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#### **Research Article**

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# Effect of caffeine on acrylamide-induced cellular hepatotoxicity

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#### **Abstract**

**Objectives:** This study aims to determine the effects of caffeine on protein oxidation and endoplasmic reticulum (ER) stress in acrylamide (AA)-induced cellular hepatotoxicity.

**Methods:** First, HepG2 cells were incubated with  $10^1$ ,  $10^3$ , and  $10^4$   $\mu$ M AA. MTT assay was used to evaluate the cell viability. The western blot method was used to measure protein carbonyl (PC) levels to evaluate protein oxidation, and glucose-regulated protein 78 (GRP78), activating transcription factor 4 (ATF4), and C/EBP homologous (CHOP) levels to evaluate ER stress. Then, according to our results, cells were incubated with 10, 50, and 200  $\mu$ M caffeine simultaneously with  $10^4$   $\mu$ M AA and cell viability, PC, GRP78, ATF4 and CHOP levels were measured with the same methods.

**Results:**  $10^3$  and  $10^4$   $\mu$ M AA caused a significant decrease in cell viability (p<0.05 and p<0.001, respectively).  $10^4$   $\mu$ M AA caused a significant increase in PC (p<0.05), GRP78 (p<0.01), ATF4 (p<0.001), and CHOP (p<0.001) levels. 50 and 200  $\mu$ M caffeine significantly reduced PC levels in AA-incubated cells (p<0.05 and p<0.01, respectively). Caffeine concentrations did not significantly change in cell viability, GRP78, ATF4, and CHOP levels (p>0.05 for all).

**Conclusions:** Our study showed that; AA, especially at higher concentrations, caused an increase in cytotoxicity, protein oxidation, and ER stress, and caffeine decreases protein oxidation without changing AA-induced cytotoxicity and ER stress.

**Keywords:** acrylamide; caffeine; hepG2 cells; protein oxidation; endoplasmic reticulum stress

#### Introduction

Caffeine is a methylxanthine alkaloid, mainly found in coffee, cocoa beans, and tea leaves. Caffeine is the most widely used psychoactive drug in the world [1]. Caffeine is a well-known non-selective adenosine receptor antagonist that blocks adenosine and adenosine receptor binding-induced sleepiness in the central nervous system [2]. Caffeine decreases oxidative stress in many conditions [1]. Caffeine has anti-inflammatory, anti-infection, and anti-fibrosis effects, and its beneficial effects are shown in neurodegenerative disease, cardiovascular disease, obesity, and diabetes. It has also been reported that caffeine has hepatoprotective effects and prevents the progression of liver diseases [3, 4, 5, 6].

Acrylamide (AA) is a synthetic substance primarily used in experimental research and in synthesizing various polymers [7, 8]. AA is also formed in foods at high temperatures during frying and baking processes [9]. Food products such as fried potatoes, bread, biscuits, and coffee contain AA [7]. In addition, AA toxicity can be seen through respiration, water pollution, or skin contact [10] and causes neurotoxicity and genotoxicity [7]. AA metabolism mainly occurs in the liver. Also, AA causes toxicity in the liver [11, 12].

One of the damage mechanisms in AA toxicity is oxidative stress [11]. Reactive oxygen species (ROS) increase affects cellular redox balance and leads to post-translational modification of proteins. Protein carbonyl (PC) is a frequently used marker for protein oxidation and oxidative stress [13].

The endoplasmic reticulum (ER) is a vital organelle for folding and post-translational modifications of proteins. Various stimuli (accumulation of modified proteins, aging, pathological diseases, oxidative stress, etc.) cause ER stress by impairing ER function. Unfolded protein response is activated in the ER to maintain homeostasis, increase folding capacity, or induce cell death as a last option [14]. Glucoserelated protein 78 (GRP78), a heat shock protein 70 family member, is an ER chaperone. Under physiological conditions, GRP78 acts as a stress sensor embedded in the ER membrane and is inactive while bound to membrane proteins. When ER stress occurs, GRP78 separates from these proteins, becomes active, and binds to misfolded proteins.

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This separation initiates the UPR cascade, including three major pathways named protein kinase R-like ER kinase (PERK), inositol requiring enzyme 1 (IRE1) and activating transcription factor 6 (ATF6) [14, 15].

In previous studies, it was reported that PERK- eukary-otic translation initiation factor  $2\alpha$  (eIF2 $\alpha$ )-activating transcription factor 4 (ATF4)-C/EBP-homologous protein (CHOP) signaling was the main pathway in AA-induced apoptotic neuronal cell death, and AA had a slight effect on IRE pathway [16, 17]. In this signaling pathway, PERK activation leads to the phosphorylation of eIF2 $\alpha$ , which reduces the amount of protein accumulated in the lumen by inhibiting protein translation and stimulates the transcription of ATF4 [18]. ATF4 is associated with amino acid metabolism and oxidation/reduction, and it activates CHOP, also called the growth arrest and DNA damage-inducible gene 153 (GADD153), which is the most important molecule for the ER stress-induced apoptotic cell death [19, 20, 21, 22].

No study investigated the effects of caffeine on AA toxicity in the liver. Oxidative and ER stresses are tightly associated with AA-induced hepatotoxicity development through complex signaling pathways involving apoptosis, inflammation and autophagy [11]. In this study, we aimed to determine the effects of caffeine on protein oxidation and ER stress in AA-induced cellular hepatotoxicity. The HepG2 cells are immortal human-derived liver hepatoma cell lines frequently preferred in chemical-induced hepatotoxicity models [11, 12]. Depending on daily coffee consumption, plasma caffeine concentrations of adults are usually between 10 and 50 µM, and it has been reported that caffeine has toxic effects at concentrations above 200 µM [23, 24, 25]. In this study, we first determined the effects of AA concentrations in HepG2 cells and then the effects of 10, 50, and 200 µM caffeine in AA-induced protein oxidation and ER stress.

#### Materials and methods

#### **Experimental design**

This study is approved by the ethics committee (TÜTF-BAEK 2018/299). HepG2 cells (ATCC) were incubated at 37 °C and 5 %  $\rm CO_2$  atmosphere. EMEM (Wisent) containing 10 % FBS (Thermo) and 1 % antibiotic and antimycotic (Thermo) was used for the cultivation.

Cells were incubated with  $10^1$ ,  $10^3$ , and  $10^4\,\mu\text{M}$  AA (Merck) to investigate the effect of AA concentrations for 24 h. The control group was simultaneously incubated with medium without acrylamide in this experiment. For the treatment, cells were incubated with 10, 50, and 200  $\mu\text{M}$ 

caffeine (Sigma) simultaneously with  $10^4~\mu M$  AA for 24 h. In this experiment, the control group was simultaneously incubated with a medium that included neither acrylamide nor caffeine. Also, a flask of cells was incubated with 12  $\mu M$  tunicamycin (Sigma), which is shown to induce ER and oxidative stress, for 24 h to show ER and oxidative stress-induced protein bands for western blot experiments [26, 27].

#### MTT assay

MTT assay was used to evaluate cell viability [28].  $10^4$  cells per well seeded into 96 well plates. After removing mediums containing treatments, 5 mg/mL MTT (Sigma) and  $100~\mu$ L EMEM were added to wells at the end of the incubations. After 4 h at the CO<sub>2</sub> incubator at 37 °C, formazan was dissolved by DMSO and Sorenson buffer, and the optical densities of wells were measured at 570/630 nm [29]. The results were expressed as percentages of the control group.

#### Western blotting

After incubations, cells were lysed with 10 mM Tris at pH 8.0 with 50 mM EDTA and %1 SDS (w/v) at 90 °C. The protein content of lysates was measured [30]. For determining PC levels, 5 µg protein containing lysate was incubated with 2,4dinitrophenylhydrazine (DNPH) (Sigma) [31] as previously described [32]. 20 µg of total protein were loaded into electrophoresis gels to determine GRP78, ATF4, and CHOP levels. Proteins of samples were loaded to electrophoresis gel and blotted to the PVDF membrane (Roche) after separation [33]. After 1 h of blocking with milk powder (Sigma) in TBST (5 %), membranes were incubated with specific antibodies (PC: 1:5,000, GRP78; 1:10,000-, ATF4; 1:2000, CHOP; 1:200, tubulin; 1:10,000) overnight at 4 °C and secondary antibodies for 1 h at laboratory conditions. Specific bands on the membrane were visualized with an electrochemiluminescence imaging system and calculated with Image-J [34]. Results were calculated by dividing specific protein results into the same sample's loading control tubulin result. Then, the results were divided into the result of the control group presented in the same experiment and on the same membrane. All results were expressed as fold change relative to the control.

#### Statistical analysis

SPSS 20 program (IBM) was used for statistical analysis. Results were calculated as a percentage of control for each MTT assay and as a fold of control for each Western blot. Data was expressed as the mean ± standard deviation of three results from three independent experiments. The One-Way ANOVA test was used to compare parameters among the groups, and then Tukev's test was used to compare groups. The p<0.05 was considered statistically significant.

#### **Results**

#### AA causes cytotoxicity at higher concentrations

The viability percentages were  $100 \pm 2 \%$  for  $10^1 \mu M$ ,  $88 \pm 5 \%$ for  $10^3 \, \mu M$ , and  $57 \pm 7 \, \%$  for  $10^4 \, \mu M$  AA-incubated cells.  $10^3 \, \mu M$ AA caused significant cytotoxicity compared to control and  $10^{1} \mu M$  AA-incubated cells (p<0.05 for both).  $10^{4} \mu M$  AA caused significant cytotoxicity compared to control, 10<sup>1</sup> AAincubated cells and 10<sup>3</sup> µM AA-incubated cells (p<0.001 for all) (Figure 1).

## AA increases protein oxidation and ER stress at $10^4 \, \mu M$

PC levels of  $10^1 \mu M$  AA-incubated cells were 1.26  $\pm$  0.15 fold,  $10^3 \, \mu M$  AA-incubated cells were 1.18  $\pm$  0.20 fold, and  $10^4 \, \mu M$ AA-incubated cells were  $2.04 \pm 0.62$  fold of control.  $10^4 \,\mu\text{M}$  AA caused a significant increase in PC levels compared to the control group (p<0.05) (Figure 2).

GRP78 levels of 10<sup>1</sup> µM AA-incubated cells were  $1.11 \pm 0.16$  fold,  $10^3 \mu M$  AA-incubated cells were  $1.16 \pm 0.33$ fold, and  $10^4 \, \mu M$  AA-incubated cells were 1.73  $\pm$  0.17 fold of control.  $10^4 \, \mu M$  AA-incubated cells had significantly higher GRP78 levels than control (p<0.01), 10<sup>1</sup> AA-incubated cells (p<0.05) and  $10^3 \mu M$  AA-incubated cells (p<0.05) (Figure 2).

ATF4 levels of  $10^1 \, \mu M$  AA-incubated cells were  $1.05 \pm 0.05$ fold,  $10^3 \, \mu M$  AA-incubated cells were 2.92  $\pm$  0.29 fold, and  $10^4 \,\mu\text{M}$  AA-incubated cells were 20.18  $\pm$  2.34 fold of control.

10<sup>4</sup> μM AA-incubated cells had significantly higher ATF4 levels than the control,  $10^1$  AA-incubated cells and  $10^3$  µM AAincubated cells (p<0.001 for all) (Figure 2).

CHOP levels of  $10^1$  µM AA-incubated cells were 1.19  $\pm$  0.10 fold,  $10^3 \,\mu\text{M}$  AA-incubated cells were 1.50  $\pm$  0.21 fold, and  $10^4 \,\mu\text{M}$  AA-incubated cells were 2.26  $\pm$  .19 fold of control. 10<sup>3</sup> μM AA-caused a significant increase in CHOP levels (p<0.05). 10<sup>4</sup> µM AA-incubated cells had significantly higher CHOP levels than the control (p<0.001), 10<sup>1</sup> AA-incubated cells (p<0.001) and  $10^3 \mu M$  AA-incubated cells (p<0.01) (Figure 2).

### Caffeine concentrations do not change cell viability in AA-incubated cells

According to our results, HepG2 cells were incubated with caffeine concentrations simultaneously with 10<sup>4</sup> µM AA, which induced cytotoxicity, oxidative, and ER stress.

The viability percentages were  $58 \pm 1\%$  for  $10^4 \,\mu\text{M}$  AAincubated cells,  $58 \pm 3\%$  for  $10^4 \mu M$  AA+  $10 \mu M$  caffeineincubated cells,  $63 \pm 3\%$  for  $10^4 \mu M$  AA+  $50 \mu M$  caffeineincubated cells, and  $62 \pm 1 \%$  for  $10^4 \mu M$  AA+ 200  $\mu M$  caffeineincubated cells. 10, 50, and 200 µM caffeine did not significantly alter cell viability in 10<sup>4</sup> µM AA-incubated cells (p>0.05 for all) (Figure 3).

# Caffeine decreases AA-induced protein oxidation at higher concentrations without affecting ER stress

PC levels of  $10^4 \, \mu M$  AA-incubated cells were  $2.04 \pm 0.10$  fold,  $10^4 \, \mu M$  AA+ 10  $\mu M$  caffeine-incubated cells were 1.80  $\pm$  0.15 fold, 10<sup>4</sup> μM AA+ 50 μM caffeine-incubated cells were  $1.64 \pm 0.12$  fold, and  $10^4 \, \mu M$  AA+ 200  $\mu M$  caffeine-incubated cells were 1.45  $\pm$  0.13 fold of control. 50  $\mu M$  caffeine caused a significant decrease in PC levels in 10<sup>4</sup> μM AA-incubated cells (p<0.05). 10<sup>4</sup> μM AA+ 200 μM caffeine-incubated cells had

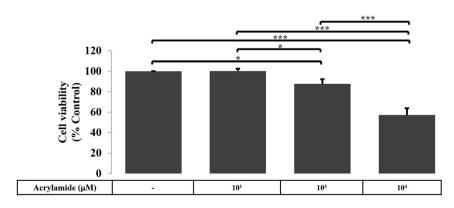


Figure 1: Effects of acrylamide on cell viability in HepG2 cells. Results are expressed as the mean  $\pm$  standard deviation of three results from three independent experiments. Data was analyzed by one-way ANOVA, and turkey post-hoc test were performed for multiple comparisons. \*: p<0.05, \*\*\*: p<0.001.

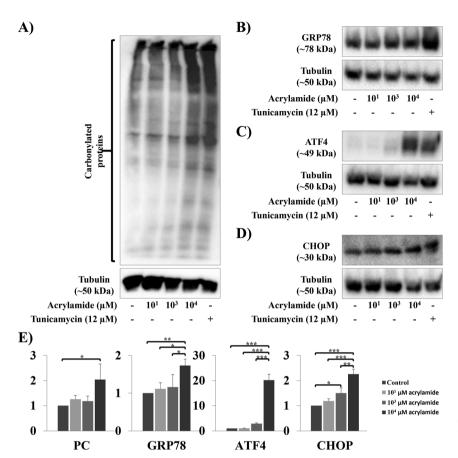
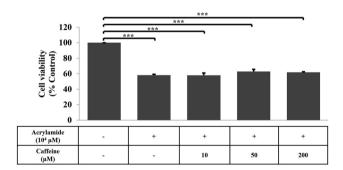


Figure 2: Effects of acrylamide on protein oxidation and ER stress in HepG2 cells. Effect of acrylamide concentrations on (A) PC levels (B) GRP78 levels (C) ATF4, and (D) CHOP levels in HepG2 cells. Tunicamycin-treated cells were used to show ER and oxidative stress-induced protein bands (not calculated). Results are expressed as the mean ± standard deviation of three results from three independent experiments (E). Data was analyzed by one-way ANOVA, and turkey post-hoc test were performed for multiple comparisons. \*: p<0.05, \*\*\*: p<0.01, \*\*\*: p<0.001.



**Figure 3:** Effects of caffeine on cell viability in acrylamide-incubated HepG2 cells. Results are expressed as the mean  $\pm$  standard deviation of three results from three independent experiments. Data was analyzed by one-way ANOVA, and turkey post-hoc test were performed for multiple comparisons. \*\*\*: p<0.001.

significantly lower PC levels than  $10^4~\mu M$  AA incubated cells (p<0.01) and  $10^4~\mu M$  AA+  $10~\mu M$  caffeine-incubated cells (p<0.05) (Figure 4).

GRP78 levels of  $10^4\,\mu\text{M}$  AA-incubated cells were  $1.74\pm0.14$  fold,  $10^4\,\mu\text{M}$  AA+  $10\,\mu\text{M}$  caffeine-incubated cells were  $1.79\pm0.22$  fold,  $10^4\,\mu\text{M}$  AA+  $50\,\mu\text{M}$  caffeine-incubated cells were  $1.75\pm0.19$  fold, and  $10^4\,\mu\text{M}$  AA+  $200\,\mu\text{M}$  caffeine-incubated cells were  $1.94\pm0.19$  fold of control.  $10,\,50,\,$  and

200  $\mu$ M caffeine treatment did not significantly alter GRP78 levels in 10<sup>4</sup>  $\mu$ M AA-incubated cells (p>0.05 for all) (Figure 4).

ATF4 levels of  $10^4 \, \mu M$  AA-incubated cells were  $19.64 \pm 4.20$  fold,  $10^4 \, \mu M$  AA+  $10 \, \mu M$  caffeine-incubated cells were  $17.84 \pm 3.24$  fold,  $10^4 \, \mu M$  AA+  $50 \, \mu M$  caffeine-incubated cells were  $19.21 \pm 5.34$  fold, and  $10^4 \, \mu M$  AA+  $200 \, \mu M$  caffeine-incubated cells were  $22.00 \pm 4.07$  fold of control. 10, 50, and  $200 \, \mu M$  caffeine treatment did not significantly alter ATF4 levels in  $10^4 \, \mu M$  AA-incubated cells (p>0.05 for all) (Figure 4).

CHOP levels of  $10^4\,\mu\text{M}$  AA-incubated cells were  $2.06\pm0.10$  fold,  $10,000\,\mu\text{M}$  AA+  $10\,\mu\text{M}$  caffeine-incubated cells were  $1.60\pm0.15$  fold,  $10^4\,\mu\text{M}$  AA+  $50\,\mu\text{M}$  caffeine-incubated cells were  $1.95\pm0.38$  fold, and  $10^4\,\mu\text{M}$  AA+  $200\,\mu\text{M}$  caffeine-incubated cells were  $2.17\pm0.09$  fold of control. 10, 50, and  $200\,\mu\text{M}$  caffeine treatment did not significantly alter CHOP levels in  $10^4\,\mu\text{M}$  AA-incubated cells (p>0.05 for all).  $10^4\,\mu\text{M}$  AA+  $200\,\mu\text{M}$  caffeine-incubated cells had significantly higher CHOP levels than  $10^4\,\mu\text{M}$  AA+  $10\,\mu\text{M}$  caffeine-incubated cells (p<0.05) (Figure 4).

#### **Discussion**

Caffeine is a plant-based substance in coffee, tea and energy drinks. Thus, it is one of the most frequently consumed

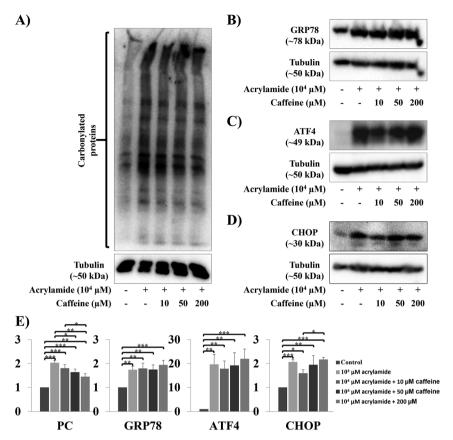


Figure 4: Effects of caffeine on protein oxidation and ER stress in acrylamideincubated HepG2 cells. Effect of caffeine concentrations on (A) PC levels (B) GRP78 levels (C) ATF4, and (D) CHOP levels in acrylamideincubated cells. Results are expressed as the mean  $\pm$  standard deviation of three results from three independent experiments (E). Data was analyzed by one-way ANOVA, and turkey post-hoc test were performed for multiple comparisons. \*: p<0.05, \*\*: p<0.01, \*\*\*: p<0.001.

chemicals worldwide [1]. AA is a water-soluble substance that people are exposed to occupationally and with food [35]. It's metabolized in the liver and has been shown to have toxic effects on the liver [11, 12, 36]. Previous studies reported the hepatoprotective effects of caffeine in alcoholic liver injury [5] and liver fibrosis [6]. Still, there is no study investigating the effects of caffeine on AA toxicity in the liver. In the present study, we aimed to examine the effects of caffeine on protein oxidation and ER stress in AA-induced cellular hepatotoxicity. Also, researchers pre-treated rat primary astrocytes [37], human astrocytoma [38, 39], neuroblastoma and glioblastoma cells [40] with caffeine as ataxia telangiectasia mutated (ATM)/ATM-Rad3-related (ATR) inhibitor for investigating the mechanism action of AA in their study. However, we could not find any study determining the effect of caffeine together with acrylamide. With the present study, we also determined the effect of caffeine together with acrylamide for the first time.

First, to evaluate the effects of AA concentrations on cytotoxicity, protein oxidation, and ER stress, cells were incubated with  $10^1$ ,  $10^3$ , and  $10^4 \,\mu$ M AA. In our study,  $10^3$  and 10<sup>4</sup> μM AA caused cytotoxicity in HepG2 cells. Our results support the studies in the literature, which showed that AA has toxic effects on HepG2 cells at high concentrations [12, 40, 41]. It is known that acrylamide weakens antioxidant defense systems by causing glutathione depletion during its metabolism, and oxidative stress increases acrylamide toxicity [7, 11]. We showed that 10<sup>4</sup> µM AA increased oxidative stress with increased PC levels in HepG2 cells, and similar to our results, the increase of oxidative stress in acrylamide-treated HepG2 cells was shown in previous reports [41, 42]. It was demonstrated that AA has upregulated ER stress pathways [16, 43]. Also, previous studies reported that AA causes an increase in ER stress, including GRP78 and CHOP levels in kupffer cells, rat liver [44] and HepG2 cells [42]. Compatible with the literature,  $10^3 \,\mu\text{M}$  and  $10^4 \,\mu\text{M}$  AA caused an increase in ER stress in HepG2 cells.

Based on our first results, we chose 10<sup>4</sup> µM AA to induce cytotoxicity, protein oxidation and ER stress in subsequent experiments to investigate the effects of caffeine concentrations. Plasma caffeine concentrations in adults are usually between 10 and 50 µM, and it has been reported that caffeine has toxic effects at concentrations above 200 µM [23, 24, 25]. Thus, to determine the effects of caffeine on AA-induced cellular hepatotoxicity, the cells were incubated with 10, 50, and 200  $\mu$ M caffeine simultaneously with 10<sup>4</sup>  $\mu$ M AA.

In our study, caffeine concentrations did not significantly affect the cell viability in AA-treated HepG2 cells. Our findings showed that our caffeine concentrations were ineffective in preventing AA-induced hepatotoxicity in HepG2 cells. On the other hand, caffeine was reported to be effective at 20 mM concentration against lipopolysaccharide-induced cytotoxicity [45]. We used caffeine concentrations compatible with human plasma caffeine concentrations, and our previous report showed that cell viability decreased to 84 % even when incubated with 200  $\mu$ M caffeine for 24 h in HepG2 cells [46]. Also, it was shown that higher caffeine concentrations are cytotoxic to HepG2 cells [47]. In light of our previous study and literature, higher caffeine concentrations may do more harm than good in AA-induced hepatotoxicity in HepG2 cells.

We found that 50 and 200  $\mu$ M caffeine caused a significant decrease in PC levels in AA-incubated HepG2 cells. The antioxidant properties of caffeine are well-known [1], and our results showed that caffeine causes an antioxidant effect against AA-induced protein oxidation. Also, Pasaoglu et al. [48] reported that 30 and 100 mg/kg/day oral caffeine treatment for 14 days causes a decrease in liver advanced oxidation protein products levels in rats, and Amer et al. [49] reported 37.5 mg/kg/day caffeine caused a decrease in liver protein carbonyl levels and malondialdehyde levels in thioacetamide-induced liver injury in rats.

Caffeine concentrations in this study did not significantly change the GRP78, ATF4, and CHOP levels in AA-treated HepG2 cells. Our findings showed that 10, 50, and 200  $\mu M$  caffeine does not affect AA-induced ER stress in HepG2 cells. We could not encounter any study investigating the effect of caffeine at concentrations similar to our research on ER stress in the liver *in vivo* or *in vitro*. However, at high concentrations, there are contradictory results regarding the effects of caffeine on ER stress in different liver-associated cell lines. Hu et al. [45] reported that caffeine is effective at 20 mM concentration against lipopolysaccharide-induced cytotoxicity and ER stress in L02, human fetal hepatocyte, cell line. On the other hand, Li et al. [50] reported that caffeine induces ER stress and apoptosis at 5 mM concentrations and above in LX-2, immortalized human hepatic stellate cell line.

As a limitation, although caffeine and acrylamide have systemic effects, our *in vitro* study investigated the effects of these chemicals in a cellular liver model using an immortal cell line.

In conclusion, we found that AA, especially at higher concentrations, caused an increase in cytotoxicity, protein oxidation, and ER stress, and caffeine decreases protein oxidation without changing AA-induced cytotoxicity and ER stress. Our results show that caffeine may be beneficial in preventing protein oxidation, but it's insufficient to prevent AA-induced hepatotoxicity.

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**Research ethics:** The ethics committee of Trakya University School of Medicine approved this study (TÜTF-BAEK 2018/299, date of approval: 17.09.2018).

**Informed consent:** Not applicable.

**Author contributions:** The authors have accepted responsibility for the entire content of this manuscript and approved its submission.

Use of Large Language Models, AI and Machine Learning Tools: None declared.

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**Data availability:** The raw data can be obtained on request from the corresponding author.

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