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OPEN Adalimumab Reduces Photoreceptor Cell Death in A Mouse Model of Retinal Degeneration

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Growing evidence suggests that inflammation is involved in the progression of retinitis pigmentosa (RP) both in patients and in animal models. The aim of this study was to investigate the effect of Adalimumab, a monoclonal anti-TNFlpha antibody, on retinal degeneration in a murine model of human autosomal recessive RP, the rd10 mice at postnatal day (P) 18. In our housing conditions, rd10 retinas were seriously damaged at P18. Adalimumab reduced photoreceptor cell death, as determined by scoring the number of TUNEL-positive cells. In addition, nuclear poly (ADP) ribose (PAR) content, an indirect measure of PAR polymerase (PARP) activity, was also reduced after treatment. The blockade of TNFlpha ameliorated reactive gliosis, as visualized by decreased GFAP and IBA1 immunolabelling (Müller cell and microglial markers, respectively) and decreased up-regulation of TNFlpha gene expression. Adalimumab also improved antioxidant response by restoring total antioxidant capacity and superoxide dismutase activity. Finally, we observed that Adalimumab normalized energetic and metabolic pattern in rd20 mouse retinas. Our study suggests that the TNF α blockade could be a successful therapeutic approach to increase photoreceptor survival during the progression of RP. Further studies are needed to characterize its effect along the progression of the disease.

Retinitis pigmentosa (RP) is a group of inherited retinal dystrophies characterized by progressive and irreversible loss of vision that in most models studied, parallels photoreceptor cell death¹⁻³. RP is the leading cause of genetic blindness in adults with an estimated incidence of 1 in 3,500 ~ 4,500 human births⁴. Patients with RP typically loose night vision in adolescence, peripheral vision in young adulthood, and central vision later in life, due to progressive, sequential loss of rod and cone photoreceptor cells. Although many therapeutic approaches have been developed to prevent photoreceptor cell death, no effective treatment is still available. More than 70 genes, including phosphodiesterase 6 (PDE6) subunit genes, have been identified to date whose mutations cause different forms of RP^{5,6}. Besides the genetic

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defect, evidence suggests that oxidative stress and neuroinflammation contribute to its progression^{7–9}. In particular, inflammatory processes including microglial activation and upregulation of inflammatory cytokines (TNF α , IL-1 β , etc.) and chemokines (MCP1, RANTES, etc.) have been described in patients as well as in animal models of RP^{10–12}.

Tumor necrosis factor alpha (TNF α) is a pleiotropic cytokine essential for the induction and maintenance of the inflammatory immune responses. It is a well characterized mediator of cellular activities including proliferation, survival, differentiation, inflammation and cell death. In the retina, it is likely secreted from activated macrophages, astrocytes, microglia and Müller glial cells. TNF α can trigger several well-characterized death-promoting (caspase-dependent and caspase-independent cell death) and survival-promoting pathways, depending upon the predominating signalling pathway in the particular cell type¹³. In the eye, TNF α appears to have a role in the pathogenesis of inflammatory diseases such as uveitis¹⁴, as well as in retinal degenerations such as diabetic retinopathy, age-related macular degeneration and, recently, in RP^{11,15-17}.

Several anti-TNF α agents (Infliximab, Adalimumab, Etarnecept, Golimumab and Certolizumab pegol) have been developed and approved for clinical use in inflammatory diseases such as rheumatoid arthritis, psoriasis and ankylosing spondylitis¹⁸. These anti-TNF α agents are antibodies against TNF α or TNF α receptor. In Ophthalmology, they are widely used as an alternative to traditional immunosuppressive treatments in non-infectious uveitis. More recently anti-TNF α agents have been used for retinal diseases such as neovascular age-related macular degeneration, diabetic macular edema and retinal vein occlusions^{19–21}.

To further explore the *in vivo* potential benefits of blocking TNF α we adopted the *rd10* mouse, a model of human autosomal recessive RP. This mouse carries a mutation on the β subunit of the cGMP PDE6 gene ($Pde6\beta$) that produces retinal degeneration².

In this study, we first analysed the early stages of retinal degeneration and the profile of TNF α expression in rd10 mice under our housing conditions. We observed a peak of TNF α gene expression and photoreceptor cell death at postnatal day (P) 18. Therefore, we studied the effect of Adalimumab, a monoclonal antibody against TNF α , on the progression of the retinal degeneration at this age. We observed that Adalimumab prevented TNF α upregulation, reduced photoreceptor cell death, slowed microglial and Müller cell activation, improved antioxidant response and ameliorated the energetic and metabolic dysfunction at P18. Based on these results we suggest that anti-TNF α therapies could be promising treatments to improve photoreceptor cell survival in humans with RP. However, further studies are needed to investigate the molecular mechanisms involved in the protective effect of anti-TNF α agents and their long-term effect.

Results

Temporal progression of retinal degeneration in rd20 **mice.** Light exposure accelerates progression of retinal degeneration in many animal models of RP^{22,23}. Hence we first investigated the time course of retinal degeneration in rd10 mice from P13 to P20 stabulated under light intensities between 98 and 131 lux. Sytox Green nuclear staining showed a decline of the number of photoreceptor cell nuclei rows in the outer nuclear layer (ONL) in rd10 mice. As expected, we did not find a significant difference between control and rd10 mice at P13. But the number of cell nuclei rows in the ONL dropped abruptly from eleven to five from P15 to P20 (Fig. 1A). Degeneration was evident at P18 (Fig. 1D).

Then, we assessed photoreceptor cell death by TUNEL staining which detects DNA breaks in both apoptotic and necrotic nuclei. From P15 to P20 there was a significant increase of TUNEL-positive cells in rd10 mice, reaching a peak at P18 (12.5 \pm 2.4 TUNEL-positive cells/normalized ONL area (five-fold increase between P15 and P18), Mann-Whitney U test, p < 0.0001) (Fig. 1B,D, and Supplementary Table S1).

It has been previously described that TNF α is upregulated in the retina of rd10 mice previous to photoreceptor cell death peak^{11,24}. Therefore, we analysed gene expression of TNF α during the time course of retinal degeneration in our housing conditions. As shown in Fig. 1C, TNF α was upregulated from P13 to P20 with the highest expression at P18 (6.1 \pm 1.2 relative expression (six-fold increase over control retinas), Mann-Whitney U test, p = 0.0002) concomitantly with the peak of TUNEL-positive cells. At P18 a significant increase of TNF α protein level was also observed in the retina of rd10 mice (mean = 65.5 pg/mg protein, confidence interval [25.1–170.9], generalized least squares model, p = 0.0034) compared to control retinas (mean = 17.1 pg/mg protein, confidence interval [15.8–18.5]).

In view of these results, P18 was chosen as the end point for the ensuing analysis of the potential protective effect of Adalimumab.

Adalimumab reduced photoreceptor cell loss and PARP activation. To determine whether Adalimumab treatment protects against photoreceptor cell death, we quantified the normalized ONL thickness ratio and the number of rows of nuclei in the ONL at P18. Normalized ONL thickness decreased slower in Adalimumab-treated rd10 retinas (reduction of 20% over control retinas) than in vehicle-treated rd10 retinas (reduction of 40% over control retinas) (Fig. 2A and Supplementary Table S1). A mixed linear model analysis also revealed that Adalimumab significantly reduced (p = 0.047)

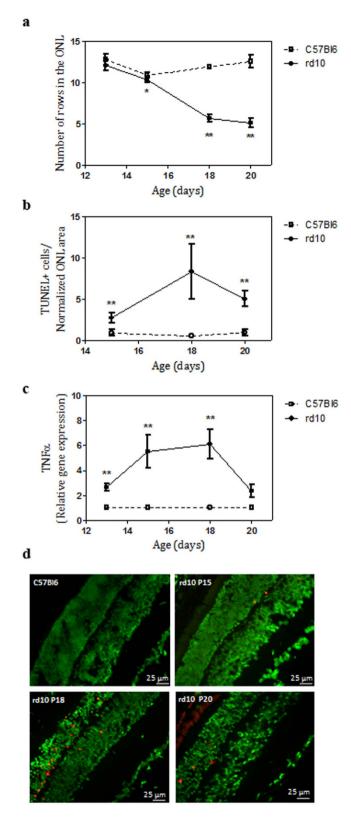


Figure 1. Time course of retinal degeneration in the rd10 mouse retina. (a) Quantitation of the number of rows of photoreceptor nuclei from P13 to P20 in control (C57Bl6) and rd10 mice. (b) Number of TUNEL-positive cells by normalized ONL thickness in control and rd10 mice from P15 to P20. (c) Relative TNF α gene expression from P13 to P20 in rd10 mice. Values are the mean \pm SEM of, at least, five retinas per group. Values that are significantly different are indicated by asterisks *p < 0.05, **p < 0.001, ***p < 0.0001 (Mann-Whitney U test). (d) Photoreceptor cell death was visualized by TUNEL (red) in Sytox Green-counterstained in retinal sections from control mice at P18 and from rd10 mice at P15, P18 and P20.

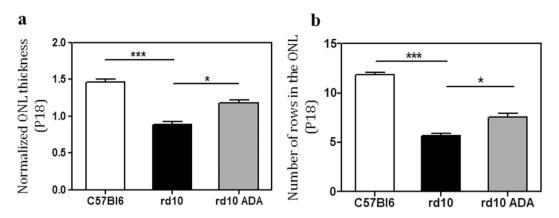


Figure 2. Adalimumab administration decreased photoreceptor cell loss in the rd10 mouse retina at P18. Bar graph illustrates the effect of Adalimumab (ADA) on normalized ONL thickness (a) and number of rows of nuclei in the ONL (b) Values are the mean \pm SEM of six retinas per group. Values that are significantly different are indicated by asterisks *p < 0.05, ***p < 0.001 (Mixed linear model analysis).

photoreceptor cell loss (7.6 \pm 0.4 number of rows of nuclei in the ONL) compared to vehicle-treated rd10 retinas (5.9 \pm 0.3 number of rows of nuclei in the ONL) (Fig. 2B and Supplementary Table S1).

We previously described that Infliximab, other monoclonal anti-TNF α antibody, significantly reduced the number of TUNEL-positive cells in Zaprinast-treated explants of porcine retina²⁵. In the current study we observed that intraperitoneal administration of Adalimumab significantly decreased (\sim 78%, p=0.048) the number of TUNEL-positive cells in the ONL at P18 (2.7 \pm 0.4 TUNEL-positive cells) compared to vehicle-treated rd10 retinas (12.5 \pm 2.4 TUNEL-positive cells) (Fig. 3A,C, and Supplementary Table S1). Therefore the blockade of TNF α was preventing photoreceptor cell loss in rd10 mice. However, the mechanisms of TNF α -induced cell death remained unclear.

Paquet-Durand *et al.* (2007) demonstrated non-apoptotic mechanisms involved in photoreceptors cell death such as the activation of PARP which led to necroptosis in a similar mouse model of RP²⁶. To assess whether TNF α inhibition prevented PARP activation, we visualized the content of PAR polymers (an indirect marker for PARP activity) and its retinal location at P18. As shown in Fig. 3B, an intense and nuclear PAR staining was found in photoreceptor cells of vehicle-treated rd10 retinas. In contrast, Adalimumab substantially decreased (up to 90%, p < 0.001) nuclear PAR staining in photoreceptor cells (2.1 \pm 0.6 PAR-positive cells), compared to vehicle-treated rd10 retinas (25.1 \pm 5.5 PAR-positive cells) (Fig. 3C). Taking together these results suggested that Adalimumab could slow down photoreceptor cell death through PARP pathway inhibition in rd10 mice. Further studies will be designed to test this hypothesis.

Adalimumab reduced reactive gliosis. Chronic microglia activation is associated with various neurodegenerative diseases including RP. We investigated whether Adalimumab treatment reduced microglial activation in rd10 mouse retinas. Microglial cells were identified by specific labelling with Iba1, a microglia/macrophage-specific calcium-binding protein. The level of migration from the inner to the outer layers of retina was employed as an indicator of microglial activation. Iba1 immunostaining revealed that microglia was located in all layers, especially in outer layers in vehicle-treated rd10 retinas. Conversely, after intraperitoneal administration of Adalimumab, microglia was preferentially located in the inner layers of the retina. A few microglial cells were observed in the outer plexiform layer but not in the ONL in Adalimumab-treated rd10 retinas (Fig. 4A). To quantify microglial behaviour, we performed a weighted analysis where values close to 1 indicate a high rate of microglial migration to outer layers. Statistical analysis confirmed that microglial activation was reduced (p = 0.006) in Adalimumab-treated rd10 retinas (0.28 \pm 0.03) compared to vehicle-treated rd10 retinas (0.43 \pm 0.02) at P18 (Fig. 4C and Supplementary Table S1).

Gliosis commonly involves upregulation of the glial fibrillary acidic protein (GFAP), an intermediate filament protein, in Müller glial cells. We also studied whether the blockade of TNF α with Adalimumab ameliorates glial activation in rd10 mice. Figure 4B shows that GFAP was upregulated in vehicle-treated rd10 retinas (p < 0.001) compared to control retinas. Adalimumab tended to decrease (up to 50%) (p = 0.27) GFAP-positive immunolabelling compared to vehicle-treated rd10 retinas at P18 (Fig. 4C).

After evaluating the effect of Adalimumab in prevention of reactive gliosis, we analysed whether Adalimumab affected gene expression of the inflammatory mediators TNF α and IL-6 by real time PCR at P18. After Adalimumab treatment, TNF α expression was significantly reduced (p = 0.006) (Fig. 4C and Supplementary Table S1). At P18 we did not observe changes in IL-6 expression neither in vehicle-treated rd10 retinas nor in Adalimumab-treated rd10 retinas (data not shown).

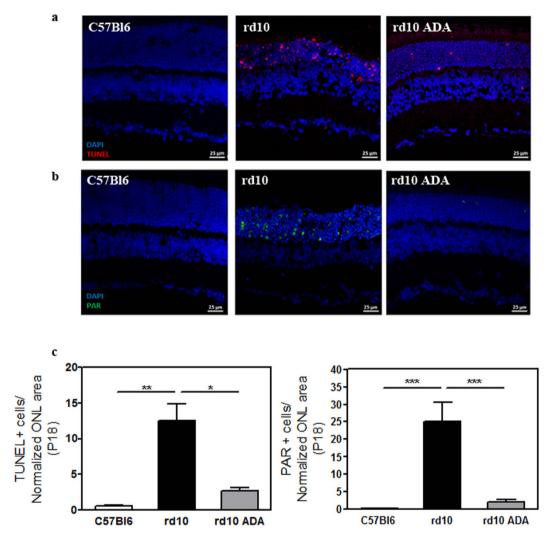


Figure 3. Adalimumab significantly reduced photoreceptor cell death in the rd10 mouse retina at P18. Representative photomicrographs of retinal sections showing (a) TUNEL-stained sections revealing dead photoreceptors in DAPI-counterstained and (b) PAR content in DAPI-counterstained retinal sections. (c) Bar graph illustrates the effect of Adalimumab (ADA) on the number of TUNEL-positive nuclei and PAR-positive cells. Values are the mean \pm SEM of, at least, five retinas in each group. Values that are significantly different are indicated by asterisks *p < 0.05, **p < 0.01, ***p < 0.001 (Mixed linear model analysis).

Adalimumab ameliorated energetic and metabolic dysfunction. It has been described a coordinated downregulation of several metabolic pathways at the onset of retinal degeneration, affecting amino acid biosynthesis, glycerol phospholipid metabolism and oxidative phosphorylation in 14-d-old Rh1P37H retinas, a Drosophila model expressing the equivalent of the most common RP-linked mutation, $RhoP23H^{27}$. First, we assessed whether rd10 mice presented a deregulated expression of enzymes of energy metabolism and mitochondrial structure and dynamics at P18 by quantitative reverse phase protein microarrays (RPPM)²⁸ using validated antibodies (Supplementary Figure S1). We analysed the expression of proteins of glycolysis (GAPDH, LDHA), electron transport and oxidative phosphorylation (OXPHOS) (NDUFS3, SDHB, CORE 2, COXII and β-F1-ATPase), mitochondrial structure (HSP60) and dynamics (OPA1), β-oxidation (HADHA), the pentose phosphate pathway (G6PDH), the antioxidant response (SOD1, SOD2 and catalase) and the expression of β-actin as reference protein (Supplementary Figure S1).

At the onset of RP, the expression of most of these markers was significantly downregulated in retinas of rd10 mice leading to a global energy depletion that might contribute to cell death (Table 1). No significant differences in β -actin were found between control $(0.64\pm0.05 \text{ arbitrary units (a.u.)/ng total protein)}$ and rd10 retinas $(0.71\pm0.06 \text{ a.u./ng total protein)}$. Hence we observed an energy depletion and mitochondrial dysfunction during photoreceptor cell death. As shown in Table 1, intraperitoneal administration of Adalimumab significantly ameliorated this phenotype in retinas from rd10 mice at P18.

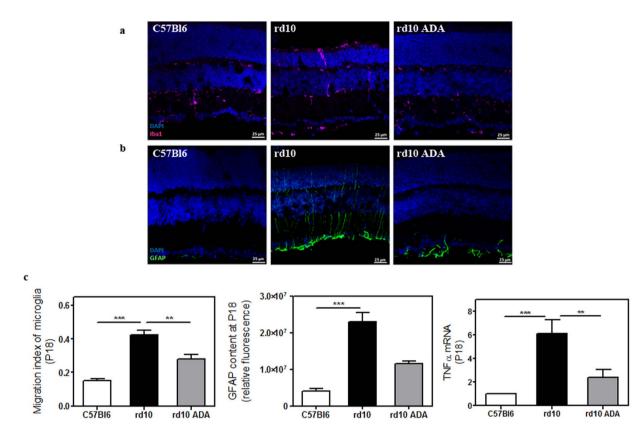


Figure 4. Adalimumab ameliorated reactive gliosis in the rd10 mouse retina at P18. Representative photomicrographs of retinal sections showing (a) Iba1-labelling to visualize microglial cells and (b) GFAP content in DAPI-counterstained sections. (c) Bar graph illustrates the effect of Adalimumab (ADA) on migration index of microglia, the corrected fluorescence of GFAP content and TNF α gene expression. Values are the mean \pm SEM of, at least, five retinas per group. Values that are significantly different are indicated by asterisks **p < 0.01, ***p < 0.001 (Mixed linear model analysis).

	Protein	С	rd10	rd10 ADA
OXPHOS	NDUFS3	3.1 ± 0.1	1.5 ± 0.2*	2.4 ± 0.2^a
	SDHB	2.1 ± 0.1	1.3 ± 0.1*	2.1 ± 0.2^a
	CORE2	1.4 ± 0.1	0.8 ± 0.1*	1.3 ± 0.1^a
	COX II	2.0 ± 0.1	0.9 ± 0.1*	2.1 ± 0.2^a
	B-F1	1.6 ± 0.1	0.9 ± 0.1*	1.8 ± 0.1^a
Glycolysis	GADPH	2.0 ± 0.2	1.2 ± 0.1*	2.1 ± 0.2^a
	LDHA	0.70 ± 0.02	0.34 ± 0.04*	0.60 ± 0.05^{a}
Pentose phosphate pathway	G6PDH	0.48 ± 0.04	0.36 ± 0.05*	$0.64 \pm 0.07^{\mathrm{a}}$
β-oxidation	HADHA	0.34 ± 0.02	0.17 ± 0.01*	0.37 ± 0.03^{a}
Mitochondrial dynamics	OPA1	1.9 ± 0.1	1.0 ± 0.1*	1.8 ± 0.2^a
Mitochondrial structure	HSP60	0.13 ± 0.01	0.05 ± 0.01*	0.14 ± 0.01^{a}

Table 1. Summary of proteomic changes in rd10 **mice with or without Adalimumab.** Note: Values are expressed as the mean \pm SEM of fluorescence intensity (arbitrary units) per ng of total protein of six retinas per group. Values that are significantly different from C57Bl6 (C) are indicated by asterisks *p < 0.05. 'a' represent statistical differences (p < 0.05) between rd10 and rd10 ADA (One way ANOVA, Newman-Keuls post-test). ADA: Adalimumab.

Adalimumab improved antioxidant response. It has been reported that inflammatory processes and oxidative stress are closely linked in several pathologies. The contribution of oxidative damage to cell death in rd10 mouse retinas has been previously described^{29–31}. In particular, anti-TNF α treatments have been demonstrated to have a positive effect on improving antioxidant response and reducing the

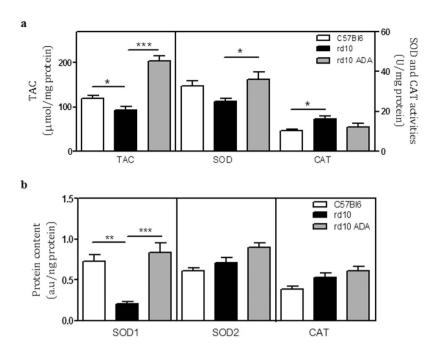


Figure 5. Adalimumab improved antioxidant response at P18. Total antioxidant capacity (TAC), superoxide dismutase (SOD) and catalase (CAT) activities (a) and protein content of SOD1, SOD2 and CAT (b) were measured. Quantitation is shown in bar graphs. Values are the mean \pm SEM of, at least, six retinas per group. Values that are significantly different are indicated by asterisks *p < 0.05, **p < 0.01, ***p < 0.001 (Mixed linear model analysis).

production of reactive oxygen species (ROS) in several pathological situations³². Anti-TNF α antibodies decrease cytokine concentrations, thus directly influencing ROS/RNS production by inflammatory cells or indirectly modulate ROS-stimulated signalling pathways activity. Therefore we evaluated whether Adalimumab improved the antioxidant response in retinas from rd10 mice. For this purpose, we measured the total antioxidant capacity (TAC), enzymatic activity of superoxide dismutase (SOD) and catalase (CAT) which play a major role in the first line of antioxidant defence.

The TAC was reduced (p = 0.031) in vehicle-treated rd10 retinas (92.6 \pm 7.8 nmol of Trolox/mg protein) compared to control retinas (119 \pm 5.7 nmol of Trolox/mg protein). Adalimumab significantly increased TAC to 204.5 \pm 11.6 nmol of Trolox/mg protein (p < 0.001) (Fig. 5A and Supplementary Table S1).

Total SOD activity seemed to decrease (p = 0.12) in vehicle-treated rd10 retinas (25 \pm 1 U/mg protein) compared to control retinas (33 \pm 2 U/mg protein). However, Adalimumab-treated rd10 retinas showed higher values of total SOD activity (p = 0.042), being similar to control retinas (36 \pm 4 U/mg protein) (Fig. 5A and Supplementary Table S1).

Analysis of the protein content of the cytosolic (SOD1) and mitochondrial (SOD2) antioxidant enzymes superoxide dismutase and CAT by RPPM suggested that the diminished TAC and SOD activities could be due to a very large reduction of the expression of SOD1 in vehicle-treated rd10 retinas $(0.20\pm0.03\,\text{a.u/ng}$ protein, p=0.004) (Fig. 5B). The content of SOD2 in vehicle-treated rd10 retinas $(0.71\pm0.06\,\text{a.u/ng}$ protein) was similar to control retinas $(0.61\pm0.04\,\text{a.u/ng}$ protein). Adalimumab increased the content of SOD1 $(0.84\pm0.11\,\text{a.u/ng}$ protein, p=0.001) and, to a lesser extent, of SOD2 $(0.90\pm0.05\,\text{a.u/ng}$ protein, p=0.087) (Fig. 5B and Supplementary Table S1).

CAT activity was higher (p = 0.011) in vehicle-treated rd10 retinas (16 ± 2 U/mg protein) than in control retinas (10 ± 1 U/mg protein). CAT content had a tendency to increase (0.53 ± 0.05 a.u/ng protein, p = 0.15) compared to control retinas (0.39 ± 0.03 a.u/ng protein). Adalimumab had a tendency to reduce CAT activity (p = 0.078) but it did not have effect on CAT content (Fig. 5A,B, and Supplementary Table S1).

Discussion

TNF α inhibitors have been widely used for the treatment of different autoimmune disease, including ocular inflammatory disorders, with promising results^{33–35}. Nevertheless, to our knowledge, we present the first study that evaluates the effect of TNF α inhibitors on the progression of the retinal degeneration in a murine model of RP. The results obtained in the present study revealed that intraperitoneal administration of Adalimumab, a recombinant human monoclonal antibody against TNF α , reduced retinal degeneration decreasing photoreceptor cell death and reactive gliosis at early stages of RP in rd10 mice.

We used the rd10 mouse model of human autosomal recessive RP. This mutant mouse mimics the disease progression seen in some human conditions and offers an amenable therapeutic window. In general, rod photoreceptors start to degenerate between P16 and P20, with a maximum cell death around P21 and P25. By P60, rods are undetectable and only cones remain^{2,36}. Under our housing conditions, we observed that photoreceptor degeneration started around P15, with maximum TUNEL-positive cells around P18. This time point of maximum cell death agreed with results recently described by Arango-Gonzalez *et al.* $(2014)^{37}$. In our study light intensity was higher (98-131 lux) than light intensity described by other authors (around $60 \text{ lux})^{38,39}$. Therefore, the early onset of photoreceptor degeneration could be explained by light conditions.

In the present study, we demonstrated that intraperitoneal injections of Adalimumab displayed a neuroprotective effect on photoreceptors survival at the onset of their death in *rd10* mice. The administration of 3 mg/kg Adalimumab from P9 to P17 contributed to preserve retinal structure reducing the loss of photoreceptors at P18. Further, Adalimumab strongly reduced the number of TUNEL-positive cells in the ONL. Based on several studies using different animal models of RP, it has been suggested that apoptosis is playing a minor role in retinal degeneration³⁷. However, non-apoptotic mechanisms seem to prevail as major contributors to cell death including accumulation of cGMP and activation of histone deacetylase (HDAC), PARP (PAR accumulation) and calpains^{40,41}.

On the other hand, TNF α is considered both apoptotic and necroptotic inducer mainly through its binding to the receptor TNFR1. TNF α binds to TNFR1 leading to the recruitment of multiple proteins that can trigger different downstream pathways as activation of the transcription factor NF-kB, MAPKs (mitogen-activated protein kinases), caspase 8 promoting cell survival, apoptosis or necroptosis, etc⁴². The two most extensively studied ways of necroptosis are initiated by TNFR1 and by PARP pathway. PARP pathway is initiated by overactivation of the DNA repair enzyme PARP-1, resulting in the massive synthesis of PAR and, eventually in necroptosis⁴³. An evident controversial exists concerning to the link between TNF α and PARP pathway during necroptosis. Previously, PARP pathway has been suggested as a key element of TNF α -mediated necroptosis⁴⁴. But other authors propose that TNF α -induced necroptosis and PARP pathway represent distinct and independent routes to necroptosis 45,46. In our previous study we found that TNF α inhibition did not prevent PAR accumulation but also increased it supporting the idea that PARP pathway was independent of $TNF\alpha$ -associated pathways in cultures of porcine retina exposed to Zaprinast²⁵. In the current study we confirmed that there was an excessive nuclear PAR content in vehicle-treated rd10 mouse retinas that was effectively prevented by Adalimumab treatment. Therefore our results suggest that PARP activation is involved in TNFα-mediated photoreceptor cell death in rd10 mice. As described above, PARP activation could be consequence of excessive cGMP accumulation that in turns would activate HDAC. PARP activation could be also consequent of the inflammatory process. We should confirm whether reduced inflammation could be the primary responsible for down regulation of PARP activity or could be secondary to reduced cell death. Thus, the mechanistic interpretation of these results is complicated and deserves further studies.

Upregulation of TNF α and activation of microglial cells are early events involved in photoreceptor death in murine models of RP^{11,47}. We corroborated microglial activation and migration to outer retina in rd10 mice. However, we did not visualize Iba1-positive cells in the ONL in retinas from rd10 mice treated with Adalimumab. Thus, inhibition of TNF α reduced excessive microglial activation and migration of microglial cells to the outer layers. Microglial activation is a hallmark of neuroinflammation. It is associated with various neurodegenerative diseases including retinal degenerations^{10,48}. Some therapeutic approaches are designed to inhibit microglial activation. For instance, minocycline attenuated photoreceptor cell death in different animal models of retinal degeneration^{49,50} and improved vision in patients with diabetic macular edema⁵¹. Around 80 clinical trials are in progress to investigate the effect of minocycline in several pathologies including retinal degenerations (ClinicalTrials.gov, Identifiers: NCT02140164 and NCT01468844).

Recent studies showed that Adalimumab decreased glial cell activation (overexpression of GFAP) and preserved retinal organization in organotypic cultures of porcine neuroretina exposed to $TNF\alpha^{52}$. Activated Müller cells can release antioxidants, growth factors and cytokines, including $TNF\alpha$, contributing to retinal regeneration or degeneration. Müller cells are activated in models of RP resulting in overexpression of GFAP, activation of ERK (extracellular signal-regulated kinase), translocation of Müller cell bodies to the outer retina and thickening of their processes^{25,53}. In the present study, Adalimumab tended to ameliorate GFAP overexpression in rd10 retinas.

Adalimumab specifically blocks the interaction of soluble TNF α , and to a lesser extent, transmembrane TNF α with its receptors, inhibiting downstream cellular signalling mechanisms²¹. We found that blocking TNF α tends to normalize the increased TNF α production, as described in other pathological situations⁵⁴. TNF α blockade prevented the activation of signalling pathways involving the activation of p38 MAPK or NF- $\kappa\beta^{55,56}$. MAPKs such as ERK, JNK (c-Jun N-terminal kinase) and p38, are involved in many inflammatory and degenerative processes including retinal cell death⁵³. Thus, we speculate that Adalimumab could attenuate the inflammatory process by interfering downstream signalling of TNF α , at the level of p38 MAPK or ERK activation. Further studies are needed to confirm this hypothesis.

Inflammation and oxidative stress are closely linked processes in several pathological situations. It has been widely suggested that oxidative stress contributes to the pathogenesis of RP in animal models and patients^{29–31}. For instance, we previously found that RP patients present a reduced antioxidant

response⁹. In the current study we assessed whether down regulation of inflammation (TNF α and reactive gliosis) induced by Adalimumab also influenced antioxidant response. Adalimumab improved antioxidant defence machinery in rd10 mouse retinas. This treatment improved the total antioxidant capacity and total SOD activity probably by increasing SOD1 content in the retina of the rd10 mouse. It has been shown that SOD1 deficient mice are more sensitive to the damaging effects derived of oxidative insults^{57,58}. Thus, SOD1 seems to be a key defender against oxidative damage in retina. It has been previously shown that Adalimumab can influence the antioxidant and oxidant status in other pathological situations such as psoriasis, rheumatoid arthritis, ankylosing spondylitis or uveitis^{32,59,60}. For instance, Adalimumab is capable of upregulating SOD activity and downregulating catalase activity in mesenchymal stem cells from skin of patients with psoriasis⁵⁹ and to restore glutathione (GSH) content and glutathione peroxidase activity in eves of a uveitis experimental model⁶⁰.

The retina is an organ with one of the highest energy demands per tissue weight due to the photoreceptor cell metabolism. Glucose is metabolized through two major metabolic pathways: (1) glycolysis to obtain energy (ATP), reducing power (NADH) and pyruvate and (2) the pentose phosphate pathway (PPP) to obtain reducing power (NADPH) and ribose for nucleic acid biosynthesis. Under aerobic conditions, pyruvate supplies ATP, precursors of certain amino acids as well as NADH through the citric acid cycle or Krebs cycle (TCA cycle). The NADH generated by the TCA cycle is fed into the mitochondrial oxidative phosphorylation (OXPHOS) pathway. During OXPHOS, electrons are transferred from electron donors to electron acceptors such as oxygen in redox reactions releasing energy and finally ATP. Photoreceptors contain a dense concentration of mitochondria in their inner segments that provides the ATP for the phototransduction processes mainly through the OXPHOS pathway^{61,62}. Part of the oxygen consumed in OXPHOS is transformed to reactive oxygen species which are converted to H₂O by antioxidant enzymes. However, a deficient antioxidant defence results in oxidative damage that initiates a cascade of events resulting in mitochondrial dysfunction and cell death⁶³. In this study we corroborated the downregulation of OXPHOS pathway observed at the onset of photoreceptor degeneration in a Drosophila model expressing the equivalent of the most common RP-linked mutation, RhoP23H²⁷. We also observed significant alterations in other energetic pathways as glycolysis, pentose phosphate pathway and β -oxidation indicating global energy depletion in rd10 retinas at P18. We also detected mitochondrial alterations in proteins related to its dynamics and structure supporting the results reported by other authors^{63,64}. Adalimumab ameliorated energetic and metabolic dysfunction in rd10 mice.

It is tempting to speculate that Adalimumab or other anti-TNF α agents could be a promising therapy for RP and other retinal degenerations. However, further studies are needed to investigate in detail the molecular mechanisms involved in the neuroprotective effect of Adalimumab and its long-term effect.

Methods

Animals and treatment. rd10 mice were used as human model of autosomal recessive retinitis pigmentosa. Wild-type C57Bl6 mice with the same genetic background as rd10 mice were used as control. Mice were kept under a 12 hours light/dark cycle, humidity and temperature controlled and with food and water supplied ad libitum. All cages were placed on the lower shelf of an IVC rack with light illuminance of 115 ± 7 lux (95% CI: 98–131). Mice were housed in the Animal Facility of *Unitat Central d'Investigació* (UCIM) of Valencia University. This study was carried out in accordance with guidelines from the European Union Guidelines for the Care (European Union Directive (2010/63/EU) and the Use of Laboratory Animals. All animal procedures and protocols were approved (A1361179906177) and monitored by the Committee of Ethics in Research of the Faculty of Medicine, University of Valencia.

To determine the profile of retinal degeneration at early stages in our housing conditions, untreated *rd10* and C57Bl6 mice were euthanized at P13, P15, P18 and P20. The eyes were rapidly removed and processed as described below.

To evaluate the effect of Adalimumab, each rd10 mouse received one intraperitoneal injection of Adalimumab (Humira, Abbot Laboratories) saline solution at 3 mg/kg every three days starting at P9 and until P17. The dose of 3 mg/kg of Adalimumab was chosen based on previous studies with murine models of rheumatoid arthritis and uveitis 60,65,66 . No apparent side effects (e.g. opportunistic infections) were detected in animals treated with Adalimumab.

Control mice received the same volume of saline (vehicle) at the same time. At P18, treated and untreated *rd10* mice and C57Bl6 mice were sacrificed. The eyes were rapidly removed and processed as described below.

ELISA assay. Retinas were homogenized in 20 mM Tris-HCl pH 7.4, 10 mM EDTA containing protease inhibitor cocktail (Complete Protease Inhibitor Cocktail; Roche, Basel, Switzerland) and $200\,\mu\text{M}$ phenylmethylsufloxifluoride (PMSF). The TNF α protein levels were estimated with a high sensitivity ELISA kit (eBioscience, Ireland, UK), according to the manufacturers' instructions. Tissue TNF α levels were expressed as pg/mg protein.

Tissue processing. To obtain retinal sections, the eyes were rapidly removed and fixed in 4% filtered paraformaldehyde for two hours at room temperature and cryoprotected in a sucrose gradient

(15-20-30%). Eyes were frozen embedded in OCT and $10\,\mu m$ sections were cut in a cryostat (Leica CM1900, Nussloch, Germany).

For biochemical determinations and gene expression analysis, retinas were isolated, placed immediately into the appropriate buffer and stored at -80 °C.

Retinal immunohistochemistry and TUNEL assay. Immunofluorescent staining procedures were performed in 10μm cryosections. Sections were post-fixed in 4% filtered paraformaldehyde (Sigma-Aldrich, Madrid, Spain) in 0.1 M phosphate buffer pH 7.4 for 15 minutes at room temperature. Sections were pre-treated with citrate buffer pH 6.0 for epitope retrieval and incubated in blocking solution containing 5% normal goat serum, 1% bovine serum albumin and 0.25% Triton X-100 for one hour. Afterwards, they were incubated with primary antibody anti-PAR (1:200, Enzo Life Science, Madrid, Spain) as an indirect marker for PARP activity, anti-Iba1 (1:300, Wako Pure Chemical Industries Ltd., Osaka, Japan) and anti-GFAP (1:400, Sigma-Aldrich, Madrid, Spain) overnight at 4°C. Sections were then incubated with the fluorescence-conjugated secondary antibody Alexa Fluor 647, 555 or 488 (Invitrogen, Life Technologies, Madrid, Spain) for one hour at room temperature. After labelling, the sections were mounted in Fluoromount-G (Southern Biotechnology, Birmingham, AL, USA) and observed under a fluorescence or confocal microscope.

To evaluate cell death, the terminal deoxynucleotidyl transferase dUTP nick and labelling (TUNEL) assay was used as previously described⁶⁷.

Microscopy and quantification. The retinal sections were examined under an Eclipse 80i microscope (NIKON Instruments, Badhoevedorp, The Netherlands) or under a confocal microscope (Leica TCS SP5 Confocal microscope, Leica Microsistemas SLU, Barcelona, Spain) belonging to the Microscopy Unit of the IIS-La Fe (Valencia, Spain). ImageJ software was used to quantify the thickness and the number of rows of photoreceptor nuclei in the ONL, the number of TUNEL, PAR and Iba1-positive cells and the corrected fluorescence of GFAP.

We measured the ONL thickness of the entire retina normalized to the thickness of the inner nuclear layer (INL) to avoid the bias derived of the angle of the sectioning plane. Because the ONL thickness and the degenerative process in the *rd10* model are not uniform and dependent upon location, we performed several measurements across the entire retina (from the nasal to the temporal retina) for each mouse. The *normalized ONL thickness ratio* is defined as the ONL thickness/INL thickness. At least five entire retinas were analysed per experimental group.

The number of TUNEL or PAR-positive cells was represented as the ratio between the number of TUNEL or PAR-positive cells in the ONL and the normalized ONL thickness ratio of each section.

To evaluate microglial activation we measured the migration index (M.I) which is defined as the number of Iba1-positive cells weighted according to the retinal layer where are located (M.I = Σ (number of Iba1-positive cells in each layer \times layer weighted factor)/total number of Iba1-positive cells in the section. The layer weighted factor was 1 for ONL, 0.5 for outer plexiform layer (OPL) and 0.25 for INL.

Corrected fluorescence of GFAP was quantified as previously described²⁵. TUNEL and PAR-positive cells, microglial migration index and the corrected fluorescence of GFAP were quantified from four non-adjacent sections of at least five retinas for each experimental group.

Isolation of total RNA and cDNA synthesis. Total RNA was isolated from frozen retina using RNeasy mini kit (Qiagen, Hilden, Germany) following the manufacturer's protocol. Then, cDNA was synthesized starting from $1\mu g$ of RNA by reverse transcription using the GeneAmp Gold RNA PCR Reagent kit (Applied Biosystems, Carlsbad, CA, USA) following manufacturer's instructions.

Quantitative real time PCR. The relative expression of $TNF\alpha$ and IL-6 was measured by real-time PCR using a commercial thermal-cycler (Applied Biosystems ViiATM 7 Real-Time PCR System; Life Technologies Corporation, Carlsbad, California, USA), TaqMan® gene expression assay (Mm00443260_g1 (TNF α) and Mm00446190_m1 (IL-6) and TaqMan® 2X PCR Master Mix (Applied Biosystems, Life Technologies Corporation, Carlsbad, California, USA). β 2 microglobulin (β 2m) gene (Mm00437762_m1) was used as housekeeping gene.

Real-time PCR was performed with 1 cycle of 2 min at 50 °C, followed by 1 cycle of denaturation of 10 min at 95 °C, continued by 40 cycles of 15 seconds denaturation at 95 °C and 60 seconds annealing at 60 °C.

Evaluation of antioxidant response. Retinas were homogenized in 5 mM phosphate buffer pH 7, 0.9% NaCl, 0.1% glucose, centrifuged at $10,000 \times g$ (total antioxidant capacity (TAC) and catalase (CAT) activity) or $1,500 \times g$ (superoxide dismutase (SOD) activity) for 15 minutes at 4 °C. Supernatants were used to determine TAC, CAT and SOD activities with commercial kits following Manufacturer's Instructions (Cayman Chemical, Ann Arbor, MI, USA). Protein concentrations were measured by the bicinchoninic acid (BCA) protein assay. For total SOD activity assay one unit of SOD was defined as the amount of enzyme needed to exhibit 50% dismutation of the superoxide radical. For CAT activity assay

one unit was defined as the amount of enzyme that will cause the formation of 1.0 nmol of formaldehyde per minute at 25 °C. SOD and CAT activities were expressed as U/mg protein. Retinal TAC levels were expressed as nmol/mg protein.

Retinal content of SOD1 (cytosolic), SOD2 (mitochondrial) and CAT enzymes were measured as described below²⁸.

Reverse Phase Protein Microarrays (RPPM). Thirty-six retinas were prepared for printing onto RPPM. Frozen retinas were grinded with liquid nitrogen and the tissue powder was extracted with 90 µl of 50 mM Tris/HCl pH 8, containing 100 mM NaCl, 1 mM DTT, 1% (v/v) Triton X100, 0.1% SDS, 0.4 mM EDTA and a cocktail of protease (Roche) and phosphatase (Sigma) inhibitors. After protein extraction, samples were centrifuged at 15,000 g for 30 min at 4 °C. The protein concentration in the supernatants was determined with the Bradford reagent (Bio-Rad Protein Assay) using BSA as standard. For printing and processing, samples were diluted in PBS to a final protein concentration of 0.75 µg/µl. Serially diluted protein extracts (0-1 µg/µl) derived from HCT-116 and OVCAR8 carcinoma cells were also prepared to assess printing quality and the linear response of protein recognition by the antibodies used²⁸. A solution of BSA (1µg/µl) was also prepared for printing as internal negative control. Approximately, 1 nl volume of each sample was spotted in duplicate onto nitrocellulose-coated glass slides (FAST Slides, Scheleicher & Schuell BioScience, Inc.) using a BioOdyssey Calligrapher MiniArrayer printer (Bio-Rad Laboratories, Inc.) equipped with a solid pin (MCP310S) at constant humidity and temperature²⁸. After printing, arrays were allowed to dry and further blocked in PBS-T containing 5% skimmed milk. After, the arrays were incubated overnight at 4°C with the indicated concentrations of the following highly specific primary monoclonal antibodies (mAbs): β-F1-ATPase (0,4μg/ml) and anti-Hsp60 (4μg/ml)²⁸; anti-LDHA (4μg/ml)⁶⁸; anti-GAPDH (1/20000; Abcam), anti-complex II subunit B (SDHB, 1/500; Invitrogen), anti-complex III subunit Core 2 (Core2, 1/1000; Mitosciences), anti-catalase (1/5000; Sigma-Aldrich), anti-complex IV subunit II (COXII, 1/100; Abcam), anti-complex I Ndufs3 (1/200; Abcam), anti-β-actin (1/5000;Sigma), anti-OPA1 (1/500; BD Transduction Lab) and the indicated concentrations of the following polyclonal antibodies: anti-HADHA (1/200; Abcam), anti-SOD1 (1/200; Santa-Cruz), anti-SOD2 (1/200; Abcam) and anti-G6PDH (1/500; Thermo Scientific). After incubation, the arrays were processed and revealed as previously described²⁸. The mean fluorescent intensity of the spots was quantified using FIJI software (N.I.H., USA) and converted into arbitrary units of expressed protein/ng of total protein in the tissue extract using the expression obtained in the linear plot of the HCT116 cell line as standard (see Supplementary Figure S1).

Statistical analyses. Statistical analysis was performed using R software (version 3.1.2). Comparisons between different age categories were performed using Mann-Whitney U. A generalized least squares model was fitted to the log-transformed TNF α protein values to assess differences between the two experimental groups and also the effect of Adalimumab (ADA). Linear models were used to assess the association between ADA and the two experimental groups with the different response variables. To account for the non-independence of observations in the case of repeated measures per sample, a random intercept was added to the linear models with the variable sample as random factor. P values for the mixed linear models were computed using Satterthwaite approximation to degrees of freedom. Estimations for each variable in a linear mixed model are interpreted similarly than estimations from an ordinary linear model, that is, in an additive way. Negative values indicated a negative effect of being rd10 or having received Adalimumab. And positive values indicated a positive effect. Some response variables were logarithmically transformed to avoid negative values for the estimated control values. Result tables from the models are shown in Supplementary Table S1. Reverse Phase Protein Microarrays were analysed by One-way ANOVA and Newman-Keuls post-test. A p-value < 0.05 was considered statistically significant.

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Author Contributions

C.M.F.C. carried out biochemical determinations, performed histological analysis and contributed to write the draft; A.H.P. provided advice and training on rd10 management, carried out some histological analysis and helped to write the draft; L.O.G. performed analysis of antioxidant response and helped to histological analysis; C.C.M. and M.S.A. performed reverse phase protein microarrays; D.H. carried out statistical analysis and helped to revise the draft; D.S. helped to obtain Adalimumab and to design the study; J.M.C. and J.M.M. participated in the design of the study and helped to revise the manuscript. E.J.d.l.R. provided advice and training on rd10 management and, helped to revise the manuscript; R.R. conceived the study, designed and supervised the study, analysed data and redacted the draft. All authors read and approved the final manuscript.

Additional Information

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