

Modulation of the Proteasome Pathway by Nano-Curcumin and Curcumin in Retinal Pigment Epithelial Cells

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Keywords

Curcumin · Nano-curcumin · Proteasome · Retinal pigment epithelium

Abstract

Introduction: Curcumin has multiple biological effects including the modulation of protein homeostasis by the ubiquitin-proteasome system. The purpose of this study was to assess the in vitro cytotoxic and oxidative effects of nano-curcumin and standard curcumin and characterize their effects on proteasome regulation in retinal pigment epithelial (RPE) cells. **Methods:** Viability, cell cycle progression, and reactive oxygen species (ROS) production were determined after treatment with nano-curcumin or curcumin. Subsequently, the effects of nano-curcumin and curcumin on proteasome activity and the gene and protein expression of proteasome subunits PA28 α , α_7 , β_5 , and β_{5i} were assessed. **Results:** Nano-curcumin (5–100 μ M) did not show significant cytotoxicity or anti-oxidative effects against H₂O₂-induced oxidative stress, whereas curcumin (≥ 10 μ M) was cytotoxic and a potent inducer of ROS production. Both nano-curcumin

and curcumin induced changes in proteasome-mediated proteolytic activity characterized by increased activity of the proteasome subunits β_2 and β_{5i}/β_1 and reduced activity of β_5/β_{1i} . Likewise, nano-curcumin and curcumin affected mRNA and protein levels of household and immunoproteasome subunits. **Conclusions:** Nano-curcumin is less toxic to RPE cells and less prone to induce ROS production than curcumin. Both nano-curcumin and curcumin increase proteasome-mediated proteolytic activity. These results suggest that nano-curcumin may be regarded as a proteasome-modulating agent of limited cytotoxicity for RPE cells.

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Introduction

Curcumin (1, 7-bis-[4-hydroxy-3-methoxyphenyl]-1, 6-heptadiene-3,5-dione) is the orange and water-insoluble pigment extract of turmeric, the rhizome of *Curcuma longa*. The therapeutic potential of curcumin is currently being tested in several clinical trials after promising preliminary observations [1]. In ophthalmology, curcumin

has been proposed as a potential therapeutic agent for several conditions including dry eye syndrome [2], diabetic retinopathy [3, 4], diabetic retinal neurodegeneration [5], age-related macular degeneration (AMD) [6], retinitis pigmentosa [7, 8], and light- and oxidative stress-induced retinal neurodegeneration [9]. On the other hand, studies have warned of the potential toxic effects of curcumin in retinal pigment epithelial (RPE) cells [10–12] and retinal endothelial cells [13].

One of the major obstacles precluding the implementation of curcumin as a therapeutic agent lies in its poor bioavailability in organs and tissues other than the gastrointestinal tract when given orally [14]. At present, various curcumin formulations with improved bioavailability, including nano-particles as drug delivery systems, are being assessed [15]. However, these different formulations do not have comparable biological properties [16]. Recently, a highly absorptive curcumin dispersed with polysaccharide nano-particles (Theracurmin® or nano-curcumin) has been developed. The oral absorption efficacy of nano-curcumin is approximately 40-fold greater than that of curcumin in both rats and humans [17], which implies that this specific formulation is taken up by the epithelial cells of the gastrointestinal tract and reaches blood levels that are deemed sufficient for bioactivity [18, 19].

The ubiquitin-proteasome system (UPS), a multi-catalytic cytoplasmic and nuclear protein complex present in all eukaryotic cells, is responsible for non-lysosomal proteolysis [20]. Intracellular proteins are tagged for proteolysis after binding of ubiquitin moieties. These are then recognized by the 19S regulatory particle of the proteasome, which, in combination with the 20S catalytic core, forms the household or “classical” proteasome [21]. Within the proteasome core, 3 different specialized catalytic subunits are responsible for proteolytic cleavage of the carboxyl end of proteins: β_1 for acidic amino acids (caspase-like), β_2 for basic amino acids (trypsin-like), and β_5 for hydrophobic amino acids (chymotrypsin-like). In response to stress and injury, the household proteasome changes its configuration into the so-called immunoproteasome, which is formed upon replacement of the constitutive subunits by the inducible subunits, β_{1i} , β_{2i} , and β_{5i} [22, 23]. Therefore, the ratio between the household (β_1 , β_2 , and β_5) and inducible subunits (β_{1i} , β_{2i} , and β_{5i}) is a marker of cellular stress [24–26].

Curcumin has been shown to modulate proteasome function by different mechanisms. First, the carbonyl carbons of curcumin interact with the hydroxyl group of the amino-terminal threonine residue of the β_5 subunit; this

results in the suppression of the protease activity of the proteasome, and particularly that of the chymotrypsin-like (β_5) subunit [27]. Second, curcumin inhibits COP9 signalosome (CSN) kinase activity [28, 29]. CSN is a protein complex that controls the stability of many proteins such as ligases [30], and it possesses structural similarities to multiple non-ATPase subunits of the 19S lid of the proteasome [31]. The ligases interact with specific ubiquitin-conjugating enzymes in the ubiquitination of substrates, and, as such, CSN functions as an interface between signal transduction and ubiquitin-dependent proteolysis [32]. Third, curcumin has been shown to inhibit ubiquitin isopeptidases, a family of cysteine proteases (deubiquitinases) responsible for the reutilization of ubiquitin by the 26S proteasome [27, 33, 34]. Curcumin contains an α,β -unsaturated ketone and 2 sterically accessible β -carbons that mediate inhibition of these enzymes [31].

A steadily increasing number of clinical trials are investigating the potential therapeutic effects of curcumin and other curcuminoid formulations. However, recent reports of curcumin-mediated retinal cytotoxicity [10–13] could imply that chronic intake of curcumin negatively affects retinal function. The aims of this study are the following: (1) to assess and characterize the cytotoxic and oxidative effects of nano-curcumin and standard curcumin in RPE cells in vitro, and (2) to investigate the in vitro effects of nano-curcumin and standard curcumin on proteasome expression and activity in RPE cells, to ascertain whether they can be used as proteasome-modulating agents in retinal disorders.

Methods

Culture, Maintenance, and Treatment of ARPE-19 Cells

Experiments were conducted using ARPE-19 cells, a human RPE cell line with structural and functional properties that are characteristic of RPE cells in vivo. Cells were cultured at 37°C in 5% CO₂ in gelatin-coated T75 cell culture flasks (Corning, Lowell, MA, USA) in Dulbecco modified Eagle medium (DMEM; Gibco Life Technologies, Carlsbad, CA, USA), low glucose, pyruvate in the presence of 1% penicillin/streptomycin, and 10% fetal calf serum. Cell growth was monitored and the medium was changed twice a week. For the passaging of cells, TrypLE Express (Invitrogen, Carlsbad, CA, USA) was added for trypsinization of the cells and cell suspensions were diluted 3-fold. For experiments, cells were cultured in 6-well plates. Upon confluence, cells were washed once with phosphate-buffered saline (PBS), serum-starved for 24 h, and then treated with different concentrations of nano-curcumin (Theracurmin, kindly provided by Dr. C. Tamura, Theravalue, Tokyo, Japan) and curcumin (kindly provided by Dr. E. Kemper, Academic Medical Center, Amsterdam, The Netherlands). Both the nano-curcumin and standard curcumin were dissolved in sterile water.

Table 1. Primer details

Gene	GenBank	Forward primer	Reverse primer	Size, bp	Temperature, °C
PSME1	NM_006263	CAGCCCCATGTGGGTGATTATC	GCTTCTCGAAGTTCTTCAGGATGAT	139	82
PSMA7	NM_002792	CCTGGAAGGCCAATGCCATAG	TTTGCCACCTGACTGAACCACTTC	149	82
PSMB5	NM_002797	CCATGATCTGTGGCTGGGATAAG	GGTCATAGGAATAGCCCCGATC	144	83
PSMB8	NM_004159	CTGGAGGCGTTGTCAATATGTACC	GCAGCAGGTCACCTGACATCTGTAC	81	76

Gene nomenclature, GenBank accession code, primer sequences, and the predicted size and temperature of the amplified product.

Protein Extraction

Cells were harvested using TrypLE Express (Invitrogen), collected in Eppendorf tubes and centrifuged for 10 min at 400 g. Supernatant was removed and the pellet was suspended in TSDG buffer (10 mM Tris, pH 7.5, 25 mM KCl, 10 mM NaCl, 1.1 mM MgCl₂, 0.1 mM EDTA, and 8% glycerol), 5 mM ATP and 1× protease inhibitor (Roche Applied Science, Penzberg, Germany).

Cells were lysed by 3 cycles of freezing in liquid nitrogen and thawing at room temperature. After centrifugation (15 min; 10,000 g), protein concentrations were determined using a Bradford protein assay (Serva, Heidelberg, Germany).

Cytotoxicity Assays of ARPE-19 Cells Treated with Nano-Curcumin or Curcumin

To assess the viability of untreated ARPE-19 cells and the possible toxic effects of nano-curcumin and curcumin, the PrestoBlue cytotoxicity assay (Invitrogen) was performed according to the manufacturer's instructions. A resazurin-based compound is converted to its reduced form in the intact mitochondria of viable cells; this causes a shift in its color and fluorescence and can thus be quantified fluorometrically or spectrophotometrically. The assay was also used to test whether nano-curcumin or curcumin (concentrations of 20 μM incubated for 3 h) affect the cytotoxicity of 0–1,000 μM hydrogen peroxide (H₂O₂) in ARPE-19 cells.

The assays were carried out in 96-well plates (10,000–25,000 cells per well). After cell adherence and subsequent washing, PrestoBlue reagent was added to each well. The plates were then incubated at 37°C for 20–30 min. After incubation, the solution containing PrestoBlue was transferred from the wells of the assay plates to new wells in a 96-well plate, and absorbance was read on a plate reader (Bio-Rad, Hercules, CA, USA), with the excitation and emission wavelengths set at 570 and 600 nm, respectively.

Cell Cycle Progression Analysis of ARPE-19 Cells Treated with Nano-Curcumin or Curcumin

In order to evaluate the effects of nano-curcumin and curcumin on cell proliferation, the Click-iT EdU Alexa Fluor 488 imaging kit (Invitrogen) was applied according to the protocol provided by the manufacturer. Briefly, ARPE-19 cells at 30–50% confluence were incubated for 24 h with EdU (5-ethynyl-2'-deoxyuridine). EdU, an analog of thymidine, is incorporated into newly synthesized DNA, and is subsequently recognized by azide dyes via a copper-mediated ("click") reaction. As a negative control, untreated cells were assessed and fluorescence per cell was detected using a FACS LSRII cell sorter (Becton Dickinson, Breda, The Netherlands).

Analysis of Reactive Oxygen Species Production of ARPE-19 Cells Treated with Nano-Curcumin or Curcumin

The production of reactive oxygen species (ROS) in ARPE-19 cells in the presence or absence of nano-curcumin or curcumin (5 or 50 μM) was determined by a FACS-based ROS detection kit (Enzo Life Sciences, Plymouth Meeting, PA, USA) using a modified protocol [35]. Untreated and unstained cells were used alongside a positive control with a ROS inducer (200 μM pyocyanine) and a negative control with a ROS inhibitor (5 mM N-acetyl-L-cysteine). ARPE-19 cells incubated with nano-curcumin or curcumin were also treated with 250 μM H₂O₂. After staining for 30 min, fluorescence per cell in the green channel was detected using a FACS LSRII.

Measurement of Activity of Proteasome Subunit Complexes after Treatment of RPE Cells with Nano-Curcumin or Curcumin

We investigated the potential role of nano-curcumin and curcumin as proteasome-modulating agents in RPE cells. Proteasome catalytic subunits β₂, β_{5i}/β₁, and β₅/β_{1i} were labeled in lysates of ARPE-19 cells treated with nano-curcumin or curcumin (5, 50, or 100 μM) with a 0.5 μM activity-based probe BODIPY-epoxomicin for 1 h at 37°C (BodipyFl-Ahx3L3VS, MV121, provided by Prof. Dr. H. Overkleeft, Institute of Chemistry, Leiden, The Netherlands) [36]. Sample buffer (350 mM Tris-HCl pH 6.8, 10% SDS, 30% glycerol, 6% β-mercaptoethanol, 0.02% bromophenol blue) was added to 20 μg of protein lysate. The samples were boiled for 5 min and loaded on a 12.5% SDS-PAGE gel. After running the proteins on the gel, fluorescence imaging was performed using a Trio Typhoon (GE Medical Systems, Little Chalfont, UK) and a 580 bandpass filter to detect the probe directly on the gel. Proteasome total activity values were normalized on the basis of the total proteasome content in cells as indicated by the levels of the α₇ subunit of the 20S proteasome (1:1,000; MCP72; Enzo Life Sciences).

Western Blot Analysis of Isolated RPE Proteasome Subunits after Treatment with Nano-Curcumin or Curcumin

Western blot analysis was performed as described previously [37]. After treatment with nano-curcumin or curcumin (5, 10, 20, or 50 μM at 24 h and 50 μM at 48 h), 20 μg of RPE cell protein was loaded on a 12.5% SDS-PAGE gel, and, after electrophoresis, was transferred to nitrocellulose membranes and semi-quantitatively analyzed. Membranes were incubated overnight or longer at 4°C in the presence of either a monoclonal antibody against the α₇ subunit of the 20S proteasome (1:1,000; BML-PW8110-0025; Enzo Life Sciences) or one of the following polyclonal antibodies:

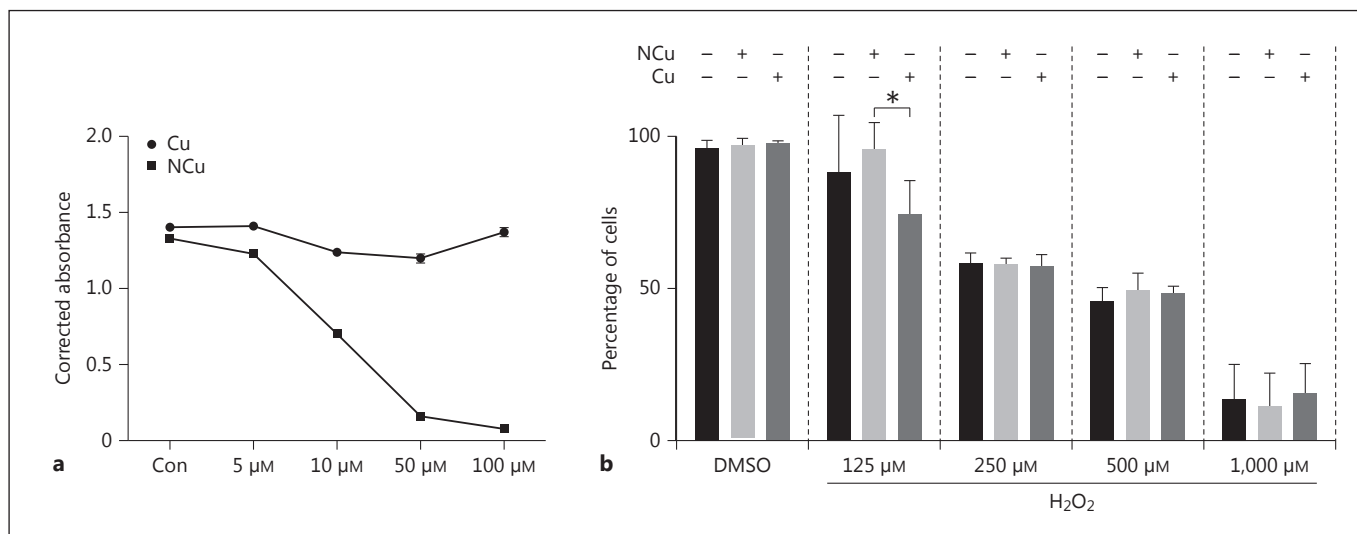


Fig. 1. Effects of nano-curcumin and curcumin on cell viability. Viability assays of ARPE-19 cells exposed to 0–100 μM of nano-curcumin (NCu) or standard curcumin (Cu) in the presence of 0–1,000 μM of H₂O₂. **a** Concentration-dependent effects of nano-curcumin and curcumin on ARPE-19 cell viability expressed as

corrected absorbance. **b** Cytotoxicity effects of 0–1,000 μM of H₂O₂ on ARPE-19 cells, untreated and after pre-treatment with nano-curcumin or curcumin (20 μM for 3 h) and expressed as mean percentage of viable cells ± standard deviation (SD) when compared to untreated cells (0.1% DMSO). * $p < 0.05$ significant change.

the anti-β₅ subunit of the 20S proteasome (1:1,000; BML-PW8895–0100; Enzo Life Sciences), or the anti-β_{5i} subunit of the 20S proteasome (1:1,000; ab3329; Abcam, Cambridge, UK). Anti-β-actin (1:10,000; a5441; Sigma-Aldrich, St. Louis, MO, USA) was used as loading control. The intensity of the bands was quantified by absorbance measurements using Odyssey (LI-COR Biosciences, Lincoln, NE, USA). Quantification was performed with Image Studio Lite v4.0 (LI-COR). Values were normalized using β-actin (1:10,000; ab8227; Abcam).

RNA Isolation and Quantification of RPE mRNA after Treatment with Nano-Curcumin or Curcumin

Real-time quantitative PCR experiments were performed to detect mRNA expression of the proteasome regulatory subunits, PA28α (PSME), α₇ (PSMA7), β₅ (PSMB5), and β_{5i} (PSMB8). Total RNA was isolated from ARPE-19 cell cultures (6 samples/experimental condition) according to the manufacturer's instructions (TRIzol; Invitrogen) after incubation with low (10 μM) and high (50 μM) concentrations of nano-curcumin and curcumin.

Total RNA (1 μg) was treated with DNase I (amplification grade; Life Technologies) and reverse-transcribed into first-strand cDNA using a Maxima® First Strand cDNA synthesis kit (Thermo Scientific, Roskilde, Denmark). Real-time qPCR was performed using a CFX96 system (Bio-Rad) as described previously [38]. Primer details are presented in Table 1. The specificity of the primers was confirmed by NCBI BLAST. The presence of a single PCR product was verified by both the presence of a single melting temperature peak and detection of a single band of the expected size on 3% agarose gel. Non-template controls were included to verify the method and the specificity of the primers. Ct values were converted to arbitrary absolute amounts ($2^{-Ct} \times 1E^{12}$).

Statistical Analysis

Data is presented as fold-change, with a fold-change of 1.0 meaning the same level as control samples. Asterisks indicate a significant change relative to the control samples. Differences between experimental conditions were calculated with one-way or two-way ANOVA, with $p < 0.05$ indicating a statistical difference. Statistical analysis of data was performed using IBM SPSS v20 (SPSS, Chicago, IL, USA).

Results

Nano-Curcumin Is Not Cytotoxic for ARPE-19 Cells, unlike Curcumin, and Does Not Protect against Oxidative Stress-Mediated Cell Death

We found no significant cytotoxicity in ARPE-19 cells exposed to 5–100 μM of nano-curcumin whereas exposure to 10–100 μM curcumin induced significant cytotoxic effects (Fig. 1a). Hardly any viable cells were found after incubation in the presence of 50–100 μM curcumin (Fig. 1a). Next, we assessed whether nano-curcumin or curcumin prevented cytotoxic effects of 125–1,000 μM H₂O₂. The highest non-lethal concentration of curcumin (20 μM) was selected, and compared to a similar concentration of nano-curcumin. As expected, exposure to H₂O₂ caused dose-dependent cytotoxicity in ARPE-19 cells (Fig. 1b). Pre-treatment with 20 μM curcumin increased the cytotoxicity of RPE cells when exposed to 125 μM

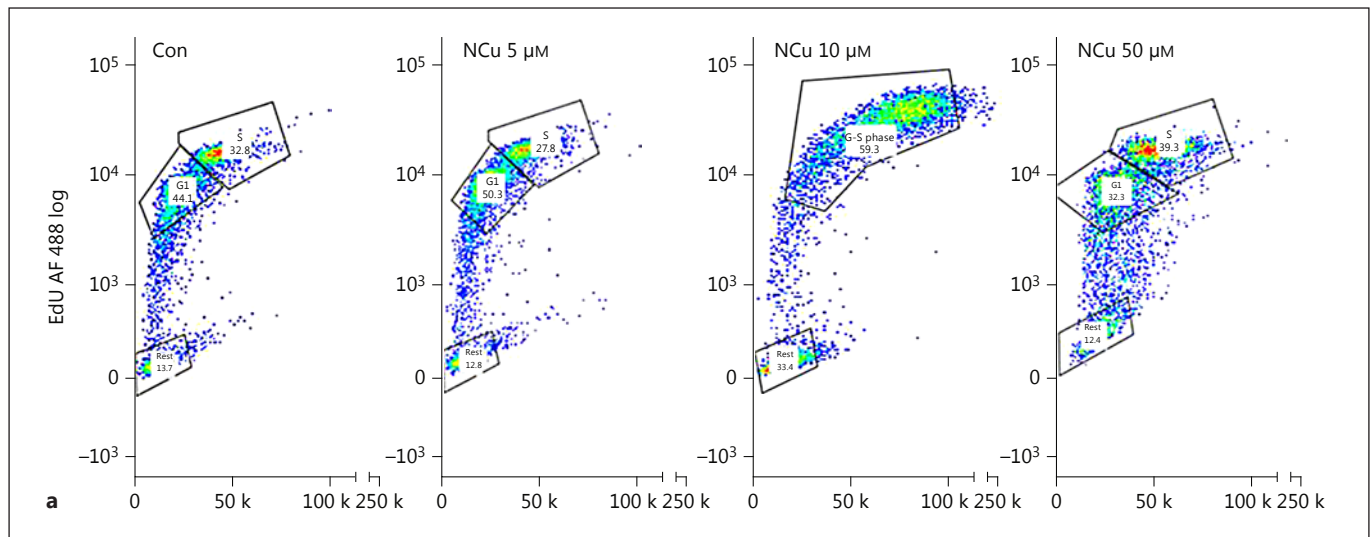
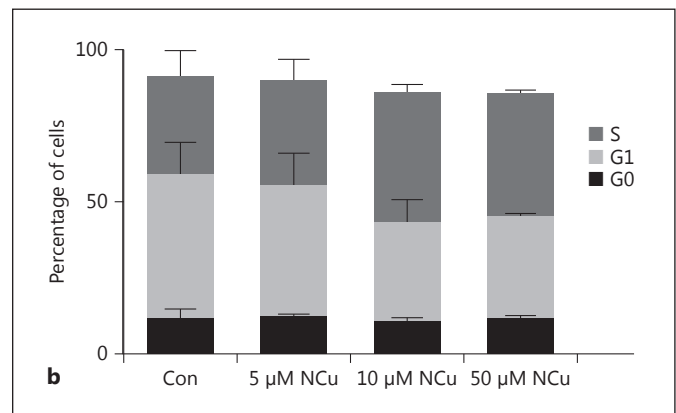


Fig. 2. Effects of nano-curcumin on cell proliferation. Cell proliferation assay of ARPE-19 cells in the presence of 0–50 μM of nano-curcumin (NCu) expressed as percentage of cells in the G0, G1, and S phases after flow cytometric analysis of fluorescence of incorporated EdU. **a** Contour plot of cells treated with 0, 5, 10, and 50 μM of nano-curcumin. **b** Percentages of cells in the different cell phases. Con, control (untreated). No significant differences were found.



H_2O_2 , in contrast to nano-curcumin which showed a mild protective effect ($p = 0.04$). This effect subsided upon higher concentrations of H_2O_2 (Fig. 1b).

Curcumin (15 μM) treatment of RPE cells was previously shown to arrest cell cycle progression with arrest in the G1 phase [12]. Cell cycle progression of ARPE-19 cells exposed to 0–50 μM nano-curcumin did not show significant changes in cell proliferation (Fig. 2).

Nano-Curcumin Is a Mild ROS Inducer and Curcumin Is a Potent ROS Inducer in ARPE-19 Cells

ROS cause damage to cells and the extracellular matrix [39]. Because of the high retinal consumption of oxygen, the retina may be particularly susceptible to oxidative damage. Accordingly, oxidative stress has been linked to several senile degenerative diseases of the retina, including AMD [40–42].

We determined the percentage of ROS-producing cells in the presence of 5 or 50 μM of nano-curcumin or curcumin, and assessed whether the oxidative effects of nano-curcumin differed from those of standard curcumin. We demonstrated strong ROS-inducing effects in cells treated with both low and high concentrations of curcumin, comparable to the pro-oxidative effects of pyocyanin which was used as a positive control (Fig. 3). These effects may explain the dose-dependent cytotoxic effects of curcumin as shown in Figure 1. Low concentrations of nano-curcumin (5 μM) did not induce significant ROS production, with ROS levels that were comparable to those obtained in unstained, untreated cells. High doses of nano-curcumin (50 μM) induced significant ROS production. These results attribute significant pro-oxidative effects to curcumin, whereas low concentrations of nano-curcumin do not lead to significant ROS production in RPE cells.

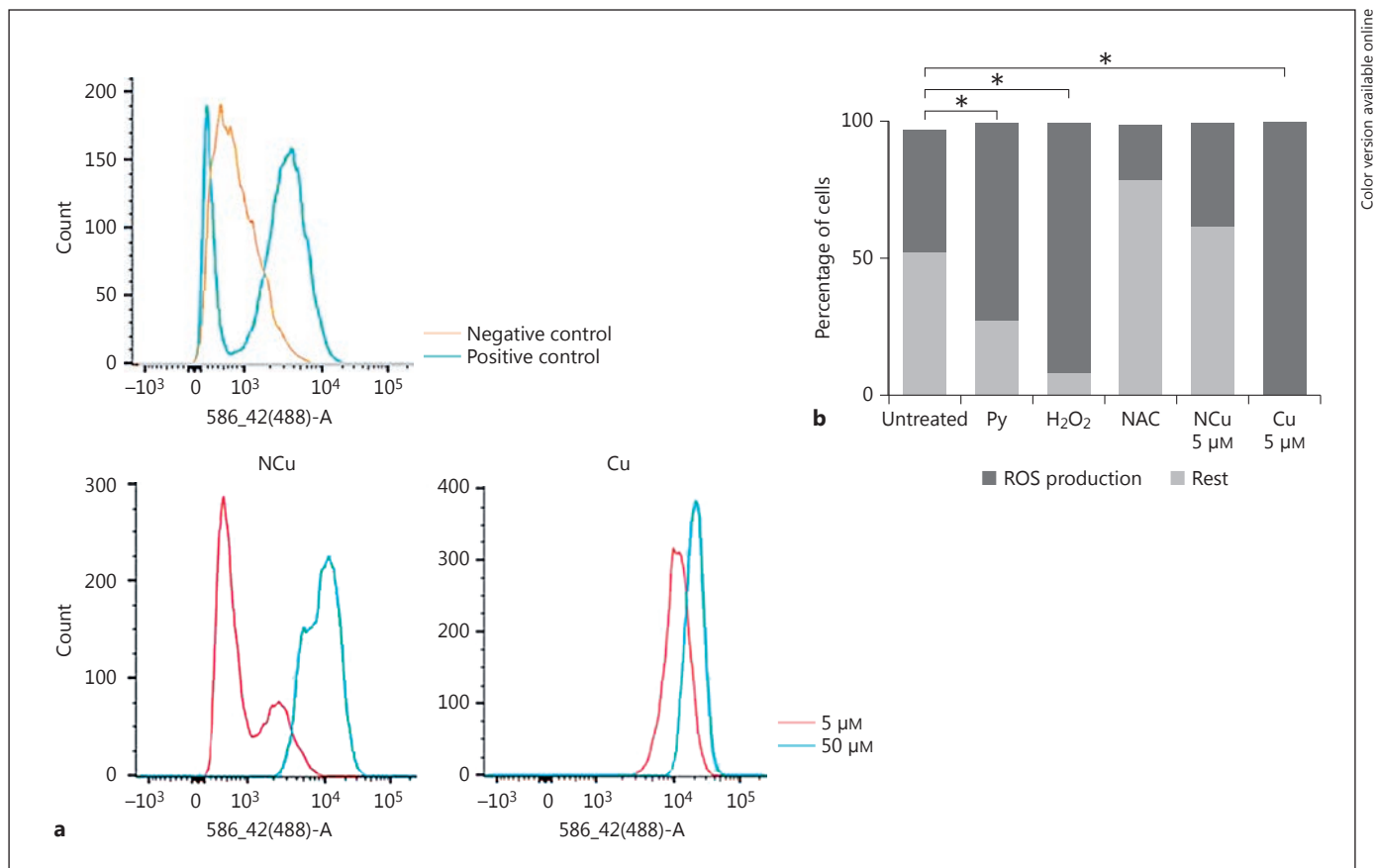


Fig. 3. Effects of nano-curcumin and curcumin on ROS production. Detection of ROS production in ARPE-19 cells treated with low (5 μ M) and high (50 μ M) concentrations of nano-curcumin (NCu) or curcumin (Cu). Cells were also treated with 200 μ M pyocyanin (general ROS inducer, positive control) and 5 mM of N-acetyl-L-cysteine (general ROS inhibitor, negative control). Cells were stained with ROS-ID™ total ROS detection reagent and analyzed using flow cytometry. Untreated cells were used as a control. Cell debris was ungated. The cells with increased levels of oxidative stress display bright green fluorescence in the presence of the ROS detection solution. **a** Histogram plots showing cells incubated in

the presence of pyocyanin (positive control), N-acetyl-L-cysteine (negative control), nano-curcumin, and curcumin. Bright green fluorescence depicting increased ROS production (as observed in the positive control) is demonstrated in cells incubated with 50 μ M of nano-curcumin and 5 and 50 μ M of curcumin. **b** Percentage of cells actively producing ROS compared to untreated cells, incubated with pyocyanin, H₂O₂, N-acetyl-L-cysteine, nano-curcumin, and curcumin. All 3 independent experiments were carried out in triplicate. Py, pyocyanin; NAC, N-acetyl-L-cysteine; ROS reactive oxygen species. * $p < 0.05$ significant change.

Nano-Curcumin and Curcumin Affect the Proteolytic Activity of the Proteasome Catalytic Subunits β_2 , β_{5i}/β_1 , and β_5/β_{1i}

In order to assess whether nano-curcumin and curcumin affect the proteolytic activity of individual proteasome subunits, a proteasome activity-based probe assay was conducted. At present, available proteasome activity probes are unable to distinguish β_1 and β_5 from their inducible counterparts, β_{1i} and β_{5i} , because the subunits run identically on gels (Fig. 4a). Nano-curcumin, at low concentrations (5 μ M), down-regulated the activity of the $\beta_5/$

β_{1i} proteasome catalytic subunits (0.57 fold-change, $p = 0.0009$). Intermediate (10 μ M) and high (50 μ M) concentrations of nano-curcumin up-regulated the activity of the β_2 (1.27 fold-change, $p = 0.03$ and 1.55 fold-change, $p = 0.0005$, respectively) and β_{5i}/β_1 proteasome catalytic subunits (1.46 fold-change, $p = 0.01$ and 1.54 fold-change, $p = 0.04$, respectively). Intermediate (10 μ M) and high (50 μ M) concentrations of curcumin down-regulated the activity of the β_5/β_{1i} proteasome catalytic subunits (0.89 fold-change, $p = 0.009$ and 0.92 fold-change, $p = 0.015$, respectively). Similar to the effects observed with nano-curcumin, high

Fig. 4. Effects of nano-curcumin and curcumin on proteasome activity. Proteasome activity labelling in ARPE-19 cell lysates. **a** After treatment with 0–50 μM of nano-curcumin (NCu) or curcumin (Cu), ARPE-19 cells were harvested and proteasomes were labeled with a proteasome activity probe. Activity of the β_2 , β_1/β_{5i} , and β_5/β_{1i} proteasome catalytic subunits was determined by fluorescence imaging. **b** Data is expressed as mean proteolytic activity per subunit complex \pm SD, normalized on the basis of the total proteasome content in cells as indicated by the levels of proteasome α_7 subunit. * $p < 0.05$ significant change; ** $p < 0.01$; *** $p < 0.001$.

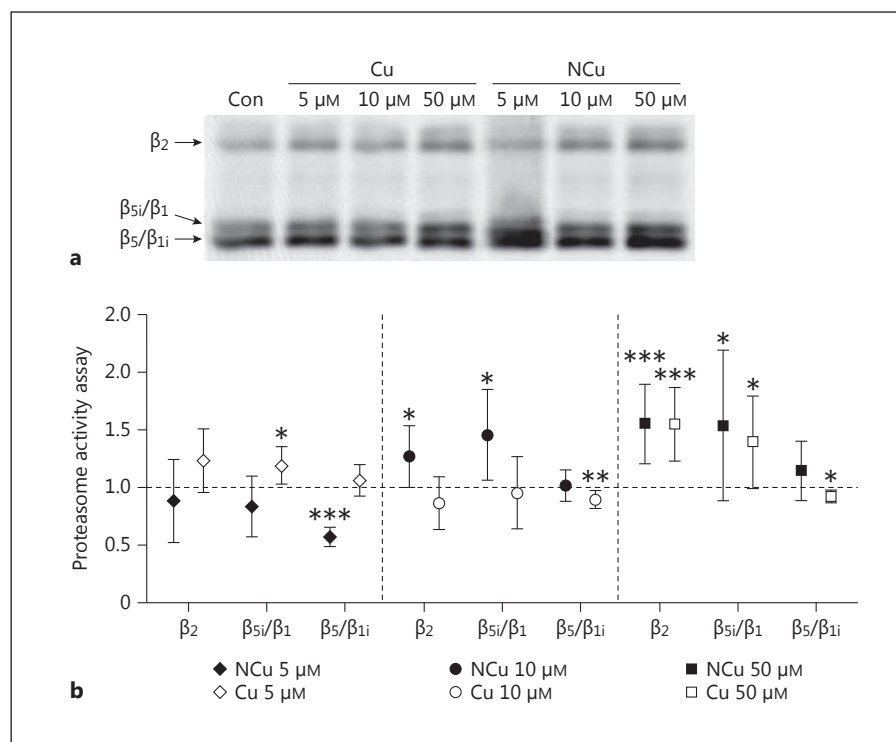
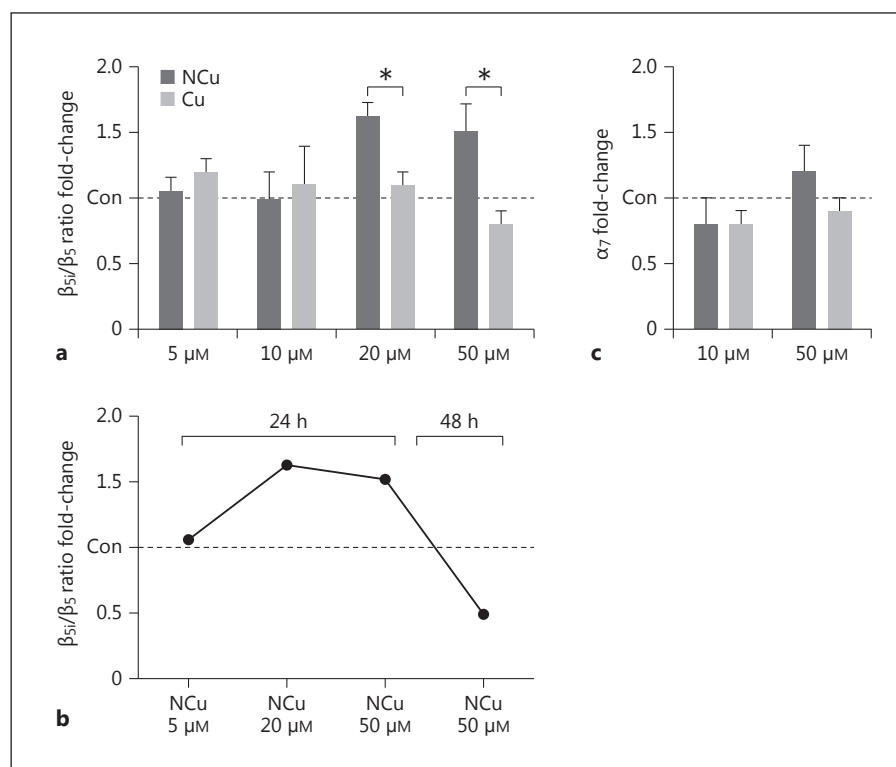


Fig. 5. Effects of nano-curcumin and curcumin on proteasome protein expression. Protein levels of proteasome subunits β_5 , β_{5i} , and α_7 were assessed by Western blot with actin expression as loading control after the incubation of ARPE-19 cells with 0–50 μM of nano-curcumin (NCu) and curcumin (Cu) at 24 h (β_5 , β_{5i} , and α_7) and 48 h (β_5 and β_{5i}). **a** β_{5i}/β_5 ratio fold-change at 24 h induced by 0–50 μM of nano-curcumin and curcumin, corrected for actin and relative to control samples. **b** Time-dependent effects in the β_{5i}/β_5 ratio fold-change induced by nano-curcumin at 24 and 48 h, corrected for actin and relative to control samples. **c** Protein levels of proteasome subunit α_7 induced by 0–50 μM of nano-curcumin and curcumin, corrected for actin and relative to control samples. Data is expressed as the mean \pm SD. * $p < 0.05$ significant change.



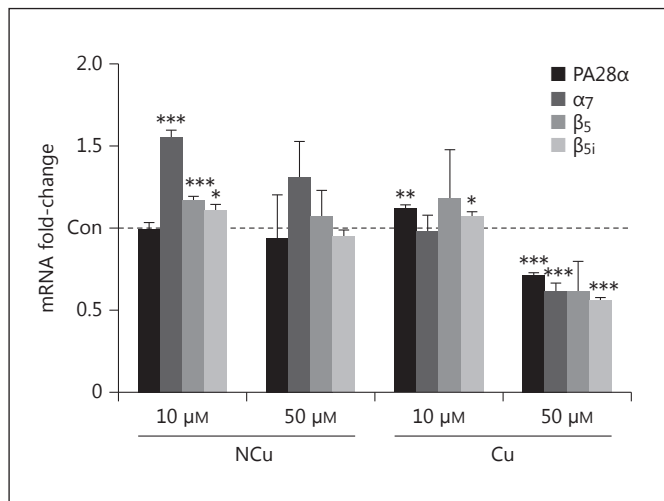


Fig. 6. Expression of proteasome-related genes after exposure to nano-curcumin and curcumin. mRNA levels of proteasome subunits PA28α, α₇, β₅, and β_{5i} after 72 h of treatment with nano-curcumin and curcumin (0, 10, and 50 μM). The mRNA expression levels of α₇ were increased upon stimulation with 10 μM of nano-curcumin, and β₅ and β_{5i} upon stimulation with 10 and 50 μM of nano-curcumin. Treatment with 10 μM of curcumin increased PA28α and β_{5i} mRNA levels. Higher concentrations of curcumin (50 μM) significantly decreased mRNA levels of PA28α, α₇, and β_{5i}. Values represent mRNA expression levels ± SD relative to untreated control cells. * $p < 0.05$ significant change; ** $p < 0.01$; *** $p < 0.001$.

(50 μM) concentrations of curcumin up-regulated the activity of the β₂ and β_{5i}/β₁ proteasome catalytic subunits (1.55 fold-change, $p = 0.002$ and 1.40 fold-change, $p = 0.035$, respectively) (Fig. 4b). These results suggest that nano-curcumin and curcumin have dose-dependent effects on proteasome-mediated proteolytic activity.

Nano-Curcumin Induces a Mild Activation of the Immunoproteasome

Protein levels of the proteasome subunits β₅, β_{5i}, and α₇ were assessed after treatment with various concentrations of nano-curcumin and curcumin. The ratio between β_{5i} and β₅ was calculated at 2 time points as a marker of immunoproteasome activation. Protein levels of the α₇ subunit of the 20S proteasome depict the total pool of proteasomes in cells. Nano-curcumin (50 and 20 μM) increased the protein levels of proteasome subunit β_{5i} (mean 1.6 fold-change, $p = 0.01$, data not shown), resulting in an increased β_{5i}/β₅ ratio when compared to the unstimulated and curcumin-treated RPE cells (Fig. 5a). This effect was both concentration- and time-dependent, as shown in Figure 5b which demonstrates that, at 48 h, the increase

in β_{5i}/β₅ ratio was overturned. Neither nano-curcumin nor curcumin had a significant effect on the protein levels of the proteasome α₇ subunit (Fig. 5c). These results suggest that nano-curcumin may induce a mild activation of the immunoproteasome.

Nano-Curcumin and Curcumin Affect the Transcription of Proteasome-Related Genes

In order to assess whether nano-curcumin affected the transcription of proteasome-related genes, qPCR of ARPE-19 cells treated for 72 h with 10 and 50 μM of nano-curcumin and curcumin was performed. Figure 6 shows that 10 μM nano-curcumin increased the gene expression of the proteasome α₇ subunit (1.56 fold-change, $p = 0.03$), the β₅ subunit (1.18 fold-change, $p = 0.004$), and the β_{5i} subunit (1.11 fold-change, $p = 0.04$). Curcumin (10 μM) increased the gene expression of the proteasome PA28α subunit (1.12 fold-change, $p = 0.01$), and the β_{5i} subunit (1.10 fold-change, $p = 0.04$). Higher concentrations of curcumin (50 μM) strongly reduced the gene expression of all proteasome-related genes ($p < 0.001$). These results show that nano-curcumin and curcumin have concentration-dependent effects on the transcription of proteasome-related genes.

Discussion

The results of this study support the pleiotropic properties of curcumin and its various formulations. A consensus regarding the effects of curcumin in the retina is yet to be reached. In this study, it was demonstrated that nano-curcumin induces changes in proteasome modulation with limited cytotoxic effects at low concentrations, suggesting that this specific formulation is a safe alternative for the retina and RPE. While some studies praise the beneficial effects of curcumin and its potential use in ophthalmic disease, others advocate that retinal function ought to be monitored during the intake of curcumin. Such contradictory findings may be explained by the hormetic properties attributed to curcumin, i.e., it is toxic at high doses but is able to exert adaptive stress responses at low doses [43, 44]. In addition, certain biological effects of curcuminoids may depend on the specific formulation of curcumin used [16, 45]. For instance, the number of methoxy groups dictates the anti-oxidative potency of curcumin [46] whereas the presence of phospholipids in curcumin formulations affects its biological activity [47]. Therefore, the use of different curcuminoid formulations may lead to contradictory results.

Our study demonstrates significant cytotoxic effects of standard curcumin in RPE cells at concentrations $>20\ \mu\text{M}$. These findings are in agreement with the data of previous studies in which curcumin exhibited pro-apoptotic effects in RPE cells at concentrations of approximately $20\ \mu\text{M}$ [10, 11]. These concentrations are in the same concentration range that is supposed to be effective in cancer cells [48–52]. On the other hand, nano-curcumin, a formulation of curcumin dispersed with polysaccharide nano-particles, showed no major cytotoxicity in RPE cells. The different cytotoxicity of these 2 formulations may be explained by the smaller size and increased solubility of nano-curcumin compared to curcumin. Curcumin, hardly soluble in water, forms particles that are $>20\ \mu\text{M}$ in diameter in aqueous solutions whereas the average diameter of nano-curcumin in aqueous solutions is 100 times smaller ($0.2\ \mu\text{M}$) [17]. Furthermore, nano-curcumin is more stable due to the formation of an amorphous state upon hydrogen bonding [53]. This suggests that monitoring of retinal toxicity is not required during concomitant intake of nano-curcumin; this may not hold true for standard curcumin.

The uptake of nano-curcumin in RPE cells was not investigated in this study. However, other studies provide evidence that RPE cells, similar to other cell lines, take up nano-curcumin. The epithelial cells of the gastrointestinal tract of rats and humans *in vivo* absorb nano-curcumin over 40 times more efficiently than standard curcumin [17, 18], and nano-curcumin has been shown to be taken up by oesophageal cancer cells and non-cancer cells *in vitro* [19].

Our findings suggest that curcumin-mediated cytotoxicity may be partly caused by increased ROS production. In contrast to curcumin, low concentrations of nano-curcumin ($5\ \mu\text{M}$) do not induce ROS production in non-oxidative conditions and have significantly fewer pro-oxidative effects when compared to curcumin. Higher concentrations of nano-curcumin, however, significantly induce ROS production. These results are in agreement with data from previous studies with other cell lines, in which curcumin was shown to possess rapid ROS-inducing effects [54–56]. In contrast, other *in vitro* ARPE-19 cell studies attribute anti-oxidative properties to curcumin in aging and H_2O_2 -mediated oxidative stress [57, 58]. According to our findings, and taking into account both the conflicting and supportive results reported in other studies, it appears that curcumin exhibits both pro- and anti-oxidative properties. Differences in experimental set-up, with regard to concentrations and stimulation times, may partly explain the contradictory findings.

Recent evidence shows that dysfunction of the UPS may contribute to the pathogenesis of several eye diseases [24, 26, 59–65]. Modulation of this pathway remains, however, an unexplored therapeutic target in retinal degenerative diseases. Curcumin has been reported to affect proteasome activity and expression in a biphasic, dose-dependent manner [31]. Low concentrations of curcumin were found to up-regulate proteasome activity but high concentrations inhibited proteasome activity [31]. Chymotrypsin-like activity was increased by 46% after treatment of human keratinocytes with curcumin (up to $1\ \mu\text{M}$ for 24 h) whereas higher concentrations of curcumin were inhibitory. At $10\ \mu\text{M}$, proteasome activity decreased to 46% of its initial value [66]. A possible explanation for this biphasic mechanism could be the fact that curcumin is both pro-oxidative and anti-oxidative. Mild or transient oxidative stress up-regulates proteasome activity and transiently enhances intracellular proteolysis whereas severe or continuous oxidative stress impairs proteasome function and decreases intracellular proteolysis [67].

Our study demonstrates that both nano-curcumin and curcumin exert concentration-dependent changes in the activity of proteasome individual subunits in RPE *in vitro*. A recently developed activity-based probe assay was used instead of the more commonly used ubiquitin-independent fluorogenic peptides and ubiquitin-dependent fluorescent reporters. Activity-based probes can detect alterations in proteasomal activity in gels, and can also be applied to visualize active proteasomes in living cells [68–70]. Unlike fluorescence-labelled tags, activity-based probes label only completely assembled and active proteasome complexes, which may explain why the results obtained by these 2 different methods are not comparable and at times inconsistent. Overall, the activity of the β_2 and β_{5i}/β_1 proteasome catalytic subunits is up-regulated with increasing concentrations of nano-curcumin and curcumin, whilst the activity of the β_5/β_{1i} proteasome catalytic subunits is down-regulated after the treatment of RPE cells with both nano-curcumin and curcumin. Short-term treatment of cells with high concentrations of both nano-curcumin and curcumin induced the expression of β_{5i} subunits, which translated in a higher number of β_{5i} subunits than β_5 subunits, i.e., the household proteasome underwent a change in configuration upon activation of the immunoproteasome. This effect cannot be explained by an increase in the number of proteasomes as the protein expression levels of the α_7 subunit remained unchanged after nano-curcumin and curcumin treatment. The pool of proteasome subunits was replenished after

treatment with low concentrations of nano-curcumin, as demonstrated by the increased mRNA expression of the α_7 , β_5 , and β_{5i} proteasome subunits. Of note, the effects on mRNA expression observed after treatment with 50 μ M curcumin, namely the down-regulation of the transcription of all proteasome-related genes, possibly reflects the potent cytotoxic effects at similar concentrations.

The pro-oxidative state induced by high concentrations of nano-curcumin and curcumin may explain the changes found in proteasome activity, and gene and protein expression. Indeed, oxidative stress has been shown to affect proteasome function [71, 72]. Notably, in the RPE, the activity of β_5 is the rate-limiting step of proteasome activity [20, 27, 65]. Changes due to nano-curcumin treatment approximate those in other conditions that have been shown to be associated with β_5 proteasome activity inhibition, namely, aging [63] and complement overactivation [73]. Changes in the ratio between the immunoproteasome and the classic proteasome are indicative of cellular stress, inflammation, and oxidative stress [24–26, 60, 73]. It has been suggested that a continuous expression of the immunoproteasome in the retina is protective against neuronal stress and promotes repair mechanisms [24]. A possible mechanism for the inactivation of the β_5 subunit lies in the chemical structure of curcumin, via inhibition of the ubiquitin isopeptidase activity located at the 19S regulatory subunit of the 26S proteasome. Curcumin belongs to a class of compounds with α,β -unsaturated ketones and 2 sterically accessible β -carbons [33]. Furthermore, curcumin has been shown to bind directly to the 20S proteasome [27]. A recent study has confirmed that both carbonyl groups of curcumin are indeed susceptible to nucleophilic attack by N-terminal threonine of the β_5 chymotrypsin-like subunit of the proteasome, thereby inhibiting its proteolytic activity [27].

The formulation of curcumin, the duration of culture, and the concentration used may account for the contra-

dictory results described hitherto in several studies. It appears that the cytotoxic profile of each formulation of curcumin depends on its bioavailability, absorption, and cellular uptake. Further in vivo studies are required to assess the molecular effects of supplementation of curcumin in relation to retinal function. It is not known whether systemic administration of curcumin reaches the RPE and the neurosensory retina at concentrations capable of inducing significant molecular effects. If so, our findings suggest that curcumin may promote the development of RPE and retinal dysfunction which, in turn, can accelerate development of age-related retinal diseases. On the other hand, our results show that 2 different formulations of curcumin produce different biological effects. Nano-curcumin may be a safer alternative in clinical trials in the future. Finally, our results attribute significant proteasome-modulating properties to both nano-curcumin and curcumin. The changes in proteasome activity incurred after nano-curcumin treatment are characterized by an activation of the immunoproteasome, with consequent changes in the protein and gene expression of proteasome-related subunits.

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Disclosure Statement

The authors declare that they have no conflict of interest.

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