crosses). Student's *t* test for the difference in abundance between each two groups results in $p < 0.05$ in all cases. **d** Western blot analyses of vitreous samples corresponding to non-diabetic participants (C1, C2), and non-PDR and PDR patients. Blots were quantified by densitometry, with bars representing the mean \pm SD

lysine coated plates (BD, Madrid, Spain). After 24 h medium was replaced and either TNF- α (10, 25 and 50 ng/ml) or IL-1 β (10 and 25 ng/ml) (Prepro, London, UK) were added for additional 72 h. To study IRBP expression, immunoblotting and real-time PCR were performed.

Statistical analysis

Results are expressed as mean \pm SD. Comparisons between groups were performed using Student's *t* test or ANOVA. Levels of statistical significance were set at $p < 0.05$.

Results

IRBP in vitreous fluid

DIGE and western blot analyses The results for DIGE proteomic analysis of IRBP are summarised in Fig. 1a–c. To further validate these results, we assessed ten PDR, ten non-PDR and ten macular hole vitreous samples in addition to the samples used in DIGE. Intravitreal IRBP concentration (densitometric arbitrary units) was significantly lower in PDR < NPDR < control (1.95 ± 1.63 , 4.47 ± 2.30 , 8.19 ± 3.41 , $p = 0.03$) (Fig. 1d). As expected, IRBP was not detected in any serum sample (data not shown). This is because synthesis of IRBP is highly restricted to the neuroretina and pineal cells.

IRBP in retina

β -Actin mRNA expression was similar between diabetic and non-diabetic retinas ($p = \text{NS}$). Thus, we calculated *IRBP* mRNA gene expression after normalising with β -actin.

IRBP mRNA levels obtained in diabetic donors were significantly lower than those obtained from non-diabetic donors (1.16 ± 0.93 vs 2.48 ± 2.01 , $p = 0.03$) (Fig. 2a).

Densitometric analysis of immunoblots showed lower IRBP levels in the retinas from diabetic donors than in

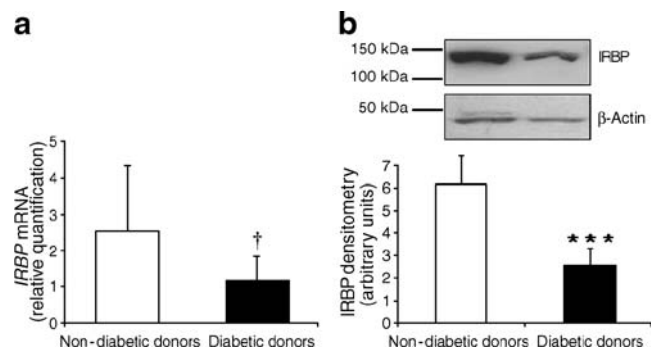
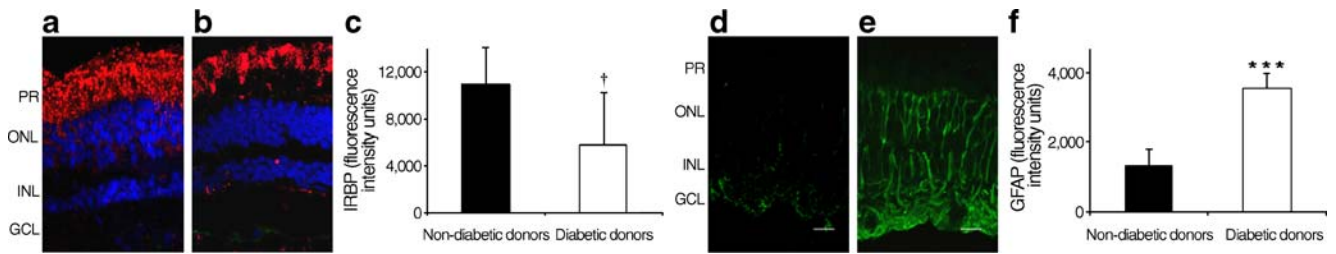


Fig. 2 **a** Real-time quantitative RT-PCR analysis of *IRBP* mRNA in human retinas from diabetic and non-diabetic donors. *IRBP* mRNA gene expression was calculated after normalising with β -actin. Bars represent the mean \pm SD, $\dagger p = 0.03$. **b** Western blot analyses of IRBP in human retina from non-diabetic and diabetic donors, with quantification by densitometry. Bars represent mean \pm SD, $***p < 0.001$



a non-diabetic donor (d) and a diabetic donor (e). Key, as above (a, b). In the diabetic retina, the Müller cells' endfeet show abundant GFAP immunofluorescence and the radial processes stain intensely throughout the inner and outer retina. Scale bar, 20 μ m. f Quantification of GFAP immunofluorescence in non-diabetic and diabetic retinas. Bars represent mean \pm SD, *** $p<0.001$

those from non-diabetic donors (2.58 ± 0.69 vs 6.14 ± 1.17 , $p<0.001$) (Fig. 2b). Laser scanning confocal images of IRBP immunofluorescence are shown in Fig. 3a–c. IRBP immunofluorescence was significantly lower in the retinas from diabetic donors than in those from non-diabetic donors ($5,787 \pm 4,531$ vs $10,963 \pm 3,787$, $p=0.04$).

Effect of diabetes in retinal neurodegeneration: apoptosis and glial activation

To assess the presence of early diabetic damage in the retinas studied, we determined the rate of apoptosis and glial activation, two well-recognised markers of retinal neurodegeneration. The percentage of apoptotic cells was higher in diabetic neuroretinas than in the age-matched non-diabetic retinas, but the differences, while having possible clinical consequences, were not statistically significant (outer nuclear layer $0.48 \pm 0.44\%$ vs $0.05 \pm 0.06\%$, $p=0.09$; inner nuclear layer 2.23 ± 2.04 vs 1.29 ± 2.31 , $p=0.34$; ganglionar cell layer $2.97 \pm 0.60\%$ vs $1.30 \pm 0.43\%$, $p=0.08$). Neuroglial activation was demonstrated in diabetic eyes by an increase of GFAP production in the neuroretina. GFAP immunofluorescence was significantly higher in diabetic than in non-diabetic retinas ($3,123 \pm 418$ vs $1,119 \pm 489$, $p<0.01$) (Fig. 3d–f).

Discussion

IRBP, a large glycoprotein highly restricted to the interphotoreceptor matrix, facilitates the transport of retinoids between the photoreceptor outer segments and RPE and therefore plays a major role in the visual cycle [7, 8]. In addition it is involved in the maintenance of photoreceptors [9, 10]. In the

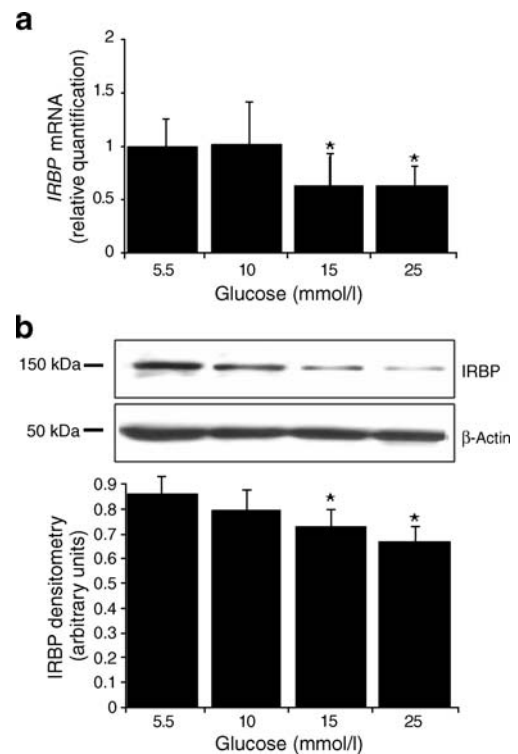
Effect of high glucose and cytokines on IRBP production in vitro

Glucose caused a dose-dependent reduction of *IRBP* mRNA expression and IRBP content in human Y79 retinoblastoma cultures (Fig. 4a, b). TNF- α and IL-1 β also decreased IRBP production by human Y79 retinoblastoma cells (Fig. 5a–d).

Effect of high glucose and cytokines on IRBP production in vitro

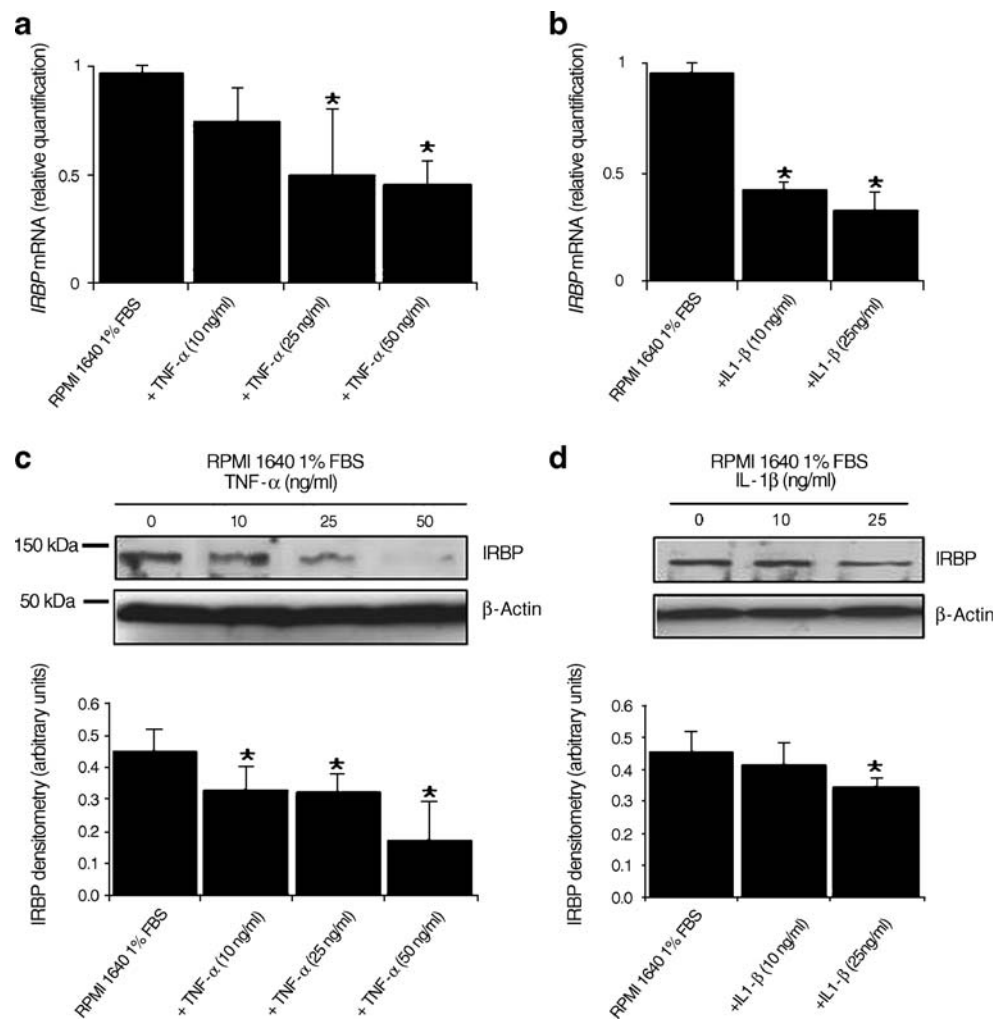
Glucose caused a dose-dependent reduction of *IRBP* mRNA expression and IRBP content in human Y79 retinoblastoma cultures (Fig. 4a, b). TNF- α and IL-1 β also decreased IRBP production by human Y79 retinoblastoma cells (Fig. 5a–d).

Effect of high glucose and cytokines on IRBP production in vitro



Effect of high glucose and cytokines on IRBP production in vitro

Fig. 5 Effect of treatment with TNF- α (10, 25 and 50 ng/ml) and IL-1 β (10 and 25 ng/ml) on *IRBP* mRNA levels (RT-PCR) (a, b) and protein content (western blot and quantification by densitometry) (c, d) in Y79 cell cultures. FBS, fetal bovine serum. Control: RPMI 1640 1% FBS. Bars represent mean \pm SD, * p <0.05 vs control



present study we provide evidence that downregulation of *IRBP* is an early event in the human diabetic retina.

One of the limiting factors in diabetic retinopathy research is the impossibility, for obvious reasons, of obtaining retinal biopsies from diabetic patients. Since retinal biopsy is not available, vitreous fluid from diabetic patients undergoing vitreoretinal surgery is currently used to indirectly explore the events taking place in the retina. However, because most vitrectomies are indicated in PDR, the results obtained in the study of vitreous fluid are only valid for advanced stages of diabetic retinopathy. In the present study, we obtained vitreous fluid from diabetic patients with mild or moderate NPDR, who underwent vitreoretinal surgery for other reasons unrelated to diabetic retinopathy. This approach enabled us to demonstrate for the first time that patients with NPDR have lower intravitreous *IRBP* levels than non-diabetic patients. Because disruption of the blood–retinal barrier is an early event in the pathogenesis of diabetic retinopathy, one of the confounding factors when interpreting results obtained in vitreous fluid measurements is that they can also reflect

serum levels, rather than local production by the retina. However, given that *IRBP* is only produced by photoreceptors and pineal cells [7, 19], intravitreous *IRBP* concentration is a reliable index of its retinal synthesis.

In addition, we found that expression and content of *IRBP* were lower in retinas from diabetic donors than in those of non-diabetic donors. Therefore, there is a low abundance of *IRBP* in the diabetic eye. Given that our diabetic donors were free of ophthalmoscopic abnormalities, it could be argued that retinas from these patients were without any diabetes-related damage. However, this is not the case, because glial activation, a characteristic feature of diabetic retinal neurodegeneration, was present. Furthermore, the higher, but not significant rate of apoptosis observed in the neuroretinas from diabetic donors strongly suggests that an *IRBP* deficit exists in very early stages of diabetic retinopathy.

The mechanisms leading to reduced amounts of *IRBP* in the diabetic eye could not be determined with the current data. *IRBP* is a dominant antigen in autoimmune uveitis [20]. Recently, moreover, proteolysis of *IRBP* has been shown to occur [21]. The loss of *IRBP* detected in the

vitreous fluid of diabetic patients could be a consequence of increased proteolytic activity due to leucocyte-released proteases [22, 23]. In fact, due to the anatomical blood–retinal barriers, IRBP has a hidden nature, but the disruption of the blood–retinal barrier that occurs in diabetic retinopathy facilitates antigen presentation and the subsequent activation of autoreactive T cells [21]. Therefore, it is possible that IRBP acts as a first trigger of the inflammatory process present in the early stages of diabetic retinopathy and that the lower levels of IRBP detected in the diabetic eye are the consequence of increased proteolytic activity secondary to inflammation. However, we found low mRNA levels of *IRBP* in the retinas of diabetic donors and, in consequence, lower production, rather than increased catabolism seems to be the primary cause of the IRBP deficit detected in the diabetic eye. In addition, high glucose levels and proinflammatory cytokines were able to downregulate *IRBP* expression and IRBP content in human Y79 retinoblastoma cells, suggesting that the diabetic milieu might play an essential role in the underproduction of IRBP in diabetic patients. Nevertheless, it should be noted that lower expression of *IRBP* was found in Y79 cells in comparison with non-transformed cells. Therefore, our results cannot be readily extrapolated to the clinical setting. As a result, further studies in primary cultures of photoreceptors are needed.

Neurodegeneration is an early event in the pathogenesis of diabetic retinopathy, antedating and participating in the microcirculatory abnormalities that occur in the condition [2, 3, 24]. Neuroretinal damage produces functional abnormalities such as colour vision defects, reduced contrast sensitivity and abnormalities in dark adaptation. These alterations can occur before microvascular lesions are detectable in ophthalmological examinations, but they tend to be predictive of PDR, which ultimately leads to degenerative changes and significant visual impairment [24–26]. The design of our study does not allow us to say whether the lower IRBP content in diabetic retinas is a cause or a consequence of retinal neurodegeneration. However, given that IRBP has neuroprotective properties, it is possible that lower production of IRBP participates in diabetic retinal neurodegeneration. In addition, since IRBP is synthesised by photoreceptors, the higher apoptosis rate detected in the photoreceptor layer could be involved in lower IRBP production, thus initiating a vicious circle that could be added to the classical pathways involved in diabetic retinopathy development.

In conclusion, lower IRBP production is an early event in the human diabetic retina and is associated with retinal neurodegeneration. Since IRBP plays a major role in the visual cycle and is essential for photoreceptor survival, an IRBP deficit could be involved in the pathogenesis of diabetic retinopathy. However, further studies are required to determine the cause of IRBP underproduction in the diabetic retina and its clinical implications.

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Duality of interest The authors declare that there is no duality of interest associated with this manuscript.

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