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Zinc attenuates ethanol-induced Sertoli cell toxicity and apoptosis through caspase-3 mediated pathways

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ABSTRACT

Ethanol enhances apoptosis in testicular germ cells. Zinc reduces ethanol-induced apoptosis of somatic cells through inhibition of caspase-mediated pathways. Little is known about the effects of ethanol on Sertoli cells and the effects of Zinc on ethanol-induced testicular injury. The hypothesis tested was that ethanol enhances apoptosis of Sertoli cells through up-regulation of caspase-3 and Zinc inhibits ethanolinduced effects. Cultured Sertoli cells (TM4) were exposed to ethanol (160 mM), Zinc (8 µM) and Zinc prior to ethanol for duration of 24 or 48 h and their effects on TM4 cell viability was then investigated by MTT assay. Caspase-3 mRNA expression was also investigated using real-time RT-PCR. Cell viability decreased and caspase-3 mRNA expression increased in cells exposed to ethanol, while exposure to Zinc showed opposite effects. Pretreatment with Zinc recovered ethanol-induced anti-proliferative effects and over-expression of caspase-3. Zinc reduced ethanol-induced Sertoli cell toxicity and apoptosis via caspase-3 mediated pathways.

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1. Introduction 24

Humans are exposed to varied environmental agents poten-25 tially hazardous to the reproductive system. Male reproduction, 26 which its dysfunction is known to contribute to more than half 27 of all cases of global childlessness [1], is highly responsive and 28 sensitive to a variety of chemical and physical agents [2]. Chronic 29 stress, obesity, urogenital trauma and inflammation, smoking and 30 alcohol abuse are associated with decreased male fertility [3,4]. 31 Alcohol has been shown to have a deleterious effect at all levels 32 of male reproductive system, including the hypothalamus, ante-33 rior pituitary gland and the testes [5]. Chronic alcohol intake in 34 experimental animals has been associated with marked degener-35 ative changes such as germ cell damage, testicular atrophy and 36

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reduction in seminiferous tubular size [6-10]. Moreover, spermatogenic failure, hypogonadism, gonadal dysfunction and infertility by triggering testicular germ cell apoptosis have been reported in humans with a history of chronic alcohol abuse [11]. However, the mechanism(s) of ethanol-induced infertility remain poorly understood [12]. Ethanol exposure also adversely affects the secretory function of testicular Sertoli cells [7], and disrupts Sertoli cellmediated blood-testes barrier [13]. Sertoli cells play a supportive and crucial role in spermatogenesis by providing physical support, nutrients and survival signals, controlling germ cell proliferation and apoptosis which is necessary for successful spermatogenesis [14-17]. Ethanol is very toxic for Sertoli cells, making them to be the primary cellular target of ethanol toxicity with marked Sertoli cell vacuolization being a morphological marker of ethanol-induced testicular injury [7,9,18–20]. 03 51

Alcohol consumption induces apoptotic cell death in different cell lines and various organs including the brain, liver, thymus and testis [14,21-23]. Within the testes, apoptosis occurs under physiologic conditions and provides to avoid overproduction of normal cells and maturation of abnormal cells. Under pathologic conditions, such as an exposure to specific environmental toxins, a greater portion of testicular cells degenerate through the apoptosis, leading to spermatogenic failure and infertility [24]. The process of

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Fig. 1. The effect of ethanol on TM4 cell viability. (A) Comparison of the effects of different concentrations of ethanol on TM4 cell viability following 24 or 48 h exposure periods. ^a Represents p = 0.005 compared to the control group (one-way ANOVA). All results are presented as mean \pm S.E.M derived from two determinations in duplicate cultures. (B) Comparison of data between 24 and 48 h exposure periods revealed no significant changes. As concentration of ethanol increased, cellular viability decreased and eventually reached to approximately 50% of control level at concentration of 160 mM following 48 h exposure period.

apoptosis is triggered and regulated by the expression of several proteins. Caspase, a highly specific class of cysteine proteases, is expressed in many mammalian cells and known to mediate a crucial stage of the apoptotic process [25,26]. Of particular interest is caspase-3, the most widely studied member of the caspase family, which is, at least partially, responsible for the degradation of DNA and cleavage of cellular proteins, leading to apoptosis [27–30]. Previous *in vitro* and *in vivo* studies have shown involvement of several proteins and pathways in ethanol-induced apoptosis of testicular cells, including Fas system, active caspases and bax-dependent signaling pathways [14,22,23]. Based on the results, ethanol appears to induce apoptosis in Leydig cells via bax-dependent caspase-3 activation [22].

It has been reported that ethanol-induced testicular damage is 73 mainly mediated by DNA damage, oxidative stress and androgen 74 suppression [7,14,23,31]. Oxidative stress is caused by imbalanced 75 between the production of reactive oxygen species (ROS) and 76 the production of antioxidant system that is responsible for their 77 neutralization and removal. Increased oxidative stress is a well-78 accepted mechanism of alcohol induced tissue injury, particularly 79 in the liver, heart, central nervous system and testis [3,5,32,33]. In 80 the testis, oxidative stress affects reproductive activity, damages 81 DNA structure and induces apoptosis, leading to fertility distur-82 bances or embryo development disorders [3,34,35]. Zinc as an 83 antioxidant agent, reduced ethanol-induced apoptosis of somatic 84 cells in different non-sexual organs [36,37]. Zinc plays important 85 roles in various biological activities but its role in spermatogenesis 86 is not well known. Zinc deficiency has been documented in alco-87 holic liver disease, skin, esophagus, gastric and kidney damages, 88 in both patient and animal [36–40]. It has been shown that Zinc 89 deprivation induces apoptosis in diverse cell lines and potentiates 90 91 death receptor-mediated apoptosis [41,42]. In this respect, testicular apoptosis after dietary zinc deficiency has also been reported 92



Fig. 2. The effect of Zinc on TM4 cell viability. (A) Comparison of the effects of different concentrations of Zinc on TM4 cell viability following 24 or 48 h exposure periods. At concentration of 8 μ M of Zinc, cell viability and proliferation increased compared to control and higher concentrations of Zinc. ^a Represents p < 0.01 compared to the control and the 64 μ M Zinc-treated groups. ^b Represents p < 0.01 compared to the control. ^c Represents p < 0.05 compared to the 64 μ M Zinc-treated group (one-way ANOVA). All results are presented as mean \pm S.E.M derived from two determinations in duplicate cultures. Comparison of data between 24 and 48 h exposure periods revealed no significant changes. (B) As concentration of Zinc increased cell viability decreased and eventually showed a trend around the control level.

[43]. Moreover, Zinc inhibits ethanol-induced hepatic apoptosis in both cell culture and mice subjected to a long-term ethanol exposure [36,37], however its protective effects on ethanol-induced testicular injury is poorly understood.

Despite of toxic effects of ethanol and protective effects of zinc on multiple cells and organs, there is little available information regarding their effects on testis and testicular somatic cells. In the present study, the effect of different concentration of ethanol and zinc (single and co-administration) on the cell viability and apoptosis (by measuring caspase-3 gene expression) in cells of the TM4 cell line, derived from mouse Sertoli cells, were investigated to determine whether an enhancement of Sertoli cell death occurs as a result of ethanol exposure and whether zinc can inhibit ethanolinduced effects.

2. Material and methods

2.1. Cell line and culture conditions

The Mus Musculus Sertoli cell line, TM4 (purchased from Pasteur Institute, Tehran, Iran) was used in all experiments. The TM4 cells were cultured in Dulbecco's Modified Eagle's Medium (DMEM)/F12/hepes media (Atocel, Austria) supplemented with 2.5% fetal bovine serum (Atocel, Austria), 5% Horse serum (Atocel, Austria), penicillin (50 units/ml) and streptomycin (50 units/ml), in 25 cm² tissue flasks maintained at 37 °C in a humidified atmosphere with 95% O₂ and 5% CO₂ in a cell incubator. The medium was routinely aspirated and replaced with fresh medium every 24–48 h. When the cultured cells achieved 70–80% confluence, the cells were transferred after short term (maximum, 1–2 min) trypsinization with 1% trypsin and subcultured twice per week. Cell viability was

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Fig. 3. Comparison of the effects of different treatments on TM4 cell viability. The experimental groups were assigned to treatment conditions that included ethanol (160 mM), Zinc (8 μ M), pretreatment and control (un-treated) groups, following 24 or 48 h exposure periods. In the pretreatment group, the cells were pretreated with Zinc (8 μ M, for 48 h), before ethanol (160 mM) exposure. Results are presented as mean ± S.E.M derived from two determinations in duplicate cultures. ^a Represents p < 0.001 compared to the ethanol group. ^c Represents p < 0.05 compared to the ethanol group. Comparison of data between 24 and 48 h exposure periods revealed no significant changes.

determined by the trypan blue staining followed by hemocytom etry and 105 cells were seeded in 5 ml fresh medium in 25 cm²
 flasks. All chemicals and reagents purchased are of molecular biol ogy grade.

125 2.2. Cell viability (MTT assay)

Cell viability was determined using the MTT (3-(4,5-126 dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide) assav 127 kit (Cayman Company, USA) according to the manufacturer's 128 protocol. In order to determine the effect of ethanol and Zinc, 129 cells were seeded to 96-well culture plate at a density of about 130 5×10^3 cells/well, and. Although the Sertoli cells have high adhe-131 sion properties, but to prevent the possible effects of unknown 132 laboratory errors a stabilization period of 24 h has been assigned. 133 After this stabilization period, cells were treated with ethanol 134 (ethanol for molecular biology, Fluka, Sigma/Aldrich, Steinheim, 135 Germany) at concentrations of 0, 20, 40, 80 and 160 mM, for 24 136 or 48 h. To determine the effect of Zinc, TM4 cells were treated 137 with 0, 8, 16, 32 and 64 µM Zinc sulfate (Sigma, USA) for 24 or 138 48 h in a final volume of $100 \,\mu$ l. At the beginning of treatments, 139 the medium was aspirated from each well and replaced with 1001 140 ethanol or Zinc containing medium, at described concentrations 141 (4 wells each and control groups of 10 wells). In control wells, the 142 medium was aspirated from each well and replaced only with a 143 free medium. After 24 h of the last treatment, a 10 μ l sample of the 144 MTT labeling reagent was added to each well, and the cells were 145 then incubated in the CO₂ incubator. After an incubation period of 146 3-4 h, 100 µl of Crystal Dissolving Solution was added to each well. 147 The absorbance and optical density (O.D.) were then measured 148 with a micro plate reader at 545-630 nm. Percent viability was 149 calculated as (O.D. of drug-treated sample/control O.D.) × 100. 150 After primary analysis, a concentration of 160 mM for ethanol 151 and 8 µM for Zinc was considered as effective concentrations. 152 Accordingly, in another set of experiments, TM4 Sertoli cells were 153 cultured and investigated in four main groups including: Ethanol 154 (160 mM), Zinc (8 µM), pretreatment and control groups in two 155 subgroups including: 24 and 48 h exposure periods. In each of 156 duplicate cultures, cells were seeded to 96-well culture plate at a 157 density of 5×103 cells/well (4 wells each group/exposure period, 158 total number of 24 wells and control group of 10 wells). Therefore 159 cells for all groups and subgroups were seeded to a single culture 160 plate, seeded at the same time, with the same density. After a sta-161 162 bilization period of 24 h, in 48 h exposure subgroups, the medium 163 was aspirated from each well and replaced with a Zinc containing



Fig. 4. Quantitative real-time PCR analysis of caspase 3 mRNA level in different treatment groups. The experimental groups were assigned to treatment conditions that included ethanol (160 mM), Zinc (8 μ M), pretreatment and control (un-treated) groups. In the pretreatment group, the cells were pretreated with Zinc (8 μ M, for 48 h), before ethanol (160 mM) exposure. Caspase 3 mRNA expression was normalized to β -actin in each sample. Vertical bars indicate the mean fold differences \pm SEM for three independent experiments in each treatment group. ^a Represents p < 0.05 compared to the control and Zinc groups.

medium for pretreatment wells and a free fresh medium for all other wells, for following 48 h. The medium was changed to the fresh related medium in the midpoint of culture. The day after that, the medium was aspirated from each well and replaced with an ethanol containing medium for wells of pretreatment and ethanol groups, with a Zinc containing medium for Zinc group and with a free medium for control group, again for following 48 h. For wells of 24 h exposure subgroups, the same protocol of 48 h exposure period was applied except that the pretreatment started one day later and extended for 48 h and ethanol and Zinc exposure was lasted only for 24 h. Therefore, pretreatment duration of Zinc for both 24 and 48 h subgroups was same and only treatment exposure time was different. MTT assay was then performed to investigate cell viability of different groups (for all wells, together simultaneously).

As the best effect of ethanol and Zinc was observed at the abovementioned concentrations, these concentrations, at 48 h exposure period, were applied for the following experiments.

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2.3. RNA extraction and reverse transcription polymerase chain
 reaction (RT-PCR)

In each set of triplicate cultures, 105 TM4 cells/well were seeded 184 in 6 well plates for 24 h, followed by a treatment according to the 185 experimental main groups: ethanol (160 mM), Zinc (8 µM), pre-186 treatment and control (one well each). Following a stabilization 187 period of 24 h, the well of pretreatment group was exposed to Zinc 188 and the other wells were exposed to free medium for following 189 48 h. The medium was changed to the fresh related medium in the 190 midpoint of culture. The day after that, the medium was aspirated 191 from each well and replaced with an ethanol containing medium 192 for wells of pretreatment and ethanol groups, with a Zinc contain-193 ing medium for Zinc group and with free medium for control group. 194 After 48 h of last treatment, the cells were trypsinized and cell pel-195 lets were used for RNA extraction with RNX-Plus (CinnaGen, Iran) 196 according to the manufacturer's instructions. Briefly, 1 ml of RNX 197 plus was added to a tube containing 10⁶ homogenized cells, and the 198 mixture was incubated at room temperature for 5 min. Chloroform 199 was added to the solution and centrifuged at 12,000 rpm for 15 min. 200 The upper aqueous phase was then transferred to a new tube and the procedure at 72 °C for 5 min. After amplification, the resulting products were separated on 2% agarose gel and stained with safe stain. The stained gel was visualized under ultraviolet light and its band width was measured by 100 bp DNA Ladder. The expected sizes of the PCR products were 250 bp (for β -actin) and 136 bp (for caspase-3).

285 rRNA→ 185 rRNA→

Denaturing agarose gel electrophoresis of total RNA, isolated from cultured mouse Sertoli cells (TM4) of different treatment groups.





RT-PCR for β -actin (left, 250 bp) and Caspase-3 (right, 136 bp) of TM4 cells.

2.4. Real time RT-PCR assay

The expression of caspase-3 gene was analyzed by real-time RT-PCR, using the mentioned specific primers and Bioneer AccuPower qPCR master Mix (South Korea) in a total volume of 25 μl according to the manufacturer's instructions. mRNA expression of Caspase-3 gene was analyzed with quantitative real-time RT-PCR using iQ5 Real-Time PCR Detection System (Bio-Rad, CA, USA). A no template control was used to test the potential contamination and primer dimer formation. The reactions were prepared in a 96-well optical plate for 10 min at 95 °C followed by 40 cycles performed with 20 s at 95 °C and 45 s at 60 °C. A melting curve analysis was conducted to confirm the specificity of the amplification reactions. Each sample was replicated three times and the threshold cycle (Ct) values were the corresponding mean. The relative expression of each mRNA was calculated using the $2^{-\Delta\Delta Ct}$ method, where Ct is the threshold cycle (analysis of relative gene expression data using real-time quantitative PCR and the 2^{-(DeltaDeltaCt} method). Relative expression levels of mRNA were normalized to β-actin and analyzed for statistical significance using one-way analysis of variance (SPSS 16). Results were expressed as mean fold differences \pm SEM for three independent experiments in each treatment group.

2.5. Statistical analysis

All data were presented as $mean \pm standard$ error for at least three separate experiments for each treatment. Statistical significance of differences between mean values was analyzed by one

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then washed in 70% ethanol and dissolved in DEPC-treated water. 204 RNA concentration and purity were determined by measuring the 205 ratio of optical density at 260 nm to that at 280 nm and 1% agarose 206 gel electrophoresis. Extracted total RNA was then used to generate 207 single-stranded cDNA with 2-step RT-PCR kit (Vivantis, Malaysia) 208 by the following reaction (according to the protocol provided by the 209 manufacturer): 10 μ g of RNA of each treatment was mixed with 1 μ l 210 of random Hexamers primer ($50 \text{ ng}/\mu l$), $1 \mu l dNTPs mix (10 mM)$ 211 and up to 10 µl of nuclease-free water. This mixture was added to 212 a cDNA synthesis mix that included 2 µl of buffer M-MuLV (10X), 213 100 unit M-MuLV reverse transcriptase and up to 10 µl of water and 214 was followed by incubation of the mixture at 42 °C for 60 min, and 215 then for 5 min at 85 °C. The synthesized cDNA was directly used as 216 template for RT-PCR by the following reaction: 5 µl of cDNA tem-217 plate was amplified by RT-PCR in 25 µl of total reaction volume 218 containing 1 μ l of each specific primer β -actin (housekeeping gene) 219 and Caspase-3 (target genes) primer sequences (takapozist, Iran). 220 Primer sequences for β-actin were 5'-TAGGCGGACTGTTACTGAGC-221 3' (sense) and 5'-GCTCCAACCAACTGCTGTC-3' (anti-sense) and 222 for Caspase-3 were 5'-GCAGCTTTGTGTGTGTGTGTGTGTC-3' (sense) and 223 5'-AGTTTCGGCTTTCCAGTCAG-3' (anti-sense). RT-PCR procedure 224 (2-steps RT-PCR, Vivantis, Malaysia) was carried out using a 225 GeneAmp 9600 PCR system (Perkin Elmer, Norwalk, CT, USA) under 226 the following conditions (briefly): initial denaturation at 94 °C for 227 2 min, followed by 30 amplification cycles, each consisting of denat-228 229 uration at 94 °C for 30 s, annealing at 60 °C for 40 s and extension at 72 °C for 50 s, with an additional extension step at the end of 230

an equal volume of isopropanol was added. The mixture was then

centrifuged at 12,000 rpm for 15 min and the resulting pellet was

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way ANOVA followed by Tukey's HSD post-hoc test using SPSS 16
 statistical analysis software (SPSS Inc., Chicago, IL). The level of significant difference was set at P < 0.05. The fold differences of gene
 expression normalized to control were presented graphically in the
 form of histograms, using Microsoft Excel computer program.

272 **3. Results**

273 3.1. Effects of ethanol on Sertoli cell viability

In order to investigate the effect of ethanol on Sertoli cell viabil-274 ity and proliferation, cultured TM4 cells were exposed to ethanol 275 at final concentrations of 0 (cells exposed to an ethanol free media, 276 as control group), 20, 40, 80 and 160 mM, for 24 and 48 h and MTT 277 assay was then carried out. In both 24 and 48 h exposure periods, 278 cell viability decreased as concentration of ethanol increased in 279 comparison to control group (Fig. 1). After 48 h exposure period, cell 280 viability significantly decreased from $100 \pm 12\%$ in control group to 281 $54.6 \pm 4\%$ in cells exposed to 160 mM ethanol (P<0.05). Ethanol 282 decreased viability and inhibited proliferation of TM4 cells in a 283 concentration-dependent manner with IC50 values of 160 mM for 284 285 24 h and 230 mM for 48 h exposure. Comparison of data between 24 286 and 48 h exposure periods revealed no significant changes (Fig. 1).

287 3.2. Effects of Zinc on Sertoli cell viability

To investigate Zinc-induced changes in Sertoli cell viability and 288 proliferation, TM4 cells were incubated with Zinc at final concen-280 trations of 0 (cells exposed to Zinc free media, as control group), 200 8, 16, 32 and $64 \,\mu\text{M}$ for 24 and 48 h and then MTT assay was 291 carried out. In both 24 and 48 h exposure periods, cell viability 292 increased in comparison to control in lower concentrations of Zinc 293 (Fig. 2). The cellular viability in control group was $\%98 \pm 10$, which 294 in cells exposed to 8μ M Zinc increased significantly to $%286 \pm 53$ 295 and $\%261 \pm 41$ following 24 and 48 h exposure periods, respec-296 tively (P < 0.001). At higher concentrations of Zinc, cell viability 297 and proliferation decreased as concentration of Zinc increased, in 298 comparison to cells exposed to 8 µM Zinc, and at the highest con-299 centration of Zinc (64 µM) cellular viability showed a trend almost 300 around the control level (Fig. 2). Comparison of data between 24 301 and 48 h exposure periods revealed again no significant changes 302 303 (Fig. 2).

3.3. Effects of pretreatment with Zinc on ethanol-induced
 changes in Sertoli cell viability

Following above mentioned primary experiments and analy-306 sis, a concentration of 160 mM for ethanol and $8 \,\mu$ M for Zinc was 307 considered as effective concentrations. In order to investigate the 308 effect of pretreatment of Zinc on ethanol-induced cell viability sup-309 310 pression, in another set of experiments, cultured TM4 cells were exposed to 8 µM Zinc for 48 h before addition of ethanol and cellu-311 lar viability was then assessed and compared with groups exposed 312 to ethanol and Zinc alone. As shown in Fig. 3, exposure to ethanol 313 inhibited cell proliferation, whereas cellular viability and prolifera-314 tion increased in Zinc group and in the group received Zinc prior to 315 ethanol, compared to the control and ethanol groups. Pretreatment 316 with Zinc inhibited ethanol-induced cell proliferation suppression 317 (Fig. 3). Comparison of data between 24 and 48 h exposure periods 318 revealed again no significant changes. 319

320 3.4. mRNA expression of caspase-3 in Sertoli cells exposed to
 321 ethanol, Zinc and Zinc prior to ethanol

Quantitative real-time RT-PCR analysis of the mRNA levels of caspase-3 was performed, in order to determine the relative levels of expression of this gene, in the experimental groups. The experimental groups were assigned to treatment conditions that included ethanol (160 mM), Zinc (8 μ M), pretreatment (cells were pretreated with Zinc before ethanol exposure) and control (untreated) groups. Expression of β -actin gene was tested for internal control of the mRNA levels. Expression of caspase-3 gene was normalized to the endogenous control to acquire the relative threshold cycle (Δ Ct) and related to the Δ Ct of un-treated condition to find the relative expression level (2- Δ Δ Ct) of treated groups. Caspase-3 mRNA levels in the un-treated cells were set at 1.00.

Caspase-3 mRNA levels was significantly increased to 2.42 ± 0.53 by treatment with ethanol (P ≤ 0.03 , Fig. 4). On the contrary, caspase-3 mRNA levels were decreased to 0.41 ± 0.16 by treatment with Zinc, but these differences were not found to be significant. As shown in Fig. 4, mRNA level of caspase-3 in the pretreatment group decreased to around control level and showed approximately same gene expression level with un-treated conditions.

4. Discussion

In the present study cell viability and proliferation as well as caspase-3 gene expression have been investigated in TM4 Sertoli cells, exposed to ethanol, Zinc or Zinc prior to ethanol, for 24 or 48 h. The main findings were that cell viability and proliferation decreased in cells exposed to ethanol while increased in cells exposed to Zinc, in a concentration-dependent manner. Pretreatment with Zinc prior to ethanol exposure decreased ethanol-induced anti-proliferative effects. There were not significant differences in Sertoli cell viability between 24 and 48 h exposure periods in experimental groups. Moreover, caspase-3 mRNA levels increased in ethanol-treated cells while decreased in Zinc-treated cells, compared to untreated group. Pretreatment with Zinc recovered ethanol-induced elevation of caspase-3 mRNA expression levels to approximately control levels. These findings suggested that ethanol shows cytotoxic and apoptotic effects in TM4 Sertoli cells, whereas Zinc shows anti-apoptotic effects and attenuates ethanol-induced cellular toxicity.

It is widely accepted that ethanol is a gonadal toxin. A growing body of evidence in humans and experimental animals indicates that ethanol abuse leads to testicular atrophy and male infertility [5,6,32,44]. It is well known that ethanol exposure disrupts the HPG axis, adversely affects the secretory function of Sertoli cells, and produces a certain degree of testicular dysfunction. Furthermore, increases in germ cell apoptosis were observed after testicular exposure to ethanol in human, laboratory animals and cell cultures [14,22,23]. Apoptosis has been indicated as an important mechanism by which ethanol caused tissue injury and enhanced toxicity of testicular germ cells [23]. In spite of intensive studies for ethanolinduced injury or apoptosis in germ cells, little is known about the effects of ethanol on isolated Sertoli cells. Since Sertoli cells have a supporting influence on spermatogenesis and do not turn over, the degradation of the Sertoli cells might be a major factor responsible for the loss of germ cells.

In the present study evaluation of cell viability via MTT assay confirmed that ethanol decreases Sertoli cell survival and exhibits an anti-proliferation effect, in a concentration- dependent manner. This finding is in agreement with the other studies which showed ethanol at concentrations comparable to those used in this study, exerts a cytotoxic effects on isolated Leydig [22], Sertoli [7,23] and germ cells [14,23]. Moreover, the ethanol concentrations used in this study at 80 and 160 mM, correspond to 360 and 720 mg/dl. In in vivo animal study in rat, the serum ethanol concentrations were reached to about 200, 300 and 500 mg/dl 3 h after subcutaneous injection of 2.5, 3 and 5 g/kg ethanol, respectively [45]. As

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previously reported, the concentrations used in this study seem 387 as clinically heavy drunken condition [22,45]. National Institute 388 on Alcohol Abuse and Alcoholism (2004) has designated a blood 389 alcohol concentration level of 80 mg/dl or higher as an indica-390 tor of heavy drinking [46]. Previous study has been reported a 391 toxic threshold for alcohol in the range of 180-200 mg/dl [45]. 392 Accordingly, maintaining blood ethanol concentrations at or above 303 200 mg/dl for four consecutive hours was the minimum condition 30/1 for triggering degenerative response. Some of the previous studies 395 have also revealed a time-dependent effect of ethanol on repro-396 ductive activity [47-50]. According to our results no significant 397 differences were found between 24 and 48 h exposure periods in 398 the experimental groups. Most of the above mentioned studies are 399 in vivo, in which hypothalamic pituitary gonadal (HPG) axis activ-400 ity is performed to regulate gonadal function. Duration dependent 401 impairment of testicular function in alcohol abusers or in experi-402 mental animals has been reported to might be due to impaired HPG 403 axis [47,48,51] which is absent in cell culture in vitro studies. 404

It is well known that caspase family concentration is critical in 405 determining the susceptibility of cells to apoptotic stimuli. Caspase-406 3, in particular, is a key player in the execution of apoptosis and has 407 408 many cellular targets that, when activated, produce the features of apoptosis. The present data have also revealed ethanol-induced up-409 regulation and increased expression of caspase-3 mRNA in Sertoli 410 cells, which has already been reported in other cells and tissues 411 exposed to ethanol [14,22,52-54]. Decreased cell viability and over-412 expression of caspase-3 mRNA in Sertoli cells exposed to ethanol 413 may be a reflection of ethanol-induced testicular injury resulting 414 in enhanced Sertoli cell apoptosis, which may be partly involved in 415 infertility associated with alcohol abuse. 416

It is well known that ethanol produces oxidative stress within 417 the testes [33,55,56]. Excessive ROS production beyond critical lev-418 els overwhelmed the antioxidant defense strategies of spermatozoa 419 in seminal plasma and resulted in increased oxidative stress. Zinc, 420 a cytoprotectant and anti-oxidant agent, coordinates intracellular 421 422 signaling pathways, stimulates cell proliferation and suppresses apoptosis [41]. The hepatoprotective effect of Zinc against ethanol 423 induced-hepatocyte injury has been well documented and seems 424 to be related to its antioxidant properties. Accordingly, it has 425 been reported that Zinc can inhibit ethanol-induced hepatocyte 426 427 apoptosis by several independent mechanisms: by an indirect antioxidative effect and probably by inhibition of caspase activa-428 tion and death receptor-mediated pathways [36,37]. Our data is the 429 first report that Zinc, inhibited ethanol induced testicular injury. 430 Ethanol exerted anti-proliferation and cytotoxic effects on Sertoli 431 cells as well as induced apoptosis, at least in part, through over-432 expression of caspase-3. However, all these parameters exhibited, 433 inhibited or recovered when the ethanol exposure was pretreated 434 with Zinc. 435

At concentrations of 8 and 16 µM, Zinc elevated cell viability and 436 437 proliferation throughout the 24 and 48 h exposure periods maximally up to 2-3 fold the control level. In addition, caspase-3 mRNA 438 expression was below the control level in cells exposed to Zinc. 439 These data suggest that Zinc not only suppressed ethanol-induced 440 anti-proliferation and apoptotic effects, but also inhibited duration-441 dependent and spontaneously-induced apoptosis, confirming the 442 previous evidences that Zinc regulates intracellular signaling path-443 ways and reduces apoptosis and changes in Zinc levels are sufficient 444 to alter susceptibility to apoptosis [38,41,57]. Irrespective to this, 445 Zinc at higher concentrations (32 and 64 µM) did not affect signif-446 icantly cellular viability of TM4 cells, suggesting that its protective 447 and anti-apoptotic effects occurs at lower concentration which is 448 in agreement with previous study showing that antioxidant agents at low concentration can inhibit or delay oxidative damage [58]. 450 451 Zinc itself at higher concentrations (100-200 µM) induces apopto-452 sis which has been reported previously in different cells and tissues

[15,59]. The role of oxidative stress and antioxidants in male fertility has been previously discussed. Accordingly, the type, dose and duration of antioxidant therapy differ substantially and should probably be adjusted to the level of oxidative stress [3]. The efficacy of Zinc supply seems to be strictly related to the dose and length of the treatment. After Zinc physiological supplementation, recovery has been observed in elderly, in cancer, in infections as well as in patients with cell disease, since it decreased oxidative stress [38,60–61]. However, long treatment or high doses of Zinc may provoke a Zinc accumulation with subsequent damage, causing toxic effects [62]. Therefore the cellular level of Zinc must be controlled within a suitable range [62,63]. Nevertheless, in the case of oxidative stress, doses of antioxidants should be higher than usual daily dose. Determination of the most preferred active doses of antioxidants like Zinc requires further research. 453

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Despite of toxic effects of ethanol and protective effects of Zinc on Sertoli cells reported in the current study, it is not possible to define with any certainty that protective effects of Zinc is generalized or is a cell type-specific phenomenon. Investigating the protective effects of Zinc on cultured primary Sertoli cells and non-Sertoli cells will be informative to show whether Zinc is specifically protects Sertoli cells, or will any cultured cell line behave similarly under these conditions.

In summary, decreased cell viability and enhanced apoptosis through over-expression of caspase-3 mRNA occurred in Sertoli cells exposed to ethanol. Pretreatment of Zinc prior to ethanol exposure attenuated ethanol-induced Sertoli cell toxicity and apoptosis through inhibition of caspase-3 mediated pathways. Since Sertoli cells are the key supportive cells for spermatogenesis and they are very receptive to modification, a better understanding of the role of Sertoli cells in ethanol-induced testicular injury would make it feasible to identify novel mechanisms of male infertility in alcoholism and lead to more effective therapeutic approaches.

Conflict of interest

The authors have no conflicts of interest to declare regarding the study described in this article and the preparation of the article.

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