

# Impact of plant derivatives on the growth of foodborne pathogens and the functionality of probiotics

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**Abstract** Numerous studies have been published on the antimicrobial and antioxidant properties of various plant components. However, there is relatively little information on the impact of such components on the enhancement of probiotics and production of antimicrobial compounds from these probiotics. Hence, this paper focuses on the influence of plant-derived components against pathogens, enhancement of cell viability and functionality of probiotics, and potential applications of such components in food safety and human health.

**Keywords** Plant components · Probiotics · Viability · Pathogens

## Introduction

Plants are rich in several functional compounds including phytochemicals, phenols, polyphenols, essential oils, and micronutrients (Cowan 1999; Tajkarimi et al. 2010). These compounds have been reported to have antimicrobial and antioxidant activity. These natural compounds have shown potential for food preservation and quality applications (Olasupo et al. 2003; Tajkarimi et al. 2010). In addition, these natural compounds play a major role in the functionality of many biological systems including microbes.

Microbes including natural microflora (gut microbiota) in human require several mineral nutrients for their survival and growth. Thus, the aforementioned natural compounds, as a good source of nutrients, could enhance the microbial and enzyme activities (Bomba et al. 2006; Sutherland et al. 2009; Yadav et al. 2011). These natural compounds contain a number of antioxidants, which make the compounds a functional food. Several antioxidants have shown to improve a population of beneficial intestinal flora especially *Lactobacillus*. Duda-Chodak et al. (2008) have shown that antioxidants such as catechin and chlorogenic acids in the range of 100–400  $\mu\text{M}$  act as a stimulatory effect on the growth of *Lactobacillus casei* (DSM 20011). These antioxidants, major components of certain fruits, and green tea serve as an oxygen scavenger and reduce the redox potential of the media. As probiotics including lactic acid bacteria grow well in low oxygen environment, catechol or other simple phenols produced from catechin and chlorogenic acid metabolism could be related to the stimulatory effect of probiotic bacteria. Phytochemical components have been also shown to stimulate the growth of lactic acid bacteria (Bomba et al. 2006). Gut microflora including probiotics is anaerobic or facultative anaerobic. Since oxygen is toxic to these microorganisms, the presence of antioxidant helps scavenge oxygen and thus reduces the availability of oxygen from the environment and promotes growth. In addition, antioxidants increase the activities of antioxidant enzymes such as superoxide dismutase and catalase. These enzymes require minerals ( $\text{Mn}^{2+}$ ,  $\text{Ca}^{2+}$ ) as cofactors for various biological activities of microorganisms (Alberto et al. 2001). The normal intestinal flora with these enzymes is able to neutralize reactive oxygen species and could thus contribute to prevent oxidative epithelial damages (Lin and Yen 1999;

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Wijeratne et al. 2005; Spyropoulos et al. 2011; del Carmen et al. 2011). Higher enzyme production could ultimately provide treatment of inflammatory diseases or post-cancer drug treatments. These enzymes also enhance the production of organic acids such as lactic and acetic acids, which act as antimicrobial agents and suppressed the growth of pathogenic bacteria (Ibrahim and Bezkorovainy 1994; Nakashima 1997; Ibrahim and Salameh 2001; Ibrahim et al. 2003; Ibrahim 2005).

Furthermore, plants rich in micronutrients such as manganese and zinc also have shown to enhance the ability of probiotics to produce organic acids. These organic acids offer the potential as a natural antimicrobial, thus improving the safety of foods and human health (Nakashima 1997; Ibrahim et al. 2003; Bomba et al. 2006; Kang and Fung 2000). In addition, plant components indirectly induce probiotics activity to promote the production of antimicrobial compounds (Ibrahim and Bezkorovainy 1994; Calomme et al. 1995; Bomba et al. 2002; Wishon et al. 2010). To date, little information is available on the indirect influence of mineral nutrients on the production of antimicrobial agents by probiotics including bifidobacteria. Kang and Fung (2000) and Zaika and Kissinger (1984) found that manganese ions ( $Mn^{2+}$ ) present in spices are strong stimulants for starter cultures. Several microorganisms require iron for growth (O'Sullivan 2001; Ibrahim 2005; Yadav et al. 2011). Bifidobacteria tend to be better at iron scavenging than other intestinal flora. Pathogens must be able to adapt to this iron-limiting environment in order to infect. Hence, growth inhibition of these competing organisms by depriving them of iron (O'Sullivan 2001) is an excellent example of the indirect effect of plant components on microorganisms. Since the early 1990s, our research group has focused on understanding the concept and mechanism to establish how these functional compounds from plants could indirectly influence the production of antimicrobial compounds contributing to improve human health (Wishon et al. 2010; Ibrahim 2005; O'Sullivan 2001; Ibrahim and Salameh 2001; Ibrahim and Bezkorovainy 1994). However, the actual mechanism for this influence has not been demonstrated comprehensively yet.

Our literature review showed that different plant components and their derivatives exhibit antibacterial activity against different pathogens or spoilage microorganisms. Most previous studies on plant products have focused mainly on antimicrobial and antioxidant properties. However, there is relatively little information pertaining to the dual impact of plant functional components on probiotics and pathogens. Plant components rich in antioxidants have not only shown to inhibit the growth of pathogens but also proven to favor the growth of probiotics. Therefore, in this review, we aimed to provide an overview on the impact of plant derivatives on the growth of foodborne pathogens and the functionality of probiotics.

## Antimicrobial properties against pathogenic bacteria

The antimicrobial properties of natural compounds derived from plants have been recognized for centuries, but only scientifically confirmed in the last 30 years (Olasupo et al. 2003). There has been increasing interest in finding new natural antimicrobials for application in food production to prevent or inhibit microbial growth and extend shelf life (Fattouch et al. 2007; Lanciotti et al. 2004). Some of the studies on antimicrobial properties of plant products against foodborne pathogens are discussed below.

Garlic (*Allium sativum*) extract has a broad range of antimicrobial activity. Allicin, an organosulfur compound present in garlic, acts as a growth inhibitor for both Gram-positive and Gram-negative bacteria including *Escherichia*, *Salmonella*, *Streptococcus*, *Staphylococcus*, *Klebsiella*, *Proteus*, and *Helicobacter pylori* (Belguith et al. 2009; Ankri and Mirelman 1999). Avato et al. (2000) have also reported that the antibacterial activity of garlic is due to the action of allicin, diallyl thiosulfinic acid, or diallyl disulfide. A recent study showed that organosulfur compounds present in garlic have higher antimicrobial activity than those of garlic phenolic compounds. Lv et al. (2011) have reported that a bactericidal property of garlic-derived organosulfur compound was much greater against *Campylobacter jejuni* compared to phenolic compound. Garlic extract has excellent antibacterial activity against *Escherichia coli*, *Salmonella*, and *Aeromonas hydrophila*. Indu et al. (2006) reported that garlic extract showed antibacterial activity against all serogroups of *E. coli*. However, enterohemorrhagic *E. coli* (serogroup O157) and enterotoxigenic *E. coli* (serogroup O8) were highly sensitive to garlic extract. Similarly, Indu et al. (2006) have also reported high antibacterial activity against *Salmonella* serotypes and *A. hydrophila* at 75 and 100 % concentrations of garlic extracts.

Antimicrobial activity of *Capsicum annum* extract against *Salmonella* Typhimurium (ATCC 14028) in beef meat was reported by Careaga et al. (2003). The minimum inhibitory concentration of *C. annum* extract to prevent the growth of *Salmonella* Typhimurium in minced beef meat was 1.5 ml/100 g. Similarly, Dorantes et al. (2000) have reported the inhibitory effect of three varieties of *C. annum* extracts against *Listeria monocytogenes* (Scott A), *Staphylococcus aureus* (FRI-S6), *Salmonella* Typhimurium (ATCC 13311), and *Bacillus cereus* (wild type). The phenolic compound, 3-hydroxycinnamic acid (coumaric acid), is responsible for antimicrobial activity in *Capsicum* (Dorantes et al. 2000).

Chinese chive that belongs to the same family as garlic is an important ingredient in Asian cooking. Ibrahim et al. (2009) have shown that crude chive extract containing sulfur compounds can be effective against the growth of *Salmonella* and could be used in food products to prevent the growth of this pathogen. The antimicrobial effect of Chinese

chive against *E. coli* and yeast (*Pichia membranaefaciens* CCRC 20859) has been also reported (Mau et al. 2001).

Caffeine (1,3,7-trimethylxanthine) is a methylated xanthine alkaloid derivative present in plant species. It has shown significant growth inhibition against *E. coli* O157:H7 at a concentration of 0.5 % (Ibrahim et al. 2006). Scientific literature suggests that antibacterial properties in coffee have been reported to arise from caffeic acid, chlorogenic acid, and protocatechnic acid (Dogasaki et al. 2002). Similarly, polyphenols (epicatechin, catechin, caffeine, chlorogenic acid, gallic acid, theobromine, theophylline, gallic acid, epigallocatechin gallate, catechin gallate, epicatechin gallate, and theaflavin) from tea have also been found to have antimicrobial activity against *Salmonella aureus* and *L. monocytogenes* in laboratory medium (Kim et al. 2004). Kim and Fung (2004) have reported the antimicrobial activity of crude water-soluble arrowroot (*Puerariae radix*) tea against *E. coli* O157:H7, *Salmonella enterica*, *L. monocytogenes*, and *S. aureus*. The antimicrobial activity of arrowroot tea was due to the presence of catechins. Catechins present in green tea extract, epigallocatechin gallate and epigallocatechin, showed strong antimicrobial activity due to the galloyl moiety present in their structures (Shimamura et al. 2007). Polyphenols extracted from green tea extract have shown inhibitory effects on Gram-positive as well as Gram-negative bacteria (Gadang et al. 2008; Perumalla and Hettiarachchy 2011). Ndi et al. (2007) have reported antimicrobial activity of the plant *Eremophila* (Myoporaceae) against Gram-positive organisms such as streptococci and staphylococci. It has been suggested that the antibacterial properties of *Eremophila* is due to the compound serrulatanes, which is structurally related to a diterpenoid quinone antibiotic called bioflorin (Ghisalberti 1994).

Essential oils (EOs) obtained from plants have been known to possess antimicrobial activity. EOs and their constituents have been used in numerous food applications and are classified as generally recognized as safe. Several previous studies have shown that EOs from cinnamon along with geraniol, lemongrass, and clove have antimicrobial effect (Aureli et al. 1992; Burt 2004; Chao et al. 2000; Kim et al. 1995). EOs of lemongrass, cinnamon, and geraniol were found to be the most effective in inhibiting the growth of *Salmonella enteritidis*, *E. coli*, and *Listeria innocua* (Raybaudi-Massilia et al. 2006). Turgis et al. (2009) have shown that mustard EOs can be used against *E. coli* O157:H7 and *Salmonella* Typhi. Allyl isothiocyanate, a non-phenolic volatile compound found in the cruciferae family at low concentration, effectively inhibits several pathogenic microorganisms (Turgis et al. 2008). In a recent study, oregano EOs showed antimicrobial effects against *E. coli*, *S. aureus*, *Bacillus subtilis*, and *Saccharomyces*

*cerevisiae* (Lv et al. 2011). Phenols and terpenes are the main two groups of constituents responsible for antimicrobial effects in oregano EOs. The major constituents of oregano EO include carvacrol (30.17 %), p-cymene (15.20 %),  $\gamma$ -terpinen (12.44 %), and thymol (8.62 %) (Lv et al. 2011). The use of natural organic compounds as food additives could also be helpful in controlling *Cronobacter* spp. in various types of foods. In a study by Lee and Jin (2008), the order of inhibition of the natural organic compounds (EOs) against *Cronobacter* spp. was thymol > eugenol > diacetyl > cinnamic acid.

Nanasombat and Lohasupthawee (2005) have reported the degree of antibacterial property of various spices tested against *Salmonella* and other enterobacteria in the order of clove > kaffir lime peels > cumin > cardamom > coriander > nutmeg > mace > ginger > garlic > holy basil > kaffir lime leaves. The antimicrobial activity in the components of spice oils of the terpenoid family is due to the presence of terpene fraction of the oils (Davidson and Branden 1981). Different types of pepper, parsley, and dill have shown antibacterial activity against natural microflora, coliforms, yeast and molds, and *S. aureus* in Kareish cheese (Wahba et al. 2010). All chilli peppers contain phytochemicals known collectively as capsaicinoids (Antonious et al. 2009; Spiller et al. 2008). Shan et al. (2007) have suggested that the antibacterial activity of a total of 46 extracts from spices and herbs was closely associated with the presence of their phenolic constituents. The authors have reported that all the tested spices have a strong antibacterial effect against *B. cereus*, *L. monocytogenes*, *S. aureus*, *E. coli*, and *Salmonella* Anatum. Oregano and thyme EOs possess significant in vitro colicidal and colistatic properties, and this bactericidal concentration of oregano EOs irreversibly damaged *E. coli* O157:H7 cells within 1 min (Burt and Reinders 2003). The bulb extracts of *Allium cepa* and *A. sativum* exhibited activity against filamentous and non-filamentous fungi. *A. sativum* showed significant inhibition of all tested bacterial (Gram-positive and Gram-negative) and fungal pathogens (Srinivasan et al. 2001). The volatile oils of black pepper, clove, geranium, nutmeg, oregano, and thyme possessed antibacterial activity against 25 different genera of tested bacteria with various degrees of growth inhibition (Dorman and Deans 2000). Nutmeg showed good antimicrobial activity against *L. monocytogenes*, whereas its effect on *E. coli* and *Salmonella* were strain dependent (Indu et al. 2006).

### Mechanisms of antimicrobial action

The possible mode of action for different plant-derived compounds as antimicrobial agents has been reviewed extensively (Burt 2004; Lanciotti et al. 2004; Smith-Palmer et al. 2001; Sofos et al. 1998; Lopez-Malo et al. 2005;

Davidson 2001). The exact mechanism of action is not yet fully understood. However, there are number of proposed mechanisms of antimicrobial action of such compounds (Holley and Patel 2005; Lanciotti et al. 2004; Proestos et al. 2008; Lambert et al. 2001; Ultee et al. 1999; Raccach 1984).

Most of the plant species (fruits and vegetables) naturally synthesized organic acids such as acetic, citric, succinic, malic, tartaric, benzoic, and ascorbic. In addition, microorganisms including lactic acid bacteria and bifidobacteria also produce acids as a result of fermentation (Ibrahim and Bezkorovainy 1994). These organic acids inhibit the growth of both bacterial and fungal cells (Brul and Cootte 1999). The inhibition of microorganisms by acetic acids is due to the decrease of intracellular pH by releasing protons from undissociated molecules in the cytoplasm causing metabolism inhibition (Olasupo et al. 2004; Hosein et al. 2011). Some organic acids such as lactic acid have been shown to alter the permeability levels of the bacterial outer membrane. The exposure to such acidic substances increases the outer membrane permeability of Gram-negative bacteria (Alakomi et al. 2000; Helander and Mattila–Sandholm 2000; Derrickson-Tharrington et al. 2005). Gill and Holley (2006) have reported the effect of trans-cinnamaldehyde (bark extract of cinnamon) on the bacterial plasma membrane causing changes in cell membrane composition and thereby deactivating bacterial growth. Helander et al. (1998) reported that the outer membranes of *E. coli* and *Salmonella* Typhimurium disintegrated when exposed to carvacrol and thymol. Similarly, Rasooli et al. (2006) have reported thickening, disruption of the cell wall, and increased roughness and lack of cytoplasm in *L. monocytogenes* following exposure to thyme essential oil. Fisher and Phillips (2008) have also mentioned the morphological changes in the *Enterococcus* species after treatment with citrus essential oils. Carvacrol and tea tree oil have been shown to increase membrane fluidity and cause the leakage of protons and potassium ions causing disruption of membrane and inhibition of adenosine triphosphate (ATP) synthesis (Ultee et al. 1999) against *S. aureus* (Halcón and Milkus 2004). Oussalah et al. (2006) have reported that the control of cellular processes such as DNA transcription, protein synthesis, and loss of enzymatic activity due to decrease in pH occurs during cell membrane disruption in the presence of various essential oils. Essential oils penetrate the cell membrane as well as the mitochondrial membrane, leading to greater cytoplasmic and  $K^+$  ion loss (Raybaudi-Massilia et al. 2006). Lambert and Hammond (1973) have likewise mentioned potassium leakage as an early indicator of membrane damage.

Phenols affect enzyme activity associated with energy production at low concentration, while they cause denaturation of protein at high concentration. Therefore, the modes

of action of phenolic compounds (EO fractions) as antimicrobial agents can be concentration dependent (Davidson 2001; Nychas 1995; Lopez-Malo et al. 2005; Sofos et al. 1998; Juven et al. 1994). Bajpai et al. (2008) reported that the antimicrobial activity is due to phenols' ability to alter microbial cell permeability, causing the loss of macromolecules, interfering with membrane function, and thereby causing deformation in structure and functionality (Rico-Munoz et al. 1987; Kabara and Eklund 1991). The antimicrobial activity of isothiocyanates derived from onion and garlic is responsible for the inactivation of extracellular enzymes through oxidative cleavage of disulfide bonds. The formation of the reactive thiocyanate radical is believed to produce the antimicrobial effect (Delaquis and Mazza 1995). Similarly, Helander et al. (1998) have reported that carvacrol, thymol, and trans-cinnamaldehyde decrease the intracellular ATP content of *E. coli* O157:H7 cells and subsequently increase extracellular ATP, indicating the disruptive action on the cell membrane.

Phenolic compounds and their essential oils containing hydroxyl group (–OH) are more inhibitory against microorganisms compared to those compounds that contain the carbonyl group. The –OH group can easily bind the active site of enzymes altering cell metabolism of microorganisms (Farg et al. 1989). In addition, the position of the –OH group in the phenolic ring structure influences antimicrobial activity. Pauli and Knobloch (1987) have reported that the alkyl substitution (o- and p-alkyl) into phenolic compounds showed strong antifungal activity. The essential oils of eugenol and thymol contribute to the inhibitory action by membrane disruption in both Gram-negative and Gram-positive bacteria (Walsh et al. 2003). Gustafson et al. (1998) have demonstrated that tea tree oil acts on the cell membrane causing cytoplasmic leakage, cell lysis, and cell death. It has also been reported that there is a rapid inhibition of the energy metabolism of *L. monocytogenes* and *Listeria sakei* in the presence of high concentrations of eugenol and cinnamaldehyde. Polyphenol compounds known as catechins have shown strong antimicrobial effects by altering the membrane morphology and disrupting the cell membrane (Ikigai et al. 1993). Other phenolic compounds such as caffeic acid and chlorogenic acid can stimulate the DNA degradation induced by Fe (III) and bleomycin (Hiramoto et al. 1996; Moran et al. 1997). Moran et al. (1997) have also reported that some antioxidants with trace amount of metals show enough prooxidative activity to cause DNA damage. Hashimoto et al. (1999) have demonstrated that gallic acid esters in epicatechin and epigallocatechin gallate have an affinity for lipid bilayers and that the effect of this affinity causes damage to the membrane structure. The actual mechanism of inhibition by all phenolic compounds is primarily the compounds' reacting with the cell membrane or inactivating essential cellular enzymes or

a combination of both (Davidson and Branden 1981). Therefore, the damage to bacterial cells occurs with the physical disruption of the target site cell membrane, dissipation of the proton motive force, or by the inhibition of associated membrane enzyme activity as suggested by Maillard (2002).

Various authors in the past have hypothesized different modes of action of spices and plant derivatives. Juven et al. (1994) have reported hydrophobic and hydrogen bonding of phenolic compounds to membrane proteins, followed by partition in the lipid bilayer. Cox et al. (2000) have shown the perturbation of membrane permeability consequent to its expansion and increased fluidity resulting in the inhibition of membrane-embedded enzymes. Membrane disruption, cell wall perturbation, and destruction of the electron transport system are a few other antimicrobial mechanisms of plant spices and their derived products (Caccioni et al. 1998; Juglal et al. 2002; Tassou et al. 2000).

Previous studies have shown that Gram-positive bacteria are more sensitive to antimicrobial action compared to Gram-negative bacteria that possess a lipopolysaccharide outer membrane, which is relatively impermeable to phenolic compounds (Smith-Palmer et al. 2001; Burt 2004; Farag et al. 1989; Lemos et al. 1990). It can be inferred that phenolic compounds can sensitize the phospholipid bilayer of the cytoplasmic membrane causing increased permeability, unavailability of vital intracellular components, and impairment of bacterial enzyme systems (Juven et al. 1994; Kim et al. 1995; Farag et al. 1989; Wendakoon and Sakaguchi 1995). Further research to demonstrate the mechanisms by which these plant-derived products cause cell death is warranted before utilizing them in food as natural preservatives or as commercial applications in the pharmaceutical industry.

### Plant components on viability and growth of probiotics

Plant species possess several functional components as growth-promoting factors for probiotics. Functional components such as phenolic compounds, antioxidants, and micronutrients have shown not only to activate the growth but also to increase bacterial populations. Some species of these probiotics are able to metabolize phenolic compounds, and these compounds also serve as oxygen scavenger, which could have increased the growth of certain probiotic bacteria (Alberto et al. 2001). The functional components derived from plant source when incorporated into the diet could lead to beneficial physiological changes in human microflora. Probiotics, the beneficial microorganism in the human gut, can therefore be potentiated by functional components of natural origin. This section summarizes the effect of such components on the enhancement of selected probiotics.

Phenolic compounds present in berries have shown to enhance the growth of lactobacilli and bifidobacteria (Tabasco et al. 2011). Alberto et al. (2001) reported that phenolic compounds such as gallic acid and catechin normally present in grapes have stimulated the growth rate and resulted in greater cell densities of *Lactobacillus hilgardii* 5W in the FT80 liquid medium. This growth stimulation could be related to the ability of *Lactobacillus* spp. to metabolize these phenolic compounds. These compounds serve as substrate for the bacteria and this could have provided the stimulatory effects. Anthocyanin pigments present in berries have also influenced the growth of *Lactobacillus acidophilus* (Pratt et al. 1960). Ávila et al. (2009) have mentioned that  $\beta$ -glucosidase present in *Lactobacillus* spp. (*Lactobacillus plantarum* IFPL722, *L. casei* LC-01) and *Bifidobacterium lactis* BB-12 can convert anthocyanin into other phenolic acid compounds with different bioavailability and bioactivity that could have contributed to the increase in growth of probiotics. Similarly, carbohydrates such as pectins and pectic-oligosaccharides have been reported to stimulate the growth of probiotic including *Bifidobacterium* spp. (Manderson et al. 2005; Olano–Martin et al. 2002).

The supplementation of probiotic food products containing *L. casei* with plant compounds containing antioxidant properties such as catechin at concentrations of 100–400  $\mu$ M and chlorogenic acid at 400  $\mu$ M has been shown to enhance the growth of *L. casei* (Duda-Chodak et al. 2008). Tea phenolic compounds including epicatechin, catechin, 3-*O*-methylgallic acid, gallic acid, and caffeic acid have been identified as inhibitory agents for *Clostridium* and *Bacteroides* spp. However, the growth of probiotics such as *Lactobacillus* spp. and *Bifidobacterium* spp. was less inhibited indicating the possible effect of phenolic compounds on the viability of probiotics (Lee et al. 2006). Similarly, Jaquet et al. (2009) reported an increase in bifidobacterial populations after drinking coffee in healthy volunteers with lower initial bifidobacterial numbers. This stimulating effect is mainly due to the presence of phenolic compounds (chlorogenic acids) and soluble fiber.

The consumption of tannin-rich pomegranate products by humans has shown the potential to improve the imbalance of intestinal bacteria caused by stress and other factors. Bialonska et al. (2009) reported that commercial extract of pomegranate byproduct at 0.01 % (v/v) significantly enhanced the growth of *Bifidobacterium breve* NRRL B-41408 and *Bifidobacterium infantis* NRRL B-41661, thereby promoting human gut health. Aqueous extraction of garlic and black peppercorns significantly enhance the growth of *Lactobacillus reuteri*. Similarly, according to Sutherland et al. (2009), the growth-promoting activities of aqueous extracts of banana, apple, and orange have been shown to enhance the growth of *L. reuteri*, *Lactobacillus rhamnosus*, and *B. lactis*.

Components present in these plant extracts such as sugars and phytochemicals including phenolic and organic acids have been shown to enhance the growth of these probiotics. Soy germ rich in isoflavones had a positive effect on the viability of probiotics mainly due to the presence of oligosaccharides, which could act as prebiotics that help promote the growth and activity of these beneficial microorganisms. De Boever et al. (2000) have reported the viability of lactobacilli when the cultures were subjected to a 2-week treatment period by adding 2.5 g/day soy germ powder to the culture medium. Michael et al. (2010) studied the effect of several plant extracts on the viability of *Lactobacillus delbrueckii* ssp. *bulgaricus* and *Streptococcus thermophilus* in nonfat yogurt. This study demonstrated that the addition of olive, garlic, onion, and citrus extracts at 0.5 and 0.1 % (w/v) maintained the viability of the yogurt cultures during refrigerated storage for 21 days. In vitro evaluation of almond skins during fermentation showed potential to be used as a novel source of prebiotics to enhance the populations of bifidobacteria (Mandalari et al. 2010).

The growth in bacterial population is attributed to the conversion of polyphenols to phenolic acids as one of the major group of phenolic metabolites by the colonic microbiota (Lafay and Gil-Izquierdo 2008). Since bifidobacteria are acetate and lactate producers, the increase in concentration of organic acids (acetate and propionate) during fermentation of almond skin might have increased the number of bifidobacteria (Mandalari et al. 2010). Roberfroid (2000) suggested fructooligosaccharides as prebiotics, as they stimulate the growth and metabolism of probiotic bacteria in the gut. Panesar and Shinde (2012) have shown good viability of *L. acidophilus* and *Bifidobacterium bifidum* in aloe vera and water blend juice (1:3)-fortified yogurt during a 28-day storage period. Germinated rough rice is another potential supplement for the growth of *L. plantarum* in food fermentation (Trachoo et al. 2006). High levels of nutrients and bioactive compounds such as proteins, amino acids, sugars, vitamins, gamma-oryzanol, gamma-amino butyric acid, tocotrienols, tocopherols, and other phytochemical substances may have promoted the growth of these bacteria (Hagiwara et al. 2004; Oh et al. 2003; Tian et al. 2004).

### Impact of plant components on probiotics and pathogens

Plant contains several functional components such as phenolic compounds, antioxidants, and minerals. Such plant-derived components have been shown to influence colonic microflora (Kelly et al. 1994) either by promoting the growth of beneficial microorganisms or by acting against pathogens. This could be due to the effect of plant extracts, which might be used in potentiating the neutralization effect

of beneficial microorganisms against enterotoxin-producing bacteria (Bomba et al. 2002). This section highlights the potential influence of plant components on probiotics and the mode of action of such probiotics against pathogens. Our main interest here is to identify the dual activity shown by these plant components. Table 1 shows the impact of various plant components on probiotics and harmful microorganisms. The impact of such functional components can be either direct or indirect (Fig. 1).

#### Direct effect

Plant components have a direct effect on the growth and viability of probiotics as well as the inhibition of harmful microorganisms. Some of the recognized functional components in selected plant species are summarized in Table 1, which provides an outline of functional components, the impact of these components on the growth and viability of probiotics, antimicrobial action against pathogens, and references. This direct effect can be attributed to the impact of functional components to enhance the growth of probiotics or antimicrobial action against pathogens (Fig. 1).

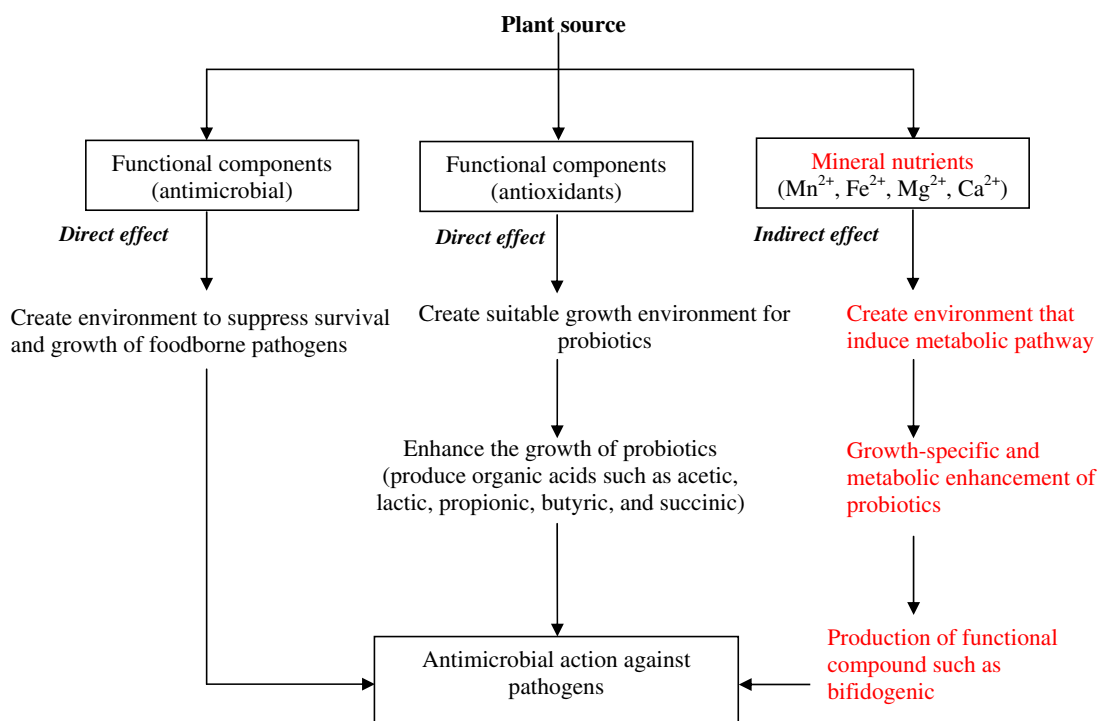
Sedighi et al. (2011) have demonstrated that almond pudding (10 % almond smash in 90 % rice extract, w/v) in infant's diet may enhance the protection against *E. coli* O157:H7 by increasing the *Bifidobacterium* spp. populations in infant's gastrointestinal system. Almond is a good source of oligosaccharide, which acts as a prebiotic to promote the growth of beneficial microorganisms (*Bifidobacterium* spp. and lactobacilli) and inhibit the growth of *Clostridium histolyticum* (Mandalari et al. 2010). The presence of phytochemicals and nutrients such as vitamin E could have attributed to the growth enhancement and suppression of *Bifidobacterium* spp. and *E. coli* O157:H7, respectively. Some spices including cinnamon and origanum have been shown to enhance the growth of lactic acid bacteria and *Bifidobacterium* spp., which has resulted in higher acid production that acts as an antimicrobial compound against pathogens including *E. coli* O157:H7 and *H. pylori* (Shelef 1984; Ibrahim et al. 2003; Ali et al. 2005; Behrad et al. 2009). Polyphenols, phytic acid, and oxalic acid are some of the important classes of antioxidant present in plants. Oxalic acid showed growth-enhancing response for *L. acidophilus*. Similarly, plant products such as apple skins, onions, tea, red wine, leafy green vegetables, and berries are rich in quercetin. A recent study by Yadav et al. (2011) showed the enhancing effect of antioxidant quercetin at 0.5–2.0 mg/5 mL toward the growth of probiotics such as *L. acidophilus* and marketed consortium of eight probiotic cultures. Salem et al. (2009) also reported the potentiality of apple skins as a media source for the production of lactic acid using *L. reuteri*. The produced lactic acid may act as an antimicrobial agent against harmful microorganisms. The components like oxalic acid and phytic acid are also known

**Table 1** Impact of plant-derived components on microorganisms

Plant species	Functional component(s)	Action on probiotics	Antimicrobial action against pathogens	References
Almond (skins)	Polyphenols and dietary fiber	Significantly increased the population of <i>Bifidobacterium</i> spp.	<i>E. coli</i> O157:H7, <i>Clostridium histolyticum</i>	Sedighi et al. (2011), Mandalari et al. (2010)
Aloe vera	Glucose, mannose, amino acids, anthraquinones, dihydroxyanthraquinones, saponins, aloin, and other phenolic compounds	Shown viability of <i>Lactobacillus acidophilus</i> and <i>Bifidobacterium bifidum</i>	<i>Staphylococcus aureus</i> , <i>Pseudomonas aeruginosa</i> , <i>Streptococcus pyogenes</i> , <i>Escherichia coli</i>	Panesar and Shinde (2012), Agarry et al. (2005), Arunkumar and Muthuselvam (2009)
Berries	Phenolics (ellagitannins, anthocyanins, catechins, and proanthocyanidins)	Increased the survival of bifidobacteria in yogurt No effect on <i>Lactobacillus rhamnosus</i>	<i>Staphylococcus aureus</i> and <i>Salmonella enterica</i> serovars	Puupponen-Pimiä et al. (2001; 2005)
Cinnamon (spice)	Cinnamaldehyde, eugenol	Presence of herbs did not affect the probiotic populations of <i>Lactobacillus</i> spp. and <i>Bifidobacterium</i> spp.	<i>Helicobacter pylori</i>	Behrad et al. (2009), Ali et al. (2005)
Coffee	Phenolic compounds (chlorogenic acids) and soluble fiber	Increased <i>Bifidobacterium</i> spp.	<i>E. coli</i> O157:H7	Jaquet et al. (2009), Ibrahim et al. (2006)
Garlic	Sugars, small proteins, Allicin, phenolics and organic acid	Enhanced the growth of <i>Lactobacillus reuteri</i>	<i>Escherichia coli</i> O157:H7 and <i>Escherichia coli</i> LF82	Sutherland et al. (2009), Shelef (1984)
Grape extract	Phenolic compounds: monomeric phenolic compounds and dimeric, trimeric, and tetrameric procyanidins Flavanoids: flavan-3-ols (catechins and proanthocyanidins)	Maximum growth of <i>Bifidobacterium breve</i> and <i>Bifidobacterium bifidum</i> observed	<i>Aeromonas hydrophila</i> , <i>Bacillus cereus</i> , <i>Enterobacter aerogenes</i> , <i>Enterococcus faecalis</i> , <i>Escherichia coli</i> , <i>Escherichia coli</i> O157:H7, <i>Mycobacterium smegmatis</i> , <i>Proteus vulgaris</i> , <i>Pseudomonas aeruginosa</i> , <i>Pseudomonas fluorescens</i> , <i>Salmonella enteritidis</i> , <i>Salmonella</i> Typhimurium, <i>Staphylococcus aureus</i> , and <i>Yersinia enterocolitica</i>	Özkan et al. (2004), Tabasco et al. (2011)
Olive leaf extracts	Polyphenol compounds (oleuropein and other secoiridoid), flavonoids: rutin flavonol, luteolin-7-glucoside	Stimulated the growth of <i>Bifidobacterium infantis</i> and <i>Lactobacillus acidophilus</i>	Exposure to olive leaf extract showed complete destruction of <i>E. coli</i> cells <i>Salmonella</i> Typhi, <i>Staphylococcus aureus</i> , <i>Vibrio cholerae</i>	Haddadin (2010), Markin et al. (2003) Bisignano et al. (1999)
Origanum (spice)	Antioxidant origanox	Little effect on the growth of <i>Bifidobacterium longum</i>	<i>E. coli</i> O157:H7	Ibrahim et al. (2003)
Pomegranate	Ellagitannins and alkaloids	Significantly enhanced the growth of <i>Bifidobacterium breve</i> and <i>Bifidobacterium infantis</i>	Pathogenic clostridia and <i>Staphylococcus aureus</i>	Bialonska et al. (2009)
Soybean, chickpea	Isoflavones (biochanin A)	No effect on <i>Bifidobacterium</i> spp.	<i>Clostridium</i> spp.	Sklenickova et al. (2010), Kramer et al. (1984)
Tea	Tea phenolics: epicatechin, catechin, 3-O-methyl gallic acid, gallic acid, and caffeic acid	<i>Bifidobacterium</i> spp. and <i>Lactobacillus</i> sp. were less severely affected Improved the survivability of bifidobacteria in yogurt during storage	<i>Clostridium perfringens</i> , <i>Clostridium difficile</i> , and <i>Bacteroides</i> spp. <i>Staphylococcus aureus</i> , <i>Salmonella</i> Typhimurium, <i>Escherichia coli</i> , and <i>Listeria monocytogenes</i>	Lee et al. (2006), Akahoshi and Takahashi (1996) Kristanti and Punbusayakul (2008)

to act as chelator for iron, thereby creating an environment to suppress the survival and growth of other microorganisms (Yadav et al. 2011). Hence, microorganisms especially pathogens that require iron for their growth cannot compete with other microorganisms without iron in the environment. Grape

extract has shown that the active compounds present in grapes promote the growth of *Bifidobacterium* spp. as well as are effective against pathogenic microorganisms (Özkan et al. 2004; Tabasco et al. 2011). The antioxidant resveratrol (0.143 mg/kg/day) present in grapes has shown to enhance



**Fig. 1** Impact of plant components on survival of pathogens and functionality of probiotics

the growth of *Bifidobacterium* and *Lactobacillus* in a rat model study after 20 days of resveratrol intake (Larrosa et al. 2009). On the other hand, resveratrol at 60  $\mu\text{g}/\text{mL}$  prevent the expression of virulence factors of *Proteus mirabilis* protecting human urothelial cell damage (Wang et al. 2006).

Tea phenolics and their derivatives have shown to inhibit the growth of pathogenic bacteria such as *Clostridium perfringens*, *Clostridium difficile*, *Bacteroides* spp., *S. aureus*, *Salmonella* Typhimurium, and *L. monocytogenes*, while probiotics such as *Bifidobacterium* spp. and *Lactobacillus* spp. were relatively unaffected, thus improving the intestinal microflora (Akahoshi and Takahashi 1996; Lee et al. 2006). Rosenthal et al. (1997, 1999) reported that tea catechins and ferulic acid also inhibit the growth of pathogenic bacteria (coliforms and *Salmonella*) but not the growth of lactic acid bacteria, indicating the role of fermentative metabolism of lactic acid bacteria. Similarly, Weisburger (1999, 2000) reported that the consumption of tea reduces the number of undesirable bacteria in the intestine but has the reverse effect on beneficial bacteria showing the dual activity of functional component catechins. Goto et al. (1999) demonstrated the effect of tea catechins (300 mg/day for 6 weeks) on fecal contents and metabolites of elderly residents and found that consumption of green tea promotes the growth of *Bifidobacterium* and *Lactobacillus* in the gut wall and decreases the survival of *S. aureus* and *Streptococcus pyogenes*. As shown in Table 1, plant-derived functional components increase the viability of probiotics such as lactic

acid bacteria and thus enhance the metabolic activity and survivability of these probiotics. Lactic acid bacteria, in turn, enhance intestinal health by producing antimicrobial substances that inhibit epithelial invasion by pathogens (Servin and Coconnier 2003). Lactobacilli also produce antimicrobial proteins known as bacteriocins that have been shown to display a wide antibacterial spectrum against Gram-positive bacteria (Jack et al. 1995). Some strains of *L. reuteri* have the ability to convert glycerol into the broad spectrum antimicrobial substance reuterin. Reuterin can inhibit the growth of Gram-negative and Gram-positive bacteria, yeasts, fungi, and protozoa and also remain active at a wide range of pH (Talarico et al. 1988, 1990; Dobrogosz and Lindgren 1994). Reuterin also affect pathogens by producing stress response thereby taking longer time to adapt to the environment (Bian et al. 2011). Rasch et al. (2007) reported that reuterin elongates the lag phase of *L. innocua* ATCC 33090 and hinders cell division irrespective of the pH of the media. As shown in Table 1, some of the plant components enhance the growth of bifidobacteria. Some of the mechanisms that have been suggested for the inhibitory action of bifidobacteria towards Gram-negative pathogens include reduction of pH by the production of organic acids, the inhibitory action of undissociated organic acid molecules, and the production of specific antimicrobial substances (Fuller 1989; Ibrahim and Bezkorovainy 1994; Servin 2004). Makras and De Vuyst (2006) have reported strong antimicrobial activity of organic acids produced by



*Bifidobacterium* strains against *S. enterica* Typhimurium and *E. coli*.

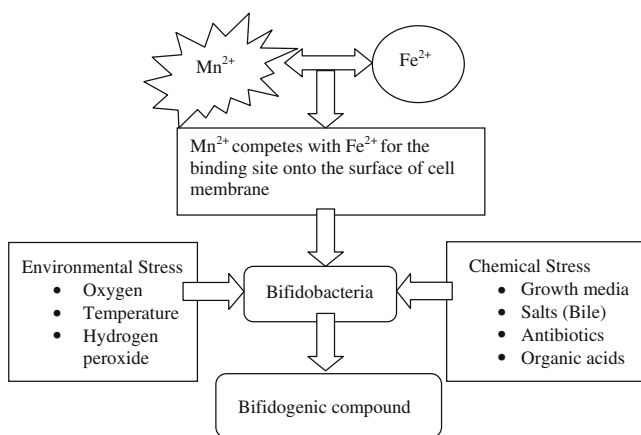
This dual effect is probably due to the inhibition of other bacteria in the intestinal tract that could have provided a niche for the enhancement of probiotics. Most of the plant components have been found to have strong antimicrobial properties against numerous pathogens. In addition to these strong antimicrobial properties, plant components also promote the growth and viability of probiotics, which produce organic acids, and other antimicrobial substances that inhibit a wide range of pathogenic microorganisms.

#### Indirect effect

Several mineral nutrients such as  $Mg^{2+}$ ,  $Mn^{2+}$ ,  $Fe^{2+}$ , and  $Ca^{2+}$  have been demonstrated to bring an impact on growth and functionality of probiotics (Boyaval 1989; Ibrahim and Bezkorovainy 1994; Wishon et al. 2010). Magnesium ( $Mg^{2+}$ ) is considered as an essential nutrient for the growth of several lactic acid bacteria such as *Lactobacillus helveticus*, *Lactobacillus lactis*, and *L. delbrueckii* (Rogosa and Mitchell 1950). Aksu and Kutsal (1986) have reported 70 % increase in lactic acid production by *L. delbrueckii* NRRL B-445 when  $MgSO_4$  was added in molasses, yeast extract, and  $(NH_4)_2HPO_4$ . Similarly,  $Mg^{2+}$  has shown to increase the survival of *Streptococcus lactis* ML3 in phosphate buffer (Thomas and Batt 1968). It has been also reported that many enzymes, which require  $Mg^{2+}$ , may also be activated in the presence of manganese ( $Mn^{2+}$ ).  $Mn^{2+}$  present in tomato juice (0.011 %) and asparagus juice has been shown to enhance the growth and acid production by lactic acid bacteria (Stamer et al. 1964). Tiwari et al. (1980) and Tewari et al. (1985) have shown increased growth of *Lactobacillus bulgaricus* with Fe-EDTA complex and higher acid production with the addition of ferric chloride ( $9 \times 10^{-5}$  mM) and cobalt chloride to paneer whey. McDonald (1957) suggested that  $Ca^{2+}$  and  $Mg^{2+}$  ions have stimulated the growth of *Lactococcus lactis* and *Lactococcus cremoris*. These enhancing effects could have resulted from an increase of free ionic form of minerals in growth media that act as an activator of different metabolic reactions such as cell division and stabilization of nucleic acids (Boyaval 1989). Therefore, the presence of these minerals has shown to create an environment that induces a new metabolic pathway. This metabolic pathway could enhance the growth-specific metabolites of probiotics that ultimately lead to the production of functional compounds. These functional compounds could have antimicrobial properties that act against pathogens (Fig. 1, indirect effect).

Some examples on how plant components such as iron ( $Fe^{2+}$ ) and manganese ( $Mn^{2+}$ ) indirectly influence the survivability of microorganisms have been described earlier (Ibrahim and Bezkorovainy 1994). Iron is an

essential nutrient for all microorganisms; however, probiotics such as bifidobacteria and lactobacilli bind the iron and hence reduce the availability to pathogenic microorganisms (Bomba et al. 2002). Spices are rich in  $Mn^{2+}$  ions and have been reported to be strong stimulators of starter culture and rapid acid production (Zaika and Kissinger 1979, 1984; Kang and Fung 2000). Ibrahim et al. (2010) have reported that the addition of  $Mn^{2+}$  to the growth culture stimulated the production of  $\alpha$ - and  $\beta$ -galactosidase activity of *L. reuteri* CF2-7F. Therefore,  $Mn^{2+}$  ions could be used for the growth of bacteria and high activity of these enzymes. Extracts of plant components such as clove, cardamom, ginger, celery seed, cinnamon, and turmeric contain  $Mn^{2+}$  that enhances acid production by *L. plantarum* (Zaika and Kissinger 1984). Stimulating the presence of  $Mn^{2+}$  has shown to stimulate the growth of *L. casei* YIT-9018 (Nakashima 1997). The production of lactic acid and other antimicrobial compounds from *Lactobacillus* spp. could act antagonistically against foodborne pathogens (Fig. 1). Ibrahim and Bezkorovainy (1994) reported that bifidobacteria have the ability to inhibit the growth of pathogenic bacteria by producing organic acids, and the antimicrobial activity of such probiotic (*Bifidobacterium longum*, NCFB 2259) is influenced by spices (Ibrahim et al. 2003). A diagram of the indirect stimulatory effect of  $Mn^{2+}$  ions (the mineral found in spices) on the production of bifidogenic compound by bifidobacteria is shown in Fig. 2. Novel functional compounds produced by bifidobacteria under low  $Fe^{2+}$  concentration have been reported by Ibrahim (2005) and O'Sullivan (2001). Iron is known to be a growth-promoting factor for bifidobacteria, and  $Mn^{2+}$  has been shown to have a strong inhibitory effect against  $Fe^{2+}$  uptake (Ibrahim and Bezkorovainy 1994; Bezkorovainy et al. 1996). When  $Fe^{2+}$  is limited in the growth environment, bifidobacteria tend to grow at a slower rate and start to produce a bifidogenic compound that has antimicrobial properties. This bifidogenic compound scavenges  $Fe^{2+}$  from the environment, thus making  $Fe^{2+}$  unavailable to pathogens (Bezkorovainy et al. 1996; Miller-Catchpole et al. 1997) and other competitive microorganisms. Ibrahim (2005) also reported that bifidobacteria could inhibit *E. coli* O157:H7. From Ibrahim's study, it was confirmed that ferrous ( $Fe^{2+}$ ) plays an important role in the production of bifidogenic compounds, and when  $Fe^{2+}$  is not available or iron uptake is inhibited, bifidobacterial cells could also be stressed by several environmental and chemical factors that ultimately lead to the production of antimicrobial bifidogenic compound (Fig. 2) (Ibrahim 2005; Morrison et al. 2001). This is how the  $Mn^{2+}$  present in most of the spices inhibits the uptake of  $Fe^{2+}$  and ultimately suppressed the growth of pathogens by producing new antimicrobial compounds.



**Fig. 2** Role of manganese ( $Mn^{2+}$ ) and iron ( $Fe^{2+}$ ) in the production of antimicrobial bifidogenic compound

Almost all bacteria, including probiotics, require iron in their natural habitat as an essential element for growth except some lactobacilli (Weinberg 1997). For this reason, lactobacilli have an advantage over other microorganisms that depend on iron. Elli et al. (2000) reported that *L. acidophilus* and *L. delbrueckii* are able to bind ferric hydroxide on their cell surface making it unavailable to pathogenic microorganisms. Functional component tannins are capable of forming complexes with polymers and minerals, making nutrients unavailable to other pathogen (Smith et al. 2005). Thus, it can be inferred that the competition by these probiotic bacteria for limited resources is an indirect way of inhibiting the growth of pathogenic microorganisms. However, the level of stimulation and inhibition of such plant components on bacteria varies depending on the bacterial species and the chemical structure of the compounds. Our literature review has shown that plant products possess bioactive components that not only promote the growth of probiotics but also inhibit the growth of pathogens directly or indirectly with various antimicrobial compounds produced from probiotics.

## Applications

Plants, their respective extracts, and functional compounds from them have been used as antibacterial agents and remedies for human illnesses. Various plant species have been known to contain naturally occurring compounds with antimicrobial activity (Kim et al. 2004; Beuchat and Golden 1989; Lopez-Malo et al. 2005). Major groups of chemicals present in plant components that are responsible for enhancing the safety and quality of food and drugs include polyphenols, quinines, flavanols/flavanoids, alkaloids, and lectins (Cowan 1999). Enhancement of food safety is one of the interests of the food industry, and the use of natural

food preservatives from plant sources is one of the ways to control microbial growth and reduce the incidence of foodborne illnesses and spoilage (Eklund 1980). These plants contain large amounts of antioxidant compounds, such as ascorbic acid, vitamin E, and phenolic compounds, and are also potentially useful in reducing oxidative damage to human cells (Jayakumar et al. 2011). Many of these plant products have been studied in the past for different purposes as antimicrobials, antioxidants, medicinal, and food preservatives on a commercial basis. Therefore, plant products are of great interest in food safety and human health.

## Food safety

Despite the use of various preservatives and methods to control food spoilage and foodborne pathogens, food safety continues to be a challenge to the entire food industry. Antibiotic-resistant foodborne pathogens have become a major health concern (Kiessling et al. 2002), as these pathogens are resistant to several food preservation techniques such as heat and acid treatment. There is a need to develop natural preservatives that could prevent the growth of such pathogens. Consumers are increasingly concerned about the safety of foods prepared with chemical preservatives. Therefore, plant-derived compounds as a natural alternative could be used to prolong the shelf life of foods and to enhance microbial safety.

The antibacterial activity of spices has attracted the interest of those who work in food safety occupations. For example, rosemary at 0.5 %w/v and its essential oil (1 % v/w) produced significant antilisterial effect on pork sausage during storage at 5 °C for 50 days (Pandit and Shelef 1994). Grohs and Kunz (2000) reported the inhibitory effect of a mixture of several aromatic herbs on the growth of meat spoiling microorganisms (*B. subtilis*, *Enterococcus* spp., *Staphylococcus* spp., *E. coli* K12, and *Pseudomonas fluorescens*) thus preserving the color and smell of fresh-portioned pork meat. Leuschner and Zamparini (2002) observed the bacteriostatic and bactericidal effect of garlic (1 %w/v) and clove (1 %w/v) against *S. enterica* and *E. coli* O157:H7, respectively, in mayonnaise. Skandamis et al. (2002) observed 1–2 log reduction of *Salmonella* Typhimurium in beef fillets stored under aerobic and modified atmosphere packing when treated with oregano essential oil. Growth of *S. aureus* was inhibited when 1 % of cloves, onion, ginger, and black and red pepper were added to meat homogenates (Nkanga and Uraih 1981). Caffeic acid, the most effective of the phenolic acids known to occur in apples, has been shown to inhibit *E. coli* O157:H7 and other foodborne pathogens (Reinders et al. 2001). Components of natural origin such as caffeic acid could be a better alternative to the currently used method for food preservation. Green tea components have been shown to enhance the shelf

life of raw, frozen, and cooked meat patties (Jo et al. 2003; Tang et al. 2001). Bañkn et al. (2007) suggested that ascorbate, green tea extract, and grape seed extracts delayed microbial spoilage, redness loss, and lipid oxidation, in low sulfite beef patties, thereby increasing shelf life. Trans-cinnamaldehyde, a major component of bark extract of cinnamon (Holley and Patel 2005), could be used as an antimicrobial agent to inactivate *E. coli* O157:H7 in apple juice and apple cider (Baskaran et al. 2010). Extracts of pepper, parsley, and dill have been shown to exhibit antimicrobial activity against natural microflora, coliforms, yeasts and molds, and *S. aureus* in Kareish cheese to prevent spoilage (Wahba et al. 2010). The compound thiosulfinates (allicin) present in garlic extract have also been shown to prevent food contamination from pathogenic microorganisms (Durairaj et al. 2009).

A combination of spices and bifidobacteria could be used to control the growth of *E. coli* O157:H7 in ready-to-eat foods and to increase the bio-safety of many consumable food products (Ibrahim et al. 2003). Plant products such as ginger enhance the growth of *L. reuteri* (Sutherland et al. 2009), which produces the broad spectrum antimicrobial compound reuterin (Axelsson et al. 1989). The antimicrobial activity of reuterin has been studied against different foodborne Gram-positive and Gram-negative pathogens in milk. Results showed that reuterin exhibited bacteriostatic and bactericidal activity against *L. monocytogenes* Ohio serotype 4b and *E. coli* O157:H7 ATCC 43894, respectively (Arques et al. 2004). A variety of metabolic products produced by lactic acid bacteria have been shown to interfere with the growth of other microbes and have been applied to food systems to prevent the growth of harmful microorganisms (Vandenbergh 1993).

Consumers are increasingly demanding healthier and safer food products that are free of chemical additives. Natural preservatives may lower processing costs and extend product shelf life. Natural preservatives could also be exploited by small food industries that do not require advanced technology such as non-thermal processing methods. Therefore, these preservatives could maintain both quality and reduce processing cost. In addition, natural preservatives may help prevent the emergence of antibiotic resistance microorganisms (Gálvez et al. 2010). Therefore, the use of natural preservatives to increase the shelf life of food is a promising method, as these substances of plant origin have antioxidant and antimicrobial properties.

#### Human health

In the last few years, considerable attention has also been given to the study on how diet can impact gut microflora. The use of dietary components to increase the metabolic activity or the number of beneficial microorganisms

(probiotics) without altering the existing gut ecosystem has been shown to be the best way to improve human health from a practical standpoint. Consumption of plant source products can have considerable potential for the prophylaxis and therapy of gut infections. Many studies in the past have shown methods to promote beneficial bacteria through the intake of dairy products incorporated with bifidobacteria (Poch and Bezkorovainy 1988; Collins and Hall 1984). Recently, attention to the impact of plant-derived components has focused largely on the beneficial physiological changes in the colonic microenvironment.

As discussed earlier, plant components can stimulate the growth of probiotics that can promote other health benefits in human health (Fukushima and Iino 2006). Parkar et al. (2008) have reported the potential influence of polyphenols on gut microecology with regard to inhibiting the growth and adhesion of the gut pathogens while at the same time enhancing the proliferation and adhesion of the probiotic, *L. rhamnosus* 299. This probiotic activity also helps restore the balance of the intestinal microflora in the host (Plummer et al. 2005). Shelef (1984) has mentioned that the consumption of spices may reduce cancer risk by producing a bacterial shift in the intestinal tract.

Up to now, studies have focused on the antibacterial effect or beneficial effect of probiotics mainly on the basis of selection of efficient strains of probiotics with regard to source, functionality, adhesion property, bile and acid resistance, enzyme activity, and their viability in food (Ouweland et al. 2002; Parvez et al. 2006; Percival 1997). Other factors that may enhance the efficacy of probiotics include gene manipulation, the combination of number of strains of microorganisms, and the combination of probiotics and synergistically acting components (Bomba et al. 2002). However, the contribution of probiotics to the improvement of microflora depends not only on the survivability of probiotics during storage conditions but also on the survivability of probiotics during transit period of acidic conditions of the stomach and during degradation by hydrolytic enzymes and bile salts in the small intestine (Playne 1994). It is also believed that the ingested probiotics pass through the feces without having adhered or multiplied in the intestinal mucosal cells (Servin and Coconnier 2003). Orrhage et al. (1991) have also reported that intake of supplements with bifidobacteria does not cause any significant change in the intestinal microbiota as bifidobacteria do not easily adhere in the gut. Adherence of probiotics to intestinal mucosal cells is believed to provide maximum probiotic effect. However, it has been reported that exogenously administered probiotics seem to pass into feces instead of adhering to the mucosal cells. Thus, Bezkorovainy (2001) has suggested a continuous ingestion of probiotic culture to obtain a continuous exogenous

probiotic effect. However, this may not seem as practical as getting a probiotic benefit from the consumed diet itself. Ouwehand et al. (2001) reported that resistance to low pH is not only important for the survival of probiotic but also for adhesion. The authors have also mentioned that the adhesion of most of the tested strains was significantly reduced once exposed to pepsin or low pH after passing through the stomach. These findings again indicate that probiotics obtained through dietary supplements may not be able to fulfill all probiotic properties as claimed. Furthermore, probiotics must survive and retain their functionality during storage as well as in the foods into which they are incorporated. Thus, increasing the number of natural microbiota including probiotic bacteria present in the human gut via consumption of natural ingredients of plant origin seemed to be the most practical approach for a healthy gut ecosystem.

However, people are not much aware of the impact of plant-based diet and its influence on existing microflora including probiotics in the human gut. The use of functional components of plant origin can modify the colonic microbiota by increasing the number of specific probiotics. This could help transform some of the phenolic compounds present in plant components and ultimately provide a wide range of implications for the health of the host, thereby producing beneficial effects. An increase in human consumption of natural ingredients of plant origin would provide a dual health benefit by (1) stimulating the growth of probiotics and (2) inhibiting the growth of harmful microorganisms. Consuming a diet rich in functional ingredients from plant sources has been shown to have a positive impact on human health. As a result, we strongly believe that such ingredients indirectly influence the functionality on the natural microbiota in the human gut, including bifidobacteria (Bialonska et al. 2009; Sutherland et al. 2009; Alberto et al. 2001; Ibrahim et al. 2010; Liu et al. 2011). Several reports have demonstrated that functional ingredients enhance the growth of lactobacillus and bifidobacteria, a major part of human microbiota. For instance, several polyphenols have been shown to enhance the proliferation and adhesion of probiotics and inhibit the growth and adhesion of gut pathogens in in vitro and in vivo studies. It is desirable to increase the number of probiotics and inhibit the growth of potential pathogens in the human gut. Therefore, consumption of a diet containing these plant-derived compounds has the potential to influence our gut microecology, improve the intestinal ecosystem, and thereby contribute to gut health benefits and the overall maintenance of human health.

## Conclusions

Plant species have been known to contain a wide variety of functional components. The impact of these components on

microorganisms can be either direct or indirect. Most of these compounds have shown antimicrobial properties against foodborne pathogens. Similarly, antioxidants present in plant compounds have also shown to stimulate the growth of probiotics. These probiotics in turn produce organic acids, which act as antimicrobial agents. On the other hand, mineral nutrients present in plants indirectly induce growth-specific metabolites of probiotics to promote the production of antimicrobial compounds. Hence, the production of these organic acids and antimicrobial compounds could contribute to improve food safety and human health.

Recently, scientists are showing interest in the relationship between nutrients and gene expression known as nutrigenomics. Understanding the impact of plant components on gene expression in probiotics such as lactobacilli and bifidobacteria could further help improve food safety and human health. Screening for the risk of various diseases and determining an individual's ideal health-promoting diet through genetic testing is a new area of research to understand human response to foods. Future research in the area of nutrigenomics could play a significant role to understand how diet affects gene expression to improve an individual's entire health. Hence, there are many opportunities to use this information to accelerate progress toward a comprehensive understanding of the functionality of plant-derived components on nutrigenomics.

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