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The viability of *Lactobacillus fermentum* CECT5716 is not essential to exert intestinal anti-inflammatory properties

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ABSTRACT

Probiotics have been used as alternative therapies in intestinal inflammatory disorders. Many studies have shown that different bacterial probiotic strains possess immuno-modulatory and anti-inflammatory properties. However, there is an increasing interest in the use of non-viable bacteria to reduce the risks of microbial translocation and infection. The aim of this study was to evaluate whether the viability of *L. fermentum* CECT5716 is essential to exert its intestinal anti-inflammatory effect. We compared the preventative effects of the viable and non-viable probiotic in the TNBS model of rat colitis. *In vitro* studies were also performed in Caco-2 and RAW 264.7 cells to evaluate the probiotic effects on IL-8, IL-1 β and nitrite production, and p44/42 and p38 MAP kinase protein expressions. *In vitro* results revealed a decrease in the stimulated production of pro-inflammatory mediators regardless of the viability of the probiotic. Likewise, both forms of the probiotic administered to colitic rats produced a significant reduction of IL-1 β and TNF- α levels and colonic iNOS expression. In conclusion, both live and dead *L. fermentum* CECT5716 have demonstrated to attenuate the inflammatory process and diminish the production of some of the inflammatory mediators. In fact, the viability of this probiotic did not affect its immuno-modulatory and anti-inflammatory properties.

INTRODUCTION

The human gut is a huge complex ecosystem where microbiota plays many critical roles in maintaining homeostasis. When the intestine microbiota is drastically altered, the process termed as dysbiosis takes place, and this has been associated with various intestine conditions like inflammatory bowel disease (IBD) or irritable bowel syndrome (IBS) ¹. Therefore, the modification of the gut microbiota following the administration of probiotics has emerged as a promising strategy to alleviate these diseases ^{2, 3}. According to the guidelines from FAO/WHO, probiotics are defined as ‘live microorganisms that, when ingested in adequate amounts, exert a health benefit on the host’ ^{4, 5}. This definition specifies that probiotic microorganisms must be ‘alive’, and this specification is supported by an extensive number of studies suggesting that to provide health benefits, probiotic microorganisms must be viable ⁶. Nevertheless, other studies have reported that non-viable forms of some probiotics can also exert beneficial effects on the host, most probably due to the capacity of human cells to recognize specific bacterial components or products, including immuno-stimulatory DNA, cell wall and membrane components like peptidoglycan or lipoteichoic acid, as well as intra- and extracellular polysaccharide products, thus promoting the immune responses, which are commonly mediated by the mucosa-associated lymphoid tissue (MALT) ⁷. Another important issue about probiotics is related to the knowledge of the mechanisms underlying their beneficial effects, which at present is also limited. It is generally accepted that these mechanisms are multifactorial and strain specific, which can be associated with the production of antimicrobial substances, the reduction of luminal pH, the competition for nutrients, the competitive exclusion of pathogen binding, the enhancement of the barrier function and the modulation of the immune system ^{8, 9}.

Different strains of *Lactobacillus fermentum* have been reported to have beneficial properties, particularly for gastrointestinal health. Thus, either *L. fermentum* BR11¹⁰ or *L. fermentum* CECT5716¹¹ or *L. fermentum* ACA-DC 179¹² have shown to exert intestinal anti-inflammatory effects in different experimental models of colitis in mice and rats. When considering *L. fermentum* CECT5716, several mechanisms have been proposed to be involved, including its ability to produce antimicrobial substances, such as bacteriocins,¹³ thus limiting the deleterious effect of potential pathogens. Also, this strain is able to produce antioxidant compounds, like glutathione, and short-chain fatty acids, which facilitate the growth of other lactobacilli species¹¹. Furthermore, several studies have reported its immuno-modulatory properties¹⁴.

The aim of this study was to evaluate whether the viability of *L. fermentum* CECT5716 is essential to exert its intestinal anti-inflammatory effect by comparing the preventative effects of the probiotic in both viable and non-viable conditions in the trinitrobenzenesulfonic acid (TNBS) model of rat colitis, a well-established model of intestinal inflammation with some resemblance to human IBD¹⁵. In addition, *in vitro* experiments in a human intestinal epithelial cell line were performed to better characterize the biological effects of this probiotic. The results revealed that the viability of *L. fermentum* CECT5716 is not a requisite to show beneficial effects in this experimental model of rat colitis, in which their immunomodulatory properties on cell mitogen activated protein kinase (MPAK) signalling pathways in epithelial cells may play a role.

MATERIALS AND METHODS

Reagents

All chemicals were obtained from Sigma-Aldrich Quimica (Madrid, Spain), unless otherwise stated. Glutathione reductase was provided by Boehringer Mannheim GmbH (Ingelheim, Germany).

Preparation of the probiotic

Lactobacillus fermentum CECT5716, a human breast milk derived strain¹³ was provided by Biosearch, S.A. (Granada, Spain), and was normally growth in MRS media at 37°C under anaerobic conditions using the Anaerogen system (Oxoid, Basingstoke, UK). For probiotic treatment, bacteria were prepared daily after their suspension in sterile phosphate-buffered saline (PBS) solution. Dead bacteria were obtained after heating the microorganisms at 95°C for 30 minutes in a thermoblock MBT250 (ETG Entwicklungs- und Technologie Gesellschaft mbH Ilmenau, Ilmenau, Germany). In order to determine the viability, inactivated probiotic was grown in MRS media at 10⁸ UFC/ml and the bacterial count was carried out at 24, 48 y 72 hours. The viability of inactivated probiotic did not exceed 0.0001%.

***In vitro* assays**

The human colonic epithelial colorectal adenocarcinoma cell line Caco-2 and the mouse macrophage RAW 264.7 cells (obtained from Cell Culture Unit of the University of Granada, Spain) were grown in Dulbecco's modified Eagle's medium (DMEM), supplemented with 10% Fetal Bovine Serum (FBS), l-glutamine (2 mmol/l), penicillin (100 units/ml) and streptomycin (1 mg/ml), and maintained at 37°C in a humidified, 5% CO₂ environment. For experiments, confluent cells in cell culture flasks were trypsinized and seeded on 24-well plates to 70-80% of confluence, and then they were pre-treated for 3 h either with live or dead bacteria suspended in DMEM at 10⁸ colony forming units (CFU)/ml. Cells were stimulated with different stimuli,

lipopolysaccharide (LPS) (100 ng/ml) (RAW 264.7) and interleukin (IL)-1 β (1 ng/ml) (Caco-2) during 24 h to evaluate IL-8, IL-1 β and nitrite levels, and for 30 min to evaluate p44/42 and p38 MAP kinase protein expression. Cytokine production was quantified by ELISA assay (R&D Systems, Abingdon, UK), whereas nitrite determination was performed by Griess assay as described elsewhere¹⁶. The p44/42 MAP kinase protein expression in Caco-2 cells was performed by immunoblotting¹⁷ (BD, Franklin Lakes, NJ USA).

In vivo studies

Experimental design

Female Wistar rats (180–200 g) were obtained from Janvier (St Berthevin Cedex, France), housed in makrolon cages and maintained in air-conditioned animal quarters, which were monitored according to the recommendations from the Federation of European Laboratory Animal Science Associations (FELASA), with a 12 h light–dark cycle, and fed standard rodent chow (Panlab A04, Panlab, Barcelona, Spain) and water ad libitum throughout the experiment. The monitoring of the rats revealed no infection with common murine pathogens during the period of the experiment. This study was carried out in accordance with the Directive for the Protection of Vertebrate Animals used for Experimental and other Scientific Purposes of the European Union (86/609/EEC) and approved by the Animal Research and Ethic Committee of the University of Granada (Spain).

The rats were randomly assigned to four groups (n = 10); two of them (non-colitic and control groups) received orally PBS solution (1 ml) and the other two (treated groups) received the probiotic orally, live or dead, at the concentration of 5×10^8 CFU suspended in 1 ml of PBS solution, by means of an oesophageal catheter, daily for 3 weeks. Two

weeks after starting the experiment, the rats were fasted overnight, and those from the control and the treated groups were rendered colitic as previously described.¹⁸ Briefly, they were anaesthetized with isoflurane and given 10 mg of TNBS dissolved in 0.25 ml of 50% ethanol (v/v) by means of a Teflon flexible cannula inserted 8 cm through the anus. Rats from the non-colitic group were administered intracolonicly 0.25 ml of PBS instead of TNBS. The body weight, water and food intake, as well as stool consistency, were recorded daily throughout the experiment. All rats were killed with an overdose of halothane 1 week after the induction of colitis, and the colon was obtained for the assessment of colonic damage.

Assessment of colonic damage

After the rats were sacrificed, the colon was removed aseptically and placed on an ice-cold plate, the colonic segment was cleaned of fat and mesentery, blotted on filter paper; each specimen was weighed, and its length measured under a constant load (2 g), and the weight/length ratio determined. The colon was scored for macroscopically visible damage on a 0–10 scale by two observers unaware of the treatment, according to the criteria previously reported,¹⁸ which takes into account the extent as well as the severity of colonic damage (Table 1). The colon was subsequently divided into different segments for biochemical determinations, which were frozen at -80 °C for myeloperoxidase (MPO) activity, tumour necrosis factor (TNF)- α , interferon (IFN)- γ and IL-1 β production, and inducible nitric oxide synthase (iNOS) expression, except one sample that was weighed and frozen in 1 ml of 50 g/l trichloroacetic acid for total glutathione content determinations. All biochemical measurements were completed within 1 week from the time of sample collection and were performed in duplicate. MPO activity was measured as previously described;¹⁹ the results were expressed as MPO units per gram of wet tissue and one unit of MPO activity was defined as that

degrading 1 μmol hydrogen peroxide/min at 25 °C. Total glutathione content was quantified in liver with the recycling assay described by Anderson,²⁰ and the results were expressed as nmol/g wet tissue. The colonic samples for TNF- α , IFN- γ and IL-1 β determination were immediately weighed, minced on an ice-cold plate and suspended in a tube with 10 mM sodium phosphate buffer (pH 7.4) (1:5 w/v). The tubes were placed in a shaking water bath (37 °C) for 20 min and centrifuged at 9,000 g for 30 s at 4 °C; the supernatants were frozen at -80 °C until cytokine assay. TNF- α , IFN- γ and IL-1 β were quantified by ELISA assay (R&D Systems, Abingdon, UK), and the results were expressed as ng/g wet tissue, respectively. iNOS expression was analyzed by Western blotting as previously described,²¹ and control of protein loading and transfer was conducted by detection of the β -actin levels (BD, Franklin Lakes, NJ USA).

Statistics

All results are expressed as the mean \pm SEM. Differences between means were tested for statistical significance using a one-way analysis of variance (ANOVA) and post-hoc least significance tests. Non-parametric data (score) are expressed as the median (range) and were analyzed using the Mann–Whitney U-test. Differences between proportions were analyzed with the chi-square test. All statistical analyses were carried out with the GraphPad Prism version 5.0 (La Jolla, CA, USA), with statistical significance set at $P < 0.05$.

RESULTS

Lactobacillus fermentum CECT5716, live or dead, inhibits the stimulated IL-8 production, p44/42 and p38 MAP kinase protein expression in Caco-2 cells.

The incubation of confluent Caco-2 cells with *L. fermentum* CECT5716 (10^8 CFU/mL), live or dead, for 3 h significantly increased the IL-8 production in comparison with basal conditions, without showing statistical significance when live or dead bacteria are considered (Figure 1). The incorporation of IL-1 β in the cell culture for 24 hours resulted in the stimulated release of IL-8, which was approximately 10-fold higher than that obtained after the incubation of the cells with the probiotics on basal conditions (Figure 1). When Caco-2 cells were previously exposed to the probiotic, live or dead, a significant inhibition of the IL-1 β -stimulated production of IL-8 was observed; however, in this case, the cell pre-treatment with live probiotic showed a higher reduction in this cytokine production than the dead bacteria (Figure 1).

The stimulatory effect of IL-1 β on Caco-2 cells was associated with increased phosphorylation of the MAP kinases, both p42/44 ERK and p38 (Figure 1). The pre-treatment of these cells with live or dead probiotic did not significantly modify the expression of these MAP kinases in basal conditions, but it showed inhibitory effects when they were stimulated with IL-1 β , showing a reduced phosphorylation of the MAP kinase p42/44 ERK and p38 when compared with stimulated cells without probiotic (Figure 1).

***Lactobacillus fermentum* CECT5716, live or dead, inhibits nitric oxide and IL-1 β production in stimulated RAW 264.7 cells.**

Similarly to what occurred with Caco-2 cells, the incubation of confluent RAW 264.7 cells with *Lactobacillus fermentum* CECT5716, live or dead, for 3 h resulted in a significant increase of the release of IL-1 β and nitric oxide when compared with those cells without probiotic, showing a significant higher increase in nitrite production when dead probiotic was administered in comparison with the values obtained with live

bacteria (Figure 2). LPS incorporation to the culture media of these macrophages for 24 hours resulted in an increased production of IL-1 β and nitric oxide in comparison with the levels obtained in basal conditions, being these significantly reduced when the cells were previously incubated with the probiotic, either live or dead; however, the reduction in nitrite production was achieved to a lesser extent when dead probiotic was used (Figure 2).

Intestinal anti-inflammatory effects of *Lactobacillus fermentum* CECT5716, live or dead, in the TNBS model of rat colitis.

The administration of the probiotic *Lactobacillus fermentum* CECT5716, live or dead, for 2 weeks before colitis induction did not result in any symptom of diarrhoea or affect weight evolution in comparison with untreated rats (data not shown). However, the administration of the probiotic to colitic rats resulted in an overall lower impact of the TNBS-induced damage compared to the untreated colitic control group when evaluated one week after the colonic insult. Thus, the intestinal anti-inflammatory effect was evidenced macroscopically by a significant reduction in the colonic damage score in comparison with that of control rats ($P < 0.05$) (Figure 3), since a significant decrease of the area of colonic necrosis and/or inflammation was observed in both colitic groups treated with the probiotic. However, this anti-inflammatory effect was not associated with significant differences on the colonic weight/length ratio among colitic groups, which was increased significantly as a consequence of the inflammatory process (Figure 3). The later contrasts with a previous study reported for this strain of *L. fermentum* in the same model of rat colitis, in which a significant reduction in the colonic weight/length ratio was obtained after probiotic treatment¹¹. This could be explained due to the different experimental conditions at the two time points. There are small quantitative differences in the inflammatory response and in the probiotic effect that

may be responsible for the lack of a significant effect after probiotic administration to colitic rats on this ratio in the present study.

Biochemically, the preventative beneficial effects showed by the probiotic, either live or dead, were evidenced by the reduction of the increased colonic MPO activity observed in the colitic control group (Figure 3), being this enzyme activity a marker of neutrophil infiltration ²². In addition, colonic inflammation was characterized by a decreased content in glutathione content, most probably as a consequence of the colonic oxidative stress induced by the inflammatory process, as previously reported in this model of experimental colitis ²³ (Figure 3). The treatment with *L. fermentum* CECT5716, live or dead, resulted in a significant increase in the colonic glutathione content, although no significant differences were observed between both treated groups (Figure 3).

Finally, the colonic inflammatory process induced by TNBS was also associated with increased levels of the pro-inflammatory cytokines TNF α and IL-1 β (Figure 4), as well as by a greater colonic iNOS expression (Figure 4) in comparison with non-colitic animals. The administration of the probiotic to colitic rats resulted in a significant reduction of both cytokine levels, and a lower colonic iNOS expression when compared to TNBS control animals, without showing differences between viable and dead probiotic (Figure 4).

DISCUSSION

In the last decade, several studies have supported the potential use of probiotics in human IBD, mainly in pouchitis and ulcerative colitis ²⁴, thus confirming the preclinical studies performed in experimental models of colitis ²⁵. Different mechanisms have been involved in these effects, and some of them could be attributed to the interaction of probiotics with other microorganisms, either members of the microbiota or potential

pathogens, which result in the restoration of the dysbiosis that characterizes these intestinal conditions²⁶. It is evident that this type of interaction is typically dependent on the viability of probiotics. Also, the presence of viable probiotics in the intestinal lumen is required to promote the production of short chain fatty acids (SCFA), which has been also proposed to contribute to the intestinal anti-inflammatory effect.²⁷ In addition, other mechanisms are related with the cross-talk between probiotics and host cells, clearly contributing to the well-known immunomodulatory properties ascribed to these beneficial bacteria. However, and in contrast to the direct effects exerted by the probiotics on the microbiota composition or SCFA production, their interaction with the host cells is not exclusively dependent on the bacterial viability, due to the capacity of immune cells to recognize specific bacterial components or products, thus promoting the corresponding immunological response⁷. Supporting this, different surface proteins extracted from the *Propionibacterium freudenreichii* ITG P20 surface have been recently characterized and could account for its immunomodulatory properties, being responsible for the induction of the regulatory cytokines IL-10 and IL-6.²⁸ In fact, the *in vitro* assays performed in the present study confirm this possibility, since in both intestinal epithelial cells and macrophages, the probiotic viability does not appear to be essential to affect cell activity, both in basal conditions and when the cells are stimulated with IL-1 β or LPS, respectively. Furthermore, the cellular mechanisms involved in these effects seem to be similar, because when the expression of the MAP kinases p42/44 ERK and p38 were evaluated in unstimulated or stimulated epithelial cells, both live or dead probiotic showed the same pattern of activities. It is well known that cell-wall components from Gram-negative such as lipopolysaccharides as well as host-derived cytokines such as IL-1 β and TNF α , increase IL-8 secretion from intestinal epithelial cells through the activation of mitogen activated protein kinase (MAPK)^{29, 30}.

In consequence, the ability of the probiotic to modulate MAPK activity can justify its effects on IL-8 production in intestinal epithelial cells.

These results confirmed that this strain of *L. fermentum* CECT5716 exhibits one the important features of potential probiotic candidates; that is, the capacity to modulate the immune response of the host, which clearly contributes to its intestinal anti-inflammatory effect, as stated in the *in vivo* experiments performed in the present study. However, it is interesting to note that these beneficial effects are not exclusively dependent on the probiotic viability, since both live and dead probiotic ameliorated colonic inflammation induced by the instillation of TNBS in rats. Previous studies have also revealed that the beneficial effects of probiotics in experimental colitis may be achieved by nonviable bacteria.^{31, 32} The capacity showed by dead probiotic to display intestinal anti-inflammatory properties would imply the participation of different bacterial components in these effects. Thus, it has been demonstrated that cell walls from Gram-positive bacteria, including *Enterococcus faecium* SF68 and *Lactobacillus casei* Shirota, can exert immunomodulatory properties, by affecting macrophage and B-cell activity^{33, 34}. Furthermore, it has been also reported that genomic DNA isolated from the probiotic preparation VSL#3 ameliorated the severity of colitis in DSS-, TNBS-induced and spontaneous colitis in IL-10 KO mice, being these effects mediated by TLR-signalling (mainly TLR-9)³¹. However, it is important to note that, in contrast, other studies have determined the viability requirements for the probiotic to exert intestinal anti-inflammatory properties in experimental colitis, like *Lactobacillus salivarius* ssp. *Salivarius* CECT5713³⁵.

Classically, the pathogenesis of IBD was mainly attributed to an exacerbated adaptive immune response against antigens present in the luminal environment of the intestine. Most recently, a novel hypothesis has proposed that this inflammatory disease of the gut

may result from a primary defect in intestinal innate immunity, which in turn could cause an imbalance between immune responses and tolerance to the gut microbiota that leads to chronic intestinal inflammation³⁶. Considering this, the immune modulatory properties of the probiotic *L. fermentum* CECT5716, as evidenced both *in vitro* and *in vivo* in the present study, seem to play a key role. Thus, in normal conditions, i.e. when the intestinal mucosa is not submitted to any offending agent, the probiotic can promote the strengthening of the immunological barrier by stimulating and maintaining the state of alert of the innate and adaptive immune system. In fact, the *in vitro* assays showed an increased production of the innate cytokines IL-8 and IL-1 β , as well as of NO, when either intestinal epithelial cells or macrophages were incubated with probiotic. A similar overproduction of cytokines has been reported for peripheral blood mononuclear cells exposed to well-established probiotic strains of lactobacilli, streptococci, *Leuconostoc* spp., and *Bifidobacterium breve*^{37,38}.

However, in an inflammatory environment, the probiotic is able to decrease the exacerbated immune response, as confirmed both *in vitro* and *in vivo*. *In vitro*, the probiotic inhibited the stimulated production of IL-8 (intestinal epithelial cells), IL-1 β and NO (macrophages). *In vivo*, probiotic treatment to colitic rats resulted in reduced colonic production of the pro-inflammatory cytokines TNF α and IL-1 β , as well as a down-regulation of colonic iNOS expression, when compared with the untreated corresponding controls. IL-8 is a chemokine that stimulates migration of neutrophils from intravascular to interstitial sites and can directly activate neutrophils and regulate the expression of neutrophil adhesion molecules³⁹. In consequence, the ability of this probiotic to decrease IL-8 production can contribute to inhibit leukocyte infiltration in the inflamed tissue in colitic rats, as evidenced by the lower colonic MPO activity values observed in treated colitic rats in comparison with the untreated colitic group. An

increase in neutrophils is a key feature in the pathogenesis of colitis in humans and animals ⁴⁰, which once activated by different stimuli, including IFN- γ , promote the release of reactive oxygen species products, such as hydrogen peroxide and hypochlorous acid, leading to a situation of oxidative stress and causing local tissue damage ⁴¹, and corroborated in the present study by a depletion of colonic glutathione content, which was partially prevented after probiotic treatment. In fact, reducing or limiting the influx of these pro-inflammatory cells has previously been demonstrated to attenuate inflammation ⁴², and this could be one of the mechanisms involved in the beneficial effect showed by this strain of *L. fermentum* CECT5716. Similarly, different studies have also reported the ability of other probiotics, like *Enterococcus faecalis*, to modulate and attenuate the inflammatory responses further to prevent inflammatory diseases, such as necrotizing enterocolitis in infants, through interaction with the expression and production of IL-8, as a result of MAPK signalling pathway inhibition ⁴³. Furthermore, in the DSS-colitis model, similarly to what occurs in patients with IBD, p38 levels are increased in the colonic tissue ⁴⁴, and when treated with p38 inhibitor, mucosal IL-1 β and TNF- α levels were reduced ⁴⁵, consistent with what has been found for *L. fermentum* CECT5716 treatment in the present study.

In conclusion, *L. fermentum* CECT5716 has demonstrated to attenuate the inflammatory process with amelioration of the production of some of the mediators involved in the inflammatory response, and this study has revealed that the viability of the probiotic was not required for its anti-inflammatory activity.

ACKNOWLEDGEMENTS

This work was supported by the Spanish Ministry of Economy and Competitiveness (SAF2011-29648) and the Junta de Andalucía (AGR-6826 and CTS 164) with funds from the European Union; F Algieri is a predoctoral fellow of Junta de Andalucía, N Garrido-Mesa is a postdoctoral fellow of Ramón Areces Foundation; ME Rodríguez-Cabezas is a postdoctoral fellow of CIBEREHD. The CIBEREHD is funded by the Instituto de Salud Carlos III.

FIGURE LEGENDS

Figure 1. A) Effects of *Lactobacillus fermentum* CECT5716, live or dead, on IL-8 production in Caco-2 cells, in basal conditions and stimulated with IL-1 β (1 ng/ml). Data are expressed as means \pm SEM. Bars with a different letter differ statistically ($P < 0.05$). B) Effects of *Lactobacillus fermentum* CECT5716, live or dead, on p44/42 and p38 MAP kinase protein expression in Caco-2 cells in basal conditions and stimulated with IL-1 β (1 ng/ml). The experiments were performed three times, with each individual treatment being run in triplicate.

Figure 2. Effects of *Lactobacillus fermentum* CECT5716, live or dead, on A) IL-8 production in RAW 264.7 cells, in basal conditions and stimulated with LPS (100 ng/ml); and B) nitrite accumulation in RAW 264.7 cells, in basal conditions and stimulated with LPS (100 ng/ml). Data are expressed as means \pm SEM. Bars with a different letter differ statistically ($P < 0.05$). The experiments were performed three times, with each individual treatment being run in triplicate.

Figure 3. Effects of *Lactobacillus fermentum* CECT5716, live or dead, on colonic A) weight/length ratio, B) macroscopic score, C) glutathione content and D) myeloperoxidase (MPO) activity in TNBS rat colitis one week after damage induction. Data (n=10 per group) are expressed as means \pm SEM; groups with a different letter differ statistically ($P < 0.05$).

Figure 4. Effects of *Lactobacillus fermentum* CECT5716, live or dead, on colonic A) TNF α production, B) IL-1 β production and C) iNOS expression in TNBS rat colitis one week after damage induction. Data (n=10 per group) are expressed as means \pm SEM; groups with a different letter differ statistically ($P < 0.05$).

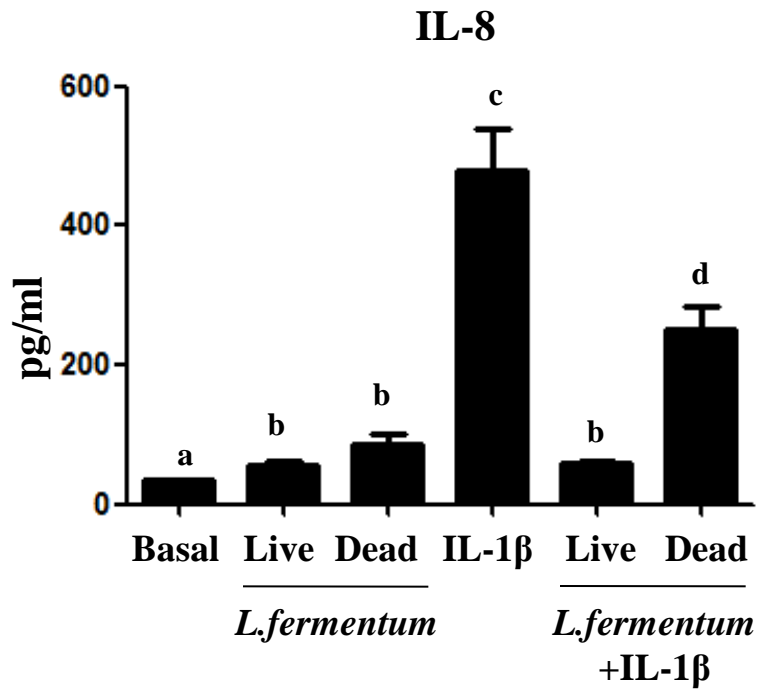
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FIGURE 1

A)



B)

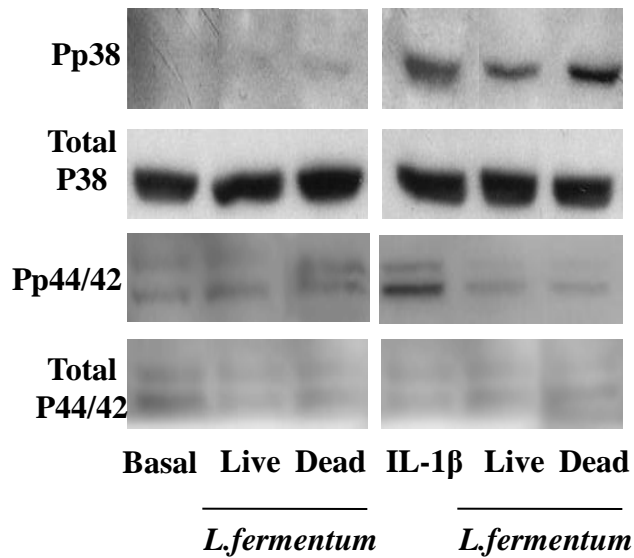
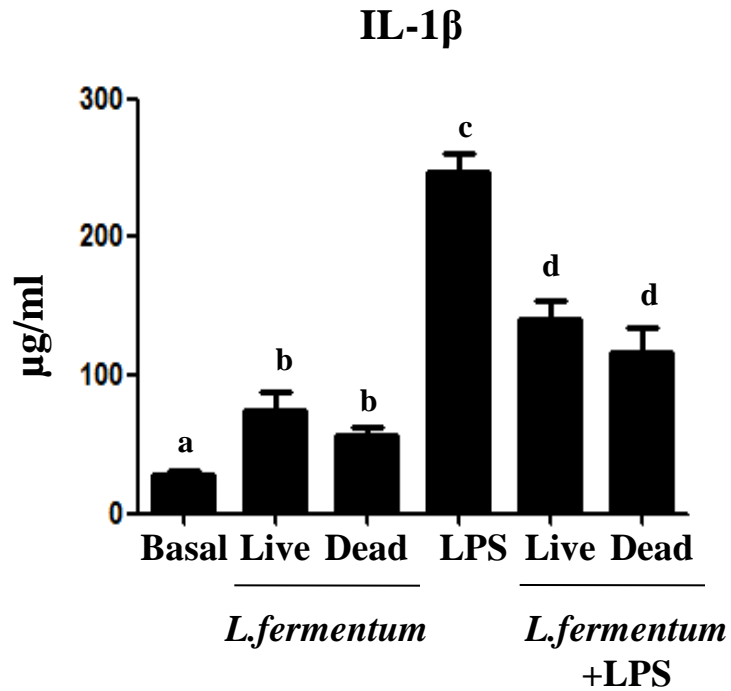


FIGURE 2

A)



B)

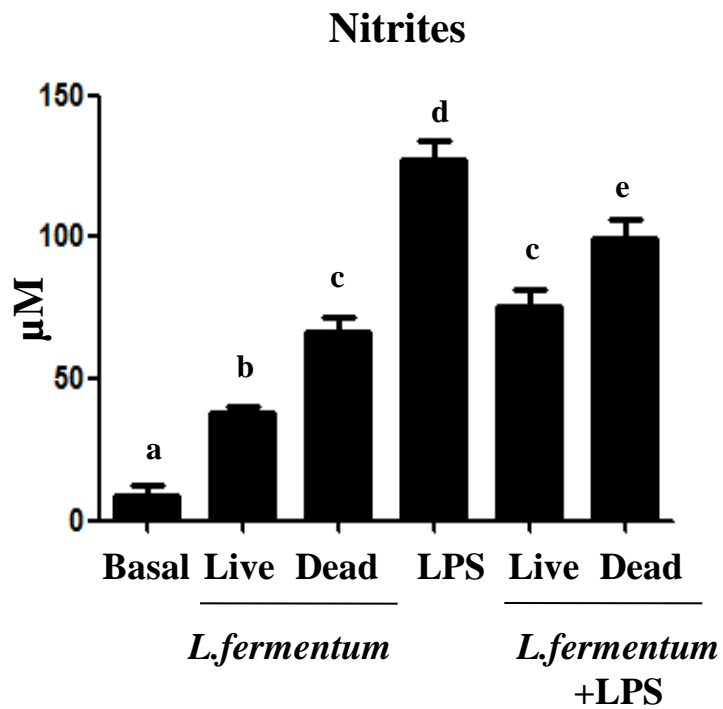


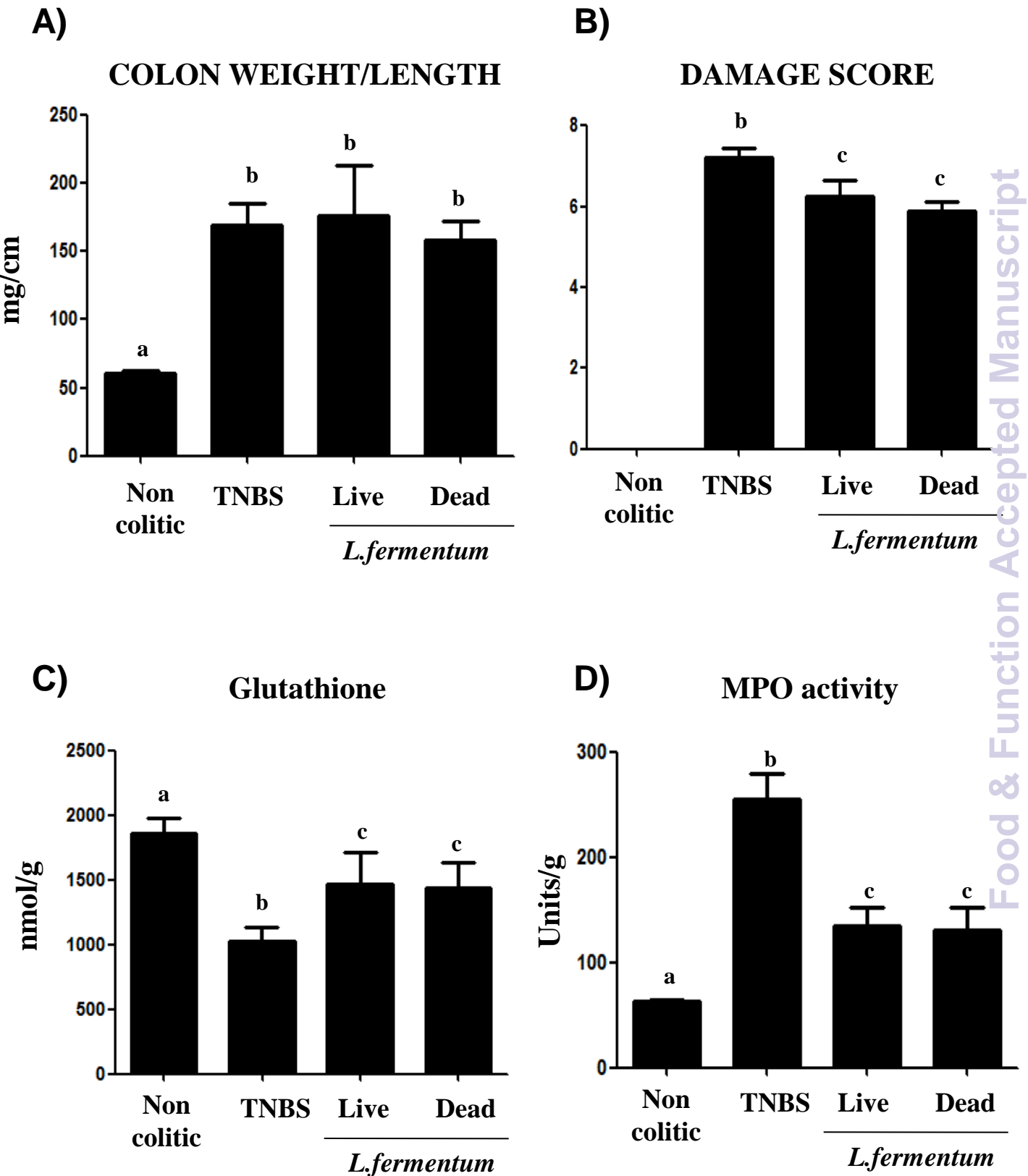
FIGURE 3

FIGURE 4

