Biochemical role of probiotics in improving cell proliferation and regulating proinflammatory cytokines-mediated oxidative damage of ethanol-induced gastric mucosal injury in rats

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ABSTRACT
In the present study the gastroprotective effect and the molecular mechanisms of probiotics in a rat model of ethanol-induced gastric injury were evaluated. Thirty-five male rats were divided into five equal groups. Group 1: (Control normal group) rats received no drugs. Group 2: (Early ulcer non-protected group) rats received absolute ethanol (0.5ml/100g rat) orally on an empty stomach and sacrificed one hour later. Group 3: (Probiotics protected group) rats received probiotic (135 mg/kg body weight/day) orally for 21 days before ethanol administration then sacrificed one hour after ethanol administration. Group 4: (Late ulcer non-treated group) rats received absolute ethanol (0.5 ml/100g rat) orally on empty stomach and sacrificed after 21 days. Group 5: (Late ulcer + Probiotics treated group) rats first administered with absolute ethanol (0.5 ml/100g rat) on empty stomach at the first day of experiment then after one hour, probiotic was administered (135 mg/kg body weight/day) for 21 days then sacrificed. The results showed a significant increase in L-MDA and obvious increase in GSH concentration and CAT activity were observed in stomach of ulcerated rats. This expression was downregulated after probiotics administration. Meanwhile, a significant down-regulation of peroxisome proliferator-activated receptor gamma (PPARγ) and transforming growth factor (TGF-β1) gene were detected in ethanol-induced rat gastric injury. This expression was upregulated after probiotics administration. Conclusively, probiotics improving gastric cell proliferation and protect gastric mucosa against injury-induced in rats via anti-inflammatory and anti-oxidative mechanisms.

1. INTRODUCTION

Gastric mucosal injury occurs when injurious and destructive factors overcome an intact mucosal defense or when the mucosal defense is impaired (Laine et al., 2008). The destructive factors such as stomach Hydrochloric Acid (HCl), ethanol, free oxygen radicals, Helicobacter pylori and Non-Steroidal Anti-inflammatory Drugs (NSAIDs) that accelerate the gastric mucosal injury leading to gastric ulceration (Al-Wajeeh et al., 2017). The acute phase of gastric ulceration involves composite action of inflammatory cytokines like tumor necrosis factor-alpha (TNF-α), interleukins (IL-6, IL-10 and IL-1β), over expression of p65 subunit of nuclear factor-kappa B (NF-kB), also the inflammation is further aggravated by the recruitment of inflammatory cells like neutrophils and mononuclear cells (Amirshahrkhi and Khalili, 2015). The severity and maintenance of gastric damage also depends on the generation of free radicals that are highly reactive intermediates which bring about macromolecular changes (Da Silva et al., 2013).

Ethanol is considered as an agent that induces extreme gastric ulcers as it stimulates severe instabilities in the gastric mucosa (Hiruma et al., 2009). Alcohol consumption has been commonly linked to gastric mucosal injury including gastritis, gastric ulcer and even gastric carcinoma (Franke et al., 2005). The mechanisms underlying ethanol-induced gastric ulcer have not been fully defined. Yet, mounting evidence has indicated that proinflammatory cytokines, oxidative stress and apoptosis play essential roles in its pathogenesis (Al Batran et al., 2013). Ethanol induces its gastrointestinal toxicity through several mechanisms such as stimulation of acid secretions (Laloo et al., 2013), proinflammatory cytokines, oxidative stress (Mei et al., 2012), invasion of activated neutrophils and apoptosis as well as exhaustion of mucosal cytoprotective moieties, including Nitric Oxide (NO) and prostaglandin E2 (PGE2) (Antonisamy et al., 2014). Activation of neutrophils is associated with an upregulated inflammatory response with increased gastric expression of (NF-kB) which controls the generation of proinflammatory cytokines including tumor necrosis factor-alpha (TNF-α). These events enlarge the
inflammatory cascade via triggering the release of other proinflammatory mediators and enhancing further recruitment of macrophages and neutrophils, thereby exacerbating the gastric insult (Sangiovanni et al., 2013). Meanwhile, oxidative stress has been implicated in the development of ethanol-induced gastric injury where an arsenal of reactive oxygen species (ROS) generated by activated leukocytes triggers mucosal damage via lipid peroxidation and via depletion of the antioxidant defenses such as reduced glutathione (GSH), glutathione peroxidase (GPs) and the total antioxidant capacity (TAC) (Liu et al., 2012). In the clinical setting, the current approach for the management of gastric ulcers focuses on the use of proton pump inhibitors and H2 receptor antagonists, administration of these drugs has been associated with several adverse effects such as nausea, constipation, gynecomastia and impotence that limit their use (Halabi et al., 2014).

Numerous studies have indicated that probiotics can be used for the treatment of gastric ulcers. The idea of using probiotics arose from the study conducted by (Elliott et al., 1998). Probiotics are live microorganisms which when administered in adequate amounts are highly benefit to health (Sanders, 2003). Probiotics was found to be beneficial in liberating toxic factors, enzymes and proteins during their intestinal transit so contribute to improve the host immune defense digestion and absorption of nutrients (Buts and De, 2006). So, the ability of probiotics to modulate gut microbiota is having therapeutic potential (Nitin et al., 2016). The possible mechanisms of protection of the gastric mucosal barrier induced by probiotics are (1) Increased levels of basal mucosal prostaglandins, (2) Increased cell proliferation/apoptosis ratio, (3) Stimulation of local immune responses, (4) Release of antioxidant substances, (5) Stimulation of the expression of gastric mucins, (6) Improvement in gastrointestinal permeability, and (7) Decreases in bacterial overgrowth (Gotteland et al., 2006). Many probiotic effects are mediated through immune regulation, through balance control of proinflammatory and anti-inflammatory cytokines. From this point, probiotics can be used as innovative tools to alleviate intestinal inflammation (Isolauri et al., 2001).

Accordingly, the present study was designed to evaluate the gastroprotective effect of probiotics in a rat model of ethanol-induced gastric mucosal injury. Parameters related to gastric oxidative stress, inflammation and healing were determined in order to elucidate some of the mechanisms behind probiotics gastroprotective impact.

2. MATERIAL AND METHODS

2.1. Experimental animals:

Thirty-five white male albino rats, 6-8 weeks old and average body weight 150-170 g were used in the experimental investigation of this study. Rats were obtained from Laboratory Animals Research Center, Faculty of Veterinary Medicine, Benha University. Animals were housed in separate metal cages, fresh and clean drinking water. Rats were kept at constant environmental and nutritional conditions throughout the period of experiment. The animals were left 2 weeks for acclimatization before the beginning of the experiment.

2.2. Chemicals and drugs:

The drugs and chemicals used in the present study were:

a. Probiotics: This product was kindly supplied from NEWDYNOVET FREE TRADE, El-montazah- Alexandria and manufactured by Multipharm. Co. USA. Probiotic is a powder added to normal saline at a dose of (0.0128×10⁹) CFU per gram of rat body weight (Gupta et al., 2013) which is equivalent to 135 mg/kg b. wt. orally dissolved in normal saline (Sanchez et al., 2015).

b. Absolute ethyl Alcohol was manufactured by Sigma-Aldrich Pharmaceutical Chemicals Co.3050 Spruce Street and St Louis; Germany.

Induction of Gastric injury:

Rats were fasted for 18 hours and allowed free access of water prior to the administration of ethanol for gastric ulcer induction. The gastric ulcers were induced in all rats except the control group by orally administered with absolute ethanol at a dose level of (0.5 ml/100 g rat) (Mei et al., 2012).

2.3. Animal grouping:

Rats were randomly divided into five main equal groups, 7 rats each, placed in individual cages and classified as follow:

Group 1: (Control Normal group): Rats received no drugs, served as control non-treated for all experimental groups.

Group 2: (Early ulcer non-protected group): Rats received absolute ethanol (0.5ml/100g rat) orally on empty stomach and sacrificed one hour after ethanol administration.

Group 3: (Probiotics protected group): Rats received probiotic (135 mg/kg body weight/day) orally for 21 days before ethanol administration. One hour after administration of ethanol the animals were sacrificed.

Group 4: (Late ulcer non-treated group): Rats received absolute ethanol (0.5 ml/100g rat) on empty stomach and were left free and sacrificed 21 days later after ethanol administration.

Group 5: (Late ulcer + Probiotics treated group): Rats first administered with absolute ethanol (0.5 ml/100g rat) on empty stomach at the first day of experiment then after one hour, probiotic was administered (135 mg/kg body weight/day) for 21 days then sacrificed.

2.4. Sampling:

Gastric tissue specimen were collected from all animal groups (control and experimental groups) once at the end of 21 days.

2.4.1. Gastric tissue for biochemical analysis:

After 21 days of treatment with probiotics the rats were sacrificed by cervical decapitation. The stomach was quickly removed, and opened along the greater curvature using a scissors. The stomach was cut into small pieces, homogenized with a glass homogenizer in 9 volume of ice-cold 0.05 mM potassium phosphate buffer (pH7.4) to make 10% homogenates. The homogenates were centrifuged at 6000 rpm for 15 minutes at 4°C then the resultant supernatant were used for the determination of L-Malondialdehyde (L-MDA) concentration and Catalase (CAT) enzyme activity.

Also, 0.2 g of stomach tissues were minced into small pieces homogenized with a glass homogenizer in 0.4 ml of 25% metaphosphoric acid (MPA) (ref. No.: 253-433-4, Sigma-Aldrich, Germany), then 1.4 ml of distilled water was added, mixed and incubated for 1 hr and centrifuged for 10 min at 20°C for subsequent biochemical analysis. Briefly, gastric tissues were cut, weighed and minced into small pieces, homogenized with a glass homogenizer in 9 volume of ice-cold 0.05 mM potassium phosphate buffer (pH7.4) to make 10% homogenates. The homogenates were centrifuged at 6000 rpm for 15 minutes at 4°C then the resultant supernatant were used for the determination of L-Malondialdehyde (L-MDA) concentration and Catalase (CAT) enzyme activity.

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Effect of probiotics administration on gastric tissue L-MDA, CAT and GSH of ethanol-induced gastric injury in male rats is presented in table (1). Gastric L-MDA concentration was significantly increased in ethanol-induced rat gastric injury in both early gastric ulcer and late ulcer in comparison with normal control group. Probiotics administration to ethanol-induced gastric injury in rats showed a significant decrease in L-MDA concentration in treated and protected groups when compared to injured- non-treated groups. While, gastric CAT activity and GSH concentration were significantly decreased in ethanol-induced rat gastric injury when compared with the normal control group. Probiotics administration to ethanol- induced gastric injury in rats significantly elevated gastric antioxidant CAT activity and GSH concentration as compared to injured- non-treated groups.

The obtained results of qPCR in table (2) revealed a significant upregulated expression of NF-κB gene in ethanol-induced gastric injury in rats at early and late ulcer when compared to the normal control group. This expression was significantly downregulated following administration of probiotics in treated and protected groups when compared with injured-non-treated groups. Meanwhile, a significant downregulated expression of PPARγ and TGF-β1 gene were observed in ethanol-induced rat gastric mucosal injury at early and late gastric ulcer as compared to the normal control group. This expression was significantly upregulated following administration of probiotics in treated and protected groups vs injured non- treated groups.

Histopathological examination of stomach mucosa of control normal rats showed normal appearance of surface epithelium, mucous neck cells, parietal cells and chief cells of mucosa in gastric tissue of (Fig. 1a). Multilocally, there was necrosis and desquamation of the surface epithelium characterized by hyper-eosinophilic, shrunken cytoplasm with pyknotic nuclei in gastric tissue of early ulcer non-protected group (Fig. 1b). Meanwhile, there were small erosions in the mucosa characterized by necrosis and loss of surface epithelium in gastric tissue of Probiotics protected group (Fig. 1c). Also, there were variable sizes areas of erosions primarily affecting the surface and deep mucosa in gastric tissue of Late ulcer non-treated group (Fig. 1d). The examined stomach of (Late ulcer + Probiotics treated group) revealed normal histological appearance of mucosal cells with rare small erosions in the surface epithelium and the erosive areas were characterized by necrosis and loss of the surface epithelium (Fig. 1e).

4. DISCUSSION

Ethanol rapidly penetrates the gastric mucosa, causing membrane injury, exfoliation of cells, erosion, and ulcer formation (Franke et al., 2005). Alcohol causes severe oxidative stress in gastric tissue, which is shown as an enhancement in lipid peroxidation that occurs via an increase in the L-MDA level and a decrease in the gastric GSH concentration (Cadirci et al., 2007). The obtained results showed significant increase in L-MDA concentration with marked decrease in CAT activity and GSH concentration in gastric tissue of ethanol-induced gastric injury in rats. Similarly, Al-Wajeeh et al. (2017) showed significant decreases in the endogenous antioxidant enzymatic activity and increased lipid peroxidation in stomach tissue of ethanol-induced gastric injury in Sprague-Dawley rats.

3. RESULTS

3000 rpm then the clean supernatant was removed and used for determination of Reduced glutathione (GSH) concentration.

2.4.2. Gastric tissue for molecular analysis:

Rats gastric tissue were immediately excised after scarification and frozen in liquid nitrogen and then stored at -80°C until used for Nuclear factor kappa B (NF-κB), peroxisome proliferator-activated receptors gamma (PPARγ) and Transforming growth factor β1 (TGF-β1) gene expression analysis by qPCR.

2.4.3. Gastric tissue for histopathological examination:

Gastric tissue specimens were taken from different parts of the stomach for histopathological examination. The specimens were preserved in 10% buffered neutral formalin. The fixed tissue were rinsed in tap water, dehydrated through graded series of alcohols, cleared in xylene and embedded in paraffin wax. 5 μm thick sections were cut and stained with hematoxylin and eosin (H&E) (Bancroft and Stevens, 1996) and then the tissues were examined by light microscopy.

2.5. Analysis:

2.5.1. Biochemical analysis:

Gastric tissue (L-MDA). CAT and GSH were determined according to the methods described by Ohkawa et al. (1997), Aebi (1984) and Beutler et al. (1963), respectively.

2.5.2. Molecular analysis

Total RNA was isolated from stomach tissue of rats using RNeasy Mini Kit (Thermo Qiagen, #74104) according to the manufacturer’s protocol. Following determination of RNA concentration and purity by Quawell nanodrop Q5000 (USA), 5 mg of total RNA from each sample was reverse transcribed using Quantscript reverse transcriptase. The produced cDNA was used as a template to determine the relative expression of (NF-κB), (PPARγ) and (TGF-β) genes using Step One Plus real time PCR system (Applied Biosystem, USA) and gene specific primers. The reference gene, β-actin, was used to calculate fold change in target genes expression. The thermal cycling conditions, melting curves temperatures, and calculation of relative expression was done. For the treated groups, assessment of 2^-ΔΔCt determined the fold change in gene expression relative to the control.

Forward and reverse primers sequence for real time PCR.

<table>
<thead>
<tr>
<th>Gene</th>
<th>Forward primer</th>
<th>Reverse primer</th>
</tr>
</thead>
<tbody>
<tr>
<td>NF-κB</td>
<td>CAGACGTTCATGACGAGA</td>
<td>GCTTCTTCCATGCCGAA</td>
</tr>
<tr>
<td>PPARγ</td>
<td>GCCCTTTGCTGACATTGAG</td>
<td>GCACGAGTGCTCTGGATG</td>
</tr>
<tr>
<td>TGF-β1</td>
<td>AAGAAGACACCAGCTCTCTA</td>
<td>TGGCTAGATGGTTTGTCTCA</td>
</tr>
<tr>
<td>β-actin</td>
<td>ACCACACTTCTCATCTC</td>
<td>CCTCAACCTTGAGATG</td>
</tr>
</tbody>
</table>

2.6. Statistical Analysis

All the data were expressed as means ±S.E. The statistical significance was evaluated by One-Way analysis of variance (ANOVA) using SPSS, 18.0 software, 2011 and the individual comparisons were obtained by Duncan’s multiple range test (DMRT). Values were considered statistically significant when p<0.05.
Dawley rats. In addition to, ethanol administration initiated gastric oxidative stress and increased the levels of lipid peroxides in a process driven by neutrophil activation. It also decreased and depleted the gastric GSH, glutathione peroxidase (GPx) and total antioxidant capacity (TAC), antioxidant defenses, which scavenge free radicals and prevent their detrimental effects (Park et al., 2008). Recently, Selmi et al. (2017) showed that ethanol intoxication induced an increase of the final products of lipid peroxidation, increase of hydrogen peroxide content as well as depletion antioxidant enzyme activities such as GPx, CAT and SOD.

In the current study a significant depletion of gastric tissue L-MDA and marked increase in GSH concentration and CAT activity were observed after administration of probiotics in gastric ulcer- induced rats. Similarly, Ejtahed et al. (2012) showed that there was a significant decrease in L-MDA levels in probiotic treated group after the consumption of probiotic yoghurt. Additionally, the concentration of L-MDA occurs in diabetic rats is significantly decreased after treatment with L. acidophilus probiotics (Harisa et al., 2009). Studies found that there is a protective effect of a probiotic mixture of 13 different bacteria and α-tocopherol on ethanol-induced gastric mucosal injury. The protective effect results from its ability to reduce ethanol-induced gastric mucosal lipid peroxidation and decrease the malondialdehyde concentration, so they may be beneficial for gastric damage induced with ethanol (Senol et al., 2011). Probiotic supplementation may lead to increasing TAC and lowering MDA, which improve antioxidant status (Zamani et al., 2019). Also, Wang et al. (2009) reported that, GSH-Px activity was increased in vivo after diet supplementation with Lactobacillus fermentum and Lactobacillus plantarum. Many Lactobacillus strains which have anti-oxidative effect were found not only causing reduction in L-MDA level but also ameliorating production of antioxidant (Ejtahed et al., 2012). Probiotics can increase CAT and SOD activities, so that it has a Potential role in the management of gastric ulcer (Khoder et al., 2016).

A significant upregulation of NF-κB gene expression level was observed in stomach of ethanol-induced gastric injury in rats as compared to control normal group. Ethanol can cause gastric epithelial injury by inducing apoptosis through the TNF-α pathway and through the formation of reactive oxygen species (ROS), which causes cellular damage through oxidative stress (Liu, 2014). At the same time, the expression of TNF-α is under the control of NF-κB signal pathway. NF-κB signaling pathway is involved in controlling the gene expression of multiple factors and plays an important role in immune response, inflammation, stress response, cell apoptosis, cancer, and ontogenesis development (Chen et al., 2001).

Moreover, Li et al. (2014) reported that, ethanol markedly initiated the over expression of nuclear factor-κB in gastric mucosa of mice exposed to ethanol. Likewise, Arab et al., (2015) showed that ethanol administration stimulate gastric inflammation by increased of NF-κB p65 expression. The mRNA expression levels NF-κB was significantly increased in the gastric mucosa after ethanol administration (Song et al., 2016). NF-κB consists of p65 and p50 subunits while NFκB-p65 subunit has been commonly regarded as a marker for NF-κB activation (Verma and Kumar 2016). NF-κB is activated when its inhibitor, IκB, is phosphorylated by oxidative stress and inflammatory cytokines. Consequently NF-κB is released which then translocate toward the nucleus to initiate transcription of target dependent inflammatory genes (Lawrence, 2009). This suggestion was confirmed by Katary and Salahuddin (2017), who reported that, ethanol consumption up regulated protein expression of NF-κB p65.

<table>
<thead>
<tr>
<th>Animal groups</th>
<th>L-MDA (nmol/g tissue)</th>
<th>CAT (U/g tissue)</th>
<th>GSH (mg/g tissue)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Group I: Normal control</td>
<td>1.31 ± 0.14</td>
<td>0.933 ± 0.04</td>
<td>4.48 ± 0.32</td>
</tr>
<tr>
<td>Group II: Early ulcer non-protected</td>
<td>2.00 ± 0.32</td>
<td>0.392 ± 0.02</td>
<td>1.19 ± 0.12</td>
</tr>
<tr>
<td>Group III: Probiotics protected</td>
<td>3.64 ± 0.34</td>
<td>0.605 ± 0.09</td>
<td>3.55 ± 0.26</td>
</tr>
<tr>
<td>Group IV: Late ulcer non-treated</td>
<td>12.29 ± 0.73</td>
<td>0.370 ± 0.03</td>
<td>0.81 ± 0.05</td>
</tr>
<tr>
<td>Group V: Late ulcer + Probiotics treated</td>
<td>5.82 ± 0.28</td>
<td>0.530 ± 0.03</td>
<td>2.61 ± 0.15</td>
</tr>
</tbody>
</table>

Data are presented as (Mean ± S.E). Mean values with different superscript letters in the same column are significantly different at (P<0.05).

<table>
<thead>
<tr>
<th>Animal groups</th>
<th>Nuclear factor kappa B (NF-κB)</th>
<th>peroxisome proliferator-activated receptors gamma (PPARγ)</th>
<th>Transforming growth factor β1 (TGF-β1)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Fold change mean</td>
<td>SEM</td>
<td>Fold change mean</td>
</tr>
<tr>
<td>Group I: Normal control</td>
<td>1.00a</td>
<td>0.07</td>
<td>1.00a</td>
</tr>
<tr>
<td>Group II: Early ulcer non-protected</td>
<td>3.78b</td>
<td>0.22</td>
<td>0.04b</td>
</tr>
<tr>
<td>Group III: Probiotics protected</td>
<td>1.60c</td>
<td>0.09</td>
<td>0.26c</td>
</tr>
<tr>
<td>Group IV: Late ulcer non-treated</td>
<td>4.66c</td>
<td>0.26</td>
<td>0.02c</td>
</tr>
<tr>
<td>Group V: Late ulcer + Probiotics treated</td>
<td>2.60d</td>
<td>0.16</td>
<td>0.24d</td>
</tr>
</tbody>
</table>

Means within the same column carrying different superscript letters are significantly different at (P<0.05).

In addition, Altura et al. (2002) stated that IκB phosphorylation (and degradation) was stimulated by ethanol. In the current study the obtained results revealed a significant down-regulation of gene expression level of NF-κB following administration of probiotics in gastric injury treated groups. Probiotic bacteria can modulate systemic inflammation, cell proliferation and also apoptosis, and such properties may be useful for future immuno-modulatory and cancer prevention (Sheil et al., 2004). Pena et al. (2005) reported that L. reuteri mediated its anti-inflammatory effects via inhibition of NF-κB signaling in human intestinal epithelial cells. Also, Iyer et al., (2008) stated that, because of the central role of NF-κB and mitogen-activated protein kinase (MAPK) signaling in inflammation, cell proliferation
and apoptosis, it was estimated that *L. reuteri* mediated anti-proliferative and pro-apoptotic effects by modulating NF-κB and MAPK signaling pathways.

A significant down regulation in expression level of PPARγ and TGF-β1 gene were observed in ethanol-induced gastric injury in rats as compared to the normal control group. Activation of PPARγ has been connected with induction of cell proliferation, apoptosis, regulation of cell differentiation, and resolution of inflammation by the inhibition of NF-κB, transactivation of nitric oxide synthase-2 (NOS2) and cyclooxygenase-2 (COX2) target genes.

The obtained results are nearly similar to the data reported by Mahmoud Awny et al. (2015), who showed a significant decrease in the gastric mRNA expression of the anti-inflammatory PPAR-γ level, with a significance up regulation in the pro-inflammatory factor (NF-κB) with I/R injury in animals. In addition, Lahiri et al. (2009), who investigated the effect of PPAR-γ activation on pro-inflammatory gene expression involved in gastric ulceration and reported that, induction of ulcer caused a significant down regulation in the expression levels of PPAR-γ in ulcer induced rats. Previous studies have showed that I/R injurious effects were mediated partially through suppressing the PPAR-γ mRNA (Matsuyama et al., 2005), which is a transcription factor, acts as a regulator of anti-inflammation, antioxidant, and phagocyte-mediated cleanup processes. PPARγ was found to interact negatively with other

![Histopathological changes of rats' stomach in control and treated rats.](image-url)
transcription factors as NF-κB, which ensures its anti-inflammatory/immunomodulatory effect (Zhao et al., 2015). TGF-β1 is well known to be a multifunctional cytokine which regulates many biological processes like cell proliferation, cell differentiation, adhesion, inter cell signaling, as well as production and the degradation of extracellular matrix proteins, so that playing an essential role during wound healing and tissue repair. These results confirm that TGF-β1 expression is part of the normal healing response of gastric tissue (Polonikov et al., 2007). TGF-β has also been reported to modulate the (COX-2) expression. Takahashi et al. (1998) in vivo and in vitro studies was reported that COX-2 protein is localized to the base of gastric ulcers in rats and that COX-2 mRNA expression is regulated positively by IL-1β and TNF-α and negatively by TGF-β1. From this point, TGF-β1 plays an important role in the process of gastric ulcer healing by it interacts with prostaglandins and COX-2. In this current study, the obtained qPCR results showed a significantly up-regulated expression of PPARγ and TGF-β1 gene expression following administration of probiotics in gastric injury treated groups. Similarly, Wu et al. (2009) showed that PPARγ over-expression protects mitochondrial membrane potential and prevents apoptosis by up-regulating the expression of the anti-apoptotic Bcl-2 family proteins. Likewise, Konturek et al., (2009) reported that Probiotic bacteria Escherichia coli Nissle or heat acute murine colitis via affecting gastric mucosal expression of IL-1b, PPARγ, HSP70 and COX-2 by up regulation. Probiotic bacteria modulate gut microbial diversity and suppress colitis via targeting myeloid cell PPARγ (Bassaganya et al., 2012). Additionally, Zhang et al. (2013) reported that Lactobacillus casei significantly increased numbers of Lactobacillus and Bifidobacterium and decreased Clostridium in the intestine, with up-regulation of PPAR-γ gene expression. Besides, Kefir, koumiss, yoghurt and the commercial probiotic preparation increased PPARα and PPAR-β/δ gene expressions (Sari et al., 2013). Furthermore, Fuji et al. (2006) reported that, the administration of B. breve can up-regulate TGF-β1 signaling and may possibly be beneficial in attenuating inflammatory and allergic reactions. TGF-β acts through the induction and maintenance of regulatory T cells expressing the forkhead box P3 (FOXP3) protein or latency associated peptide (LAP). Regulatory T cells (Tregs) include different types of cells, such as CD4+CD25+ T lymphocytes and Tr1 lymphocytes. Tregs and Tr1 lymphocytes play an essential role in the control of local inflammation in animal models (Pronio et al., 2008).

5. CONCLUSIONS

These findings suggest that oral treatment with probiotics showed a significant gastroprotective effects in ethanol-induced rat gastric injury as confirmed by antioxidant and anti-inflammatory activities. Also, the gastroprotective effect of probiotics might be mediated by adjustment of inflammatory mediators and increasing antioxidants defense mechanism. Moreover, probiotics administration may have the potential as an alternative treatment for gastric ulcer because of its cytoprotective and anti-inflammatory role in improving gastric mucosal healing and cell proliferation.

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6. REFERENCES


42. Pena, J.; Rogers, A.; Ge, Z.; Ng, V.; Li, S.; Fox, J.; & Versalovic, J. 2005. Probiotic Lactobacillus spp. Diminish Helicobacter hepaticus-Induced Inflammatory Bowel Disease in Interleukin-10-Deficient Mice. Infection And Immunity, 73(2), 912-920.


45. Sanchez, E.; Nieto JC; Boullasa A; Vidal S; Sancho EJ; Rossi G Sancho-Bru P; Oms R; Mirelis B; Juárez C; Guarner C;Soriano G. 2015. VSL#3 probiotic treatment decreases bacterial translocation in rats with carbon tetrachloride-induced cirrhosis. Official Journal of the International Association for the Study of the Liver 35(3):735-45.


