Additive effect of alpha-tocopherol and ascorbic acid in combating ethanol-induced hepatic fibrosis

P Prathibha, S Rejitha, R Harikrishnan, S Syam Das, P A Abhilash, M Indira

Department of Biochemistry, University of Kerala, Kariavattom, Thiruvananthapuram 695 581, Kerala, India

Objective: To investigate the efficacy of combined administration of alpha-tocopherol (AT) and ascorbic acid (AA) in reducing ethanol-induced hepatotoxicity.

Methods: Rats were maintained for 90 days and grouped as follows: I – control rats, II – ethanol, III – alpha-tocopherol, IV – ethanol + alpha-tocopherol, V – AA, VI – ethanol + ascorbic acid, VII – alpha-tocopherol + ascorbic acid, VIII – ethanol + alpha-tocopherol + ascorbic acid. At the end of the experimental period, markers of hepatic function, oxidative stress, and the expression of markers of inflammation and fibrosis were assayed.

Results: The markers of hepatic function, lipid peroxidation products, protein carbonyls, and the expression of nuclear factor kappa B, tumor necrosis factor alpha, transforming growth factor beta 1, cytochrome P4502E1, and collagen Type I were elevated after ethanol administration. All these parameters were reduced in the ethanol group administered AT and AA in combination. The activities of antioxidant enzymes which were reduced by ethanol administration were enhanced on combined administration of AT and AA. The reduction in hepatic fibrosis was almost 20% more in AT and AA co-administered group compared with AT and AA alone treated groups.

Discussion: Combined administration of fat soluble AT and water soluble AA was beneficial against ethanol-induced hepatotoxicity. This may be due to their different subcellular localizations.

Keywords: Ethanol, CYP2E1, NF kappa B, TGF-beta1, TNF-alpha, Reactive oxygen species, Alpha-tocopherol, Ascorbic acid

Introduction

Alcoholism has been a major health concern affecting various sections of society across the globe. Alcohol abuse causes serious clinical manifestations on almost all organs of the body. The liver is the major target organ of alcohol-induced injury. The susceptibility of the liver to alcohol-induced toxicity is due to both the high concentrations of alcohol found in portal blood (versus systemic), as well as the metabolic consequences of ethanol. Alcoholic liver disease is a spectrum of disease states that includes steatosis (fatty liver), steatohepatitis, and in severe cases, fibrosis and/or cirrhosis. Metabolism of alcohol to acetaldehyde takes place mainly in hepatocytes, a step catalyzed by alcohol dehydrogenase or cytochrome P4502E1 (CYP2E1) which leads to the release of reactive oxygen species (ROS). Generation of ROS also results from the imbalance in the oxidant–antioxidant status of the cells, especially in the liver. Excessive production of ROS can damage lipids, proteins, and DNA by inhibiting its normal function and thus leads to oxidative stress and decreased antioxidant levels in many tissues. ROS can diffuse from their site of generation and cause impairment to the structural and functional integrity of the cells causing tissue damage. Hepatic fibrosis is characterized by the excessive deposition of extra-cellular matrix (ECM) component like collagen in the hepatic stellate cells (HSCs). ROS generated in hepatocytes plays a key role in the development of hepatic fibrosis by activating collagen production in HSC which further leads to damage of cellular macromolecules.

Chronic alcoholism is associated with high risk of micronutrient deficiency. Micronutrients like alpha-tocopherol (AT), ascorbic acid (AA), carotenoids, and trace elements are important factors implicated in the defense against oxidative injury, and a deficit in any one of these elements can result in functional impairment of the overall antioxidant system.
being hydrophobic, is an important component of biological membranes and decrease in its concentration causes structural and functional damage to the cells. Unlike other fat soluble vitamins, it is not accumulated in the body to toxic levels since they get eliminated via increased metabolism.6

AA, a hydrophilic antioxidant, acts as a free radical scavenger and reduces the capability of ethanol to interact with essential molecules.7 AA protects lipid peroxidation in hydrophobic compartments either by scavenging lipid peroxides or by reducing tocopherol radicals to tocopherols.8 Numerous studies have been conducted to elucidate the antioxidant potential of AT9,10 and AA.11,12 It has been reported that AT, AA, and selenium when administered in combination imparts protection to ethanol-induced duodenal mucosal injury.13 But hardly any work is known regarding the impact of combined action of AT and AA on ethanol-induced hepatotoxicity and on fibrosis.

Hence, the major objective of this study was to understand the impact of the combination of AT and AA on ethanol-induced oxidative stress, expression of NF kappa B (nuclear factor kappa B), tumor necrosis factor alpha (TNF-alpha), transforming growth factor beta1 (TGF beta1), CYP2E1 on fibrosis.

Materials and methods
Male albino rats (Sprague Dawley strain) weighing between 100 and 140 g, bred and reared in our animal house were used for the experiment. Weight-matched animals were selected. A total of 48 rats were divided into eight groups of six rats each.

Group I (CN): Control rats.

Group II (E): Ethanol (4 g/kg body weight/day).

Group III (AT): alpha-tocopherol (250 mg/kg body weight/day).

Group IV (E + AT): Ethanol (4 g/kg body weight/ day) + alpha-tocopherol (250 mg/kg body weight/day).

Group V (AA): Ascorbic acid (200 mg/kg body weight/day).

Group VI (E + AA): Ethanol (4 g/kg body weight/ day) + ascorbic acid (200 mg/kg body weight/day).

Group VII (AT + AA): alpha-tocopherol (250 mg/kg body weight/day) + ascorbic acid (200 mg/kg body weight/day).

Group VIII (E + AT + AA): Ethanol (4 g/kg body weight/day) + alpha-tocopherol (250 mg/kg body weight/day) + ascorbic acid (200 mg/kg body weight/ day).

Animals were housed in polypropylene cages. The cages were kept in a room that was maintained between 28 and 32°C. The light cycle was 12 hours of light and dark. The study protocol was approved by the Institutional Animal Ethics Committee (IAEC -KU-14/2009-2010-BC-MI (23)). The animals were handled as per laboratory animal welfare guidelines.

Rats were fed with animal feed (The composition of feed is as follows: Moisture-10%, Crude protein-22%, Crude fibre-0.3%, Crude fat-3-4%, Nitrogen free extract-60%, Color-1% and phosphorous-0.5%) (Ashirvad Pvt Ltd, India). Food and water were given ad libitum. Ethanol was purchased from M/s Merck Ltd, Mumbai, India. Ethanol was diluted with distilled water (1:1) and given orally by gastric intubation. AT was purchased from M/s Sigma Aldrich, USA and AA was purchased from M/s Merck Ltd, Mumbai, India. AT was dissolved in coconut oil and AA was dissolved in distilled water and all of them were administered orally by gastric intubation. The dose of AA (200 mg/kg body weight) was taken from previous studies conducted in our laboratory12 and the dose of AT was decided based on the dose-dependent study in which it was found that AT at a dose of 250 mg/kg body weight is effective in preventing oxidative stress. Control, AT, AA, and AT + AA groups were administered glucose solution equivalent to the calorific value of ethanol in group II. The duration of the experiment was 90 days. At the end of the experimental period, the animals were fasted overnight and killed. The liver was dissected out and cleaned with ice cold phosphate buffer saline, blotted dry, and immediately transferred to ice cold containers for various biochemical evaluations. Blood was collected in clean, dry test tubes and allowed to clot for 30 minutes at room temperature. The clear serum was removed after centrifugation at 2000 g for 10 minutes and used immediately for the assay of various parameters.

Biochemical analysis
The activity of alanine aminotransferase (ALT) and aspartate aminotransferase (AST) was analyzed by the method of Reitman and Frankel.14 Gamma-glutamyl transferase (GGT) was analyzed by the method of Szasz.15 Malondialdehyde (MDA) was estimated by the method of Hiroshi Ohkawa.16 Hydroperoxides (HP) were estimated by the method of Mair and Hall.17 Conjugated dienes (CD) were estimated by the method of Recknagel and Ghoshal.18 Protein carboxyls were estimated by the method of Abraham and Hall.17 Total collagen was estimated by the procedure of Chandrakasan et al.20 Hydroxyproline was estimated by the procedure of Woessner21 and tissue protein was estimated by the method of Lowry et al.22 Superoxide dismutase (SOD) was assayed by the method of Kakkar et al.23 Catalase was assayed by the method of Maehly and Chance.24 The activity of glutathione reductase (GR) was determined by the method of David and Richard25 and the activity of glutathione peroxidase (GPx) was determined by the method of Lawrence and Burk.26 as modified by Agergaard and Jensen.27 The reduced glutathione

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Table 1 Primer sequences used for RT-PCR analysis

<table>
<thead>
<tr>
<th>Genes</th>
<th>Primer sequences</th>
<th>Melting temperature in (°C)</th>
</tr>
</thead>
<tbody>
<tr>
<td>GAPDH (Glyceraldehyde-3 phosphate dehydrogenase)</td>
<td>Forward: 5′-TGA CAA CCT CAC GTG TGT CA-3′</td>
<td>63.7</td>
</tr>
<tr>
<td></td>
<td>Reverse: 5′-GGC ATG GAC TGT GGT CAT GA-3′</td>
<td>64.7</td>
</tr>
<tr>
<td>CYP2E1 (cytochrome P4502E1)</td>
<td>Forward: 5′-GCC ACC CTC ATC GTC ATA TC-3′</td>
<td>59.4</td>
</tr>
<tr>
<td></td>
<td>Reverse: 5′-GCA GCC AAT CAG AAA TGT GG-3′</td>
<td>60.0</td>
</tr>
<tr>
<td>NFκB (nuclear transcription factor kB)</td>
<td>Forward: 5′-CAC CAA AGA CCC ACC ACC TCA CC-3′</td>
<td>58.9</td>
</tr>
<tr>
<td>TNF-alpha (tumor necrosis factor alpha)</td>
<td>Forward: 5′-GCC TAG TTT GCC TGG TGG AA-3′</td>
<td>59.4</td>
</tr>
<tr>
<td></td>
<td>Reverse: 5′-TAGG CCG CAT TCA AGT CAT AGT-3′</td>
<td>59.4</td>
</tr>
<tr>
<td>Collagen type I</td>
<td>Forward: 5′-GAG CCA CGA GAT CTA TG-3′</td>
<td>61.4</td>
</tr>
<tr>
<td></td>
<td>Reverse: 5′-AAG CCT GTA TTC GCT CTT CT-3′</td>
<td>62.4</td>
</tr>
</tbody>
</table>

(GSH) was determined by the method of Patterson and Lazarow.28

Total RNA isolation
Total RNA was isolated from the liver using TRIzol Reagent (M/s Sigma Aldrich, USA) by the method described by Chomczynski and Sacchi.29

Reverse transcription-polymerase chain reaction
The isolated RNA was used for reverse transcriptase-polymerase chain reaction (RT-PCR) to study the expression of glyceraldehyde-3-phosphate dehydrogenase (GAPDH), CYP2E1, NF kappa B, TGF beta1 and collagen Type I. Total tissue RNA (2 μg) was primed with 0.05 μg oligodT and reverse transcribed by omisscript reverse transcriptase using a cDNA synthesis kit (Qiagen). The PCR was carried out using eppendorf thermocycler. Primer sequences are given in Table 1. Primer sequences for GAPDH, CYP2E1, NF kappa B, TNFalpha, TGF beta1, and collagen were taken from the previous reports.30,31 The PCR contained 10 mM Tris (pH 8.3), 50 mM KCl, 1.5 mM MgCl2, dNTP at 20 mM each and gene specific primers at 0.5 mM each and 0.025 units/μl taq polymerase. After an initial denaturation step at 94°C, 35 amplification cycles were performed. Each cycle included an initial denaturation step at 94°C for 45 seconds, annealing at 56°C for NF kappa B, 55°C for TGF beta1, 61°C for TNF-alpha, 55°C for CYP2E1, 62°C for GAPDH, and 53°C for collagen Type I. A final extension step of 5 minutes at 72°C was performed in order to complete the PCR. The amplified product was electrophoresed on 2% agarose gel containing ethidium bromide. Then the gels were subjected to densitometric scanning (Bio Rad Gel Doc, CA, USA) to determine the OD of each and then normalized against an internal control GAPDH (using quantity one imaging software).

High-pressure liquid chromatography analysis

Analysis of AT in serum
Hundred microliters of serum was deproteinized with 100 μl of ethanol and extracted with 600 μl of chloroform. The extract was shaken for 5 minutes before centrifuging at 12,000 rpm for 8 minutes. The organic layer was extracted and evaporated to dryness under a nitrogen atmosphere. HPLC analysis was done by the Shimadzu Prominence SCL-20AHT (Shimadzu, Kyoto, Japan) system and the separation of AT was done by isocratic gradient elution using a Luna 5S NH2 100A column (Phenomenex, Torrance, CA, USA). The mobile phase was HPLC water (eluent A, pH 2.5) and methanol (eluent B) in 1:1 ratio. The total flow rate was 1.0 ml per minute and the time of analysis was 15 minutes. The detector’s wavelengths were set at 268 nm. The injection volume was 20 μl and the temperature of the column was thermostated at 40°C.

Analysis of AA in serum
One part serum with four parts 6% metaphosphoric acid was mixed in a polypropylene storage vial. The vial contents were vortexed and centrifuged at 10,000 g for 15 minutes at 4°C and the supernatant was used for analysis. High-pressure liquid chromatography (HPLC) analysis was done by the Shimadzu Prominence SCL-20AHT (Shimadzu, Kyoto, Japan) system and the separation of AA was done by isocratic gradient elution using a Luna 5S NH2 100A column (Phenomenex, Torrance, CA, USA). The mobile phase was HPLC water (eluent A, pH 2.5) and methanol (eluent B) in 1:1 ratio. The total flow rate was 1.0 ml per minute and the time of analysis was 15 minutes. The detector’s wavelengths were set at 268 nm. The injection volume was 20 μl and the temperature of the column was thermostated at 40°C.

Histopathological analysis
For histopathological studies, the liver was fixed in Bouin’s fixative and sections were taken in the microtome. The sections were stained using hematoxylin and eosin. The pathological changes were examined using a sensitive light microscope.

Statistical analysis
The results were analyzed using the statistical programme – SPSS/PC+, version 11.5 (SPSS Inc., Chicago, IL, USA). A one-way analysis of variance
was employed for comparison among the six groups. Duncan’s post hoc multiple comparison tests of significant differences among groups were determined. \( P < 0.05 \) was considered to be significant.

**Results**

**Biochemical analysis**

The activities of specific liver toxicity markers like ALT, AST, and GGT (Table 2) were found to be significantly elevated in ethanol-administered group when compared with control group and this was reduced to near normal levels in AT and AA-administered group, but greater reduction was observed in ethanol groups administered in combination with both AT and AA. The activities of scavenging enzymes catalase, SOD, GPx and GR (Table 4) were found to be reduced in ethanol-administered group and this was elevated in AT and AA-administered groups, but greater elevation was observed in ethanol group administered in combination of both AT and AA.

The levels of lipid peroxidation products – HP, CD, and MDA, in the liver were elevated in the ethanol group when compared with the control group (Table 3). Administration of AT along with ethanol reduced the levels of HP by 40.71%, CD by 28.11%,

<table>
<thead>
<tr>
<th>Table 2</th>
<th>Activity of toxicity marker enzymes in the liver and serum of rats</th>
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<tbody>
<tr>
<td></td>
<td>ALT (( \mu \text{M of pyruvate} / \text{min/mg protein} ))</td>
</tr>
<tr>
<td></td>
<td>Liver</td>
</tr>
<tr>
<td>CN</td>
<td>15.12 ± 0.58*</td>
</tr>
<tr>
<td>E</td>
<td>51.31 ± 2.00*</td>
</tr>
<tr>
<td>AT</td>
<td>15.28 ± 0.59*</td>
</tr>
<tr>
<td>E + AT</td>
<td>22.99 ± 0.89*</td>
</tr>
<tr>
<td>AA</td>
<td>15.91 ± 0.61*</td>
</tr>
<tr>
<td>E + AA</td>
<td>24.19 ± 0.94*</td>
</tr>
<tr>
<td>AT + AA</td>
<td>16.35 ± 0.63*</td>
</tr>
<tr>
<td>E + AT + AA</td>
<td>19.66 ± 0.76*</td>
</tr>
</tbody>
</table>

Values are expressed as mean ± SEM of six rats in each group. Values not sharing a common superscript differ significantly at \( P < 0.05 \) within rows.

<table>
<thead>
<tr>
<th>Table 3</th>
<th>Lipid peroxidation products, protein peroxidation products, total collagen, and hydroxyproline content in the liver</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>MDA (( \mu \text{Mol/100 g wet tissue} ))</td>
</tr>
<tr>
<td>CN</td>
<td>0.46 ± 0.01*</td>
</tr>
<tr>
<td>E</td>
<td>0.96 ± 0.03*</td>
</tr>
<tr>
<td>AT</td>
<td>0.43 ± 0.01*</td>
</tr>
<tr>
<td>E + AT</td>
<td>0.58 ± 0.02*</td>
</tr>
<tr>
<td>AA</td>
<td>0.44 ± 0.01*</td>
</tr>
<tr>
<td>E + AA</td>
<td>0.58 ± 0.04*</td>
</tr>
<tr>
<td>AT + AA</td>
<td>0.47 ± 0.01*</td>
</tr>
<tr>
<td>E + AT + AA</td>
<td>0.51 ± 0.02*</td>
</tr>
</tbody>
</table>

Values are expressed as mean ± SEM of six rats in each group. Values not sharing a common superscript differ significantly at \( P < 0.05 \) within rows.

<table>
<thead>
<tr>
<th>Table 4</th>
<th>Activities of scavenging enzymes Catalase, SOD, GPx, GR, and GSH in the liver of rats</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Catalase (U*/mg tissue)</td>
</tr>
<tr>
<td>CN</td>
<td>63.54 ± 2.48*</td>
</tr>
<tr>
<td>E</td>
<td>41.76 ± 1.63*</td>
</tr>
<tr>
<td>AT</td>
<td>63.81 ± 2.49*</td>
</tr>
<tr>
<td>E + AT</td>
<td>54.82 ± 2.14*</td>
</tr>
<tr>
<td>AA</td>
<td>64.49 ± 2.52*</td>
</tr>
<tr>
<td>E + AA</td>
<td>55.76 ± 2.17*</td>
</tr>
<tr>
<td>AT + AA</td>
<td>64.57 ± 2.52*</td>
</tr>
<tr>
<td>E + AT + AA</td>
<td>60.03 ± 2.34*</td>
</tr>
</tbody>
</table>

*Velocity constants.

**Enzyme concentration required to inhibit chromogen produced by 50% in 1 minute.

Values are expressed as mean ± SEM of six rats in each group. Values not sharing a common superscript differ significantly at \( P < 0.05 \) within rows.
Expression of NF-kappa B was analyzed in the cytoplasmic fraction of liver by agarose gel electrophoresis and the intensities of the bands were compared with that of the intensities of GAPDH bands expressed in the samples. Intensities of the bands were quantified using Bio-Rad gel doc and plotted. The results presented are average of quadruplicate experiments ± SEM statistically significant at \( P < 0.05 \).

Expression of TNF-alpha was analyzed in the cytoplasmic fraction of liver by agarose gel electrophoresis and the intensities of the bands were compared with that of the intensities of GAPDH bands expressed in the samples. Intensities of the bands were quantified using Bio-Rad gel doc and plotted. The results presented are average of quadruplicate experiments ± SEM statistically significant at \( P < 0.05 \).

Expression of CYP2E1 was analyzed in the cytoplasmic fraction of liver by agarose gel electrophoresis and the intensities of the bands were compared with that of the intensities of GAPDH bands expressed in the samples. Intensities of the bands were quantified using Bio-Rad gel doc and plotted. The results presented are average of quadruplicate experiments ± SEM statistically significant at \( P < 0.05 \).

Expression of TGF beta1 was analyzed in the cytoplasmic fraction of liver by agarose gel electrophoresis and the intensities of the bands were compared with that of the intensities of GAPDH bands expressed in the samples. Intensities of the bands were quantified using Bio-Rad gel doc and plotted. The results presented are average of quadruplicate experiments ± SEM statistically significant at \( P < 0.05 \).

Expression of collagen type I was analyzed in the cytoplasmic fraction of liver by agarose gel electrophoresis and the intensities of the bands were compared with that of the intensities of GAPDH bands expressed in the samples. Intensities of the bands were quantified using Bio-Rad gel doc and plotted. The results presented are average of quadruplicate experiments ± SEM statistically significant at \( P < 0.05 \).
and MDA by 39.58%, and the administration of AA along with ethanol reduced the levels of HP by 34.87%, CD by 27.27%, and MDA by 39.58% when compared with ethanol group. But greater reduction was observed in the group administered a combination of both AT and AA, whereas the level of HP was reduced by 48.76%, CD by 39.68%, and MDA by 46.87% when compared with ethanol-administered rats.

The levels of protein carbonyls (Table 3) in the liver were elevated in the ethanol-administered group and this was reduced in all AT and AA administered groups with the best reduction being shown by the ethanol group administered AT and AA in combination.

The content of collagen and hydroxyproline (Table 3) in the liver was elevated in the ethanol-treated animals. This was reduced in all antioxidant-administered groups in comparison with ethanol-administered group. AT reduced the levels by 23 and 34%, respectively. AA reduced the levels by 25 and 30%, respectively. Ethanol group supplemented with a combination of AT and AA showed 45% reduction in toxicity in the content of both collagen and hydroxyproline when compared with ethanol-administered group.

The activities of scavenging enzymes catalase, SOD, GPx and GR (Table 4) were found to be reduced in ethanol-administered group and this was elevated in AT and AA-administered groups, but greater elevation was observed in ethanol group administered in combination of both AT and AA.

The mRNA expressions of NF kappa B (Fig. 1A), TNF-alpha (Fig. 1B), CYP2E1 (Fig. 1C), TGF

Figure 1 (continued)
beta1 (Fig. 1D), and collagen Type I (Fig. 1D) were done using RT-PCR. In ethanol-treated rats, there was significant increase in the expression of PCR products compared with control rats and those treated with AT or AA alone or both in combination. The intensities of these were reduced significantly in all the antioxidant groups administered with ethanol, the reduction observed being more significant in the combination group.

HPLC analysis (Figs. 2 and 3) of AT and AA in serum showed that the concentration of AT and AA was reduced in the ethanol-treated group. This reverted to near normal levels in all antioxidant-administered groups but the more significant effect was seen in the ethanol group administered with both vitamins, the reduction observed being more significant in the combination group.

The histopathological architecture of the liver in CN (Fig. 4A), AT (Fig. 4C), AA (Fig. 4E), and AT + AA (Fig. 4G) showed normal cell structure. Ethanol-administered group (Fig. 4B) exhibited extensive hepatic cell damage as indicated by ballooning of hepatocytes, vacuolization, dilation of sinusoids and steatosis. These alterations were reduced in group supplemented with AT or AA along with ethanol (Figs. 4D and 4F), but maximum reduction in the damage were exhibited by ethanol group co-administered in combination with both AT and AA (Fig. 4H).

**Discussion**

Metabolism of ethanol to cytotoxic aldehyde and then in turn to acetate can lead to the production of ROS which perturbs the cell membrane resulting in the leakage of specific marker enzymes to circulation. In this study also, the specific markers of liver toxicity such as GGT, ALT, and AST were elevated in serum and liver tissue of ethanol-treated group indicating the extent of hepatic injury and this was reduced to near normal levels in ethanol groups administered with both AT or AA alone. This is in agreement with earlier findings. But greater reduction in
the toxicity was observed in ethanol group co-administered in combination with both AT and AA which was also in accordance with the findings of Yanardag et al.\textsuperscript{33} who showed that vitamins E, C, and selenium when supplemented in combination reduced the levels of specific liver toxicity markers.

Ethanol-induced liver injury is associated with increased oxidative stress and free radical-mediated tissue damage. Metabolism of ethanol results in elevated production of free radicals and depletion of these toxic species is associated with cell survival during ethanol intoxication. The characteristic
features of ethanol-induced liver diseases are progression of oxidative stress, alterations in the expression of liver inflammatory markers and hepatic fibrosis. Consistent with this, in this study it was observed that chronic ethanol consumption led to elevated levels of lipid peroxidation products viz. MDA, HP, CD, and protein carbonyls. This is in agreement with earlier studies wherein ethanol administration resulted in enhanced oxidative stress and lipid peroxidation.34 This was reversed in AT and AA-administered groups, but significant reduction in lipid peroxidation was observed in ethanolic rats administered a combination of both AT and AA. This is in agreement with the studies of Upansai et al.35 which showed that combined administration of AT and AA inhibits lipid peroxidation induced by lead which may be due to its antioxidant property. This observation was further supported by findings of Shalan et al.7 who proved the protective efficacy of AT, AA, and selenium against ethanol-induced toxicity.

Free radical-scavenging enzymes such as SOD and catalase are known to be the first-line cellular defense against oxidative damage and earlier reports have shown that they get reduced after ethanol abuse.36 In our study also, the activities of these enzymes were reduced after chronic ethanol intake and this was brought to near-normal levels in ethanol group administered with AT or AA. But more pronounced result was observed in ethanol group administered a combination with both AT and AA which may be due to their additive antioxidant activities. GSH is an antioxidant, preventing damage to important cellular components caused by ROS. The levels of GSH get decreased after chronic ethanol intake as reported by the earlier studies.37,38 This may be due to its utilization in scavenging the ROS mediated by ethanol metabolism which in turn leads to oxidation of GSH. The activities of GSH-dependent enzymes like GPx and GR was also found to be decreased after chronic ethanol administration which may be due to either free radical-dependent inactivation of enzyme or depletion of its co-substrates, that is GSH and NADPH.38 Consistent with these findings our result also exhibited decreased activities of GSH, GPx, and GR after chronic ethanol intake. This was brought to near-normal levels in ethanol group supplemented with either AT or AA with more significant alterations were shown by ethanol group administered a combination with both AT and AA.

Ethanol intake leads to the activation of various transcription factors. In order to evaluate the expression of transcription factors involved in ethanol-induced oxidative stress, the expression of CYP2E1, NF kappa B, TNF-alpha, and TGF beta1 was done. The CYP2E1 expression and activities are found to be elevated after chronic ethanol consumption.39 The induction of CYP2E1 in alcoholics is a means by which ethanol generates oxidative stress.40 This activates lipid peroxidation and excessive generation of hydroxyl ethyl radical resulting in decreased antioxidant levels. In this study, there was increased expression of CYP2E1 on alcohol administration which was reversed by the administration of AT and AA. But significant reduction was observed when AT and AA were administered together to ethanolic rats. Treatment with antioxidant vitamins in combination reduced the hepatotoxicity of the liver. This may be one of the factors for the observed reduced hepatotoxicity in animals co-administered AT and AA. This is in accordance with Carlett et al.41 who showed that pre-treatment with substances that reduce the activity of CYP2E1 or inactivate ROS could reduce the hepatotoxic effect of ethanol.

NF kappa B, a key factor associated with cellular stress in the liver resides in cytosol as dimer in complex with inhibitory kappa B (IκB) molecule. Activation of the dimer due to oxidative stress upon phosphorylation, dissociates IκB molecule, exposing NF kappa B which further leads to its nuclear translocation and DNA binding on nucleus. Elevated expression of NF kappa B has been reported in ethanol-induced liver inflammation.42 In accordance with this, this study also showed elevated expression of NF kappa B in ethanol-treated group. However, all ethanol groups supplemented with either AT or AA exhibited a decreased expression, but the reduction was more significant in the ethanol group administered a combination of AT and AA. NF kappa B is responsible for activation of other cytokines such as TNF-alpha with its involvement in fibrogenesis. Chronic ethanol consumption results in elevated levels of free radicals which may have activated NF kappa B leading to increased production of TNF-alpha. TNF-alpha is a pro-inflammatory cytokine and Kupffer cells are the major sites for TNF-alpha expression which is elevated during ethanol consumption.43 Consistent with this, in this study the levels of TNF-alpha were found to be elevated in the ethanol-administered group which was brought down in all antioxidant-administered groups with significant reduction being seen in the ethanol group administered a combination of AT and AA.

TGF beta1 mainly produced by macrophages acts as a growth factor in the stimulation and deposition of ECM. They also activate HSCs to proliferate and produce collagen which leads to hepatic fibrosis and liver injury.44 TGF beta1 enhances HSCs proliferation45 by increasing the expression of platelet-derived growth factor, an activating cytokine. HSC or Kupffer cells once activated, produce increased...
quantities of TGF beta1. In this study, it was found that the ethanol exposed group exhibited an increased mRNA expression of pro-inflammatory cytokine, TGF beta1. The mRNA levels of TGF beta1 were reduced to a greater extent in the ethanol group administered a combination of AT and AA when compared with other groups.

Hepatic fibrosis is characterized by deposition of ECM proteins such as collagen, mainly types I, III, and IV. HSCs are major collagen producing cells; they undergo activation during the development of fibrosis. This activation increases the capacity to synthesize more collagen by HSC. When animals are fed with ethanol, the acetaldehyde conversion may result in the development of hepatic fibrosis which in turn is accompanied by increased levels of hydroxyproline, a marker for collagen metabolism and hepatic fibrosis. These results in turn suggest that the formation of acetaldehyde adduct can cause hepatic fibrosis. George and Chandrakasan showed the activation of lipid peroxidation and MDA to be an important factor in the development of hepatic fibrosis. Our studies also demonstrated an elevation in the levels of collagen, its expression and hence a subsequent increase in the concentration of hydroxyproline. This may be due to increased oxidative stress and activation of NF kappa B which resulted in enhanced fibrosis. Reduction in the concentration of collagen and hydroxyproline and expression of collagen was shown by the ethanol group administered a combination of both AT and AA compared with AT or AA alone treated groups.

It has been reported earlier that chronic alcoholism leads to micronutrient deficiency and our results confirmed that AT and AA levels were reduced significantly in the ethanol-treated group and this was brought to near-normal levels in all the groups administered AT and AA in combination or alone. Hence, the enhanced levels of both hydrophilic antioxidant AA and hydrophobic AT may have provided the microenvironment in the hepatic cell for the reduction of oxidative stress and fibrosis.

Histopathological studies of the liver also indicated that after chronic ethanol exposure extensive hepatocellular damage was observed as evidenced by ballooning of hepatocytes, dilation of sinusoids, steatosis, and vacuolization. This was almost reversed in alcoholic rats supplemented AT or AA. But more significant effect was exhibited by ethanol group supplemented in combination of both AT and AA. This observation reinforced our biochemical findings that a combination of AT and AA provide better ameliorating effect than either AA or AT alone.

AT and AA are the major antioxidants that are thought to impart a protective effect by either reducing or preventing oxidative damage. Lipid soluble AT prevents lipid peroxidation chain reactions in cellular membranes by interfering with the propagation of lipid radicals. AA is a water-soluble antioxidant found in the cytosol and extracellular fluid that can interact directly with free radicals, thus preventing oxidative damage. Due to their different subcellular locations, a combination of AT and AA has been shown to have a better antioxidant effect than either of the two alone.

Conclusion
Our biochemical and histopathological studies confirm that chronic ethanol consumption leads to oxidative stress and hepatotoxicity. The increased production of ROS caused lipid peroxidation, protein oxidation, fibrosis and increased expression of NF kappa B, TNF-alpha, TGF beta1, CYP2E1, and collagen Type I. But co-administration of AT and AA was more effective than either AT or AA alone in combating ethanol-induced hepatotoxicity. The reduction in fibrosis was more than 20% in AT + AA group than in AT or AA alone supplemented group. The mechanism of action seems to be down-regulating the free radical production by reducing the expression of CYP2E1. This further led to reduced NF kappa B activation and decreased the expression of pro-inflammatory cytokines. Lipid soluble, AT prevents lipid peroxidation chain reactions in cellular membranes by interfering with the propagation of lipid radicals and water-soluble antioxidant AA found in the cytosol and extracellular fluid interacts directly with free radicals, thus preventing oxidative damage. Owing to their different subcellular locations, a combination of AT and AA proved to be more advantageous in combating ethanol-induced hepatotoxicity than administration of either AT or AA alone.

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