



Effect of extracellular vesicles of *Lactobacillus rhamnosus* GG on the expression of CEA gene and protein released by colorectal cancer cells

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Received: January 2021, Accepted: November 2021

ABSTRACT

Background and Objectives: Almost all living cells secret nano-sized structures enclosed by the lipid bilayer called extracellular vesicles (EVs) into their extracellular milieu. These EVs play important roles in several physiological processes as a cargo delivery system. In probiotics, EVs are the main communication tool with the host. The present study aimed to assess the effect of EVs originated from *Lactobacillus rhamnosus* GG on the *Carcinoembryonic antigen (cea)* gene expression and protein (CEA) synthesis in the SW480 and HT-29 cell lines.

Materials and Methods: Different concentrations of *Lactobacillus rhamnosus* GG EVs were applied on the SW480 and HT-29 cell lines. The MTT assay, Real-Time PCR, and ELISA analysis methods were exploited to explore the cell viability and the expression level of the *cea* gene in comparison with the β -actin gene as the control.

Results: The two concentrations of 80 and 100 μ g/ml of *Lactobacillus rhamnosus* GG EVs considerably affected the anti-proliferation and increased the amount of both CEA mRNA and protein (p < 0.05).

Conclusion: Our findings showed that EVs of *Lactobacillus rhamnosus* GG could induce the gene expression and protein synthesis of CEA. Also, they reduced the cell proliferation of HT29 and SW480. Thus, probiotics such as EVs of *Lactobacillus rhamnosus* GG could be useful for preventing colorectal cancer.

Keywords: Extracellular vesicles; Lactobacillus rhamnosus GG; Probiotic; Carcinoemberyonic antigen family

INTRODUCTION

Infectious microorganisms, especially Gram-negative bacteria, apply the carcinoembryonic antigen-related cell adhesion molecules (CEACAMs) family members to attach to the intestine and colonize there. Bacterial agents such as *Escherichia coli, Salmonel*- *la* Typhimurium, and *Haemophilus influenzae* interact with the CEA protein expressed on the surface of intestinal epithelial cells via mannose-specific lectin (1) leading to the uptake of infectious agents and protection from diseases. The *cea* gene family from the immunoglobulin (Ig) superfamily has a remarkably varied array of highly glycosylated glycoproteins (2,

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3). The *cea* product, CEA, is a glycoprotein with a size of 180-kDa that is highly presented on the surface of normal cells such as intestinal cells and numerous tumoral (e.g., colon cancer, stomach cancer, pancreas cancer, ovarian cancer, and lung cancer) (4). Previous phylogenetic analysis in humans shows that the CEA family includes 29 genes 21 of which are protein-coding located on chromosome 19 (region 19q13.2-19q13.4) and organized in contiguous clusters (5).

Intestinal bacteria located in the human digestive tract are responsible for digesting food into absorbable nutrients (6, 7). These bacteria are also in charge of preventing the localization of intestinal epithelium by pathogens (8). Therefore, probiotics as living microorganisms are vastly discussed that can help the health of the host if delivered in adequate amounts (9) by both anti-infective and anti-carcinogenic properties. They confer these protective effects through binding to, competitively inhibiting, and degrading the mutagens, boosting the host innate and adaptive immunity, inducing the beneficial gut microbe, and improving the metabolic activity (10).

Lactobacillus and Bifidobacterium species are among the beneficial probiotics present in the gut. Lactobacillus rhamnosus GG is reported to prevent infectious diseases by producing antimicrobial agents such as bacterial metabolites, prebiotics, lantibiotics, and extracellular vesicles (EVs) (9, 11). Generally, the Lactobacilli species are Gram-positive rod bacteria that are microaerophile or facultative anaerobe (9).

Most of the living cells release nano-sized subcellular structures enveloped by lipid bilayer membranes called extracellular vesicles with a typical diameter of 30-100 nm (12). Many different cell types secrete EVs to their extracellular environment such that they can be found in most biological fluids (13). EVs have proved to be physiologically active and able to exert various biological functions regardless of the presence of bacterial cells (14). EVs can regulate the recipient cell physiology and gene expression through initiating some cell signaling pathways and/or intercellular translocation of proteins, lipid molecules, and RNA cargo (15, 16). Today, we know that both Gram-negative and Gram-positive bacteria produce EVs from their outer membrane and peptidoglycan cell wall, respectively (17). EVs are also called outer-membrane vesicles (OMVs) in Gram-negative bacteria or membrane vesicles (MVs) in Gram-positive bacteria (18). Having thick cell

walls, the exact process of generation and release of EVs by Gram-positive bacteria is a mystery yet (19). However, their effective interaction with cellular membranes and satisfactory ability to produce large aggregates at high concentrations have been shown in different studies (20, 21). In the present study, the EVs derived from *L. rhamnosus* GG are surveyed for their role in modifying the expression and production of the *cea* gene in colorectal cancer cells and their effect has been compared with that of *L. rhamnosus* GG alone.

MATERIALS AND METHODS

Bacterial culture and EVs isolation. Lactobacillus rhamnosus GG, PTCC1637, was obtained from the Iranian Research Organization for Science and Technology (IROST), Tehran, Iran. The bacteria were cultured in Man Rogosa Sharp (MRS) broth at 37°C for 24 h. For isolation of EVs, 300 ml of condition medium was collected from the cultured L. rhamnosus GG. First, the collected medium was centrifuged for 10 min at 10000 g to eliminate the cell debris and dead cells. Then, the supernatant was passed from a 0.22 mm filter (GVS filter technology, UK), and concentrated by filtration using a Centricon Plus-70 (Millipore, MA, USA). Finally, the concentrate was ultra-centrifuged at 100000 g for 60 min and the protein amount was determined at 230 nm using Nanodrop 2000 spectrophotometer (Thermo-Scientific, US) in order to evaluate the concentration of EV.

Transmission electron microscopy. In accordance with our previous work, the morphology and size of EVs were evaluated using a transmission electron microscope (TEM) (LEO906, Germany).

Cell culture. Two types of human colon cancer cell lines (SW480 and HT 29) were provided from the Pasture Institute (Tehran, Iran). Both cell lines were cultured in high glucose DMEM culture media containing 10% fetal bovine serum (FBS), 100 U/ml penicillin, and 100 mg/ml streptomycin at 37°C and 5% CO_2 . All reagents for cell culture were provided by the Iranian Inocolon Institute.

Cell viability test. MTT assay was applied to explore the role of EV in the viability of HT29 and SW480 cells. Briefly, 5000 cells were exposed to

different EV concentrations, including 5, 10, 20, 40, 60, 80, 100, 150, and 200 μ g/ml. Also, 1×10^6 CFU of *L. rhamnosus* GG and PBS were tested as negative controls. Incubation was performed for 24 hours. After removing the medium of wells, 100 μ l of fresh DMEM containing 10 μ l of 3-(4,5-dimethylth-iazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) (5 mg/ml) (Sigma-Aldrich, Germany) was added to each one and incubated at 37°C and 5% CO₂ for further 4 h. Afterward, the culture medium was substituted with 100 μ l dimethyl sulfoxide (Sigma-Aldrich, Germany). ELISA reader (Biorad, USA) was used to record the optical density of each well at 570 nm.

Gene expression analysis. Real-time PCR was used for evaluating the effect of EVs in changing the cea gene expression level. Briefly, 80, 100, 150, 200 μ g/ml, 1 × 10⁶ CFU of *L. rhamnosus* GG, and PBS were used to treat 5×10^5 W480 and HT29 cells for 24 h. Then, the RNX-Plus kit (Cinnagen, Iran) was used according to the manufacturer's protocol for mRNA purification. The concentration of mRNA was measured using the Nanodrop spectrophotometer. Also, mRNA integrity was investigated by electrophoresis on gel agarose 1%. The Bioneer kit (Takara, Japan), containing M-MLV reverse transcriptase and both random hexonucleotides and oligo dT primers, was used for synthesizing cDNA. In the following, further real-time PCR was performed using specific primers (as depicted in Table 1) and Corbett Rotor-Gene 6000 real-time PCR cycler (Qiagen Corbett, Hilden, Germany) to evaluate the relative expression of cea and β -actin genes. The PCR procedure contained 3 min initialization time at 94°C, 40 cycles for denaturation each of 30 s at 94°C, 30 s annealing time related to each gene, and final extension for 30 s at 72°C. Rest 2009 (Oiagen, USA) was used to calculate the relative expression.

ELISA analysis. For determining the impact of EVs on the secretion of CEA protein, the ELISA assay was carried out using the Carcinoembryonic Antigen Human ELISA kit (Abcam, USA). In summary, 1×10^6 of each SW480 and HT-29 cells were exposed

to four various concentrations including 80, 100, 150, and 200 μ g/ml of EVs, 1 × 10⁶ CFU of *L. rhamnosus* GG, and PBS for 24 h. Then, the culture medium of each well was collected and 100 µl from each culture medium was added to a well coated with specific human anti-CEA antibody. After incubation for 150 min, washing steps were performed according to the kit protocol, and 100 µl of 1× biotinylated CEA detection antibody was added to the wells and shaken for 1 h at room temperature. Then, 100 µl 1× HRP-streptavidin solution was added to each well and incubated for 45 min at room temperature. Next, 100 µl TMB substrate was poured into each well and after incubation for 30 min, the reaction was stopped with 50 µl stop solution and the optical density of each well was recorded at 450 nm using the ELISA reader. The quantity of each sample was calculated according to the standard curve prepared with standard solutions using different concentrations that were available within the kit.

Statistical analysis. Results were analyzed with the nonparametric Mann-Whitney test using SPSS.11 software (SPSS, Chicago, IL, USA) considering p-value < 0.05 as statistically significant. Rest 2009 software (Qiagen, USA) was used to statistically analyze the relative gene expressions.

RESULTS

Transmission electron microscopy analysis. Fig. 1 depicts the TEM image of the EVs derived from *L. rhamnosus* GG. EVs appear in the form of round shapes in a dark field with a diameter of 50-150 nm.

Cytotoxicity impacts of extracellular vesicles from *Lactobacillus rhamnosus* GG. MTT assay was implemented to evaluate the cytotoxic properties of EVs on the tested colorectal cancer cells. As shown in Figs. 2 and 3, among all studied concentrations, the values of 60, 80, 100, 150, and 200 µg/ml of EVs caused substantial inhibitory effects on both Sw480 and HT29 after 12 and 24 h incubation, re-

Table 1.	The sequence	of primer	pairs.
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Gene name	Forward sequence	Reverse sequence	Annealing temperature (°C)
сеа	AGGCCAATAACTCAGCCAGT	GGGTTTGGAGTTGTTGCTGG	59
β-actin	TCATGAAGATCCTCACCGAG	CCGACACGCTAAGACTGC	56



Fig. 1. The electron microscopy image of the extracellular vesicle from *Lactobacillus rhamenosus*. Arrows show the example of vesicles.



Fig. 2. The viability percentage of Sw480 cells in different concentration of extracellular vesicle from *Lactobacillus rhamenosus* after 12 h and 24 h. * and ** indicate *p* value less than 0.05 and 0.1, respectively.

spectively, in comparison with the negative control (p < 0.05). Whereas only the Sw480 cells were influenced by the significant cytotoxicity of EVs at concentrations of 10, 20, and 40 µg/ml (p < 0.05). The IC₅₀ calculated for Sw480 cells and HT29 cells were 158.9 and 244.66 µg/ml, respectively. According to the present results, the growth of neither of tested cell lines was inhibited by *L. rhamnosus* GG.

Gene expression analysis. Real-time PCR was performed to determine the effect of probiotic EVs on the *cea* gene expression level. In this regard, four selected cytotoxic concentrations of EVs (80, 100, 150, and 200 μ g/ml) were examined. The results obtained from the relative expression test showed a significant increase in the expression of *cea* gene after exposure to 150 and 200 μ g/ml of EVs in SW480 cells (2.89-



Fig. 3. The viability percentage of HT-29 cells in different concentration of extracellular vesicle from *Lactobacillus rhamenosus* after 12 h and 24 h. * and ** indicate p value less than 0.05 and 0.1, respectively.

and 3.1-fold change, respectively) (Fig. 4) and HT29 cells (3.97- and 3.53-fold change, respectively) (Fig. 4) in comparison with the negative control (p < 0.05). On the other hand, treating cells with *L. rhamnosus* GG had no significant effect on the *cea* gene expression.

ELISA analysis. ELISA was used to investigate the effect of EVs on the CEA secretion. CEA concentration increased after exposure of SW480 cells to 150 and 200 µg/ml of EVs (2.8 and 3.01 ng/ml, respectively) compared to the negative control (0.7 ng/ ml) (p < 0.05) (Fig. 5a). Also, a significant increase was observed in the amount of CEA in the culture medium of HT29 cells exposed to 150 and 200 µg/ml of EVs (6.43 and 7.5 ng/ml, respectively) compared to the negative control (2.34 ng/ml) (p < 0.05) (Fig. 5b). In contrast, direct treating with *L. rhamnosus* GG did not change the amount of CEA secretion from both cell lines.

DISCUSSION

Previous studies have verified that some probiotic microorganisms can trigger cell signaling against cancer cells. These probiotics especially contain *Lactobacillus* and *bifidobacterium* species including *L. casei, L. reutei, L. lactis,* and *L. rhamnosus.* The beneficial effect of probiotic bacteria is the result of their direct and/or indirect interaction with cells via secreting bioactive components such as bacteriocins and polysaccharides (22). EV formation and their active function in delivering different elements have been frequently reported in Gram-negative bacteria and showed to be involved in their infection process. In contrast, EV formation is less studied and reported Gram-positive bacteria. Despite the differences in the cell structure of the Gram-positive and Gram-negative bacteria, there are many morphological similarities in the MVs from Gram-positive bacteria and OMVs from Gram-negative bacteria (18).

Either of pathogenic and non-pathogenic microorganisms interacting with the colonic mucosa bind to mucosal cells through the interconnection between host cell membrane receptors and specific bacterial adhesive agents. The host receptors called carcinoembryonic antigen-related cell adhesion molecules or CEACAMs are expressed on the apical membrane



Fig. 4. Ratio of *cea* gene expression mRNA after treating with different concentration of extracellular vesicle from *Lactobacillus rhamenosus* (80, 100, 150, and 200 μ g/ml) in Sw480 cells and HT-29 * indicates *p* value less than 0.05.

of the polarized colonic epithelial cells. A subfamily of these adhering molecules is Glycosylphosphatidylinisotol (GPI) anchored CEACAMs that act as epithelial cellular receptors and interact with the adhesins expressed by some Gram-negative bacteria cells (1, 2). CEACAM receptors also contribute to the innate immune defence of host and are believed to exert a protective effect against microbial attack to the colon (23). Naghibalhosaini et al. have shown in their study that E. coli can significantly inhibit or decrease the CEA production in human colon-originated cell lines (LS-180 and HT29/219). They also highlighted that Caco-2 cells co-culture with E. coli leads to a considerable decrease the CEA production, however not statistically significant. They also showed a significant impact of lipid A treatment on decreasing the CEA production from cancer cells. They argued that E. coli and lipid A effect in a time- and dose-dependent manner, respectively (24). Another study by Sun et al. has reported the interaction between bacterial adhesins of Lactobacillus Plantarum strains and colonic epithelial cells through the d-mannosyl residue of CEA (25). However, its exact mechanism is not recognized yet.

To the best of the author's knowledge, the present study is the first of its kind focusing on the increasing effect of EVs from *Lactobacillus rhamnosus* GG on CEA release from human cancer cells. The current results showed that EVs derived from *L. rhamnosus* can increase CEA secretion from HT29 and SW480 cancer cell lines in comparison with the control



Fig. 5. CEA amount after treating with different concentration of extracellular vesicle from *Lactobacillus rhamenosus* (80, 100, 150, and 200 μ g/ml) in Sw480 cells (a) and HT-29 (b). * indicates *p* value less than 0.05.

group. Our findings also showed that the proliferation of HT29 and SW480 was suppressed with 80 to 200 μ g/ml concentrations of purified *L. rhamnosus* GG EVs. Furthermore, the effect of purified EVs from *L. rhamnosus* GG on the up-regulation of *cea* gene expression and its protein (CEA) production was notable in this study.

CONCLUSION

Overall, in the present study, probiotic EVs were isolated and their anti-proliferative effect on two colon cancer cell lines (HT29 and SW480) was investigated for the first time. Our findings showed that the gene expression within the recipient cells was altered by these probiotic-derived EVs. The increased expression of *cea* in cells was triggered following the exposure to 150 and 200 μ g/ml of probiotic EVs. Finally, since the field of studying EVs from Gram-positive bacteria is young, many essential questions are still unanswered and understanding the exact mechanisms and features of these vesicles require further investigations.

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