



## Antioxidant effect of phenolic compounds (PC) at different concentrations in IEC-6 cells: A spectroscopic analysis

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### ABSTRACT

Phenolic compounds (PC) have been proposed as natural antioxidant agents that protect cells against oxidative stress-related diseases. Nonetheless, their low bioavailability forecasts controversy about mechanisms on their *in vivo* scavenging activity against reactive oxygen species (ROS). It has been proposed that PC reduce directly ROS concentration. An alternative or complementary action of PC could be the activation of the cell's antioxidant pathway, involving the regulation of gene expression, like that initiated by the Nrf2 transcription factor. To date there is not enough experimental data to support or discard this possibility. In the present study, we evaluated the use of several PC to prevent peroxidation of macromolecules and to elicit the activation of the Nrf2 transcription factor in H<sub>2</sub>O<sub>2</sub>-stressed IEC-6 enterocytic cell line. Synchrotron microspectroscopy demonstrated that PC compounds protected proteins, lipids and nucleic acids against oxidation induced by H<sub>2</sub>O<sub>2</sub>. Immunofluorescence results showed that treatment with quercetin (Qc), catechin (Cat) and capsaicin (Cap) induced the translocation of Nrf2 into the nucleus, at the same level as did H<sub>2</sub>O<sub>2</sub> treatment, thus mimicking the action of the endogenous cell response to peroxidation. Even though the detailed mechanism still needs to be elucidated, we demonstrated the activation of Nrf2 by PCs in response to oxidative stress.

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## 1. Introduction

Phenolic compounds (PC) are natural antioxidant metabolites found in plant foods, known for their ability to scavenge reactive oxygen species (ROS) and thus protect the cells against oxidative stress-related diseases. However, due to their low bioavailability, there is controversy about how they can act as antioxidant compounds, thus the direct *in vivo* ROS-scavenging activity is in doubt [1,2]. It has been proposed that the antioxidant effect of PC may be related to an indirect interaction of the cells with ROS. In this indirect mechanism, PC seem to act as mild prooxidants that can activate a series of biochemical pathways of the cellular antioxidant system [3,4]. This mild prooxidant effect has been related to the capacity of PC to chelate metal ions and produce ROS, semiquinones and quinones which are potentially cytotoxic. Some prooxidative effects have been observed in insects that consume high levels of phenols, due to the presence of non-absorbed metal ions in their

gastrointestinal tract. Data available on mammals gave different results, which make it difficult to infer what happens in these cases [5]. This mild prooxidant effect of PC can be beneficial because it can trigger the antioxidant defense mechanism, resulting in the protection of cells [5]. In a previous work by our research team, the antioxidant effects of six different PC against H<sub>2</sub>O<sub>2</sub>-generated oxidative stress were evaluated in isolated rat enterocytes, observing that catechin and capsaicin presented high antioxidant effect, while quercetin showed a mild prooxidant effect [6]. These results indicated that at the same concentration, PC may present both antioxidant and prooxidant properties, however the mechanism involved in these differential properties is still unknown. It is proposed that PC may present a direct activity by neutralizing ROS (ROS-scavenging activity) or an indirect activity in which PC exert the activation of antioxidant pathways within the cell like the Nrf2-Keap1 pathway. The effect of different PC concentrations on their antioxidant or prooxidant activities is not well known. For instance, it has been reported that quercetin, eugenol and polyphenols from green tea extract exert antioxidant and prooxidant activities in cancer cells, depending on their concentration [7–9]. To gain

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information on the effect of PC concentration on their antioxidant properties, the aim of the current study was to evaluate the protective effect of different concentrations of catechin, capsaicin and quercetin on H<sub>2</sub>O<sub>2</sub>-induced oxidative stress, using an enterocyte-derived cell line (IEC-6) as model. To evaluate the antioxidant or prooxidant effects of PC and H<sub>2</sub>O<sub>2</sub>, we used Synchrotron radiation-based Fourier transform infrared microspectroscopy (FTIRM), which allowed us to observe the state of lipids, carbohydrates and proteins at single cell level [6,10], the data obtained were compared to standard spectrophotometric techniques such as conjugated dienes (CD) and thiobarbituric acid reactive substances (TBARS). Furthermore, we used immunofluorescence to analyze the activation of the Nrf-Keap-1 pathway in the same experimental conditions. This study reveals how different concentrations of PC affect their protective activity over IEC-6 cells, how this protective activity is exerted and how this is related to the PC structure.

## 2. Materials and methods

### 2.1. Chemicals and reagents

Catechin [Cat], quercetin [Qc], capsaicin [Cap], chloroform, cyclohexane, thiobarbituric acid, trichloroacetic acid, tetramethoxypropane (TMP), sodium bicarbonate, H<sub>2</sub>O<sub>2</sub>, trypan blue, penicillin-streptomycin, hydrochloric acid and IEC-6 cell line (ATCC: CRL-1592) were purchased from Sigma-Aldrich (St. Louis, Missouri, USA). Methanol was purchased from Hycel (Jalisco, Mexico). Primary anti-Nrf2 antibody and secondary antibody Alexa Fluor 594 were purchased from Thermo-Fisher Scientific (Waltham, Massachusetts, USA). Fetal bovine serum (FBS), Dulbecco's modified Eagle's medium (DMEM) and trypsin-like enzyme Tryple Express™ were purchased from Gibco (Gaithersburg, Maryland, USA).

### 2.2. Cell culture

The IEC-6 enterocytic cell line derived from rat (*Rattus norvegicus*) small intestine was used in the present work. After arrival, cells were maintained at -80 °C until used. For the culture, cells were thawed at 37 °C for 1 min in a water bath. Subsequently the cells contained in the 1.5 mL vial were transferred to a T-25 culture flask, containing 3 mL of DMEM (10% de FBS y 1% de PEST) and incubated at 37 °C in a 5% CO<sub>2</sub> atmosphere. DMEM culture medium was changed every other day until cells reached between 85% and 90% of confluence. Before their use in any experiment, cell culture at optimal confluence were trypsinized with 1 mL of Tryple Express™ 1x for 10 min. Afterwards, 1 mL of DMEM medium was added to stop the enzyme activity, and the content of the flasks were transferred into 1.5 mL plastic conic tubes and centrifuged at 820 x g for 5 min. Supernatant was discarded and the pellet containing cells resuspended in 500 µL of DMEM medium. A sample of 50 µL of the cell suspension was collected and mixed (1:1) with 0.4% trypan blue and used for cell counting and viability (Neubauer chamber, examined with an upright microscope at 20x objective, Leica CME Leica Microsystems, Wetzlar, Germany) [11,12].

### 2.3. PC treatment

For all the treatments, 24-well culture plates were used, inoculating 100,000 cells per well, to allow adherence, cells were grown overnight before applying the treatments into the plate. Three PC were used in this study: Cap, Cat and Qc at three different concentrations for the FTIRM, CD and TBARS experiments: 50, 100 and 250 µM, while only a 100 µM concentration was used for immunofluorescence microscopy analysis. Hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>) was used to induce oxidative stress at a final concentration of

500 µM. Three different experiments were used for each polyphenol: (1) only PC was added to the medium, (2) PC added to the medium before H<sub>2</sub>O<sub>2</sub> and (3) PC added to the medium after H<sub>2</sub>O<sub>2</sub>. Cells were exposed to H<sub>2</sub>O<sub>2</sub> and PC for 30 min each. Control treatments and those treated with just one compound were first pre-incubated 30 min with DMEM and then washed with PBS (pH 7.4), centrifuged at 820 x g for 5 min and then treated with DMEM media added with the corresponding compound for 30 min. In the case of cells being treated with PC before or after H<sub>2</sub>O<sub>2</sub> they were incubated with the first compound for 30 min with DMEM added media, washed, centrifuged and incubated with DMEM media added with the second compound for 30 min. After finishing the corresponding treatment, cells were centrifuged at 820 x g for 5 min and the supernatant was discarded. Finally, cells were washed twice with PBS (pH 7.4) and centrifuged at 820 x g for 5 min.

### 2.4. Fourier transform infrared microspectroscopy (FTIRM)

For FTIRM analyzes, cells obtained during the PC treatments were washed with PBS 1x and centrifuged at 820 x g for 5 min. After centrifugation the supernatant was discarded, and cells were fixed in paraformaldehyde 4% in PBS (PFA-PBS) for 20 min. Fixed cells were centrifuged for 10 min at 820 x g, supernatant discarded and the cells washed with deionized water, and finally resuspended in 200 µL of water [10]. Fixed cells were analyzed using the FTIRM end station in the line ID-21 at the European Synchrotron Radiation Facility (ESRF) in Grenoble, France using a Thermo Nicolet Continuum microscope (Thermo Scientific, Madison, WI, U.S.A.) coupled to a Thermo Nicolet Nexus FTIR spectrometer (Thermo Scientific, Madison, WI, U.S.A.). The IR microscope is equipped with a 32X objective, a motorized sample stage and a liquid nitrogen-cooled 50 µm mercury cadmium telluride detector. For observation, 20 µL of fixed cells (approximately 20000 cells) were left to dry in BaF<sub>2</sub> windows of 1 mm height. Spectra was recorded over the range of 4000 to 900 cm<sup>-1</sup>, the spectral resolution was set to 6 cm<sup>-1</sup> and 120 scans per spectrum were collected. Spectra of 30 individual cells were recorded for each treatment.

### 2.5. Infrared spectra analysis

Infrared spectra obtained by FTIRM were analyzed using Unscrambler X software (CAMO Software, Norway). Raw spectra were pre-processed first by a vector normalization and then by a second derivative using Savitsky-Golay of second polynomial order with 21 smoothing points [6,13,14]. Raw and second derivative analysis of all results are presented in supplementary files 1 and 2, respectively. Processed spectra were used for calculation of ratios (proportions) from IR signals at different spectral bands. Effects of oxidative stress in cells has been related to the displacement or reduction of some IR bands. The following bands were evaluated for distinct phenomena: For lipid oxidation: 1740 cm<sup>-1</sup>/2960 cm<sup>-1</sup>, lipid saturation: 2920 cm<sup>-1</sup>/2960 cm<sup>-1</sup> and lipid unsaturation: 3012 cm<sup>-1</sup>/ 2960 cm<sup>-1</sup>, protein aggregation: 1630 cm<sup>-1</sup>/1650 cm<sup>-1</sup> [15–19].

### 2.6. Spectrophotometric measures: determination of lipid peroxidation by conjugated dienes (CD) and determination of thiobarbituric acid reactive substances (TBARS)

Conjugated Dienes (CD) and thiobarbituric acid reactive substances (TBARS), which measure the presence of primary and secondary products of lipid peroxidation respectively, were measured as previously reported, using approximately 300,000 cells from the PC treatments, as described in our previous work [6]. For the CD analysis, PC treated, and control cells were lysed by sonication and then lipids extracted with chloroform. The extracted lipids were

then dissolved in cyclohexane, and immediately measured with a UV-spectrophotometer at 233 nm (Thermo Scientific, Madison, WI, USA) using cyclohexane as blank. The obtained absorbance was converted to conjugated dienes concentration using an extinction coefficient of  $27000 \text{ M}^{-1}\text{cm}^{-1}$  and the results were expressed as conjugated dienes concentration ( $\mu\text{M}$ ) per million cells. For TBARS analysis, PC treated, and control cells were lysed by sonication. Samples and calibration standards were mixed with TBA reagent (20% TCA, 0.5% TBA and 2.5 N HCl) and the mixture heated for 45 min in a boiling water bath. After cooling, the solution was centrifuged, and the absorbance of the supernatant was measured at 532 nm using the reaction mixture as a blank. Tetramethoxypropane (TMP, 1.25–20  $\mu\text{M}$ ) was used as standard and results were expressed as malonaldehyde (MDA) concentration ( $\mu\text{M}$ ) per million of cells.

### 2.7. Immunofluorescence microscopy analysis

An immunofluorescence microscopy analysis was carried out to evaluate the activation of the Nrf2-Keap1 pathway. IEC-6 cells were grown in culture dishes as described in section 2.3. Then an initial population of  $\approx 10,000$  cells were inoculated into square glass slides ( $1 \text{ cm} \times 1 \text{ cm}$ ), allocated inside wells of a 24-wells culture plate, mixed with 1 mL of DMEM (10% de FBS y 1% de PEST) and left overnight to allow adherence. Subsequently cells were treated as described in the section 2.4, using a concentration of 100  $\mu\text{M}$  of each PC [6]. After PC treatments, DMEM was removed from each well and the cells were rinsed three times with PBS 1x to remove excess of DMEM. Cells were fixed with a 4% PFA-PBS solution for 20 min and rinsed with PBS 1x twice. Subsequently, anti-Nrf2 primary antibody was added (50  $\mu\text{L}$  1:200) and incubated for 30 min. Incubation was stopped washing 3 times with PBS 1x. Then, the cells were incubated with the appropriate Alexa Fluor 594-conjugated secondary antibody-(50  $\mu\text{L}$  1:400) for 30 min. At the end of the incubation, the slides were washed 3 times with PBS 1x. and counterstained in the dark for 15 min with DAPI, and washed three times with PBS 1x. Finally, slides were mounted using glycerol 70% and observed with a Leica DM2000 (Leica fluorescence microscope, Leica microsystem, Mexico) using the RGB, blue (DAPI) and fluorescent (fluorescein) filters with 20x and 40x targets (Wetzlar, Hesse, Germany) to acquire images of the treated cells.

### 2.8. Statistical analysis

All assays were carried out by triplicate and results are presented as averaged  $\pm$  standard deviations (SD). CD and TBARS results were analyzed using a one-way analysis of variance (ANOVA). When ANOVA showed significant differences, a Tukey's post hoc test was applied. Statistical significance was regarded as  $P < 0.05$ . In the case of IR spectra, proportions mentioned in section 2.6 were used. For immunofluorescence microscopy, a frequency chart with SD were used to compare results from the different treatments.

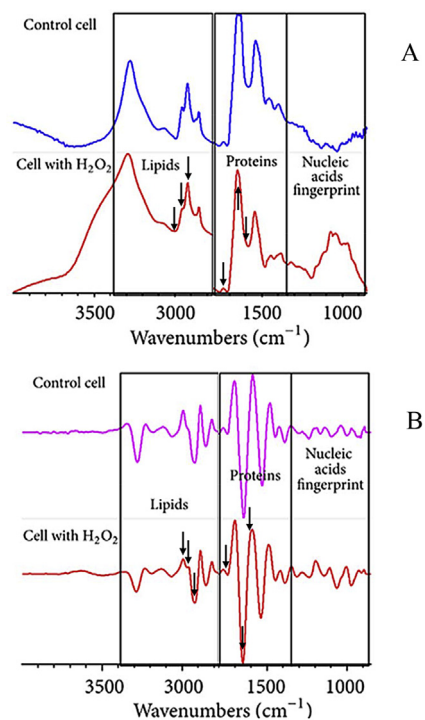
## 3. Results and discussion

In a previous study carried out by our research group, the effect of six different PC (Cap, Cat, Qc, gallic acid, caffeic acid and chlorogenic acid) against  $\text{H}_2\text{O}_2$ -induced oxidative stress in primary culture small intestine enterocyte was evaluated [6]. Results showed that some PC presented a protective effect when administered before or after the oxidative stimulus ( $\text{H}_2\text{O}_2$ ). From the six PC analyzed Cap and Cat showed the best protective effect, while phenolic acids showed weaker protective activities. When administered at 100  $\mu\text{M}$  in the absence of  $\text{H}_2\text{O}_2$ , all six PC behaved as mild pro-oxidants and this effect was more evident for Qc and phenolic acids. In order to describe the mechanism by which this PC present

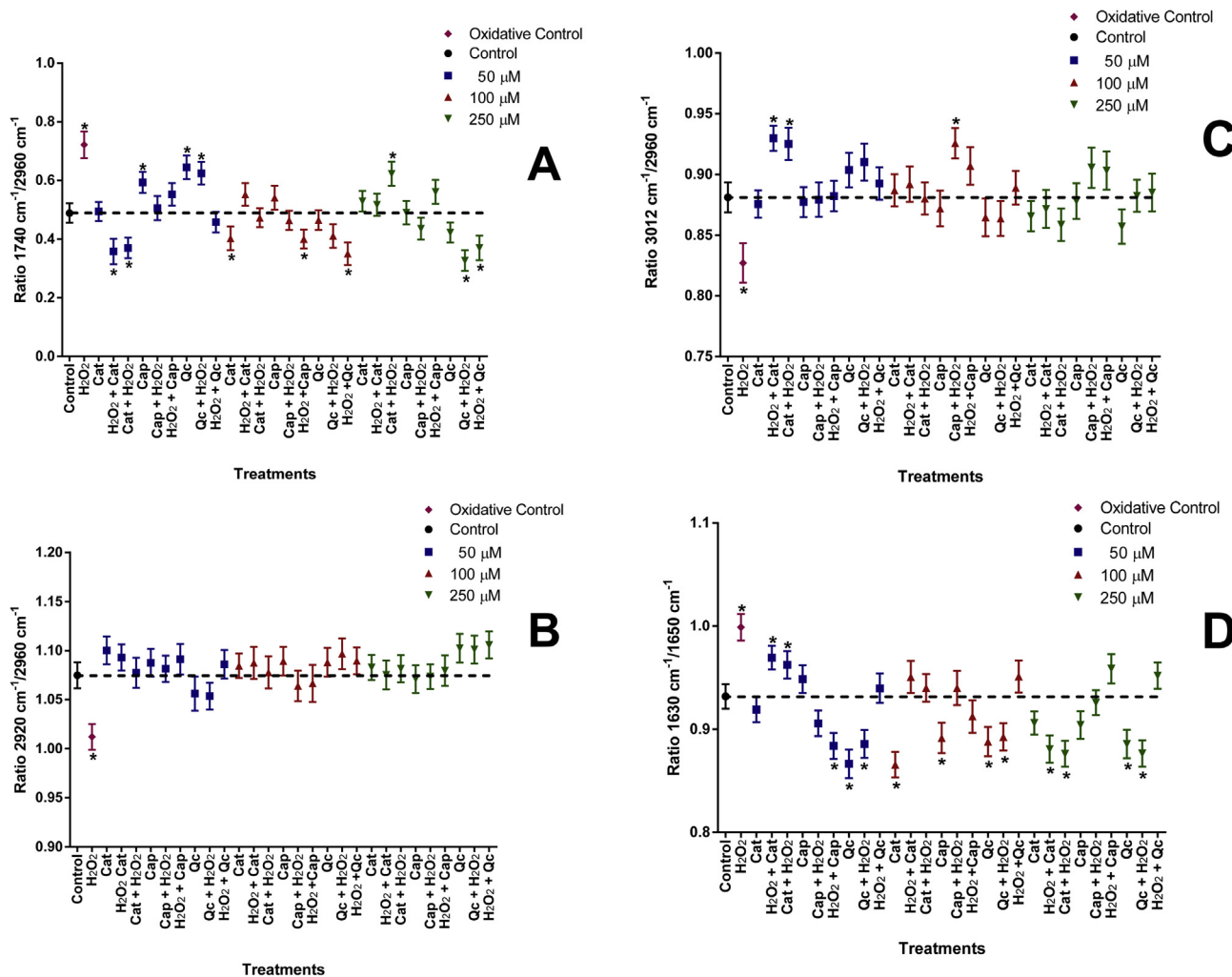
their protective effect, in the present study the effect of Cap, Cat and Qc at three different concentrations was evaluated, to determine if there is a dose-dependent effect of these PC against the  $\text{H}_2\text{O}_2$ -induced stress using a rat enterocyte cell line (IEC-6) as model.

### 3.1. Oxidative stress and antioxidant activity evaluated by infrared spectral analysis

Infrared spectroscopy was used to obtain information on the oxidation of proteins and lipids in IEC-6 cells treated with  $\text{H}_2\text{O}_2$  and/or three concentrations of Cat, Cap and Qc, by comparison of the FTIR spectra of control cells versus spectra from PC-treated cells [6]. Two sections of the FTIR spectra were analyzed. The first section, in the range of  $1750\text{--}1470 \text{ cm}^{-1}$ , known as the protein region, shows changes in aldehyde, amide I and II bands. The second section in the  $3200\text{--}2800 \text{ cm}^{-1}$  range, known as lipid region, shows changes in  $\text{CH}_2$  and  $\text{CH}_3$  alkyl chains and olefinic bond bands. Fig. 1 shows the FTIR absorbance spectra (1A) and second derivative spectra (1B) from control and  $\text{H}_2\text{O}_2$ -treated cells, to illustrate how raw and processed spectra data are seen. In order to evaluate the protective effect of PC, different FTIRM signal ratios related with lipid and protein oxidation, were analyzed:  $1740 \text{ cm}^{-1}/2960 \text{ cm}^{-1}$  (lipid oxidation),  $2920 \text{ cm}^{-1}/2960 \text{ cm}^{-1}$  (lipid saturation),  $3012 \text{ cm}^{-1}/2960 \text{ cm}^{-1}$  (lipid unsaturation) and  $1630 \text{ cm}^{-1}/1650 \text{ cm}^{-1}$  (protein aggregation). Fig. 2A shows the  $1740 \text{ cm}^{-1} / 2960 \text{ cm}^{-1}$  ratio results, which allowed us to evaluate the effect of oxidative stress on lipid oxidation (MDA production) [15]. From the analysis of this figure, it is possible to observe a significant difference between the  $\text{H}_2\text{O}_2$



**Fig. 1.** FTIRM infrared spectra (A) and second derivative analysis (B). (A) control treatment (DMEM 1 h) upper spectra and  $\text{H}_2\text{O}_2$ -treated cells (DMEM 30 min + 30 min 500  $\mu\text{M}$   $\text{H}_2\text{O}_2$  for 30 min) lower spectra. Spectra were obtained with a spectral resolution set to  $6 \text{ cm}^{-1}$ . (B) Average second derivative spectra using Savitsky-Golay with 21 smoothing points and a second polynomial order. (A) control treatment (DMEM 1 h) upper spectra and  $\text{H}_2\text{O}_2$ -treated cells (DMEM 30 min + 30 min 500  $\mu\text{M}$   $\text{H}_2\text{O}_2$  for 30 min) lower spectra. Three major zones of interest were observed: Fingerprint and nucleic acid zone ( $1470\text{--}900 \text{ cm}^{-1}$ ), protein zone ( $1720\text{--}1470 \text{ cm}^{-1}$ ), and lipid zone ( $3200\text{--}2800 \text{ cm}^{-1}$ ). The last two zones were used in this work. Arrows indicate the main points of analysis at  $1630 \text{ cm}^{-1}$ ,  $1650 \text{ cm}^{-1}$ ,  $1740 \text{ cm}^{-1}$ ,  $2920 \text{ cm}^{-1}$ ,  $2960 \text{ cm}^{-1}$  and  $3012 \text{ cm}^{-1}$ .



**Fig. 2.** Ratios between chemical bonds related to the lipid peroxidation observed with the FTIR spectra data. (A) 1740 cm<sup>-1</sup>/2960 cm<sup>-1</sup> ratio assigned to lipid oxidation, (B) 2920 cm<sup>-1</sup>/2960 cm<sup>-1</sup> ratio assigned to lipid saturation, (C) 3012 cm<sup>-1</sup>/2960 cm<sup>-1</sup> ratio assigned to lipid insaturation and (D) 1630 cm<sup>-1</sup>/1650 cm<sup>-1</sup> ratio assigned to protein aggregation. The dotted line indicates the mean value of the control cells for each ratio, all ratios were compared to the control ratio in each case, while their significance was measured via Anova. \* = Statistically differences respect to control treatment.

treatment (oxidative control), control treatment and PC treatments. H<sub>2</sub>O<sub>2</sub> treatment showed the highest ratio value, followed by Qc (free and Qc + H<sub>2</sub>O<sub>2</sub>) and Cap at low concentration (50 μM), and 250 μM Cat + H<sub>2</sub>O<sub>2</sub>. These results seem to indicate that the oxidative control and treatments with lower concentrations of Cap and Qc induced a higher production of MDA, a by-product of lipid peroxidation characterized by the presence of an aldehyde group (1740 cm<sup>-1</sup>). Similar results (increased MDA production) were observed for 250 μM Cat treatment before H<sub>2</sub>O<sub>2</sub>-induced oxidative stress, even though to a lesser degree than the oxidative control (H<sub>2</sub>O<sub>2</sub>), suggesting a low protective effect of Cat at this concentration. There is a series of treatments, mainly at 100 and 250 μM PC, with or without H<sub>2</sub>O<sub>2</sub>, without a statistical difference when compared to the control cells, which indicates that these treatments generated the same amount of MDA as the control. Interestingly, there was a third group of treatments which showed lower MDA formation (lower 1740 cm<sup>-1</sup>/2960 cm<sup>-1</sup> ratio) than the control cells. These treatments were 50 μM Cat and 250 μM Qc before or after H<sub>2</sub>O<sub>2</sub>. The data shown in Fig. 1 seems to indicate that the production of MDA may be triggered by low Cap and Qc concentrations (50 μM), in agreement with our previous work that showed higher production of MDA in Qc treated cells when compared to

control cells [6]. Interestingly, Qc showed a protective effect (lower MDA production) at the highest concentration (250 μM). Similar results have been reported in murine sperm cells [20] and PC12 cells [21] under oxidative stress, where higher Qc concentrations (100–500 μM) inhibited MDA production better than lower concentrations (<100 μM). The effects of Cat and Cap in the production of MDA were less dependent of their concentration, in agreement with previous results, which showed concentration-independent effects of Cap in an animal model [23] and of Cat in Caco-2 cells [24]. Interestingly, Cat which, similar to Qc, has a catechol group at ring B, but lacks the presence of the double bond and oxo group in ring C and therefore, has a less rigid structure than Qc, didn't show a prooxidant activity against parasites unless its concentration was high enough to be considered toxic to the organism [22]. The two hydroxyl groups in an ortho position in Cap seem to favor its scavenging activity when used against parasites or *in vitro*, showing lower oxidative stress [22,25]. These results suggest a Structure-Activity Relationship (SAR) between the structure of the PC and their protective effect against H<sub>2</sub>O<sub>2</sub>-induced oxidative stress.

The 2920 cm<sup>-1</sup>/2960 cm<sup>-1</sup> ratio evaluates the stability of lipids, associated to the proportion between primary and secondary



carbons present in the lipid structure [6,15]. The  $2920\text{ cm}^{-1}/2960\text{ cm}^{-1}$  ratio results are shown in Fig. 2B, where it is possible to observe that the oxidative control ( $\text{H}_2\text{O}_2$  treatment) exhibited a lower ratio than the control, indicating a significant effect on lipid saturation related to the loss of integrity in the lipid skeleton and the formation of new shorter aliphatic chains. No significant difference was observed between any PC treatment and the control, independently of the PC concentration. These results may indicate that the three studied PC can protect against the effects of  $\text{H}_2\text{O}_2$  on lipid saturation and integrity, and that this effect is neither dose nor structure dependent. The unsaturated lipids are targets for ROS. Thus, the third evaluated ratio,  $3012\text{ cm}^{-1}/2960\text{ cm}^{-1}$  (Fig. 2C), describes the effect of peroxidation on the decrease of unsaturated lipids inside the cell [15]. In agreement with previous results [6],  $\text{H}_2\text{O}_2$ -treated cells showed the lowest  $3012\text{ cm}^{-1}/2960\text{ cm}^{-1}$  ratio, indicating the highest oxidation of unsaturated lipids. Treatments with  $50\text{ }\mu\text{M}$  Cat before and after  $\text{H}_2\text{O}_2$  and  $100\text{ }\mu\text{M}$  Cap before  $\text{H}_2\text{O}_2$  reduced the unsaturated lipid peroxidation when compared to the control, indicating that these PC decrease the  $\text{H}_2\text{O}_2$ -induced basal oxidation of unsaturated lipids in the interior of the cells. Similar results have been reported when the effect of Cat was evaluated against ethanol-induced lipid peroxidation in rat livers [26]. The rest of the treatments presented a  $3012\text{ cm}^{-1}/2960\text{ cm}^{-1}$  ratio without significant differences with control cells, indicating that, they avoided  $\text{H}_2\text{O}_2$ -induced damage to unsaturated lipids.

Fig. 2D shows the  $1630\text{ cm}^{-1}/1650\text{ cm}^{-1}$  ratio, related to protein aggregation caused by protein oxidation as a product of oxidative stress [15]. In agreement with previous results [6],  $\text{H}_2\text{O}_2$ -treated cells showed the highest  $1630\text{ cm}^{-1}/1650\text{ cm}^{-1}$  ratio, indicating the highest protein aggregation due to oxidative stress. The treatments of Cat +  $\text{H}_2\text{O}_2$  and  $\text{H}_2\text{O}_2$  + Cat at  $50\text{ }\mu\text{M}$  showed intermediate values between control and  $\text{H}_2\text{O}_2$ -treated cells, indicating that Cat at low concentration showed smaller protective effects against protein aggregation. Interestingly, several treatments reduced protein aggregation compared to basal levels in control cells: all Qc concentrations, used alone or before  $\text{H}_2\text{O}_2$ ;  $50\text{ }\mu\text{M}$  Cap before or after  $\text{H}_2\text{O}_2$ ;  $100\text{ }\mu\text{M}$  Cap before  $\text{H}_2\text{O}_2$ ;  $100\text{ }\mu\text{M}$  Cat without  $\text{H}_2\text{O}_2$ ; and  $250\text{ }\mu\text{M}$  Cat with or without  $\text{H}_2\text{O}_2$ . Maurya et al., observed that Cat at low concentrations ( $20\text{ }\mu\text{M}$ ) showed a higher protein aggregation in erythrocyte hemoglobin under oxidative stress, probably due to a prooxidant effect [27]. It has been reported that Qc presented a protective effect against protein aggregation at concentrations above  $20\text{ }\mu\text{M}$ , however, its protective mechanism has not been described [28].

To evaluate the effect of the PC treatments on the relation between lipids and proteins, the  $2920\text{ cm}^{-1}/1650\text{ cm}^{-1}$  ratio was used to compare the  $\text{CH}_2$  from lipid skeletons with those of amide I bonds from protein (data not shown). Results showed that aside from the treatments  $\text{H}_2\text{O}_2$  + Cap and  $\text{H}_2\text{O}_2$  + Qc at  $50\text{ }\mu\text{M}$ , no significant differences were observed between all other treatments, including  $\text{H}_2\text{O}_2$ , and control, meaning that the relation between lipids and proteins was not affected by  $\text{H}_2\text{O}_2$ -induced oxidative stress. The differences observed between the control and the  $\text{H}_2\text{O}_2$ +Cap and  $\text{H}_2\text{O}_2$  + Qc  $50\text{ }\mu\text{M}$  treatments can indicate a better protection of lipids by a low concentration of Cap and Qc after the addition of  $\text{H}_2\text{O}_2$ , which is in agreement with TBARS results.

In summary we have seen that protection against lipid peroxidation by PC is dose dependent. Higher concentrations tend to exert better protective effect, except for Cat +  $\text{H}_2\text{O}_2$  at  $250\text{ }\mu\text{M}$ . Interestingly, at low concentration ( $50\text{ }\mu\text{M}$ ) PC seem to exert a prooxidant effect against lipids. The presence of the three PC protected the lipid skeleton and lipid unsaturation against  $\text{H}_2\text{O}_2$ -induced damage, regardless of its concentration. Finally, protein aggregation mediated by  $\text{H}_2\text{O}_2$ -induced oxidative stress was suppressed by all PC, except low Cat concentration, in agreement with previously reported results [27].

### 3.2. Evaluation of lipid oxidation by conjugated dienes (CD) and thiobarbituric acid reactive substances (TBARS)

To compare the FTIRM results, with standard spectrophotometric biochemical assays, primary and secondary lipid oxidation products were measured by CD and TBARS techniques respectively. Fig. 3A shows the results of CD from cells treated with Cap, Cat and Qc at three concentrations ( $50$ ,  $100$  and  $250\text{ }\mu\text{M}$ ) in the presence and absence of  $\text{H}_2\text{O}_2$ . These results showed that cells treated with  $\text{H}_2\text{O}_2$  alone produced the highest CD content with an average of  $4.72\text{ }\mu\text{M}$  of CD / million cells, while control cells showed the lowest CD content with an average of  $2.20\text{ mM}$  of CD / million cells (Fig. 3A). Interestingly, all IEC-6 treated cells showed the same trend on their CD values than primary cultures of rat enterocyte cells [6], but with lower values, probably because IEC-6 cells suffered less stress during manipulation.

Data in Fig. 3A, shows that all PC treatments presented CD values similar to those of the control, in agreement with results obtained with primary enterocyte cells [6]. When the effect of the PC concentration was evaluated, a non-statistical dose dependent increase on CD values was observed for all treatments (PC alone, before or after the addition of  $\text{H}_2\text{O}_2$ ) as the concentration increased from  $50$  to  $250\text{ }\mu\text{M}$ .

When evaluating the effect of PC on the generation of MDA, through TBARS (Fig. 3B), it was possible to observe that as in the case of CD,  $\text{H}_2\text{O}_2$ -stressed cells showed the highest MDA content, followed by PC treatments and finally positive control, which showed the lowest MDA content. MDA content of IEC-6 cells showed lower values compared to primary cultures of rat enterocyte cells [6], indicating once again that IEC-6 cells suffered less stress during manipulation.

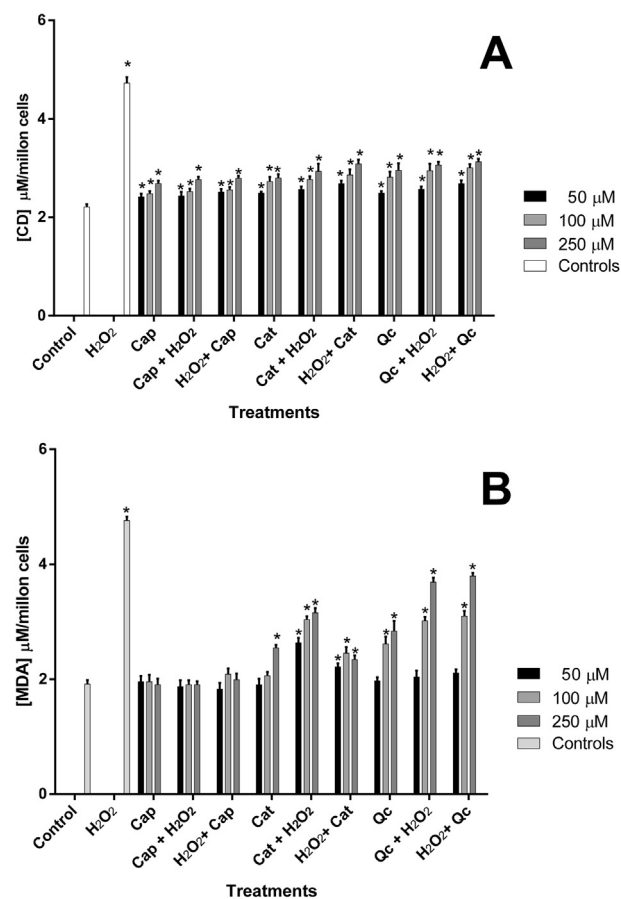
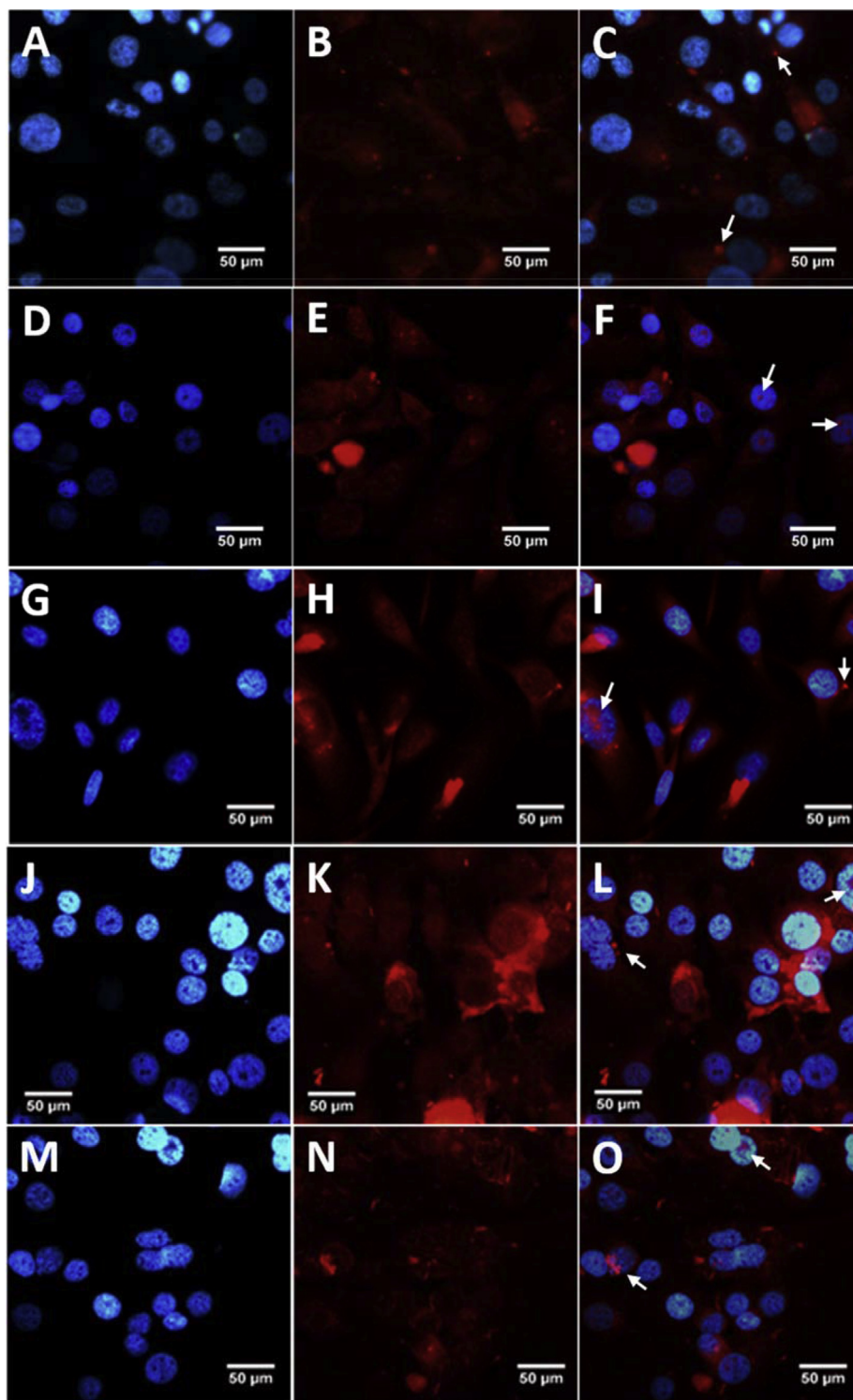


Fig. 3. Spectrophotometric assays to assess lipid peroxidation on IEC-6 treated cells. (A) CD assay. (B) TBARS assay. Values presented are averages + standard deviations of 6 individual measurements. \* = Statistically differences respect to control treatment.



**Fig. 4.** Immunofluorescence of transcription factor Nrf2 in IEC6 cells. Images A to C, are of cells without any treatment (A) Nuclei with DAPI counterstaining. (B) Location of Nrf2. (C) Co-localization of labels from (A) and (B). Images D to F, are of cells treated with H<sub>2</sub>O<sub>2</sub> (500 µM) (D) Nuclei with DAPI counterstaining. (E) Location of Nrf2. (F) Co-localization of labels from (D) and (E). G to I Qc 100 µM (G) Nuclei with DAPI counterstaining. (H) Location of Nrf2. (I) Co-localization of labels from (G) and (H). J-L, treatment with Qc (100 µM) administered prior to H<sub>2</sub>O<sub>2</sub> (500 µM) treatment. (J) Nuclei with DAPI, (K) Anti-Nrf2 fluorescence, (L) Overlap of fluorescence signals from (J) and (K). M – O, treatment with Qc (100 µM) administered after H<sub>2</sub>O<sub>2</sub> (100 µM); (M) DAPI nuclei counterstaining (N) Anti-Nrf2 fluorescence, (O) Overlap of fluorescence signals, DAPI stain and anti Nrf2 fluorescent labelling co-localize. White arrows in the merger images indicates some points in which Nrf2 is detected.

Fig. 3B shows the effect of the PC concentration on the production of MDA. It is possible to observe a dose-dependent increase on MDA concentration as Qc concentration increases, being this effect higher in the presence of H<sub>2</sub>O<sub>2</sub>. Similar results have been observed in liver cells [29] and in erythrocytes [30] treated with Qc at concentrations greater than 100 µM. This increase in MDA values shows that Qc

presents higher prooxidant effect as its concentration in the cell increases [30] and that its protective effect against H<sub>2</sub>O<sub>2</sub> is better at lower concentrations, since at higher concentrations, it might present a prooxidant effect. In the case of Cat, a significant dose-dependent increase in MDA concentration was observed for Cat alone and administered before the H<sub>2</sub>O<sub>2</sub>. This type of behavior

indicates that Cat has a slight prooxidant effect at concentrations higher than 100  $\mu\text{M}$  and that it is more effective to rescue cells from an oxidative stress-state than to prevent it. This effect has been previously reported when Cat was administered to endothelial cells [31] and HepG2 cells [32]. Finally, in the case of the Cap treatment, no significant variations in the concentration of MDA were observed after increasing Cap concentration, which indicates that Cap was the best protector against the formation of lipid oxidation products. Similar effect of Cap on the generation of MDA was observed in murine liver cells at concentrations below 100  $\mu\text{M}$  [33].

In summary, all PC, at all concentrations analyzed in the present study, showed no significant effects on CD content when administered by themselves and a complete protection against the  $\text{H}_2\text{O}_2$ -induced CD increment, which means that the protective effect is similar among different PC used. The TBARS assay showed that Cap presented the best protective effect. These results are in agreement with those reported with the FTIRM, in which, independently of the Cap concentration, the production of MDA was lower when compared with the other two PC, indicating that this PC shows a better protective effect against  $\text{H}_2\text{O}_2$ -induced lipid peroxidation.

### 3.3. Immunofluorescence for PC treatments

The immunofluorescence experiments were carried out to evaluate if the Nrf2-Keap-1 pathway was activated by the interaction of PC (alone or in the presence of  $\text{H}_2\text{O}_2$ ) with IEC-6 cells. PC can exert an antioxidant effect on the cells through two mechanisms: a direct or an indirect effect. In the case of a direct effect, the PC exerts its antioxidant activity through neutralization or stabilization of ROS [34]. While in an indirect effect, the PC do not interact directly with the ROS, but activate different biochemical pathways that increase the cell's antioxidant defenses, such as the Nrf2-Keap1 pathway which is common in all cell types. The exact mechanism by which a PC activates the indirect mechanism is still not well known [35], but it has been suggested that a mild prooxidant effect from PC within the cell can trigger it [34]. The binding of the Nrf2-Keap1 complex allows the ubiquitination of Nrf2, for its subsequent proteolysis, thus maintaining the levels of Nrf2 low and hindering its translocation to the nucleus. Inactivation of Keap1 by cysteine oxidation allows Nrf2 to translocate to the nucleus where it acts as a transcription factor for the Antioxidant Response Element (ARE) triggering the transcription of Phase II and antioxidant enzymes [36].

In order to test the hypothesis of activation of Nrf2 translocation by PC, IEC-6 cells were incubated with 100  $\mu\text{M}$  solutions of Qc, Cap and Cat, in the presence or absence of  $\text{H}_2\text{O}_2$ . Fig. 4 (A–I) shows the translocation of Nrf2 to the nucleus in the presence of Qc. Similar results were reported for Cap and Cat. Fig. 4 A–C are control cells, in which Nrf2 is located in the cytoplasmic area which indicates that without an oxidative stimuli Nrf2 is not translocated to the nucleus. Also, in Fig. 4B we can see how the lack of translocation allowed us to see the nucleus of the cells without the need for DAPI staining. Fig. 4 D–F correspond to the oxidative control cells, in which 500  $\mu\text{M}$   $\text{H}_2\text{O}_2$  was added. Fig. 4E and F shows that Nrf2 has been translocated to the cell nucleus, due to the oxidative stimulus. This translocation is necessary to activate the antioxidant and cell-protective response. Fig. 4 G–O show the effect of 100  $\mu\text{M}$  Qc treatments, in which loss of integrity in some nucleus can be observed through DAPI staining (Fig. 4G, J and 4M) when compared with the control cells (4A–C). Fig. 4H, K and 4N also show Nrf2 translocation to the nucleus in a similar pattern to the one observed for  $\text{H}_2\text{O}_2$ -treated cells (4D–F). Finally, the merged Fig. 4I, L and 4O shown that most of Nrf2 has been translocated to the nuclei, which indicates a robust activation of the Nrf2-Keap1 pathway. Similar results were observed for 100  $\mu\text{M}$  Cat and Cap treatments (data not shown). Similar effects on the translocation of Nrf2 have been reported for Qc (25  $\mu\text{M}$ ) in neurons [37], Cat (50  $\mu\text{M}$ ) in human mammary epithelium cells [34,38], Cap

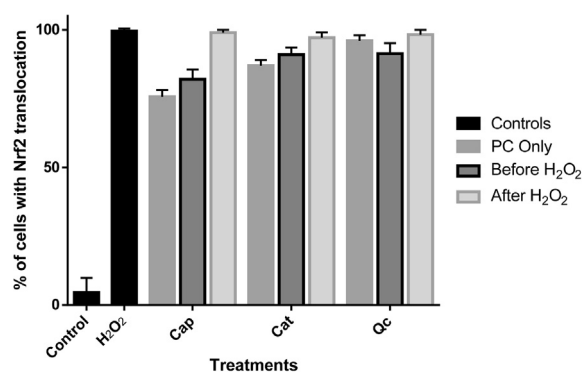


Fig. 5. Percentage of cells with translocation of Nrf2 to the nucleus. The negative and positive controls are shown in black. Meanwhile PC treatments are shown for every combination of treatments used: PC Only, PC before the addition of  $\text{H}_2\text{O}_2$  and after addition of  $\text{H}_2\text{O}_2$ . Every treatment shows its own SD.

(200  $\mu\text{M}$ ) in HepG2 cells [39], and for other phytochemicals such as isothiocyanates [40], resveratrol [41], curcumin [35] and others [35,37] in different cell systems.

Fig. 5 shows the percentage of Nrf2 translocation to the nucleus, considering cells exposed to  $\text{H}_2\text{O}_2$  as the positive control (100% translocation of Nrf2 to the nucleus), and DMEM-treated cells as the negative control (4.5% translocation). All PC treatments produced a clear translocation of Nrf2 to the nucleus, with Cap showing the lowest Nrf2 translocation percentage with 76% and  $\text{H}_2\text{O}_2$  + Cap the highest with 99%. These results show that the exposition to IEC-6 cells to Cap, Cat and Qc exerted a clear translocation of Nrf2 to the nucleus.

From the FTIRM, CD, TBARS and immunofluorescence results, we can conclude that Cat, Cap and Qc induced significant antioxidant effects in IEC-6 cells than can be mediated through the translocation of Nrf2 and activation of cellular antioxidant response although a direct antioxidant effect cannot be excluded, especially in lipid peroxidation.

## 4. Conclusion and perspective

After analyzing the FTIRM spectra, and standard spectrophotometric biochemical assays for lipid oxidation (CD and TBARS) results, it was concluded that, treatments with low concentration (50  $\mu\text{M}$ ), presented a mild oxidative stress effect, being this effect higher for Qc. Higher concentrations (100 and 250  $\mu\text{M}$ ) were more effective against  $\text{H}_2\text{O}_2$ -induced oxidative stress. Thus, higher PC concentrations showed a better antioxidant effect. Immunofluorescence studies in IEC-6 cells showed that all treatments with 100  $\mu\text{M}$  PC produced a translocation of Nrf2 to the nucleus, suggesting the activation of the Nrf2-Keap1 pathway. While these results expand our knowledge about how PC concentration can affect its antioxidant effect within enterocytes, there is still doubts about how the Nrf2-Keap1 pathway is activated and whether the concentration of PC affects the quantity of phase II enzymes that are produced by the cell. Finally, if there is an indirect antioxidant effect of the PC, the question arises whether this is the only mechanism, or there is also a direct scavenging activity at the same time.

### Declaration of competing interest

The authors declare that there is no conflict of interests regarding the publication of this paper.

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## Appendix A. Supplementary data

Supplementary data to this article can be found online at <https://doi.org/10.1016/j.saa.2019.117570>.

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