

Effects of Fermented Edible Seeds and Their Products on Human Health: Bioactive Components and Bioactivities

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Abstract: There is a long history of using fermentation in food production. Edible seeds, such as certain beans and cereal grains, are important in the human diet and provide many health benefits. Various microbes, such as lactic acid bacteria, molds, and yeasts, considered as generally recognized as safe (GRAS) microbes, are commonly used to ferment edible seeds and their products. Fermentation can change bioactive components and produce new bioactivities. In order to highlight the importance of fermentation on bioactive components and bioactivities in edible seeds, this review, therefore, summarizes recent relevant studies and discusses fermentation procedures and influences of fermentation on their bioactive components and bioactivities. Overall, fermented edible seeds and their products contain enhanced bioactive components, especially γ -aminobutyric acid and natural phenolics, and they possess versatile bioactivities, such as antioxidant and anti-cancer effects, and, therefore, can be recommended as an important part of the human diet, or they can be developed into functional foods to help in the prevention of certain chronic diseases.

Keywords: anti-cancer effect, anti-hypertensive effect, antioxidant effect, beans, bioactive peptides, cereal grains, fermentation, natural phenolics, vitamins, γ -aminobutyric acid

Introduction

Fermentation, an ancient food biotechnology, is still commonly employed in the food industry to extend shelf-life as well as to improve nutritional and sensory qualities of foods (Bourdichon and others 2012). This technology mainly includes natural and inoculated fermentations. The former utilizes the endophytic microbes naturally occurring on plant-based foods to perform the fermentation process, such as to make the traditional broad bean paste in China, while the latter is commonly carried out by inoculation of generally recognized as safe (GRAS) microbes into plant-based foods or milk-based foods. Therefore, an understanding of the fermentation methods can be of importance to efficiently perform the fermentation process to develop fermented foods with health benefits.

Edible seeds, such as edible beans and cereal grains, have been reported to possess many health benefits (Saleh and others 2013; Hayat and others 2014; Rebello and others 2014). Moreover, recent studies suggest that many fermented edible seeds and their

products contain enhanced bioactive components and exhibit various bioactivities. In order to better understand the fermentation procedures and to highlight the importance of the effects of fermented edible seeds and their products on health, we searched for relevant original English articles based on ISI Web of Science from 2000 to the present. In this review, we first summarize and classify the fermentation procedures for edible seeds and their products, then discuss the influences of fermentation on the various bioactive components, such as vitamins, γ -aminobutyric acid (GABA), natural phenolics, and bioactive peptides, and finally we highlight the possible benefits of fermented edible seeds and their products, such as antioxidant, anti-hypertensive, and anti-cancer effects. Therefore, this article provides a comprehensive review and updated information about the bioactive components and bioactivities of fermented edible seeds and their products.

Fermentation procedures

Based on the source of microbes involved in the fermentation process, there are natural and inoculated fermentations. In addition, fermentation can be divided into solid-state fermentation (SSF) and liquid-state fermentation (LSF) according to the water content in the system. Edible seeds commonly need to be pretreated before fermentation, such as by soaking, cracking, milling, sieving, and cooking. In many cases, these processing methods are combined together. Next, we summarize and discuss some basic principles involved in the fermentation process.

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Pretreatment of edible seeds

Before the fermentation process, edible seeds should be pretreated. Fermentation commonly requires soaked, cracked, and cooked seeds or milled flour to start SSF or LSF. Of course, seeds cannot be cooked or autoclaved before natural fermentation, since this will partly or completely kill microbes existing on seeds, leading to the failure of subsequent fermentation. On the other hand, it is better to reduce microbial populations existing on seeds before inoculated fermentation, since they would compete with and inhibit the growth of inoculated microbes during the fermentation process. For example, studies found that LAB-fermented soybeans and lentil also contain generally the same amount of aerobic mesophilic bacteria compared to LAB after 48 or 96 h of fermentation, suggesting the possible competitive relationship between aerobic mesophilic bacteria and LAB during fermentation (Fernandez-Orozco and others 2007; Torino and others 2013). Therefore, edible seeds can be dried by hot air, such as at 40 to 60 °C for 24 h, or cooked/autoclaved, before the inoculated fermentation process (Duenas and others 2005; Lee and others 2008; Limon and others 2015). Overall, pretreatment of edible seeds can enhance the efficiency of fermentation.

The starter culture and inoculum

Many diverse microbes have been used as the starter culture for fermenting edible seeds and their products, mainly GRAS microbes, such as food-grade bacteria, fungi, and yeasts. Lactic acid bacteria (LAB) are the most common bacteria used in edible seed fermentation, such as *Lactobacillus* (*Lb.*) *acidophilus*, *Lb. brevis*, *Lb. bulgaricus*, *Lb. casei*, *Lb. fermentum*, *Lb. johnsonii*, *Lb. paracasei*, *Lb. plantarum*, *Lb. reuteri*, *Lb. rhamnosus*, *Lb. rossiae*, *Lb. zeae*, *Lactococcus* (*Lc.*) *lactis*, *Bifidobacterium* (*Bb.*) *animalis*, *Bb. infantis*, *Streptococcus* (*Sc.*) *thermophilus*, and *Weissella* (*W.*) *paramesenteroides* (Coda and others 2010; Dordevic and others 2010; Hole and others 2012; Ko and others 2013; Lai and others 2013; Rizzello and others 2013; Choi and others 2014a; Zhao and Shah 2014; Jhan and others 2015; Zhang and others 2015c). In addition, *Bacillus* (*B.*) *subtilis* has also been commonly used to ferment edible seeds (Torino and others 2013; Wang and others 2014; Limon and others 2015). Additionally, fungi (molds), including *Aspergillus* (*A.*) *oryzae*, *A. awamori*, *A. sojae*, *A. niger*, *Agrocybe* (*Ac.*) *cylindracea*, *Cordyceps* (*C.*) *militaris*, *Coprinus* (*Cr.*) *cinereus*, *Grifola* (*G.*) *frondosa*, *Ganoderma* (*Gd.*) *austral*, *Gd. neo-japonicum*, *Gd. lucidum*, *Lentinus* (*L.*) *edodes*, *Monascus* (*M.*) *purpureus*, *M. ruber*, *Rhizopus* (*R.*) *oryzae*, *R. microsporus*, *R. oligosporus*, *R. oryzae*, and *Thamnidium* (*T.*) *elegans* (Rhyu and others 2000; Lee and others 2004, 2006, 2008; Fernandez-Orozco and others 2007; Cai and others 2012; Salar and others 2012; Choi and others 2014c; Starzynska-Janiszewska and others 2014; Subramaniam and others 2014; Xiao and others 2014; Gamboa-Gomez and others 2016), and yeasts, such as *Issatchenkia* (*I.*) *orientalis*, *Saccharomyces* (*S.*) *cerevisiae*, and *S. boulardii* (Rekha and Vijayalakshmi 2008; Dordevic and others 2010; Fan and others 2010), have also been employed to ferment edible seeds and their products. The diversity of starter cultures provides multiple choices for fermentation, and combinations of different starter cultures may enhance fermentation efficiency, but this topic needs further investigation.

The inoculum amount of a starter culture is an important factor for the fermentation process. For LAB fermentation, inoculation of 1% to 10% (bacteria (mL)/sample (mL or g)) of the starter culture (10^8 cfu/mL) has been commonly employed in SSF and LSF of edible seeds and their products, with 10^6 to 10^7 cfu/mL LAB in original samples (Duenas and others 2005; Torino and others

2013; Gan and others 2016b). For *B. subtilis* and fungal fermentation, inoculation of 5% (bacteria (mL)/sample (mL)) of the starter culture (10^5 /g sample) has been mainly employed in SSF of edible seeds (Fernandez-Orozco and others 2007; Torino and others 2013; Limon and others 2015). Overall, the inoculum of starter cultures can vary due to different starter culture organisms and substrates, and optimization of the inoculum should be necessary to perform the fermentation efficiently.

Fermentation conditions

There are several factors, such as fermentation temperature, time, humidity, and other conditions, affecting the fermentation efficiency. Natural fermentation has been reported to control the temperature at 30, 37, or 42 °C (Elyas and others 2002; Granito and Alvarez 2006; Limon and others 2015), which is probably associated with the main microbes carried by different seeds. In addition, the temperature is commonly controlled at 37 °C for LAB fermentation (Frias and others 2005; Torino and others 2013), while fermentation using *B. subtilis* and fungi has mostly been employed at 30 °C (Fernandez-Orozco and others 2007; Lee and others 2008), probably due to the optimum growth at this temperature. For fermentation time, several hours to several days has been reported, while 48 and/or 96 h is most commonly used for edible seed fermentation. In addition, it is better to control the fermentation humidity at 90% to 95% if possible (Fernandez-Orozco and others 2007; Lee and others 2008), which can provide a relatively moist air condition for the growth of microbes. SSF of edible seeds and bean milk fermentation are generally performed quiescently, while LSF of edible seeds is commonly carried out by continuous shaking/stirring, with a speed of 200 to 450 rpm (Duenas and others 2005; Frias and others 2005; Torino and others 2013; Gan and others 2016b), which can accelerate the growth of microbes, increase the interaction between microbes and substrates, and enhance the efficiency of fermentation. Similarly, adding sugars (1% to 2%) to the fermentation system, such as glucose or sucrose (Jhan and others 2015; Gan and others 2016c), can provide an extra energy source to accelerate the growth of microbes. Moreover, fermentation can be performed under aerobic or anaerobic conditions, dependent on the species of microbes involved. For example, it is better to perform LAB fermentation in an anaerobic or microaerophilic environment. Overall, the fermentation condition is critical for the efficiency of fermentation and needs to be optimized for fermenting different products.

Treatment after fermentation

After fermentation, fermented edible seeds are commonly sterilized prior to further application. Autoclaving at 121 °C for 15 min coupled with subsequent freeze-drying is a common treatment (Gan and others 2016b). In addition, hot air drying, such as at 70 °C for 3 to 4 h, has also been used to treat fermented seeds (Elyas and others 2002). On the other hand, direct freeze-drying of fermented samples has also been reported (Frias and others 2005; Lee and others 2008; Jhan and others 2015), which can be better to retain nutritional and bioactive components. It should be mentioned that the step of sterilization after fermentation may be associated with potential food safety problems, which will be discussed in the final part of this review.

Table 1–Influences of fermentation on vitamins in edible seeds and their products.

Edible seeds and their products	Inoculated microbes	Vitamins	Vitamin content		Unit	References		
			Nonfermented samples	Fermented samples				
Kidney bean (<i>Phaseolus vulgaris</i>)	Natural fermentation	Vitamin B1	0.20	0.13 to 0.17	mg/100 g DW	(Granito and others 2002)		
		Vitamin B2	0.12	0.14 to 0.16	mg/100 g DW			
Lupin (<i>Lupinus albus</i> L. var. Multolupa)	Natural fermentation	Vitamin C	6.48	N.D.	mg/100 g DW	(Frias and others 2005)		
		α -Tocopherol	0.19	0.25	mg/100 g DW			
		γ -Tocopherol	20.1	18.9	mg/100 g DW			
		δ -Tocopherol	0.25	0.14	mg/100 g DW			
	<i>Lb. plantarum</i> CECT 748	Natural fermentation	Vitamin C	6.48	N.D.	mg/100 g DW		
			α -Tocopherol	0.19	0.08	mg/100 g DW		
			γ -Tocopherol	20.1	2.35	mg/100 g DW		
		<i>B. subtilis</i> (BCRC 14716) and <i>Lb. delbrueckii</i> sp. bulgaricus (BCRC 14008)	δ -Tocopherol	0.25	0.04	mg/100 g DW		
			Vitamin C	5.41	192	mg/g		
			Vitamin E	0.08	0.38	mg/g		
Red bean (<i>Phaseolus radiates</i>)	<i>B. subtilis</i> (BCRC 14716) and <i>Lb. delbrueckii</i> sp. bulgaricus (BCRC 14008)	Vitamin C	5.41	192	mg/g	(Jhan and others 2015)		
		Vitamin E	0.08	0.38	mg/g			
Soybean (<i>Glycine max</i> cv. Merit)	<i>A. oryzae</i> 2094 ^T (ATCC 1011)	α -Tocopherol	1.00	0.16	mg/100 g DW	(Fernandez-Orozco and others 2007)		
		β -Tocopherol	0.25	0.32	mg/100 g DW			
		γ -Tocopherol	4.10	12.8	mg/100 g DW			
		δ -Tocopherol	1.65	15.8	mg/100 g DW			
		<i>R. oryzae</i> CECT 2340 (ATCC 24563)	α -Tocopherol	1.00	0.32		mg/100 g DW	
			β -Tocopherol	0.25	0.35		mg/100 g DW	
			γ -Tocopherol	4.10	10.9		mg/100 g DW	
			δ -Tocopherol	1.65	16.2		mg/100 g DW	
		<i>B. subtilis</i> CECT 39 ^T (ATCC 6051)	α -Tocopherol	1.00	0.79		mg/100 g DW	
			β -Tocopherol	0.25	0.57		mg/100 g DW	
	γ -Tocopherol		4.10	15.4	mg/100 g DW			
	δ -Tocopherol		1.65	13.4	mg/100 g DW			
	<i>Lb. plantarum</i> CECT 748 ^T (ATCC 14917)	Natural fermentation	α -Tocopherol	1.00	0.01		mg/100 g DW	
			β -Tocopherol	0.25	0.22		mg/100 g DW	
			γ -Tocopherol	4.10	3.34		mg/100 g DW	
			δ -Tocopherol	1.65	5.93		mg/100 g DW	
		<i>Lb. plantarum</i> CECT 748 ^T (ATCC 14917)	Natural fermentation	α -Tocopherol	1.00		0.01	mg/100 g DW
				β -Tocopherol	0.25		0.21	mg/100 g DW
				γ -Tocopherol	4.10		3.41	mg/100 g DW
			<i>Lb. plantarum</i> CECT 748 ^T (ATCC 14917)	Natural fermentation	δ -Tocopherol		1.65	6.07
α -Tocopherol					1.00	0.01	mg/100 g DW	
β -Tocopherol					0.25	0.22	mg/100 g DW	
Soybean	Natural fermentation	γ -Tocopherol	4.10	3.33	mg/100 g DW			
		δ -Tocopherol	1.65	6.17	mg/100 g DW			
		α -Tocopherol	1.00	0.01	mg/100 g DW			
		β -Tocopherol	0.25	0.21	mg/100 g DW			
	<i>G. frondosa</i> GCMCC 5.00248	Natural fermentation	γ -Tocopherol	4.10	3.43	mg/100 g DW		
			δ -Tocopherol	1.65	6.17	mg/100 g DW		
			α -Tocopherol	1.00	0.01	mg/100 g DW		
		<i>G. frondosa</i> GCMCC 5.00248	Natural fermentation	Thiamin	3.27	10.8	mg/L	
				Niacin	3.84	8.74	mg/L	
				Riboflavin	0.81	0.54	mg/L	
Soymilk	<i>Lb. casei</i> Zhang	Vitamin B6	~2.00	20.9	mg/L	(Li and others 2012)		
		<i>Bb. animalis</i> V9 (CP001892)	Vitamin B6	~2.00	18.1		mg/L	
		<i>Lb. acidophilus</i> NCFM	Vitamin B6	~2.00	17.9		mg/L	
	<i>Lb. rhamnosus</i> GG	Natural fermentation	Vitamin B6	~2.00	14.2	mg/L	(Zhao and Shah 2014)	
			Vitamin B6	~2.00	13.4	mg/L		
		<i>Bb. animalis</i> BB12	Natural fermentation	Vitamin B6	~2.00	13.9		mg/L
				Vitamin C	431	16.2 to 227		μ g/g
	<i>Lb. acidophilus</i> CSCC 2400	Natural fermentation	Thiamin	35.9	41.3 to 61.1	μ g/g		
			Riboflavin	196	N.D.	μ g/g		
			α -Tocopherol	237	73.2 to 214	μ g/g		
<i>Lb. paracasei</i> CSCC 279		Natural fermentation	Vitamin C	431	70.2 to 256	μ g/g		
			Thiamin	35.9	4.10 to– 14.1	μ g/g		
			Riboflavin	196	179	μ g/g		

(Continued)

Table 1–Continued.

Edible seeds and their products	Inoculated microbes	Vitamins	Vitamin content		Unit	References	
			Nonfermented samples	Fermented samples			
Buckwheat groats (<i>Fagopyrum esculentum</i> Moench)	<i>Lb. zeae</i> ASCC 15820	α -Tocopherol	237	186 to 304	$\mu\text{g/g}$	(Malgorzata and others 2015)	
		Vitamin C	431	182 to 315	$\mu\text{g/g}$		
		Thiamin	35.9	7.20 to 16.6	$\mu\text{g/g}$		
		Riboflavin	196	185 to 186	$\mu\text{g/g}$		
		α -Tocopherol	237	343 to 361	$\mu\text{g/g}$		
	<i>Lb. rhamnosus</i> WQ2	Vitamin C	431	27.9 to 183	$\mu\text{g/g}$		
		Thiamin	35.9	13.8 to 30.1	$\mu\text{g/g}$		
		Riboflavin	196	N.D.	$\mu\text{g/g}$		
		α -Tocopherol	237	278 to 377	$\mu\text{g/g}$		
		Thiamin	9.70	10.9	$\mu\text{g/g}$		
	<i>R. oligosporus</i> NRRL 2710	Pyridoxine	1.65	1.70	$\mu\text{g/g}$		(Starzynska-Janiszewska and others 2016)
		Pyridoxine	120	330	$\mu\text{g/g}$		
		Vitamin C	50	110	$\mu\text{g/g}$		
		α -Tocopherol	0.73	1.10	$\mu\text{g/g}$		
		γ -Tocopherol	106	127	$\mu\text{g/g}$		
δ -Tocopherol		2.93	3.45	$\mu\text{g/g}$			
Thiamin		0.67	1.62	mg/kg DW			
Riboflavin		0.15	1.01	mg/kg DW			

A., *Aspergillus*; B., *Bacillus*; Bb, *Bifidobacterium*; G., *Grifola*; Lb., *Lactobacillus*; R., *Rhizopus*; N.D., not detected. DW, dry weight.

Influences of Fermentation on Bioactive Components Vitamins

Vitamins are important essential nutrients. Based on solubility, they can be divided into water-soluble and fat-soluble groups. The former mainly includes the vitamin B groups and vitamin C, and the latter mainly contains vitamin A, vitamin D and E groups, and vitamin K. Edible seeds, such as edible beans and cereal grains, are good natural sources of some vitamins, mainly the vitamin B and E groups (Fardet 2010; Hayat and others 2014), while fermentation has distinct influences on different vitamins in edible seeds and their products.

The vitamin B group mainly include thiamin (vitamin B1), riboflavin (vitamin B2), niacin (vitamin B3), pantothenic acid (vitamin B5), pyridoxine (vitamin B6), biotin (vitamin B7), folic acid (vitamin B9), and cobalamin (vitamin B12). Fermentation has been reported to exhibit varying effects on vitamin B members in edible seeds and their products (Table 1). Especially, fermentation by fungi and LAB, such as *G. frondosa*, *R. oligosporus*, and *Lb. acidophilus*, has been found to increase thiamin, riboflavin, niacin or pyridoxine in soymilk and buckwheat groats (Zhao and Shah 2014; Malgorzata and others 2015; Yang and others 2015; Starzynska-Janiszewska and others 2016). This suggests that some microbes may possess the capacity of producing B group vitamins, consistent with previous studies that some LAB and fungi can synthesize B vitamins (Strzelczyk and Leniarska 1985; Martens and others 2002; LeBlanc and others 2011; Capozzi and others 2012). The biosynthetic pathways of the main vitamin B group members in microbes have been reported (Magnusdottir and others 2015), and Figure 1A shows the proposed synthetic pathway of thiamin, one important B vitamin which is increased in some fermented edible seeds and their products (Table 1).

Vitamin C (also known as ascorbic acid) generally occurs at low levels in edible seeds, and fermentation can cause opposite effects on its content in different edible seeds and their products (Table 1). Fermentation was found to reduce ascorbic acid in lupin seeds and soymilk (Frias and others 2005, Zhao and Shah 2014), while increasing it in red bean and buckwheat groats (Jhan and

others 2015, Malgorzata and others 2015). This discrepancy may be associated with the degradation or biosynthesis of ascorbic acid by microbes in the fermentation process, since different microbes possess the capacity of decomposing or synthesizing it (Bremus and others 2006; Linster and Van Schaftingen 2007). Several LAB have been reported to degrade ascorbic acid into simple organic acids, such as acetic and lactic acids (Montano and others 2013), while yeasts, such as *S. cerevisiae*, have been demonstrated to biosynthesize L-ascorbic acid from L-galactose (Hancock and others 2000). In addition, metabolically engineered yeasts have been reported to synthesize L-ascorbic acid from D-glucose (Branduardi and others 2007), using the synthetic pathway (Figure 1B) found in plants. On the other hand, although *R. oligosporus*-fermented buckwheat groats and *B. subtilis* combined with *Lb. bulgaricus*-fermented red bean have been reported to increase ascorbic acid compared to unfermented samples (Jhan and others 2015; Malgorzata and others 2015), whether these microbes can biosynthesize ascorbic acid needs further investigation.

E vitamins include tocopherols and tocotrienols, each with α , β , γ , and δ homolog. In most edible seeds, γ -tocopherol is the most abundant vitamin E, with much higher content than other tocopherols (Table 1). Fermentation has been reported to differently influence the contents of tocopherols in edible seeds and their products (Table 1), probably associated with microbes involved in the fermentation process. Fungi, such as *A. oryzae*, *R. oryzae*, and *R. oligosporus*, have been found to increase tocopherols in soybean and buckwheat groats (Fernandez-Orozco and others 2007; Malgorzata and others 2015; Starzynska-Janiszewska and others 2016). *B. subtilis* was found to significantly increase γ and δ tocopherols in soybeans (Martens and others 2002), while LAB exhibited different influences on E vitamins (Frias and others 2005; Fernandez-Orozco and others 2007; Zhao and Shah 2014). These results suggest that some microbes may biosynthesize tocopherols during the fermentation process and a possible biosynthetic pathway of γ -tocopherol was proposed (Figure 1C) based on a previous study (Rippert and others 2004). On the other hand, the influence of fermentation on tocotrienols has scarcely been investigated.

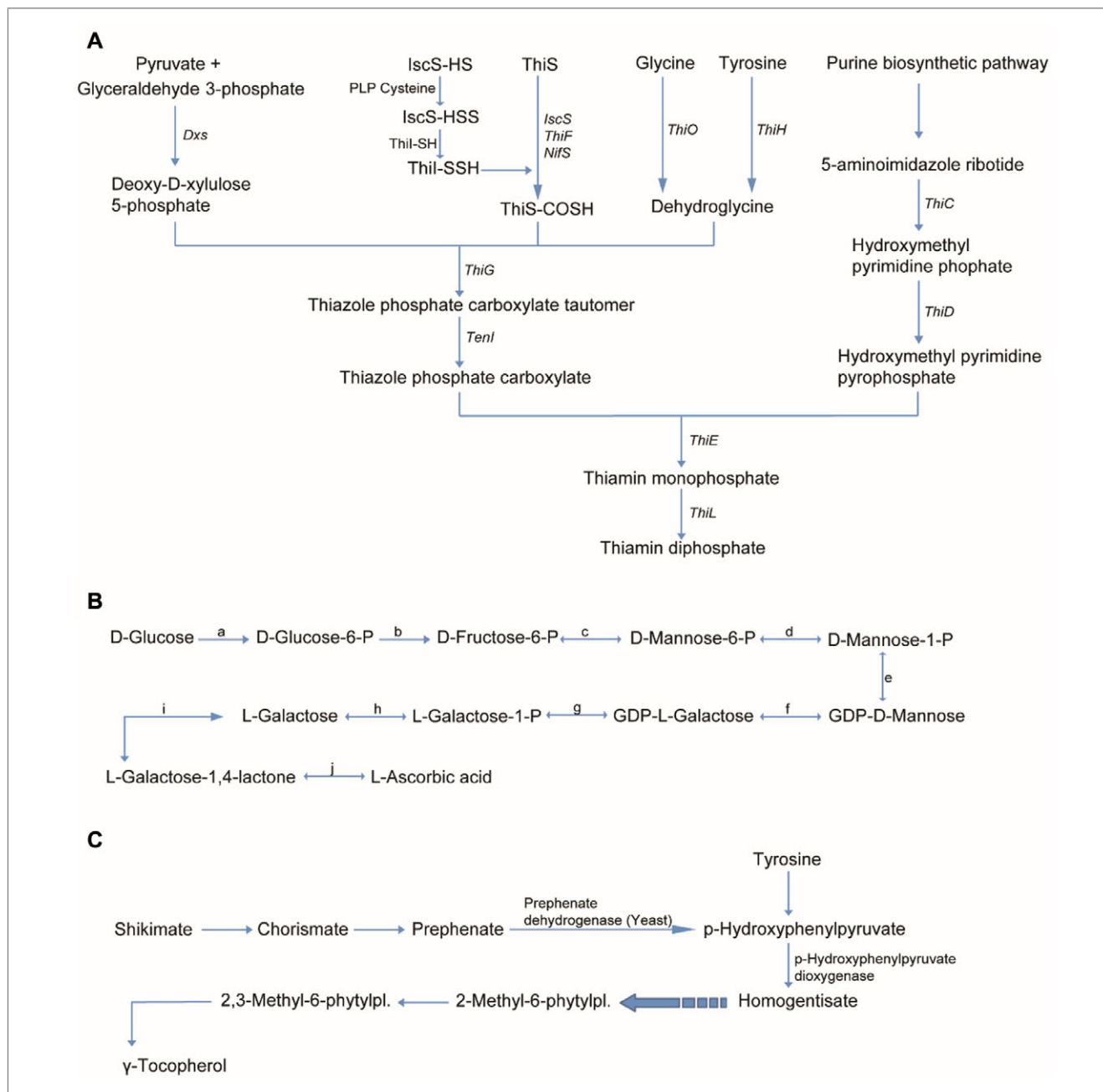


Figure 1—Proposed biosynthetic pathways of typical vitamins by microbes. (A) Proposed biosynthetic pathway of thiamin (vitamin B1), modified from Du and others (2011). Abbreviations: *Dxs*, 1-deoxy-D-xylulose 5-phosphate synthase; *IscS*, cysteine desulfurase; *NifS*, sulfur donor; *TenI*, transcriptional regulator TenI; *ThiC*, hydroxymethyl pyrimidine synthase; *ThiD*, hydroxymethyl pyrimidine (phosphate) kinase; *ThiE*, thiamin phosphate synthase; *ThiF*, adenyltransferase; *ThiG*, thiazole synthase; *ThiH*, thiazole synthase; *ThiI*, sulfur transferase; *ThiL*, thiamin phosphate kinase; *ThiO*, glycine oxidase; *ThiS*, sulfur carrier protein. (B) Proposed biosynthetic pathway of L-ascorbic acid (vitamin C), cited from Branduardi and others (2007). The enzymes a to h are as follows. a, hexokinase; b, glucose-6-phosphate isomerase; c, mannose-6-phosphate isomerase; d, phosphomannomutase; e, mannose-1-phosphate guanylyltransferase; f, GDP-mannose-3,5-epimerase; g, GDP-L-galactose phosphorylase; h, L-galactose 1-phosphate phosphatase; i, L-galactose dehydrogenase; j, L-galactono-1,4-lactone dehydrogenase. (C) Proposed biosynthetic pathway of γ -Tocopherol (vitamin E) based on a previous study (Rippert and others 2004).

In general, fermentation has distinct influences on various vitamins in edible seeds and their products, which can be associated with microbe-mediated biosynthesis or degradation of vitamins. In light of the health benefits of vitamins, it is promising to employ vitamin-producing microbes or genetically engineered microbes with enhanced vitamin-producing capacity to develop fermented edible seeds and their products rich in vitamins.

GABA

GABA, a nonprotein amino acid, is an important inhibitory neurotransmitter in the mammalian nervous system, and it plays a critical role in the regulation of blood pressure and many other physiological functions (Lee and Pan 2012; Diana and others 2014). It can be produced in plants, microorganisms, and mammals, and it is widely distributed in various foods (Jannoey and



Figure 2—Main bioactivities of fermented edible seeds and their products.

others 2010). Considering the health benefits of GABA, the need for GABA-rich foods is increasing, and various food processing methods, such as fermentation, have been employed to enhance its content (Dhakal and others 2012).

Edible seeds are natural sources of GABA, and many studies reported that fermentation can further increase GABA content in many edible seeds and their products (Table 2), such as adzuki bean, chickpea, faba bean, kidney bean, lentil, soybean, amaranth, buckwheat, millet, oat, quinoa, rice, rye, spelt, wheat, and/or their products. A variety of microorganisms has been employed in the fermentation process to enhance GABA content, and LAB are the most commonly used (Table 2). In addition, fungi and molds, such as *A. oryzae*, *R. oryzae*, and *M. purpureus*, are also used to enhance GABA content in edible seeds (Table 2).

Compared to germination where GABA is mainly synthesized by the seed itself, fermentation can accumulate GABA with the help of microorganisms. GABA is synthesized via glutamate decarboxylase (GAD)-mediated decarboxylation of glutamic acid, and GAD plays a central role in the synthesis of GABA (Battaglioli and others 2003). During the fermentation process, microbes can hydrolyze proteins and release free amino acids, such as glutamic acid, which can be used as the substrate for the synthesis of GABA by GABA-producing microbes. Many microbes have been reported

to produce GAD to synthesize GABA, such as LAB and fungi (Dhakal and others 2012; Wu and Shah 2016). In addition, it is also possible that GABA can be synthesized by the endogenous GAD of seeds, since its concentration has been reported to be increased in control samples without inoculated microbes (Rizzello and others 2008; Coda and others 2010). Besides, many factors, such as fermentation temperature, pH, and fermentation time, as well as different media additives, can influence the production of GABA, as previously reviewed by Dhakal and others (2012), and not further discussed herein. Overall, fermentation is a valuable bioprocessing strategy for producing GABA-rich products.

Natural phenolics

Natural phenolics, having at least one phenol group, and existing widely in the plant kingdom, are an important category of phytochemicals. They can be classified into different subgroups based on their chemical structures, such as phenolic acids, flavonoids, and proanthocyanidins. In plants, natural phenolics may exist in soluble and bound forms. Soluble phenolics include free and conjugated forms, with the latter conjugated with organic acids or sugar groups, and are synthesized through a multi-enzyme complex localized on the cytoplasmic surface of the endoplasmic reticulum, and they subsequently transported to intracellular vacuoles and

Table 2–Influences of fermentation on GABA content in edible seeds and their products.

Edible seeds and their products	Inoculated microbes	GABA content		Unit	References
		Nonfermented samples	Fermented samples		
Chickpea (<i>Cicer arietinum</i>)	<i>Lb. plantarum</i> C48	18.0	615	mg/kg DW	(Coda and others 2010)
	<i>Lc. lactis</i> subsp. <i>lactis</i> PU1	18.0	1031	mg/kg DW	
Faba bean (<i>Vicia faba</i> L.)	<i>Lb. plantarum</i> VTT E-133328	~50.0	~1000	mg/kg DW	(Coda and others 2015)
Kidney bean (<i>Phaseolus vulgaris</i>)	<i>B. subtilis</i> CECT 39 ^T (ATCC 6051)	3.20	2.61 to 2.69	mg/g DW	(Limon and others 2015)
	<i>Lb. plantarum</i> CECT 748 ^T (ATCC 14917)	2.27	6.76 to 9.90	mg/g DW	
Lentil (<i>Lens culinaris</i>)	Natural fermentation	2.27	7.54 to 10.6	mg/g DW	(Torino and others 2013)
	<i>Lb. plantarum</i> CECT 748 ^T (ATCC 14917)	~4.40	~6.00 to 7.16	mg/g DW	
	Natural fermentation	~4.40	~7.50 to 10.4	mg/g DW	
Soybean (<i>Glycine max</i>)	<i>B. subtilis</i> CECT 39 ^T (ATCC 6051)	~3.20	~4.00 to 6.54	mg/g DW	(Coda and others 2010)
	<i>Lb. plantarum</i> C48	24.0	57.0	mg/kg DW	
	<i>Lc. lactis</i> subsp. <i>lactis</i> PU1	24.0	39.0	mg/kg DW	
Amaranth (<i>Amaranthus hypocondriacus</i>)	<i>A. oryzae</i> FMB S46471 and <i>L. brevis</i> GABA100	~0.15	~3.60 to 6.60	g/kg DW	(Kim and Ji 2014)
	<i>Lb. plantarum</i> C48	12.0	654	mg/kg DW	(Coda and others 2010)
Buckwheat (<i>Fagopyrum esculentum</i>)	<i>Lc. lactis</i> subsp. <i>lactis</i> PU1	12.0	816	mg/kg DW	(Coda and others 2010)
	<i>Lb. plantarum</i> C48	40.0	643	mg/kg DW	
Millet (<i>Panicum miliaceum</i>)	<i>Lc. lactis</i> subsp. <i>lactis</i> PU1	40.0	562	mg/kg DW	(Coda and others 2010)
	<i>Lb. plantarum</i> C48	15.0	110	mg/kg DW	
Oat (<i>Avena sativa</i>)	<i>Lc. lactis</i> subsp. <i>lactis</i> PU1	15.0	231	mg/kg DW	(Cai and others 2014)
	<i>Lb. plantarum</i> C48	27.0	185	mg/kg DW	
	<i>Lc. lactis</i> subsp. <i>lactis</i> PU1	27.0	206	mg/kg DW	
	<i>A. oryzae</i> var. <i>effuses</i> 3.2825	57.1	59.0 to 331	μg/g DW	
	<i>A. oryzae</i> 3.5232	57.1	59.0 to 435	μg/g DW	
Quinoa (<i>Chenopodium quinoa</i>)	<i>R. oryzae</i> 3.275	57.1	59.0 to 126	μg/g DW	(Coda and others 2010)
	<i>Lb. plantarum</i> C48	78.0	415	mg/kg DW	
	<i>Lc. lactis</i> subsp. <i>lactis</i> PU1	78.0	176	mg/kg DW	
Rice (<i>Oryza sativa</i>)	<i>Lb. plantarum</i> C48	12.0	24.0	mg/kg DW	(Jannoey and others 2010)
	<i>Lc. lactis</i> subsp. <i>lactis</i> PU1	12.0	30.0	mg/kg DW	
	<i>M. purpureus</i> CMU 001	N.M.	1.24 to 28.4	mg/g DW	
	<i>Lc. lactis</i> RO50	N.M.	120 to 150	mg/100 g DW	
Rye (<i>Secale cereale</i>)	Several LAB	N.M.	11.7 to 168	mg/kg DW	(Saikusa and others 2016)
	<i>Lb. plantarum</i> C48	12.0	39.0	mg/kg DW	(Rizzello and others 2008)
	<i>Lc. lactis</i> subsp. <i>lactis</i> PU1	12.0	75.0	mg/kg DW	
Spelt (<i>Triticum spelta</i>)	<i>Lb. plantarum</i> C48	3.00	72.0	mg/kg DW	(Coda and others 2010)
	<i>Lc. lactis</i> subsp. <i>lactis</i> PU1	3.00	259	mg/kg DW	
White wheat (<i>Triticum aestivum</i> Appulo cv)	Several LAB	N.M.	12.7 to 101	mg/kg DW	(Rizzello and others 2008a)
Common wheat (<i>Triticum aestivum</i>)	<i>Lb. plantarum</i> C48	7.00	63.0	mg/kg DW	(Coda and others 2010)
Durum wheat (<i>Triticum durum</i>)	<i>Lc. lactis</i> subsp. <i>lactis</i> PU1	7.00	70.0	mg/kg DW	(Coda and others 2010)
	<i>Lb. plantarum</i> C48	19.0	75.0	mg/kg DW	
Adzuki bean milk	<i>Lc. lactis</i> subsp. <i>lactis</i> PU1	19.0	84.0	mg/kg DW	(Liao and others 2013)
	<i>Lc. lactis</i> subsp. <i>lactis</i> and <i>L. rhamnosus</i> GG	~28.0	~68.0	mg/100 mL	
Black bean milk	<i>Lb. brevis</i> FPA 3709	1.07	1.34 to 4.04	mg/mL	(Ko and others 2013)
Chickpea milk	<i>Lb. plantarum</i> M-6	~140	~150 to 537	mg/L	(Li and others 2016)
Soy milk	A mixture of LAB	~100	~130 to 180	mg/g DW	(Tsai and others 2006)
	<i>Lb. casei</i> Zhang	~24.5	~43.0	mg/L	(Li and others 2012)
	<i>Bb. animalis</i> V9 (CP001892)	~24.5	~45.6	mg/L	
	<i>Lb. acidophilus</i> NCFM	~24.5	~36.0	mg/L	
	<i>Lb. rhamnosus</i> GG	~24.5	~39.0	mg/L	
	<i>Bb. animalis</i> BB12	~24.5	~39.5	mg/L	
	<i>Lb. casei</i> Shirota	~24.5	~45.0	mg/L	
Soybean paste	Natural fermentation	~5.00	~5.00 to 80.0	μmol/g DW	(Jung and others 2016)
Oat flakes beverage	<i>Lb. plantarum</i> LPO9	~4.00	~5.00	mg/kg	(Nionelli and others 2014)
Rice-based sunsik	A mixture of microorganisms	26.0	66.0	mg/100 g DW	(Koh and others 2014)
Wheat-based dosa	<i>Lb. plantarum</i> MNZ	~0.00	~0.00 to 115	mg/kg	(Zareian and others 2015)
Wholemeal wheat	Several LAB	N.M.	11.0 to 259	mg/kg DW	(Rizzello and others 2008)

A., *Aspergillus*; B., *Bacillus*; Bb, *Bifidobacterium*; Lb., *Lactobacillus*; R., *Rhizopus*; N.D., not detected. DW, dry weight.

stored there, while bound phenolics are formed by the secretion of vacuolar soluble phenolics to the cell wall, where they can covalently bind with cell wall macromolecules, such as polysaccharides and proteins, as a component of plant cell walls (Agati and others 2012). Edible seeds, such as edible beans and cereal grains, have been reported to contain a variety of natural phenolics (Deng and others 2012; Gan and others 2016d). Phenolics mainly exist in the pigmented seed coats, primarily as flavonoids and proanthocyanidins (Gan and others 2016a), and many edible seeds contain a substantial level of bound phenolics, generally with much higher content than in common fruits and vegetables (Gan and others 2016a).

Recent studies indicate that fermentation can change phenolic composition and distribution of edible seeds and their products. Most studies report that fermentation increases the total phenolics content (TPC) of soluble phenolics, while having different effects on specific phenolic compounds (Table 3). This can be partly due to the metabolism of soluble phenolic compounds by microbes during the fermentation process, since many microbes, such as LAB and fungi, are able to produce different enzymes, such as β -glucosidase, esterase, and tannase, to metabolize soluble phenolics and/or phenolic polymers into free forms or degraded products (Rodriguez and others 2009; Hole and others 2012; Huynh and others 2014). For instance, fermentation of soybean and soy products by many microbes has been found to convert soybean isoflavone glucosides (such as daidzin) into their aglycones (such as daidzein). Our work (Gan and others 2016b) found that natural and LAB-mediated fermentation increased soluble TPC in mottled cowpea, accompanied with a significant increase of soluble catechin, which was speculated to be associated with the degradation of proanthocyanidins in the bean coat. Additionally, soluble phenolics can be released from bound phenolics during the fermentation process. Our work (Gan and others 2016b) also indicated that fermentation significantly increased soluble TPC in small runner bean, partly due to the release of bound phenolics. This may be related to microbe-mediated decomposition of cell wall components and subsequent release of bound phenolics, since many microbes possess various enzymes, such as cellulase, feruloyl esterase, glucosidase, xylanase, pectinase, and proteinases, which can degrade the cell wall matrix (Huynh and others 2014). Taken together, microbe-mediated metabolism of soluble phenolics and release of bound phenolics may both contribute to the increase of soluble TPC in some fermented edible seeds and their products. However, several studies also found that fermentation did not increase, or may even reduce, soluble TPC in some edible seeds and their products, such as lentil (Torino and others 2013), black soybean (Lee and others 2008), wheat (Subramaniam and others 2014), soymilk (Rekha and Vijayalakshmi 2008), buckwheat groats (Malgorzata and others 2015), and black rice bran (Yoon and others 2015), probably associated with the degradation of some phenolics during the fermentation process.

Although edible seeds contain substantial levels of bound phenolics, the influence of fermentation on TPC and phenolic composition of their bound phenolics has been not much investigated. Our recent work (Gan and others 2016b) found that fermentation exhibited different influences on bound TPC in selected edible beans. On one hand, bound TPC was found reduced in naturally fermented small runner bean and LAB-fermented lentil, small rice bean, and small runner bean, suggesting the release of bound phenolics during fermentation. On the other hand, bound TPC was found significantly increased after fermentation of some other beans, implying that bound phenolics were not released from

cell walls, while becoming more available for extraction. In addition, the influence of fermentation on specific bound phenolic compounds has also been scarcely studied. We found that natural fermentation of mottled cowpea increased the contents of bound ferulic and *p*-coumaric acids and reduced the content of bound protocatechuic acid, while LAB-mediated fermentation did not evidently change them (Gan and others 2016b). Another study reported that bound ferulic acid and *p*-coumaric acid increased in LAB-fermented barley, but decreased in LAB-fermented oat groats, probably releasing from cell wall matrix and become soluble phenolics (Hole and others 2012). These limited results suggest that fermentation can, overall, improve the bioavailability of bound phenolics in edible seeds and may play an important role in gut health, therefore, more studies are needed to further investigate the specific influence of fermentation on bound phenolics in different edible seeds.

Bioactive peptides

Edible seeds, such as edible beans and cereal grains, are rich in proteins, which can be hydrolyzed into small-molecule peptides during the fermentation process. Recent studies have found that some fermented edible seeds and their products can produce bioactive peptides (Table 4), mainly antioxidant and angiotensin-converting-I-enzyme (ACE) inhibitory peptides, which are discussed below.

Antioxidant peptides exhibit various antioxidant activities, such as reducing, free radical-scavenging, inhibition of lipid peroxidation, and metal ion chelation properties, which are mainly associated with the intrinsic characteristics of peptides, such as their amino acid composition, structure, and hydrophobicity. Table 4 summarizes the main antioxidant peptides derived from fermented edible seeds and their products, and some intrinsic features and antioxidant mechanisms of antioxidant peptides have been proposed based on previous studies. (1) They, in general, contain 2 to 20 amino acids, all with molecular weight lower than 6.0 kDa (Coda and others 2012). (2) The existence of amino acids, such as A (alanine), C (cysteine), H (histidine), K (lysine), L (leucine), M (methionine), P (proline), V (valine), W (tryptophan), and Y (tyrosine), may contribute to the antioxidant activity of peptides (Sarmadi and Ismail 2010). (3) Amino acid residues, including C, D (aspartic acid), E (glutamic acid), H, K, M, R (arginine), W, and Y, can be associated with the chelating activity of antioxidant peptides (Wu and others 2014). (4) Hydrophobic amino acids can enhance the solubility of peptides in the oil environment, therefore facilitating the interaction with lipophilic radical species and polyunsaturated fatty acids (PUFAs) (Coda and others 2012). (5) The sulfur group (SH group) of cysteine and of methionine exhibits antioxidant activity, since it is able to neutralize reactive free radical species to form stable oxidation products, cystine, and methionine sulfoxide, respectively (He and others 2012). (6) Histidine-containing peptides exhibit strong radical-scavenging activity due to decomposition of the imidazole ring (He and others 2012). (7) Histidine mainly acts as a chelator of metal ions at the amino terminus of peptides, while acting as an effective scavenger against various radicals at the carboxyl terminus (Coda and others 2012). (8) Peptides with aromatic amino acid residues, including F (phenylalanine), Y and W, are good donors of hydrogen and can efficiently scavenge free radicals due to their conjugated double-bond structure of a benzene ring (Zhang and others 2014). These characteristics of antioxidant peptides can be helpful for predicting other potential antioxidant peptides, and they can also provide a reference for the chemical synthesis of antioxidant peptides.

Table 3—Influences of fermentation on phenolic composition in edible seeds and their products.

Edible seeds and their products	Inoculated microbes	Extracts	TPC		Unit	Changes of main phenolic compounds	References
			Nonfermented samples	Fermented samples			
Chickpea (<i>Cicer arietinum</i> L.)	<i>R. oligosporus</i> NRRL 2710	80% Ethanol	93.0	117 to 631	mg GAE/100 g DW	N.M.	(Sanchez-Magana and others 2014)
	<i>R. oligosporus</i> NRRL 2710	NaOH hydrolysis	128	143 to 205	mg GAE/100 g DW	N.M.	
	<i>C. militaris</i> SN-18	80% Methanol	7.08	12.8	mg GAE/g DW	Shikimic acid (↑), chlorogenic acid (↑), daidzein (↑), genistein (↑), biochanin A (↑), rutin (↑), <i>p</i> -coumaric acid (↔), syringic acid (↓), ferulic acid (↓), daidzin (↓), glycitin (↓), genistin (↓), luteolin (↓)	(Xiao and others 2014)
Common bean (<i>Phaseolus vulgaris</i>)	<i>C. militaris</i> SN-18	80% Ethanol	7.36	13.3	mg GAE/g DW	N.M.	
	<i>C. militaris</i> SN-18	Water	6.07	10.5	mg GAE/g DW	N.M.	
	<i>Lb. plantarum</i> DSM 20174 and <i>R. microsporus</i> var. chinensis	PBS	1.60	1.61	mg TAE/g DW	N.M.	(Starzynska-Janiszewska and others 2014)
Black cow gram (<i>Labiab purpureus</i>)	<i>R. oligosporus</i> NRRL2710	80% Acetone	1.49	1.69	mg TAE/g DW	N.M.	
	Natural fermentation	70% Acetone	~2.85	~2.00 to 2.50	mg GAE/g DW	N.M.	(Camboa-Gomez and others 2016)
	<i>R. oligosporus</i> NRRL2710	70% Acetone	~2.50	~1.35 to 2.10	mg GAE/g DW	N.M.	
Mottled cowpea (<i>Vigna unguiculata</i>)	Natural fermentation	80% Methanol	190	287	mg GAE/100 g DW	N.M.	(Can and others 2016b)
	<i>Lb. paracasei</i> 279	NaOH + HCl hydrolysis	190	277	mg GAE/100 g DW	N.M.	
	Natural fermentation	80% Methanol	69.9	85.6	mg GAE/100 g DW	N.M.	
Mottled cowpea (<i>Vigna unguiculata</i>)	<i>Lb. paracasei</i> 279	80% Methanol	69.9	76.8	mg GAE/100 g DW	N.M.	
	Natural fermentation	80% Methanol	69.9	90.7	mg GAE/100 g DW	N.M.	
	<i>Lb. paracasei</i> 279	80% Methanol	400	626	mg GAE/100 g DW	N.M.	
Mottled cowpea (<i>Vigna unguiculata</i>)	Natural fermentation	80% Methanol	400	752	mg GAE/100 g DW	N.M.	
	<i>Lb. plantarum</i> WCFS1	NaOH + HCl hydrolysis	400	747	mg GAE/100 g DW	N.M.	
	Natural fermentation	80% Methanol	103	186	mg GAE/100 g DW	N.M.	

(Continued)

Table 3—Continued.

Edible seeds and their products	Inoculated microbes	Extracts	TPC		Unit	Changes of main phenolic compounds	References
			Nonfermented samples	Fermented samples			
Kidney bean (<i>Phaseolus vulgaris</i>)	<i>Lb. paracasei</i> 279	NaOH + HCl hydrolysis	103	105	mg GAE/100 g DW	Protocatechuic acid (↔), ferulic acid (↔), <i>p</i> -coumaric acid (↔)	(Limon and others 2015)
	<i>Lb. plantarum</i> WCFS1	NaOH + HCl hydrolysis	103	126	mg GAE/100 g DW	Protocatechuic acid (↔), ferulic acid (↔), <i>p</i> -coumaric acid (↓)	
	Natural fermentation	Water	20.7	20.2 to 21.2	mg GAE/g DW	Sinapoyl aldaric acid (↑), ferulic acid (↑), catechin and its derivatives (↓), <i>p</i> -hydroxybenzoic acid (↓), sinapoyl methylaldaric acid (↓), naringenin and its derivatives (↓)	
Speckled kidney bean (<i>Phaseolus vulgaris</i>)	Natural fermentation	80% acetone	0.25	0.42	mg TAE/g WW	N.M.	(Oboh and others 2009)
	Natural fermentation	NaOH hydrolysis	0.32	0.17	mg TAE/g WW	N.M.	
	<i>B. subtilis</i> CECT 39 ^T (ATCC 6051)	Water	15.9	31.1 to 35.9	mg GAE/g DW	Ferulic acid and its derivatives (↑), <i>p</i> -hydroxybenzoic acid (↑), hydroxycinnamic compounds (↑), catechin (↓), <i>p</i> -coumaric acid (↓)	
Lentil (<i>Lens culinaris</i>)	<i>Lb. plantarum</i> CECT 748 ^T (ATCC 14917)	Water	20.7	17.8 to 22.0	mg GAE/g DW	Sinapoyl aldaric acid (↑), catechin and its derivatives (↓), <i>p</i> -hydroxybenzoic acid (↓), sinapoyl methylaldaric acid (↓), naringenin and its derivatives (↓), ferulic acid (↓), hydroxycinnamic compounds (↓)	(Limon and others 2015)
	Natural fermentation	80% Methanol	248	441	mg GAE/100 g DW	N.M.	
	<i>Lb. paracasei</i> 279 <i>Lb. plantarum</i> WCFS1 Natural fermentation	NaOH + HCl hydrolysis	248 248 106	489 509 110	mg GAE/100 g DW mg GAE/100 g DW mg GAE/100 g DW	N.M. N.M. N.M.	
Lentil (<i>Lens culinaris</i>)	<i>Lb. paracasei</i> 279 <i>Lb. plantarum</i> WCFS1 Natural fermentation	Water	106 106 ~31.0	103 106 ~28.0 to 31.0	mg GAE/100 g DW mg GAE/100 g DW mg GAE/g DW	N.M. N.M. N.M.	(Torino and others 2013)

(Continued)

Table 3–Continued.

Edible seeds and their products	Inoculated microbes	Extracts	TPC		Unit	Changes of main phenolic compounds	References
			Nonfermented samples	Fermented samples			
Mung bean (<i>Vigna radiata</i>)	<i>B. subtilis</i> CECT 39 ^T (ATCC 6051)	Water	~24.0	~35.0 to 36.0	mg GAE/g DW	N.M.	(Can and others 2016b)
	<i>Lb. plantarum</i> CECT 748 ^T (ATCC 14917)	Water	~31.0	~27.5 to 29.0	mg GAE/g DW	N.M.	
	Natural fermentation	80% Methanol	220	308	mg GAE/100 g DW	N.M.	
	<i>Lb. paracasei</i> 279	NaOH + HCl hydrolysis	220	341	mg GAE/100 g DW	N.M.	
	<i>Lb. plantarum</i> WCFS1		220	325			
	Natural fermentation		81.1	101			
	<i>Lb. paracasei</i> 279	Acidified 80% methanol	81.1	61.3	-	Catechin (↑), eriodictyol (↑), taxifolin (↑), <i>p</i> -hydroxyphenylacetic acid (↑), eriodictyol (↑), 7-O-glucoside (↓)	
	<i>Lb. plantarum</i> WCFS1		81.1	65.4			
	<i>Lb. plantarum</i> CECT 748 ^T (ATCC 14917)		N.M.	N.M.			
	Pigeon pea (<i>Cajanus cajan</i>)	<i>C. militaris</i> SN-18	80% methanol	1372	2972	μg GAE/g DW	
<i>C. militaris</i> SN-18		80% ethanol	1480	2696	μg GAE/g DW	Chlorogenic acid (↑), vanillic acid (↑), sinapic acid (↑), rutin (↑), luteolin (↑), kaempferol (↑), shikimic acid (↔)	(Xiao and others 2015d)
<i>C. militaris</i> SN-18		80% acetone	1649	3124	μg GAE/g DW	Chlorogenic acid (↑), vanillic acid (↑), sinapic acid (↑), rutin (↑), luteolin (↑), kaempferol (↑), shikimic acid (↔)	(Obob and others 2009)
<i>C. militaris</i> SN-18		Water	1146	5680	μg GAE/g DW	Shikimic acid (↑), chlorogenic acid (↑), vanillic acid (↑), sinapic acid (↑), rutin (↑), luteolin (↑), kaempferol (↑)	(Jhan and others 2015)
Red bean (<i>Phaseolus radiatus</i>)	Natural fermentation	80% acetone	0.24	0.46	mg TAE/g WW	N.M.	(Jhan and others 2015)
	Natural fermentation	NaOH hydrolysis	0.58	0.34	mg TAE/g WW	N.M.	
	<i>B. subtilis</i> BCRC 14716 and <i>Lb. delbrueckii</i> sp. <i>bulgaricus</i> BCRC 14008	Water	2.30	3.25	mg GAE/g DW	N.M.	

(Continued)

Table 3–Continued.

Edible seeds and their products	Inoculated microbes	Extracts	TPC		Unit	Changes of main phenolic compounds	References
			Nonfermented samples	Fermented samples			
Small rice bean (<i>Vigna umbellata</i>)	<i>B. subtilis</i> BCRC 14716 and <i>Lb. delbrueckii</i> sp. bulgaricus BCRC 14008	50% Ethanol	2.59	3.63	mg GAE/g DW	N.M.	
	Natural fermentation	80% Methanol	338	588	mg GAE/100 g DW	N.M.	(Can and others 2016b)
	<i>Lb. paracasei</i> 279		338	692	mg GAE/100 g DW	N.M.	
	<i>Lb. plantarum</i> WCFS1		338	666	mg GAE/100 g DW	N.M.	
	Natural fermentation	NaOH + HCl hydrolysis	79.6	128	mg GAE/100 g DW	N.M.	
	<i>Lb. paracasei</i> 279		79.6	62.7	mg GAE/100 g DW	N.M.	
	<i>Lb. plantarum</i> WCFS1		79.6	70.7	mg GAE/100 g DW	N.M.	
	Natural fermentation	80% Methanol	211	360	mg GAE/100 g DW	N.M.	(Can and others 2016b)
	<i>Lb. paracasei</i> 279		211	469	mg GAE/100 g DW	N.M.	
	<i>Lb. plantarum</i> WCFS1		211	449	mg GAE/100 g DW	N.M.	
Small runner bean (<i>Phaseolus coccineus</i>)	Natural fermentation	NaOH + HCl hydrolysis	219	132	mg GAE/100 g DW	N.M.	
	<i>Lb. paracasei</i> 279		219	93.8	mg GAE/100 g DW	N.M.	
	<i>Lb. plantarum</i> WCFS1		219	132	mg GAE/100 g DW	N.M.	
	<i>Lb. plantarum</i> KFRI 00144; <i>Lb. delbrueckii</i> subsp. <i>latis</i> KFRI 01181; <i>Bb. thermophilum</i> KFRI 00748; <i>Bb. breve</i> K-101	80% Ethanol	N.M.	N.M.			(Pyo and others 2005)
	Natural fermentation	80% Methanol	2.98	4.98 to 5.68	mg CE/g DW	N.M.	(Fernandez-Orozco and others 2007)
	<i>A. oryzae</i> 2094 ^T (ATCC 10111)	80% Methanol	2.98	3.54	mg CE/g DW	N.M.	
	<i>R. oryzae</i> CECT 2340 (ATCC 24563)	80% Methanol	2.98	3.43	mg CE/g DW	N.M.	
	<i>B. subtilis</i> CECT 39 ^T (ATCC 6051)	80% Methanol	2.98	12.5	mg CE/g DW	N.M.	
	<i>Lb. plantarum</i> CECT 748 ^T (ATCC 14917)	80% Methanol	2.98	5.40 to 5.51	mg CE/g DW	N.M.	

(Continued)

Table 3–Continued.

Edible seeds and their products	Inoculated microbes	Extracts	TPC		Unit	Changes of main phenolic compounds	References
			Nonfermented samples	Fermented samples			
Black soybean (<i>Glycine max</i>)	<i>Lb. plantarum</i> CECT 748 ^a (ATCC 14917)	Acidified 80% methanol	N.M.	N.M.	-	Daidzein (↑), genistein (↑), glycitin (↑), naringenin (↑), kaempferol diglucoside (↑), <i>p</i> -hydroxyphenylacetic acid (↑), daidzin (↓), genistin (↓), glycitin (↓), malonyl and acetyl daidzin (↓), malonyl and acetyl genistin (↓), malonyl glycitin (↓), eriodictyol 7-O-glucoside (↓), kaempferol glucoside (↓)	(Landete and others 2015)
	Natural fermentation	80% Methanol	219	415	mg GAE/100 g DW	N.M.	(Gan and others 2016b)
	<i>Lb. paracasei</i> 279		219	395	mg GAE/100 g DW	N.M.	
	<i>Lb. plantarum</i> WCFS1	NaOH + HCl hydrolysis	219	384	mg GAE/100 g DW	N.M.	
	Natural fermentation		42.5	69.9	mg GAE/100 g DW	N.M.	
	<i>Lb. paracasei</i> 279		42.5	67.0	mg GAE/100 g DW	N.M.	
	<i>Lb. plantarum</i> WCFS1		42.5	54.0	mg GAE/100 g DW	N.M.	
	<i>R. oryzae</i> BCRC 30894,	80% Methanol	N.D.	2088 to 2545	μg GAE/g DW	Daidzein and genistein (↑), daidzin and genistin (↓)	(Cheng and others 2013)
	<i>R. oligosporus</i> NTU-5,						
	<i>R. oligosporus</i> BCRC 31996						
	<i>B. subtilis</i> BCRC 14715	80% Methanol	15.9	23.4	mg GAE/g DW	N.M.	(Juan and Chou 2010)
	<i>B. subtilis</i> BCRC 14715	80% Ethanol	17.8	22.7	mg GAE/g DW	N.M.	
	<i>B. subtilis</i> BCRC 14715	80% Acetone	26.6	40.4	mg GAE/g DW	N.M.	
	<i>B. subtilis</i> BCRC 14715	Water	6.04	12.4	mg GAE/g DW	N.M.	
	<i>A. awamori</i>	Methanol	~16.4	~27.2	mg GAE/g DW	Cyaniding 3-glucoside (↑)	(Lee and others 2008)
<i>A. oryzae</i> BCRC 30222	Methanol	~16.4	~17.8	mg GAE/g DW	Cyaniding 3-glucoside (↑)		
<i>A. sojae</i> BCRC 30103	Methanol	~16.4	~16.6	mg GAE/g DW	Cyaniding 3-glucoside (↑)		
<i>R. zygosporus</i> BCRC 31158	Methanol	~16.4	~20.0	mg GAE/g DW	Cyaniding 3-glucoside (↑)		
<i>Rhizopus</i> sp. No. 2	Methanol	~16.4	~25.0	mg GAE/g DW	Cyaniding 3-glucoside (↔)		
Natural fermentation	80% Methanol	414	547	mg GAE/100 g DW	N.M.	(Gan and others 2016b)	
<i>Lb. paracasei</i> 279		414	538	mg GAE/100 g DW	N.M.		
<i>Lb. plantarum</i> WCFS1		414	582	mg GAE/100 g DW	N.M.		

(Continued)

Table 3–Continued.

Edible seeds and their products	Inoculated microbes	Extracts	TPC		Unit	Changes of main phenolic compounds	References
			Nonfermented samples	Fermented samples			
African yam bean (<i>Sphenostylis stenocarpa</i> Harms)	Natural fermentation	NaOH + HCl hydrolysis	65.7	91.7	mg GAE/100 g DW	N.M.	(Oboh and others 2009)
	<i>Lb. paracasei</i> 279		65.7	78.3	mg GAE/100 g DW	N.M.	
	<i>Lb. plantarum</i> WCFS1		65.7	85.4	mg GAE/100 g DW	N.M.	
	Natural fermentation	80% Acetone	0.33	0.47	mg TAE/g WW	N.M.	
African locust bean (<i>Parkia biglobosa</i>)	Natural fermentation	NaOH hydrolysis	0.26	0.20	mg TAE/g WW	N.M.	(Oboh and others 2008)
	Natural fermentation	80% Acetone	0.61	0.68	mg TAE/g DW	N.M.	
Barley (<i>Hordeum vulgare</i> L.)	<i>Lb. johnsonii</i> LA1, <i>Lb. reuteri</i> SD2112, <i>Lb. acidophilus</i> LA-5	50% Methanol	N.M.	N.M.	-	Ferulic acid (↑), <i>p</i> -hydroxybenzoic acid (↑)	(Hole and others 2012)
	<i>Lb. rhamnosus</i>	NaOH hydrolysis 70% ethanol	N.M.	N.M.	-	Ferulic acid (↑), <i>p</i> -coumaric acid (↑)	
Buckwheat (<i>Fagopyrum esculentum</i>)	<i>S. cerevisiae</i>	70% ethanol	16.4	18.5	mg GAE/g DW	N.M.	(Dordevic and others 2010)
	<i>Lb. rhamnosus</i>	70% ethanol	50.7	59.4	mg GAE/g DW	N.M.	
Maize (<i>Zea mays</i> subsp. <i>mays</i>)	<i>S. cerevisiae</i>	70% ethanol	50.7	53.2	mg GAE/g DW	N.M.	(Dey and Kuhad 2014)
	<i>T. elegans</i> CCF-1456	Water	0.81	~1.00 to 4.00	mg GAE/g DW	N.M.	
Oat (<i>Avena sativa</i> L.)	<i>T. elegans</i> CCF-1456	65% ethanol	327	340 to 382	μmol GAE/g DW	N.M.	(Salar and others 2012)
	<i>A. oryzae</i> NCIM 1212, <i>A. awamori</i> MTCC No. 548, <i>R. oligosporus</i> NCIM 1215, <i>R. oryzae</i> RCK2012	Water	2.06	~2.30 to 6.40	mg GAE/g DW	N.M.	
<i>A. oryzae</i> var. <i>effuses</i>	<i>A. oryzae</i> var. <i>effuses</i>	80% ethanol	~105	~212	mg GAE/g DW	Chlorogenic acid (↑), caffeic acid (↑), <i>p</i> -coumaric acid (↑), ferulic acid (↑)	(Cai and others 2012)
		80% ethanol	~105	~186	mg GAE/g DW	Chlorogenic acid (↑), caffeic acid (↑), <i>p</i> -coumaric acid (↑), ferulic acid (↑)	
		80% ethanol	~105	~155	mg GAE/g DW	Chlorogenic acid (↓), caffeic acid (↑), <i>p</i> -coumaric acid (↑), ferulic acid (↔)	
<i>A. oryzae</i> var. <i>effuses</i> 3-2825, <i>A. oryzae</i> 3-5232, <i>R. oryzae</i> 3-275		80% ethanol	0.50	0.80 to 1.80	mg GAE/g DW	N.M.	(Cai and others 2014)

(Continued)

Table 3–Continued.

Edible seeds and their products	Inoculated microbes	Extracts	TPC		Unit	Changes of main phenolic compounds	References
			Nonfermented samples	Fermented samples			
Brown rice (<i>Oryza sativa</i>)	<i>C. militaris</i> SN-18	Water	5.88	14.1	mg GAE/g DW	Gallic acid (↑), <i>p</i> -hydroxybenzoic acid (↑), caffeic acid (↑), <i>p</i> -coumaric acid (↑), ferulic acid (↑), vanillin (↑), avenanthramide 2c,p,f (↑), luteolin (↑), apigenin (↑)	(Xiao and others 2015a)
			10.9	16.8	mg GAE/g DW		
			10.1	15.0	mg GAE/g DW		
			12.4	19.7	mg GAE/g DW		
Rye (<i>Secale cereale</i>)	<i>A. oryzae</i> NCIM 1212, <i>A. awamori</i> MTCC No. 548, <i>R. oligosporus</i> NCIM 1215, <i>R. oryzae</i> RCK2012	80% methanol	0.79	~3.20 to 7.80	mg GAE/g DW	N.M.	(Dey and Kuhad 2014)
			13.2	18.4	mg GAE/g DW		
Wheat (<i>Triticum aestivum</i> Linn.)	<i>S. cerevisiae</i>	70% ethanol	13.2	16.2	mg GAE/g DW	N.M.	(Dordevic and others 2010)
			36.9	41.6	mg GAE/g DW		
Wheat (<i>Triticum durum</i>)	<i>C. militaris</i>	70% ethanol	54.2	66.4	mg GAE/g DW	Ferulic acid (↑), <i>p</i> -coumaric acid (↑), syringic acid (↑), vanillic acid (↔), caffeic acid (↔)	(Zhang and others 2012)
			16.2	20.7	mg GAE/g DW		
Wheat (<i>Triticum spp.</i>)	<i>C. militaris</i>	70% acetone	16.2	18.4	mg GAE/g DW	N.M.	(Dordevic and others 2010)
			0.81	~2.30 to 11.6	mg GAE/g DW		
Wheat (<i>Triticum spp.</i>)	<i>A. oryzae</i> NCIM 1212, <i>A. awamori</i> MTCC No. 548, <i>R. oligosporus</i> NCIM 1215, <i>R. oryzae</i> RCK2012	70% ethanol	27.4	22.6	mg GAE/g DW	N.M.	(Subramaniam and others 2014)
			56.0	49.7	mg GAE/g DW		
Adlay (<i>Coix lachryma-jobi</i>)	<i>Gd. neo-japonicum</i> (KUM61076)	Water	27.4	32.6	mg GAE/g DW	N.M.	(Dey and Kuhad 2014)
			56.0	61.4	mg GAE/g DW		
			27.4	17.7	mg GAE/g DW		
			56.0	32.9	mg GAE/g DW		
Bambara groundnut (<i>Vigna subterranea</i>)	<i>B. subtilis</i> BCRC 14718	Methanol	8.58	13.3	mg GAE/g DW	N.M.	(Wang and others 2014)
			0.29	0.27	mg TAE/g WW		
Bambara groundnut (<i>Vigna subterranea</i>)	<i>Lb. plantarum</i>	Methanol	8.58	11.9	mg GAE/g DW	N.M.	(Oboh and others 2009)
			0.60	0.75	mg TAE/g WW		
Bambara groundnut (<i>Vigna subterranea</i>)	Natural fermentation	80% Acetone	0.29	0.27	mg TAE/g WW	N.M.	(Oboh and others 2009)
			0.29	0.27	mg TAE/g WW		
		NaOH hydrolysis					(Continued)

Table 3–Continued.

Edible seeds and their products	Inoculated microbes	Extracts	TPC		Unit	Changes of main phenolic compounds	References
			Nonfermented samples	Fermented samples			
Chestnut (<i>Castanea crenata</i>)	<i>B. subtilis</i> BCRC 14718	Methanol	13.0	19.7	mg GAE/g DW	N.M.	(Wang and others 2014)
Lotus seed (<i>Nelumbo nucifera</i>)	<i>Lb. plantarum</i>	Methanol	13.0	15.3	mg GAE/g DW	N.M.	
	<i>B. subtilis</i> BCRC 14718	Methanol	17.5	28.7	mg GAE/g DW	N.M.	
Walnut (<i>Juglans regia</i>)	<i>Lb. plantarum</i>	Methanol	17.5	24.6	mg GAE/g DW	N.M.	
	<i>B. subtilis</i> BCRC 14718	Methanol	22.8	33.9	mg GAE/g DW	N.M.	
Mung bean milk	<i>Lb. plantarum</i>	Methanol	22.8	28.6	mg GAE/g DW	N.M.	(Gan and others 2016c)
	<i>Lb. plantarum</i> WCFS1	THF	29.1	30.7 to 33.6	mg GAE/100 mL	N.M.	
Soymilk	<i>Lb. acidophilus</i> B4496 + <i>S. boulardii</i>	Acidic methanol	12.6	15.9 to 17.4	mg GAE/100 mL	Vitexin and isovitexin (↔)	(Rekha and Vijayalakshmi 2008)
	<i>Lb. bulgaricus</i> CFR 2028 + <i>S. boulardii</i>	Water	26.6	23.4	mg GAE/100 mL	Daidzein (↑), genistein (↑), daidzin (↓), genistin (↓)	
	<i>Lb. casei</i> B1922 + <i>S. boulardii</i>	Water	26.6	11.9	mg GAE/100 mL		
	<i>Lb. plantarum</i> B4495 + <i>S. boulardii</i>	Water	26.6	12.8	mg GAE/100 mL		
	<i>Lb. helveticus</i> B4526 + <i>S. boulardii</i>	Water	26.6	12.8	mg GAE/100 mL		
	<i>Lb. rhamnosus</i> CRL981	Water	26.6	12.6	mg GAE/100 mL		
	<i>Sc. thermophilus</i> CCRC 14085 and <i>Bb. infantis</i> CCRC 14603	Acidified acetonitrile solution	N.M.	N.M.	–	Daidzein (↑), genistein (↑), daidzin (↓), genistin (↓)	(Marazza and others 2009, 2012)
	<i>Lb. acidophilus</i> CSCC 2400	80% Methanol	11.3	12.6	mg GAE/g DW	N.M.	(Lai and others 2013)
	<i>Lb. paracasei</i> CSCC 279	50% Acetone	10.6	12.7	mg GAE/g DW	N.M.	(Zhao and Shah 2014)
	<i>Lb. zeae</i> ASCC 15820	Water	4.60	5.96	mg GAE/g DW	N.M.	
	<i>Lb. rhamnosus</i> WQ2	80% Methanol + Hexane	47.0	47.4 to 59.5	mg GAE/100 mL	Gallic acid (↑), ferulic acid (↑), chlorogenic acid (↑), daidzein (↑), glycitein (↑), genistein (↑), catechin (↓), daidzin (↓), glycitin (↓), genistin (↓),	
	<i>Lb. plantarum</i> WCFS1	80% Methanol + Hexane	47.0	47.1 to 57.9	mg GAE/100 mL		
	<i>Lb. plantarum</i> WCFS1	THF	28.1	30.0 to 32.1	mg GAE/100 mL	N.M.	(Gan and others 2016c)
		Acidic methanol	14.4	18.9 to 20.5	mg GAE/100 mL		

(Continued)

Table 3–Continued.

Edible seeds and their products	Inoculated microbes	Extracts	TPC		Unit	Changes of main phenolic compounds	References
			Nonfermented samples	Fermented samples			
Black soymilk	<i>Sc. thermophilus</i> S10	80% Ethanol	~62.5	89.7	mg GAE/100 g DW	Daidzein (↑), genistein (↑), daidzin (↓), genistin (↓)	(Lee and others 2015)
Soy germ	<i>A. niger</i> M46	70% Methanol	~12.0	~33.0 to 38.0	mg GAE/g DW	Daidzein (↑), daidzin (↓), glycytin (↓), genistin (↓), acetyl daidzin (↓), acetyl glycytin (↓)	(Sheih and others 2014)
Soy whey	<i>Lb. plantarum</i> BI-6	80& Ethanol	52.8	64.7	mg GAE/g	Daidzein (↑), genistein (↑), daidzin (↓), genistin (↓)	(Xiao and others 2015b)
Buckwheat groats	<i>R. oligosporus</i> NRR1 2710	80% methanol	7.19	5.15	mg RUE/g DW	Rutin (↓)	(Malgozrata and others 2015)
	<i>R. oligosporus</i> ATCC 64063	50% Acetone	6.18	7.47	mg GAE/g DW	N.M.	(Starzynska-Janiszewska and others 2016)
Oat flakes	<i>Lb. plantarum</i> LP09	PBS (0.1 M, pH 7.4)	3.00	4.66	mg GAE/g DW	N.M.	(Nionelli and others 2014)
Oat groats	<i>Lb. johnsonii</i> LA1, <i>Lb. reuteri</i> SD2112, <i>Lb. acidophilus</i> LA-5	80% Methanol	0.46	0.62	mmol GAE/L	N.M.	(Hole and others 2012)
	<i>Lb. reuteri</i> SD2112, <i>Lb. acidophilus</i> LA-5, <i>Lb. plantarum</i> WCF51, <i>Lb. fermentum</i> NCDO 1750	50% Methanol	N.M.	N.M.	–	Caffeic acid (↑), ferulic acid (↑), sinapic acid (↑)	
Black rice bran	<i>B. subtilis</i> KU3	NaOH hydrolysis	N.M.	N.M.	–	Ferulic acid (↓), <i>p</i> -coumaric acid (↓)	
		Fermented supernatant	172	139	mg GAE/g DW	N.M.	(Yoon and others 2015)

A., Aspergillus; B., Bacillus; Bb., Bifidobacterium; C., Cordyceps; Cd., Ganoderma; Lb., Lactobacillus; R., Rhizopus; S., Saccharomyces; Sc., Streptococcus; T., Thiammidium; N.M., not measured; DW, dry weight; WW, wet weight; CE, catechin equivalent; GAE, gallic acid equivalent; TAE, tannic acid equivalent; RUE, rutin equivalent.

Table 4–Bioactive peptides in fermented edible seeds and their products.

Edible seeds and their products	Inoculated microbes	Bioactive peptides	Information of peptides	References
Chickpea (<i>Cicer arietinum</i>)	<i>Lb. plantarum</i> C48 and <i>Lb. brevis</i> AM7	Lunasin-like polypeptides	Name: Leucoagglutinating phytohemagglutinin; MW: 29.5 kDa; AA sequence: MASSKFTVLFVLLTHANSSNDIYFNFRQNETLILQRDASVSSGQRLRLTNL-NGNGEPRVGSIGRAFYSAPIQIWDNTTGTVASFATSTFNQVPPNINAGPADGLAFALVPGSQPKDKGGFLGFGSNNFHTVAVEFDTLNKKDWDPTERHIGIDVNSIRSKTTRWDFVNGENAEVLIYDSNTLLVAVSLVYPSQKTSFVSDTVDLKSVLPEWVSVGFSA TTGINKGNVETNDVLSWFSFASKLSDGTTSEGLNLANLNLKIL Name: Pathogenesis related protein; MW: 16.9 kDa; AA sequence: MGVFTFEQE-TASTVPPAKLYKAMVKDADAIIPKAVDAIKTVETVEGGGPGTIKKLT-SIGKVSVKYQTKGDAKPNKEVQEGKAKGDALFKAIEGYVLANPNYN Name: Seed linoleate 9S-lipoxygenase-3; MW: 97.6 kDa; AA sequence: MFSGVTLGNRGHKIKGTVLLMRKNVDLNSLTTVGGVIGQGGFDL-GSTVDNLTAFILGRSVLQJISA TKPDA TGKGLGKATFLGIISSLPTL LAGQSAF KHFWDGDDMGIPGAFYIKFMQTEFLVSLTLDIPNHGSIYVCSWYINAK HHKIDRIFANQTYLPSETPAPLHYREELNLRGDTGERKEWERIYDYV YNDLGNPDSGENHARPYLGGSETYPYRRGRTGRKTRKDPNSESRSDYVYLP RDEAFGLHSSDFLTGLKAVSQNVVPALESVFFDLNFTNPFDSFDEVHGLYE GGKILPTNLSQISPLVLEIRTDGENTLKYPPPKVIQVSRSGWMTDEEFAREM LAGVNPVICLOEFPFRSKLDSQIYGDHTSKISKEHLEPNLEGLTVEEAIONKILF LLDHDSIMPYLRRINSTSKAYA TRTILFLNQNQLKPLAIELSLPHPPQGDHGA VSYVYQPALEGVSSWILLAKAVIVNDSCYHQLVSHWLNTHAVVEPFIATNR HLSLHPYKLYPHYRDTMINSLSARLSLVNDGGIEKTEFLWGRYSMEMSSKV YKNWVTEQALPADLIRGMAIEDPSSPGVKLVVDEYPAVDGLIWIWIKTWV QDYVSLYTSDEKLRODSELQWVKELVEVGHGDKKNEPWWPKMQTREDLIEV CSIVWITASALHAAVNFGQSYGGILNLRPTLRRFMPEKSAEFEELVKSPOKAY LKTTIPKFTLIDL SVLEISRHASDELYLGERDNPWTSDKRALEAFKFGNKLAEI EKLLTORNDEKLRNHRGVPVEPPTLLYPSKSEGLTRGIPNSI!	(Rizzello and others 2015)
Kidney bean (<i>Phaseolus vulgaris</i>)	<i>Lb. plantarum</i> C48 and <i>Lb. brevis</i> AM7	Lunasin-like polypeptides	Name: Subtilisin inhibitor 1; MW: 10.4 kDa; AA sequence: QEQCTNSQEQNVPLPRNYKQALNTTPTKTSWPELVGVTAEQAE TKIKEEMVD-VQIQVSPHDSFVTDADYNPKRVLYVDES NKVTRTPSIGName: Legumin A2; MW: 59.6 kDa; AA sequence: MATKLLALSFCFLGGCFALREQEQNECQLERLNALEPDNRIESEGGLI-ETWNPNNKQFRACGVALSRA TLQHNA LRRPYNSNAPOEIFIQOQNGYFGVMVFP GCPTEFEEQSEGEGRYDRHQVYRFRGDIIV/PTGIVFWMYNDQDTP VIAVSLDIRSSNINLDOMPRFRYLAGNHEOELRYHQOQGGKOEQENEGNI FSGFRDFLEDAFVNRHIVDRLQRNEDEEKAIVKYGGLSISPEKQARHQ RSGRQEEDEDEERQPRHORGRQEEDEEERQPRHORRRRGEDEEKKERR GSQKGRRQDNGLEETVCTAKLRINIGPSSPDIYNPEAGRIKTYTSLDLPYLRW LKLSAEHGLHKNAMEVPHYNLANSIYALKGRARLQVYVNGNGNTVFDGELEA GRALTVPQNYAAAKSLDRFSYVAFKTNDRAGIARLAGTSSVINNLPDVAATFNLQRNEARQLKSNPKFLV PARQSENRA SA Name: Phaseolin; MW: 47.5 kDa; AA sequence: MMRARVPLLLGILFLASLSASFATLSLREESQDNPFYFNSDNSWNTLFKN-QYGHIRVLQRFDQQSKRQLONLEDYRVEFRSKPETLLLQQADAELLVVRGSA ILVLVKPDRREYFLLTSDNIPFSDHQIPAGIFYLVNPDQKDLRIQLAMPVNNPQ IHEFFLS TEAQOQSYLQEFSKHILEASFNKFEENRVLFEFEEQEQGVVINDSEIKEL SKHAKSSRSKLSKQDNTIGNFNGLTERTDNSLNVLJSSIEME EGALFVPHYYSKAI VILVNEGEAHVELVGPKNKETLEYESYRAELSKDDVFPVIPAAYVAIKATSNVN FTGFGINANNRNLGAKTNDVISSIGRALDGGKVLGLTFSSG GDEVMKLINKOSGSYFVDAHHHQEQKGRKGFVY N.M.	(Rizzello and others 2015)
Lentil (<i>Lens culinaris</i>)	<i>Lb. plantarum</i> C48 and <i>Lb. brevis</i> AM7	Lunasin-like polypeptides		(Rizzello and others 2015)

(Continued)

Table 4–Continued.

Edible seeds and their products	Inoculated microbes	Bioactive peptides	Information of peptides	References
Mung bean (<i>Vigna radiata</i>)	<i>Lb. plantarum</i> B1-6	ACE-inhibitory peptides	N.M.	(Wu and others 2015)
Navy bean (<i>Phaseolus vulgaris</i>)	Different LAB	ACE-inhibitory peptides	N.M.	(Rui and others 2015)
Pea (<i>Pisum sativum</i>)	<i>Lb. plantarum</i> 299v	ACE-inhibitory peptides	MW: 1.59 kDa; AA sequence: KEDDEEEEOGEEE	(Jakubczyk and others 2013)
	<i>Lb. plantarum</i> C48 and <i>Lb. brevis</i> AM7	Lunasin-like polypeptides	Name: Proviclin; MW: 31.5 kDa; AA sequence: DNAEIKLLEEHEKETHRRRLDRKROQSEKNVIVKVKKQKQIEL-SKNAKSSKKSVSRSSEPNKSSDPYNSQYKGFETPKKNPQLQDLDFVNVYEIK EGSWLPHYNSRAIVTVNEKGDVELVCGORNENQQGLREDDDEEEQREETKNO VQSYKAKLTPGDV FVIPAGHPVA VRASNLLNLLFGINAENQRNIFLAGEDNVISQ IQKQKDLTPGSAQEVDRLENNQKSYFANAQQOQRETRSQEIKEHLYSILGAF Name: Seed linoleate 9S-lipoxygenase-2; MW: 97.1 kDa; AA sequence: MFPNVTLNKGHKIRGTVYLMRKNVLDNFNTIVSIGGNVHGVIDSGINI-IGTLDGLTAFGRSVSLQISATKSDANGKGVKGDFTLEGVLA S LPTLGAGESAFNI HFEWDEHMGIP121GAFYIKNYMQVEFFKSLTLEDVPHNHTIRFVNSWVYNSKLYK SPRIFFANKSYLPSETPSLVKYREEELQTLRGDGTGERKLHERIYDYVYNDLGNPDHGE HILARPILGSSTHPYPRRGRTGRYPTRKDPNSEKPAETETVPRDENFGHLKSSDFLA YGIKVSQCWVPAFESAFDLNFTPNFEDSFQDVRNLFEGGKILPLDVI STLSPVVKFEFRT DGEVLIKFTPPHVRVSKSAWMTDEEFAREMLAGVNPCMIRGLQEPFKSNLDPAEY GDHTSKISVDVNLIDGCTIDEALASGRFLIDYHDTFPLLRINETSAKAYATRILFLKE NGTLKPVAIELS LPHPDGDKSFGVSKVILPADEGVESTIWLAKAYV VVNDSCYHQL MSHWLNTHAVIEPEVIATNRQLSVVHPINKLAPHYRDTMMNINALARDLSLINANG LIERSLPSKYAVEMSSAVYKYVWFDTQALPNDLKRMMAVKDS SSPYGLR LIIEDYPYAVDGLIEWTAIKTWQDYVSYLYATDNDKNDSELSQHWKVEV VEKGGDLKDKPWWPKLQTFDELVEVCTIIWTASALHAAVNFQOYPYGGI ILNRP T LSRLLPEEGTA EYDEMVKSQAYLRITTPKFQTLIDLSVIEL SRHASDEVYLGQRENPHWTS DSKALQAFQKFGNKLAEIAEAKLTNKNNDPSL YHRVGPVQLPYTLHPSSKEGLTFRGIPNSISI Name: Seed biotin-containing protein SBP65; MW: 59.5 kDa; AA sequence: MASEQLSRRENITTKIQAEDSVPRQRTTHFELRETHELGNFQSLPRNENQAY LDRGARAPLSANVSESYLDRARVPLINANIPEHRVREKEDFGVYRDMGKFMQESK GGNKSLAEDRETLDTRSRMVTGTPHIKESGKQVVEERERARERAMEEEKR LTMEEISKYRNQAQOSALEALSAAQEKYERAKQATNELRNTTQAAQEKGEAAQ AKDATFEKTQQGYEMTGDVSNARSARTASEKAAQKNITLTKTQQGYEATRTDYS NAARTAAEYATPAAEKARCVAVQAKDV TLETGKTAEEKAKCAAIAKVAVDLK EKATVAGWTASHYATQLTVDGTRAAANAVEGAVGYVAPKASELAASKSVETVKGL AASAGETAKEFTARKEESWREYEAKRASQLOEGEELPSTGGIGKVLPSGER TQAQTNLQEKYQKGS DILGAVTETVSDIGSSMIKPIDNANTKVKEHGTTIT PKGQDAGGVLD AIGETIAEIAHTTKVIVVGEDDEVEKSMQKNIGSDSHSIDRAK HEGYRAPKNNVS Name: Albumin-1 C; MW: 13.9 kDa; AA sequence: MASVKLASLIVLFATLGMFLTKN-VGAISCNVCSPFDIPPCGSPLCRIPAGLVIGNCRNPYGVF LRTNDEHPNLCESDADCRKKKSGTFCGHYPNDIEYGWCFASKSEAEVFSKITPKDLKLSVSTA	(Rizzello and others 2015)

(Continued)

Table 4–Continued.

Edible seeds and their products	Inoculated microbes	Bioactive peptides	Information of peptides	References
Grass pea (<i>Lathyrus sativus</i>)	<i>Lb. rhamnosus</i> BGT10 <i>Lb. plantarum</i> C48 and <i>Lb. brevis</i> AM7 Different LAB	Antioxidant peptides Lunasin-like polypeptides Cancer preventive peptide	N.M. N.M. Name: Lunasin; Amino acid sequence: with immunoreactive epitope RGD D D D D D D D D D	(Stanisavljevic and others 2015) (Rizzello and others 2015) (Rizzello and others 2012) (Tsai and others 2006)
Soybean (<i>Glycine max</i>)	A mixture of LAB	ACE-inhibitory peptides	N.M.	(Nakahara and others 2012)
Soy milk	Natural fermentation	ACE-inhibitory peptides	Amino acid sequence: SY and GY	(Nakahara and others 2012)
Soy protein	<i>Lb. casei</i> spp. pseudoplantarum Different LAB	ACE-inhibitory peptides Cancer preventive peptide	N-terminal AA sequence: LIVTQ	(Vallabha and Tiku 2014)
Amaranth (<i>Amaranthus</i>)	Different LAB	Cancer preventive peptide	Name: Lunasin; Amino acid sequence: with immunoreactive epitope RGD D D D D D D D D D	(Rizzello and others 2012)
Barley (<i>Hordeum vulgare</i> L.)	Different LAB	Cancer preventive peptide	Name: Lunasin; Amino acid sequence: with immunoreactive epitope RGD D D D D D D D D D	(Rizzello and others 2012)
Rapeseed (<i>Brassica napus</i>)	<i>B. subtilis</i> 10160	Antioxidant peptides	N.M.	(He and others 2012)
Rye (<i>Secale cereal</i>)	A mixture of LAB	Antioxidant peptides	Amino acid sequence: VFVDEGLEVLGWRPVFNVSVVGRNAK (MW: 2.98 kDa); RLSLPAGAPVTVAVSP (MW: 1.54 kDa); NANGELCPNMMGCSQWGYCGLSEFCNGCSGACGCEK (MW: 4.03 kDa); LCPVHRAADL (MW: 1.10 kDa); PAEMVAAALDR (MW: 1.48 kDa); KVALMSAGSMH (MW: 1.13 kDa); DLADIPQQRLMAGLALVAVTVFLK (MW: 2.82 kDa); KNGSIFNSPSATAATIIHGHNYSGLAYLDFVTSK (MW: 3.58 kDa); GTIFFSQEGDGPTSVTGSVGLKPLGHFVHALGDTTNGCMSTGPHFNPTGK (MW: 5.34 kDa)	(Coda and others 2012)
Rye malt sourdoughs	Different LAB	ACE-inhibitory peptides	Amino acid sequences: LQP, LLP, VPP, and IPP	(Hu and others 2011)
Rye malt and flour	<i>Lb. reuteri</i>	Taste-active γ -glutamyl-dipeptides	Amino acid sequence: γ -EV, γ -EM, γ -EE, γ -EL, γ -EF, γ -EI	(Zhao and Gänzle 2016)

(Continued)

Table 4—Continued.

Edible seeds and their products	Inoculated microbes	Bioactive peptides	Information of peptides	References
Spelt (<i>Triticum spelta</i>)	A mixture of LAB	Antioxidant peptides	Amino acid sequence: AIAGAGVLSGYDQLQLFFGK (MW: 2.17 kDa); GNQEKVLELVQR (MW: 1.41 kDa); PAGSAAGAAP (MW: 0.77 kDa); EALEAMFL (MW: 0.92 kDa); AAGAAAAARSAGCCGR (MW: 1.39 kDa); ITFAAYRR (MW: 1.00 kDa); HPVPPKKK (MW: 0.91 kDa)	(Coda and others 2012)
Wholemeal wheat sourdough	A mixture of LAB	ACE-inhibitory peptides	Amino acid sequence: DPVAPLQSRGPEI (MW: 1.15 kDa); PVAPQLSRGILL (MW: 1.15 kDa); ELEIVMASPP (MW: 1.08 kDa); QILLPRPGQAA (MW: 1.16 kDa); PVAPLQSRGPE (MW: 1.15 kDa); PRSGNVGESGL (MW: 1.15 kDa); VAFPSRPTPR (MW: 1.08 kDa); DIIIPD (MW: 1.16 kDa); PRSGNVGESGLID (MW: 1.30 kDa); DPVAPLQSRGPEI (MW: 1.15 kDa); DPVAPLQSRGPEIP (MW: 1.26 kDa); PVAPLPRKGS (MW: 1.02 kDa); DPVAPLQSRGPE (MW: 1.02 kDa); SFTAGARTFNFDENPCDYFOGGKIKAT (MW: 2.98 kDa)	(Rizzello and others 2008)
Whole wheat	A mixture of LAB	Antioxidant peptides	Amino acid sequence: MAPAAVAAAEAGSK (MW: 1.24 kDa); DNIPIVIR (MW: 0.94 kDa)	(Coda and others 2012)
Wheat (Kamut)	A mixture of LAB	Antioxidant peptides	Amino acid sequence: YEWEPTVPNFDAKDYTDM (MW: 2.26 kDa); GVSNAAVVAGGH (MW: 1.04 kDa); DAQEFKR (MW: 0.89 kDa); PPGPGPPPPGAAAGRGGG (MW: 1.70 kDa); HKEMQAIIFYIMFIN (MW: 2.00 kDa); TGGGSTSSSSSSSLGGGASRGSVVEAAPATQGAANA-PAVPVVDTQEAGIR (MW: 5.02 kDa); DTAAGYVAPPDPAVSTGDTGLAGAEAPHESAVMSGAAAAAAPPGEAYTR (MW: 4.92 kDa)	(Coda and others 2012)
Wholemeal wheat	Different LAB	Cancer preventive peptide	Name: Lunasin; Amino acid sequence: with immunoreactive epitope RGDGDDDDDDDD	(Rizzello and others 2012)
Wheat germ	<i>Lb. plantarum</i> LB1 and <i>Lb. rossiae</i> LB5	Anti-fungal peptides	Amino acid sequence: VLHEPLF (MW: 0.85 kDa); YNNPIYVTENGIAEGNNKSLPITEAL (MW: 2.95 kDa); ALKAAAPSPA (MW: 0.82 kDa); ALIIVMLFGR (MW: 1.24 kDa); AAAAVFLSLLAVGHCAAADFNATDADADFAGNGVDFNSSDAAVYWGWPWKAR (MW: 5.29 kDa)	(Rizzello and others 2011)
Wheat germ	<i>B. Subtilis</i> B1	Antioxidant peptides	N.M.	(Niu and others 2013)
Casein miso paste based on rice and soybean	<i>A. oryzae</i> and <i>S. cerevisiae</i>	ACE-inhibitory peptides	Amino acid sequence: VPP and IPP	(Inoue and others 2009)
Peanut meal	<i>B. subtilis</i>	Antioxidant peptides	Amino acid sequence: YP	(Zhang and others 2014)
Walnut protein meal	<i>B. subtilis</i> GIM 1.135	Antioxidant peptides	N.M.	(Wu and others 2014)

The amino acid sequences of peptides were expressed by the manner of single-letter. Common amino acid abbreviations: A, alanine; C, cysteine; D, aspartic acid; E, glutamic acid; F, phenylalanine; G, glycine; H, histidine; I, isoleucine; K, lysine; L, leucine; M, methionine; N, asparagine; P, proline; Q, glutamine; R, arginine; S, serine; T, threonine; V, valine; W, tryptophan; Y, tyrosine. Other abbreviations: A., *Aspergillus*; B., *Bacillus*; Lb., *Lactobacillus*; S., *Saccharomyces*; N.M., not measured; MW, molecular weight; AA, amino acid.

In addition, fermentation can also produce peptides with ACE-inhibitory activity in edible seeds and their products (Table 4), with some features of ACE-inhibitory peptides summarized below. In general, ACE-inhibitory peptides have a low molecular weight. Many ACE-inhibitory peptides consist of 2 to 12 amino acids, and peptides with molecular weight about 0.8 to 0.9 kDa exhibit the highest ACE-inhibitory activity in LAB-fermented soymilk (Tsai and others 2006). Rizzello and others (2008) stated that most ACE-inhibitory peptides in LAB-fermented whole-meal wheat sourdough had molecular weights less than 1.6 kDa. In addition, the existence and position of some special amino acids in peptides can also influence the ACE-inhibitory activity. Peptides containing VAP (valine-alanine-proline) epitope were reported to possess strong ACE-inhibitory effect (Rizzello and others 2008). Vallabha and Tiku (2014) suggested that glutamine at the C-terminal had a more potent effect than other residues, and leucine (a branch-chained amino acid) at the N-terminal was more effective than glycine to bind with ACE. Ondetti and Cushman (1982) suggested that C-terminal arginine with a positive charge on the ϵ -amino group significantly contributed to ACE-inhibitory activity. Therefore, understanding these characteristics can be helpful for predicting potential ACE-inhibitory peptides from natural sources.

Besides, fermented edible seeds and their products also contain other bioactive peptides, such as taste-active peptides (Zhao and Gänzle 2016) and lunasin-like peptides (Rizzello and others 2015), which may possess anti-cancer effects (Rizzello and others 2012). In general, fermented edible seeds and their products are good sources of various bioactive peptides. However, bioactive peptides have only been reported in a limited number of fermented edible seeds up to now, and the identification of specific bioactive peptides, including their amino acid sequences, has also been scarcely reported. Therefore, more studies are needed to further explore bioactive peptides and their identities in fermented edible seeds and their products.

Other bioactive components

Many other bioactive components have been found in fermented edible seeds and their products, especially in yeast-fermented rice. A variety of bioactive compounds have been reported in *Monascus* spp.-fermented rice, such as monascin, ankaflavin, rubropunctatin, monascorburin, rubropunctamine, monascorburamine, two furanoisophthalides, monapurfluores A and B, monascopyridine A, B, C, and D, monasfluores A and B, xanthomonasin A and B, monascumic acid, and monacolin K (Wild and others 2003; Akihisa and others 2005a,b; Chen and Hu 2005; Hsu and others 2010), which exhibited different bioactivities, such as anti-cancer, anti-inflammatory, and hypolipidemic effects. Similarly, *Monascus* spp.-fermented monascol waxy corn and soybean also contained increased contents of monacolin K (also known as mevinolin and lovastatin) compared to unfermented samples (Pyo and Lee 2007; Kongbangkerd and others 2014), which was positively associated with the increased antioxidant effect of samples. In addition, fermented wheat germ has been found to produce benzoquinones with anti-cancer effect (Comin-Anduix and others 2002; Mueller and others 2011, Rizzello and others 2013).

Overall, fermented edible seeds and their products contain various bioactive components, especially vitamins, GABA, natural phenolics, and bioactive peptides, and these bioactive components endow the fermented products with versatile bioactivities, as discussed in the following section. In the future, bioactive com-

ponents in other fermented edible seeds and their products can be explored to provide a basis for the development of fermented functional foods.

Bioactivities of fermented edible seeds and their products

A large number of studies demonstrate that fermented edible seeds and their products exhibit manifold bioactivities (Figure 1), such as antioxidant, anti-hypertensive, and anti-cancer effects.

Antioxidant effect

The influence of fermentation on the antioxidant effect of edible seeds and their products has been extensively investigated *in vitro*. Various antioxidant effects, such as free radical-scavenging, and reducing and metal-chelating effects, have been reported to be increased in the hydrophilic extracts of most fermented edible seeds and their products compared to unfermented samples (Table 5). The increase of antioxidant effect is mainly due to the increased antioxidant levels in fermented samples, mainly antioxidant phenolics and peptides (Table 3 and 4). However, the influence of fermentation on antioxidant effect in the lipophilic and bound extracts of edible seeds and their products has been scarcely investigated. Unlike the hydrophilic extract commonly prepared by polar solutions, such as ethanol and methanol water solutions, the lipophilic extract needs to be prepared by nonpolar solutions, such as *n*-hexane or tetrahydrofuran, and the bound extract needs to be hydrolyzed by acidic, alkaline, or enzymatic solutions. Our previous study found that fermented and nonfermented soybean and mung bean milks exhibited much higher ABTS and DPPH free radical-scavenging effect in their lipophilic extracts than their hydrophilic extracts, and fermentation had different effects on antioxidant effect in their lipophilic and hydrophilic extracts (Gan and others 2016c). In addition, we also found that some edible beans exhibited substantial antioxidant effects in their bound extracts, while fermentation had varying effects on antioxidant effect in their bound extracts (Gan and others 2016b). These results suggest that ignoring the lipophilic or bound extracts may significantly underestimate the total antioxidant effect in some fermented edible seeds and their products.

On the other hand, fermentation was also reported to decrease antioxidant effect in the hydrophilic extract of some edible seeds (Table 5), such as lentil (Gan and others 2016b), black cow gram (Gan and others 2016b), common bean (Gamboa-Gomez and others 2016), soybean (Fernandez-Orozco and others 2007; Gan and others 2016b), wheat (Dordevic and others 2010), buckwheat (Dordevic and others 2010), and buckwheat groats (Malgorzata and others 2015). The reduction of antioxidant effect may be partly associated with the reduction of reduced glutathione and related antioxidant enzyme activities in these fermented samples, such as fermented soybeans (Fernandez-Orozco and others 2007), and may also be associated with the degradation of antioxidant phenolics into phenolic compounds with lower antioxidant effect in these fermented samples. This may explain why fermentation generally increased TPC in these samples, but reduced their antioxidant effect. Overall, fermentation has different influences on antioxidant effect in edible seeds and their products, probably associated with different fermentation methods, different antioxidant components in samples as well as different extraction and evaluation methods for antioxidant effect.

Anti-hypertensive effect

Hypertension is one of the most important risk factors for cardiovascular diseases. Blood pressure is tightly controlled by the

Table 5—Influences of fermentation on antioxidant effect in edible seeds and their products.

Edible seeds and their products	Inoculated microbes	Extracts	Antioxidant assay	Antioxidant effect		Unit	References
				Nonfermented samples	Fermented samples		
Chickpea (<i>Cicer arietinum</i> L.)	<i>R. oligosporus</i> NRRL 2710	80% Ethanol	ABTS	14.8	17.2 to 68.4	$\mu\text{mol TE/g DW}$	(Sanchez-Magana and others 2014)
			ORAC ABTS	30.3 18.2	34.6 to 100 19.1 to 28.7	$\mu\text{mol TE/g DW}$ $\mu\text{mol TE/g DW}$	
Common bean (<i>Phaseolus vulgaris</i> L., Bayo Victoria)	<i>C. militaris</i> SN-18	80% Methanol	ORAC	21.9	25.4 to 45.9	$\mu\text{mol TE/g DW}$	(Xiao and others 2014)
			DPPH	5.56 ^a	1.21 ^a	mg/mL	
		80% Ethanol	ABTS	3.61 ^a	0.94 ^a	mg/mL	(Gamba-Gomez and others 2016)
			RP	14.6 ^a	4.15 ^a	mg/mL	
			DPPH	4.67 ^a	1.35 ^a	mg/mL	
			ABTS	3.34 ^a	0.98 ^a	mg/mL	
			RP	12.8 ^a	5.61 ^a	mg/mL	
			DPPH	N.D. ^a	3.08 ^a	mg/mL	
			ABTS	5.25 ^a	3.97 ^a	mg/mL	
			RP	10.2 ^a	5.38 ^a	mg/mL	
DPPH	~2.00 ^a	~7.00 to 10.4 ^a	$\mu\text{g}/\mu\text{L}$				
Common bean (<i>Phaseolus vulgaris</i> L., Pinto durango)	<i>R. oligosporus</i> (NRRL2710)	70% Acetone	LDL	~0.28 ^a	~0.48 to 1.48 ^a	$\mu\text{g}/\mu\text{L}$	(Starzynska-Januszewska and others 2014)
			HRSP	~3.30 ^a	~0.70 to 4.20 ^a	$\mu\text{g}/\mu\text{L}$	
		70% Acetone	DPPH	~2.10 ^a	~7.40 to 9.70 ^a	$\mu\text{g}/\mu\text{L}$	(Can and others 2016b)
			LDL	~0.41 ^a	~1.15 to 1.74 ^a	$\mu\text{g}/\mu\text{L}$	
			HRSP	~0.50 ^a	~2.20 to 4.70 ^a	$\mu\text{g}/\mu\text{L}$	
			ABTS	16.4	18.8	$\mu\text{mol TE/g DW}$	
			DPPH	3.73	4.70	$\mu\text{mol TE/g DW}$	
			FRAP	14.9	13.8	$\mu\text{mol Fe (II)/g DW}$	
			ABTS	12.9	10.9	$\mu\text{mol TE/g DW}$	
			FRAP	5.13	6.87	$\mu\text{mol Fe (II)/g DW}$	
Common bean (<i>Phaseolus vulgaris</i>)	<i>Lb. plantarum</i> DSM 20174 and <i>R. microsporius</i> var. chinensis	Phosphate buffer	ABTS	16.4	18.8	$\mu\text{mol TE/g DW}$	(Starzynska-Januszewska and others 2014)
			DPPH	3.73	4.70	$\mu\text{mol TE/g DW}$	
		80% Acetone	FRAP	14.9	13.8	$\mu\text{mol Fe (II)/g DW}$	(Can and others 2016b)
			ABTS	12.9	10.9	$\mu\text{mol TE/g DW}$	
			FRAP	5.13	6.87	$\mu\text{mol Fe (II)/g DW}$	
			ABTS	3.52	4.45	$\mu\text{mol TE/g DW}$	
			FRAP	14.9	17.7	$\mu\text{mol Fe (II)/g DW}$	
			ABTS	12.9	12.5	$\mu\text{mol TE/g DW}$	
			FRAP	5.13	5.94	$\mu\text{mol Fe (II)/g DW}$	
			ABTS	3.52	4.13	$\mu\text{mol TE/g DW}$	
FRAP	14.9	18.7	$\mu\text{mol Fe (II)/g DW}$				
Black cow gram (<i>Labiab purpures</i>)	<i>Lb. paracasei</i> 279	80% Methanol	ABTS	12.9	12.5	$\mu\text{mol TE/g DW}$	(Can and others 2016b)
			FRAP	5.13	5.94	$\mu\text{mol Fe (II)/g DW}$	
		80% Methanol	ABTS	3.52	4.13	$\mu\text{mol TE/g DW}$	(Can and others 2016b)
			FRAP	14.9	18.7	$\mu\text{mol Fe (II)/g DW}$	
			ABTS	12.9	12.5	$\mu\text{mol TE/g DW}$	
			FRAP	5.13	5.94	$\mu\text{mol Fe (II)/g DW}$	
			ABTS	3.52	4.13	$\mu\text{mol TE/g DW}$	
			FRAP	14.9	18.7	$\mu\text{mol Fe (II)/g DW}$	
			ABTS	12.9	12.5	$\mu\text{mol TE/g DW}$	
			FRAP	5.13	5.94	$\mu\text{mol Fe (II)/g DW}$	

(Continued)

Table 5–Continued.

Edible seeds and their products	Inoculated microbes	Extracts	Antioxidant assay	Antioxidant effect		Unit	References
				Nonfermented samples	Fermented samples		
Mottled cowpea (<i>Vigna unguiculata</i>)	Natural fermentation	80% Methanol	ABTS	3.52	4.18	μmol TE/g DW	(Gan and others 2016b)
			FRAP	23.1	29.6	μmol Fe (II)/g DW	
	NaOH + HCl hydrolysis	ABTS	23.9	28.0	μmol TE/g DW		
		FRAP	8.93	17.8	μmol Fe (II)/g DW		
	<i>Lb. paracasei</i> 279	80% Methanol	ABTS	5.35	12.9	μmol TE/g DW	
			FRAP	23.1	40.1	μmol Fe (II)/g DW	
	<i>Lb. plantarum</i> WCF51	NaOH + HCl hydrolysis	ABTS	23.9	29.4	μmol TE/g DW	
			FRAP	8.93	8.59	μmol Fe (II)/g DW	
	80% Methanol	ABTS	5.35	5.55	μmol TE/g DW		
		FRAP	23.1	38.1	μmol Fe (II)/g DW		
NaOH + HCl hydrolysis	ABTS	23.9	28.9	μmol TE/g DW			
	FRAP	8.93	9.81	μmol Fe (II)/g DW			
Kidney bean (<i>Phaseolus vulgaris</i>)	Natural fermentation	Water	ABTS	5.35	6.08	μmol TE/g DW	(Limón and others 2015)
			ORAC	~120	~130	mmol TE/g DW	
	Water	ORAC	~170	~500 to 550	mmol TE/g DW		
		ORAC	~120	~115 to 130	mmol TE/g DW		
	<i>B. subtilis</i> CECT 39 ^T (ATCC 6051)	80% Methanol	FRAP	11.5	16.9	μmol Fe (II)/g DW	
			ABTS	13.1	19.1	μmol TE/g DW	
	<i>Lb. plantarum</i> CECT 748 ^T (ATCC 14917)	NaOH + HCl hydrolysis	FRAP	6.17	6.92	μmol Fe (II)/g DW	
			ABTS	4.89	4.70	μmol TE/g DW	
	80% Methanol	FRAP	11.5	24.0	μmol Fe (II)/g DW		
		ABTS	13.1	24.0	μmol TE/g DW		
NaOH + HCl hydrolysis	FRAP	6.17	5.99	μmol Fe (II)/g DW			
	ABTS	4.89	4.23	μmol TE/g DW			
Lentil (<i>Lens culinaris</i>)	<i>Lb. plantarum</i> WCF51	80% Methanol	FRAP	11.5	21.9	μmol Fe (II)/g DW	(Torino and others 2013)
			ABTS	13.1	22.6	μmol TE/g DW	
	NaOH + HCl hydrolysis	FRAP	6.17	6.26	μmol Fe (II)/g DW		
		ABTS	4.89	4.08	μmol TE/g DW		
	Water	ORAC	~0.26	~0.25 to 0.30	mmol TE/g DW		
		ORAC	~0.15	~0.21 to 0.23	mmol TE/g DW		
	Natural fermentation	Water	ORAC	~0.26	~0.27 to 0.30	mmol TE/g DW	
			FRAP	11.2	10.5	μmol Fe (II)/g DW	
	<i>B. subtilis</i> CECT 39 ^T (ATCC 6051)	80% Methanol	ABTS	12.8	11.0	μmol TE/g DW	
			FRAP	6.30	8.72	μmol Fe (II)/g DW	
<i>L. plantarum</i> CECT 748 ^T (ATCC 14917)	NaOH + HCl hydrolysis	FRAP	11.2	10.5	μmol Fe (II)/g DW		
		ABTS	12.8	11.0	μmol TE/g DW		
Natural fermentation	Water	FRAP	6.30	8.72	μmol Fe (II)/g DW		
		ABTS	11.2	10.5	μmol Fe (II)/g DW		

(Continued)

Table 5--Continued.

Edible seeds and their products	Inoculated microbes	Extracts	Antioxidant assay	Antioxidant effect		Unit	References
				Nonfermented samples	Fermented samples		
Lupin (<i>Lupinus albus</i>)	<i>Lb. paracasei</i> 279	80% Methanol	ABTS	4.50	5.62	$\mu\text{mol TE/g DW}$	
			FRAP	11.2	16.1	$\mu\text{mol Fe (II)/g DW}$	
Mung bean (<i>Vigna radiata</i>)	<i>Lb. plantarum</i> WCFS1	NaOH + HCl hydrolysis	ABTS	12.8	13.3	$\mu\text{mol TE/g DW}$	
			FRAP	6.30	4.47	$\mu\text{mol Fe (II)/g DW}$	
Lupin (<i>Lupinus albus</i>)	Natural fermentation	80% Methanol	ABTS	4.50	3.40	$\mu\text{mol TE/g DW}$	(Frias and others 2005)
			FRAP	11.2	14.5	$\mu\text{mol Fe (II)/g DW}$	
Mung bean (<i>Vigna radiata</i>)	<i>Lb. plantarum</i> CECT 748	NaOH + HCl hydrolysis	ABTS	12.8	12.5	$\mu\text{mol TE/g DW}$	(Xiao and others 2015d)
			FRAP	6.30	5.16	$\mu\text{mol Fe (II)/g DW}$	
Mung bean (<i>Vigna radiata</i>)	<i>C. militaris</i> SN-18	80% methanol	ABTS	4.50	3.63	$\mu\text{mol TE/g DW}$	
			DPPH	71.4	55.1 to 71.8	$\mu\text{mol TE/g DW}$	
Red bean (<i>Phaseolus radiatus</i>)	<i>B. subtilis</i> (BCRC 14716) and <i>Lb. delbrueckii</i> sp. bulgaricus (BCRC 14008)	80% ethanol	ABTS	~1000	~3750	$\mu\text{g VcE/g DW}$	
			FRAP	~7.30	~8.50	$\mu\text{mol Fe (II)/g DW}$	
Red bean (<i>Phaseolus radiatus</i>)	Natural fermentation	80% acetone	Iron-chelating	~400	~10.5	$\mu\text{g EDTA-2Na/g DW}$	
			DPPH	~650	~750	$\mu\text{g VcE/g DW}$	
Red bean (<i>Phaseolus radiatus</i>)	Natural fermentation	80% acetone	ABTS	~1120	~3500	$\mu\text{g VcE/g DW}$	
			FRAP	~6.20	~7.00	$\mu\text{mol Fe (II)/g DW}$	
Red bean (<i>Phaseolus radiatus</i>)	Natural fermentation	80% acetone	Iron-chelating	~440	~750	$\mu\text{g VcE/g DW}$	
			DPPH	~4.60	~10.0	$\mu\text{mol EDTA-2Na/g DW}$	
Red bean (<i>Phaseolus radiatus</i>)	Natural fermentation	80% acetone	ABTS	~1050	~1055	$\mu\text{g VcE/g DW}$	
			FRAP	~1560	~3350	$\mu\text{g VcE/g DW}$	
Red bean (<i>Phaseolus radiatus</i>)	Natural fermentation	80% acetone	FRAP	~9.30	~10.0	$\mu\text{mol Fe (II)/g DW}$	
			RP	~680	~850	$\mu\text{g VcE/g DW}$	
Red bean (<i>Phaseolus radiatus</i>)	Natural fermentation	80% acetone	Iron-chelating	~3.70	~9.20	$\mu\text{mol EDTA-2Na/g DW}$	
			DPPH	~600	~1460	$\mu\text{g VcE/g DW}$	
Red bean (<i>Phaseolus radiatus</i>)	Natural fermentation	80% acetone	ABTS	~1050	~3750	$\mu\text{g VcE/g DW}$	
			FRAP	~5.00	~16.0	$\mu\text{mol Fe (II)/g DW}$	
Red bean (<i>Phaseolus radiatus</i>)	Natural fermentation	80% acetone	RP	~310	~1280	$\mu\text{g VcE/g DW}$	
			Iron-chelating	~3.50	~4.70	$\mu\text{mol EDTA-2Na/g DW}$	
Red bean (<i>Phaseolus radiatus</i>)	Natural fermentation	80% acetone	DPPH	84.6 ^a	22.4 ^a	mg/mL	(Jhan and others 2015)
			FRAP	N.D. ^a	> 1000 ^a	mg/mL	
Small rice bean (<i>Vigna umbellata</i>)	Natural fermentation	50% Ethanol	Iron-chelating	56.0 ^a	16.6 ^a	mg/mL	
			DPPH	N.D. ^a	75.3 ^a	mg/mL	
Small rice bean (<i>Vigna umbellata</i>)	Natural fermentation	50% Ethanol	FRAP	19.8	21.3	$\mu\text{mol Fe (II)/g DW}$	(Can and others 2016b)
			ABTS	20.0	21.5	$\mu\text{mol TE/g DW}$	
Small rice bean (<i>Vigna umbellata</i>)	Natural fermentation	50% Ethanol	FRAP	6.04	11.5	$\mu\text{mol Fe (II)/g DW}$	
			ABTS	20.0	21.5	$\mu\text{mol TE/g DW}$	

(Continued)

Table 5–Continued.

Edible seeds and their products	Inoculated microbes	Extracts	Antioxidant assay	Antioxidant effect		Unit	References
				Nonfermented samples	Fermented samples		
Small runner bean (<i>Phaseolus coccineus</i>)	<i>Lb. paracasei</i> 279	80% Methanol	ABTS	4.60	7.50	$\mu\text{mol TE/g DW}$	(Can and others 2016b)
			FRAP	19.8	29.7	$\mu\text{mol Fe (II)/g DW}$	
	<i>Lb. plantarum</i> WCF51	NaOH + HCl hydrolysis	ABTS	20.0	25.7	$\mu\text{mol TE/g DW}$	
			FRAP	6.04	4.26	$\mu\text{mol Fe (II)/g DW}$	
	80% Methanol	NaOH + HCl hydrolysis	ABTS	4.60	3.48	$\mu\text{mol TE/g DW}$	
			FRAP	19.8	29.4	$\mu\text{mol Fe (II)/g DW}$	
	Natural fermentation	80% Methanol	ABTS	20.0	25.7	$\mu\text{mol TE/g DW}$	
			FRAP	6.04	5.11	$\mu\text{mol Fe (II)/g DW}$	
	80% Methanol	NaOH + HCl hydrolysis	ABTS	4.60	4.05	$\mu\text{mol TE/g DW}$	
			FRAP	11.1	11.7	$\mu\text{mol Fe (II)/g DW}$	
	80% Methanol	NaOH + HCl hydrolysis	ABTS	12.2	14.8	$\mu\text{mol TE/g DW}$	
			FRAP	20.7	9.16	$\mu\text{mol Fe (II)/g DW}$	
80% Methanol	NaOH + HCl hydrolysis	ABTS	13.2	5.63	$\mu\text{mol TE/g DW}$		
		FRAP	11.1	18.5	$\mu\text{mol Fe (II)/g DW}$		
<i>Lb. paracasei</i> 279	80% Methanol	ABTS	12.2	20.3	$\mu\text{mol TE/g DW}$		
		FRAP	20.7	5.46	$\mu\text{mol Fe (II)/g DW}$		
<i>Lb. plantarum</i> WCF51	80% Methanol	ABTS	13.2	4.04	$\mu\text{mol TE/g DW}$		
		FRAP	11.1	18.9	$\mu\text{mol Fe (II)/g DW}$		
NaOH + HCl hydrolysis	80% Methanol	ABTS	12.2	18.9	$\mu\text{mol TE/g DW}$		
		FRAP	20.7	9.08	$\mu\text{mol Fe (II)/g DW}$		
Soybean (<i>Glycine max</i>)	Natural fermentation	80% Methanol	ABTS	13.2	6.27	$\mu\text{mol TE/g DW}$	(Fernandez-Orozco and others 2007)
			ABTS	63.0	44.6 to 57.6	$\mu\text{mol TE/g DW}$	
	<i>A. oryzae</i> 2094 ^T (ATCC 1011)	80% Methanol	ABTS	63.0	38.0	$\mu\text{mol TE/g DW}$	
			ABTS	63.0	46.7	$\mu\text{mol TE/g DW}$	
	<i>R. oryzae</i> CECT 2340 (ATCC 24563)	80% Methanol	ABTS	63.0	131	$\mu\text{mol TE/g DW}$	
			ABTS	63.0	54.5 to 57.4	$\mu\text{mol TE/g DW}$	
	<i>B. subtilis</i> CECT 39 ^T (ATCC 6051)	80% Methanol	ABTS	63.0	2.22 to 3.40	$\mu\text{mol TE/g DW}$	
			ABTS	63.0	2.22 to 3.40	$\mu\text{mol TE/g DW}$	
	<i>Lb. plantarum</i> CECT 748 ^T (ATCC 14917)	80% Methanol	PRTC	2.54	4.59	$\mu\text{mol TE/g DW}$	
			PRTC	2.54	6.77	$\mu\text{mol TE/g DW}$	
	Natural fermentation	80% Methanol	PRTC	2.54	9.23	$\mu\text{mol TE/g DW}$	
			PRTC	2.54	3.60 to 4.25	$\mu\text{mol TE/g DW}$	
<i>A. oryzae</i> 2094 ^T (ATCC 1011)	80% Methanol	FRAP	7.08	4.37	$\mu\text{mol Fe (II)/g DW}$		
		FRAP	7.08	4.37	$\mu\text{mol Fe (II)/g DW}$		

(Continued)

Table 5—Continued.

Edible seeds and their products	Inoculated microbes	Extracts	Antioxidant assay	Antioxidant effect		Unit	References	
				Nonfermented samples	Fermented samples			
Black soybean (<i>Glycine max</i>)	<i>Lb. paracasei</i> 279	NaOH + HCl hydrolysis	ABTS	9.53	4.62	$\mu\text{mol TE/g DW}$	(Juan and Chou 2010)	
			FRAP	2.10	3.50	$\mu\text{mol Fe (II)/g DW}$		
		80% Methanol	ABTS	1.24	1.79	$\mu\text{mol TE/g DW}$		
			FRAP	7.08	4.64	$\mu\text{mol Fe (II)/g DW}$		
		NaOH + HCl hydrolysis	ABTS	9.53	3.47	$\mu\text{mol TE/g DW}$		
			FRAP	2.10	3.39	$\mu\text{mol Fe (II)/g DW}$		
	<i>Lb. plantarum</i> WCFS1	80% Methanol	ABTS	1.24	1.76	$\mu\text{mol TE/g DW}$		
			FRAP	7.08	4.48	$\mu\text{mol Fe (II)/g DW}$		
		NaOH + HCl hydrolysis	ABTS	9.53	3.58	$\mu\text{mol TE/g DW}$		
			FRAP	2.10	2.86	$\mu\text{mol Fe (II)/g DW}$		
		80% Methanol	ABTS	1.24	1.91	$\mu\text{mol TE/g DW}$		
			DPPH	1.56 ^a	0.83 ^a	mg/mL		
Black soybean (<i>Glycine max</i>)	<i>B. subtilis</i> BCRC 14715	80% Methanol	DPPH	1.48 ^a	0.91 ^a	mg/mL		
			DPPH	0.94 ^a	0.65 ^a	mg/mL		
		Acetone	DPPH	2.28 ^a	1.65 ^a	mg/mL		
			Water	4.52 ^a	2.17 ^a	mg/mL		
		80% Methanol	Iron-chelating	8.97 ^a	4.57 ^a	mg/mL		
			Iron-chelating	12.0 ^a	8.30 ^a	mg/mL		
	<i>B. subtilis</i> BCRC 30222	80% Acetone	Iron-chelating	16.1 ^a	14.2 ^a	mg/mL		
			Water	1.95 ^a	0.70 ^a	mg/mL		
		Methanol	DPPH	1.95 ^a	1.39 ^a	mg/mL		
			DPPH	1.95 ^a	1.24 ^a	mg/mL		
		Methanol	DPPH	1.95 ^a	1.58 ^a	mg/mL		
			DPPH	1.95 ^a	2.11 ^a	mg/mL		
Black soybean (<i>Glycine max</i>)	<i>A. awamori</i>	Methanol	Iron-chelating	2.68 ^a	1.19 ^a	mg/mL		
			Iron-chelating	2.68 ^a	1.54 ^a	mg/mL		
		Methanol	Iron-chelating	2.68 ^a	1.67 ^a	mg/mL		
			Iron-chelating	2.68 ^a	1.82 ^a	mg/mL		
		Methanol	Iron-chelating	2.68 ^a	3.11 ^a	mg/mL		
			FRAP	17.5	13.6	$\mu\text{mol Fe (II)/g DW}$		
	<i>A. oryzae</i> BCRC 30222	80% Methanol	ABTS	15.4	10.6	$\mu\text{mol TE/g DW}$		
			FRAP	3.75	9.15	$\mu\text{mol Fe (II)/g DW}$		
		NaOH + HCl hydrolysis	ABTS	2.59	5.21	$\mu\text{mol TE/g DW}$		
			FRAP	17.5	16.8	$\mu\text{mol Fe (II)/g DW}$		
		Black soybean (<i>Glycine max</i>)	<i>Rhizopus</i> sp. No. 2	80% Methanol	ABTS	15.4	10.6	$\mu\text{mol TE/g DW}$
					FRAP	3.75	9.15	$\mu\text{mol Fe (II)/g DW}$
NaOH + HCl hydrolysis	ABTS			2.59	5.21	$\mu\text{mol TE/g DW}$		
	FRAP			17.5	16.8	$\mu\text{mol Fe (II)/g DW}$		
Natural fermentation	80% Methanol			ABTS	15.4	10.6	$\mu\text{mol TE/g DW}$	
				FRAP	3.75	9.15	$\mu\text{mol Fe (II)/g DW}$	
	NaOH + HCl hydrolysis		ABTS	2.59	5.21	$\mu\text{mol TE/g DW}$		
			FRAP	17.5	16.8	$\mu\text{mol Fe (II)/g DW}$		

(Continued)

Table 5–Continued.

Edible seeds and their products	Inoculated microbes	Extracts	Antioxidant assay	Antioxidant effect		Unit	References
				Nonfermented samples	Fermented samples		
Barley (<i>Hordeum vulgare</i>)	<i>Lb. plantarum</i> WCF51	NaOH + HCl hydrolysis	ABTS	15.4	11.8	$\mu\text{mol TE/g DW}$	(Dordevic and others 2010)
			FRAP	3.75	5.88	$\mu\text{mol Fe (II)/g DW}$	
	80% Methanol	ABTS	2.59	3.55	$\mu\text{mol TE/g DW}$		
		FRAP	17.5	17.8	$\mu\text{mol Fe (II)/g DW}$		
	NaOH + HCl hydrolysis	ABTS	15.4	12.2	$\mu\text{mol TE/g DW}$		
		FRAP	3.75	6.27	$\mu\text{mol Fe (II)/g DW}$		
	<i>Lb. rhamnosus</i>	70% ethanol	ABTS	2.59	3.65	$\mu\text{mol TE/g DW}$	
			FRAP	15.6	20.0	$\text{nmol Fe (II)/mg DW}$	
	<i>S. cerevisiae</i>	70% ethanol	FRAP	15.6	19.8	$\text{nmol Fe (II)/mg DW}$	
			DPPH	76.7	63.4 ^a	$\mu\text{g/ml}$	
Buckwheat (<i>Fagopyrum esculentum</i>)	<i>Lb. rhamnosus</i>	70% ethanol	FRAP	49.4	51.5	$\text{nmol Fe (II)/mg DW}$	
			DPPH	76.7	66.3 ^a	$\mu\text{g/ml}$	
Maize (<i>Zea mays</i> subsp. <i>mays</i>)	<i>S. cerevisiae</i>	70% ethanol	FRAP	49.4	49.8	$\text{nmol Fe (II)/mg DW}$	
			ABTS	4.85	~3.00 to 13.0	$\mu\text{mol TE/g DW}$	
Oat (<i>Avena sativa</i> L.)	<i>A. oryzae</i> NCIM 1212, <i>A. awamori</i> MTCC No. 548, <i>R. oligosporus</i> NCIM 1215, <i>R. oryzae</i> RCK2012	Water	DPPH	1.84	~2.00 to 5.80	$\mu\text{mol TE/g DW}$	(Salar and others 2012)
			ABTS	~15.5	~18.0 to 24.0	$\mu\text{mol TE/g DW}$	
Oat (<i>Avena sativa</i> L.)	<i>T. elegans</i> CCF-1456	65% Ethanol	DPPH	~12.0	~12.5 to 14.0	$\mu\text{mol TE/g DW}$	(Cai and others 2014)
			CVA	~1.60	~5.50 to 7.00	Q1 200, μC	
Oat (<i>Avena sativa</i> L.)	<i>A. oryzae</i> var. <i>effuses</i> 3.2825, <i>A. oryzae</i> 3.5232, <i>R. oryzae</i> 3.275	80% Ethanol	ORAC	~240	~350 to 430	$\mu\text{mol TE/g DW}$	(Dey and Kuhad 2014)
			ABTS	4.90	~6.00 to 20.5	$\mu\text{mol TE/g DW}$	
Oat (<i>Avena sativa</i> L.)	<i>A. oryzae</i> NCIM 1212, <i>A. awamori</i> MTCC No. 548, <i>R. oligosporus</i> NCIM 1215, <i>R. oryzae</i> RCK2012	Water	DPPH	2.98	~2.50 to 9.50	$\mu\text{mol TE/g DW}$	(Xiao and others 2015a)
			DPPH	9.13 ^a	5.86 ^a	mg/ml	
Oat (<i>Avena sativa</i> L.)	<i>C. militaris</i> SN-18	water	ABTS	3.62 ^a	1.18 ^a	mg/ml	
			Iron-chelating	16.7 ^a	8.10 ^a	mg/ml	
Oat (<i>Avena sativa</i> L.)	80% methanol	Iron-chelating	DPPH	3.71 ^a	2.78 ^a	mg/ml	
			DPPH	1.80 ^a	0.83 ^a	mg/ml	
Oat (<i>Avena sativa</i> L.)	80% ethanol	Iron-chelating	RP	6.83 ^a	4.32 ^a	mg/ml	
			DPPH	15.5 ^a	2.90 ^a	mg/ml	
Oat (<i>Avena sativa</i> L.)	80% ethanol	DPPH	2.96 ^a	2.77 ^a	mg/ml		

(Continued)

Table 5–Continued.

Edible seeds and their products	Inoculated microbes	Extracts	Antioxidant assay	Antioxidant effect		Unit	References
				Nonfermented samples	Fermented samples		
Brown rice (<i>Oryza sativa</i>)	<i>A. oryzae</i> NCIM 1212, <i>A. awamori</i> MTCC No. 548, <i>R. oligosporus</i> NCIM 1215, <i>R. oryzae</i> RCK2012	80% acetone	ABTS RP Iron-chelating DPPH ABTS RP Iron-chelating ABTS	2.11 ^a 6.23 ^a 8.03 ^a 2.54 ^a 1.65 ^a 4.49 ^a 0.88 ^a 4.30	0.94 ^a 3.75 ^a 4.71 ^a 2.26 ^a 0.85 ^a 3.41 ^a 0.56 ^a ~7.50 to 18.0	mg/mL mg/mL mg/mL mg/mL mg/mL mg/mL mg/mL μmol TE/g DW	(Dey and Kuhad 2014)
Black rice bran	<i>B. subtilis</i> KU3	Fermented supernatant	DPPH DPPH	1.15 0.58 ^a	~2.40 to 4.80 0.70 ^a	μmol TE/g DW mg/mL	(Yoon and others 2015)
Rye (<i>Secale cereale</i>)	<i>Lb. rhamnosus</i>	70% ethanol	CBA FRAP	52.9 ^a 8.94	47.8 ^a 13.9	mg/mL nmol Fe (II)/mg DW	(Dordevic and others 2010)
Wheat (<i>Triticum durum</i>)	<i>S. cerevisiae</i>	70% ethanol	FRAP	8.94	10.7	nmol Fe (II)/mg DW	
Wheat (<i>Triticum spp.</i>)	<i>Lb. rhamnosus</i>	70% ethanol	FRAP	12.2	15.1	nmol Fe (II)/mg DW	
	<i>S. cerevisiae</i>	70% ethanol	FRAP	12.2	12.3	nmol Fe (II)/mg DW	
	<i>A. oryzae</i> NCIM 1212, <i>A. awamori</i> MTCC No. 548, <i>R. oligosporus</i> NCIM 1215, <i>R. oryzae</i> RCK2012	Water	ABTS	3.85	~8.00 to 19.5	μmol TE/g DW	(Dey and Kuhad 2014)
	<i>G. gargal</i>	Methanol	DPPH DPPH	1.29 57.6 ^a	~3.70 to 8.60 0.56 ^a	μmol TE/g DW mg/mL	(Posternsky and Curvetto 2014)
	<i>G. sordulenta</i>	Methanol	RP DPPH	55.0 ^a 57.6 ^a	0.55 ^a 5.80 ^a	mg/mL mg/mL	
	<i>G. frondosa</i>	Methanol	RP DPPH	55.0 ^a 57.6 ^a	4.20 ^a 0.81 ^a	mg/mL mg/mL	
	<i>Gd. australe</i> (KUM60813)	Ethanol	DPPH	12.4	15.0	mg/mL mg AAE/g DW	
	<i>Gd. neo-japonicum</i> (KUM61076)	Ethanol	FRAP ABTS DPPH	2.19 8.74 12.4	1.05 9.73 21.7	mg Fe (II)/g DW mg TE/g DW mg AAE/g DW	
	<i>Gd. lucidum</i> (VITA GL)	Ethanol	FRAP ABTS DPPH	2.19 8.74 12.4	2.41 9.47 6.61	mg Fe (II)/g DW mg TE/g DW mg AAE/g DW	
Wheat (<i>Triticum aestivum</i> Linn.)	<i>C. militaris</i>	70% Ethanol	FRAP ABTS DPPH	2.19 8.74 12.4	0.80 14.2 0.06 ^a	mg Fe (II)/g DW mg TE/g DW mg/mL	(Zhang and others 2012)

(Continued)

Table 5--Continued.

Edible seeds and their products	Inoculated microbes	Extracts	Antioxidant assay	Antioxidant effect		Unit	References
				Nonfermented samples	Fermented samples		
Adlay (<i>Coix lachryma-jobi</i>)	<i>B. subtilis</i> BCRC 14718	70% Acetone	HRSP	0.21 ^a	0.11 ^a	mg/mL	(Wang and others 2014)
			Iron-chelating RP	0.61 ^a	0.26 ^a	mg/mL	
Chestnut (<i>Castanea crenata</i>)	<i>B. subtilis</i> BCRC 14718	Water	DPPH	0.52 ^a	0.58 ^a	mg/mL	(Wang and others 2014)
			Iron-chelating RP	<0.05 ^a	<0.05 ^a	mg/mL	
			HRSP	0.12 ^a	0.07 ^a	mg/mL	
			Iron-chelating RP	0.29 ^a	0.21 ^a	mg/mL	
			DPPH	0.23 ^a	0.13 ^a	mg/mL	
			Iron-chelating RP	0.38 ^a	0.32 ^a	mg/mL	
Lotus seed (<i>Nelumbo nucifera</i>)	<i>B. subtilis</i> BCRC 14718	Methanol	HRSP	0.76 ^a	0.61 ^a	mg/mL	(Wang and others 2014)
			Iron-chelating RP	0.79 ^a	0.44 ^a	mg/mL	
			DPPH	>0.80 ^a	>0.80 ^a	mg/mL	
			Iron-chelating RP	3.12	1.19	mg/mL	
			Iron-chelating DPPH	10.1	6.15	mg/mL	
			Iron-chelating DPPH	3.12	2.24	mg/mL	
Walnut (<i>Juglans regia</i>)	<i>B. subtilis</i> BCRC 14718	Methanol	Iron-chelating DPPH	10.1	8.09	mg/mL	(Wang and others 2014)
			Iron-chelating DPPH	2.43	1.89	mg/mL	
			Iron-chelating DPPH	6.83	5.73	mg/mL	
			Iron-chelating DPPH	2.43	2.31	mg/mL	
			Iron-chelating DPPH	6.83	6.51	mg/mL	
			Iron-chelating DPPH	2.93	1.67	mg/mL	
Soymilk	<i>B. subtilis</i> BCRC 14718	Methanol	Iron-chelating DPPH	15.8	11.3	mg/mL	(Wang and others 2006)
			Iron-chelating DPPH	2.93	1.83	mg/mL	
			Iron-chelating DPPH	15.8	12.0	mg/mL	
			Iron-chelating DPPH	1.92	0.80	mg/mL	
			Iron-chelating DPPH	7.70	6.83	mg/mL	
			Iron-chelating DPPH	1.92	0.89	mg/mL	
Walnut (<i>Juglans regia</i>)	<i>B. subtilis</i> BCRC 14718	Methanol	Iron-chelating DPPH	7.70	6.71	mg/mL	(Wang and others 2006)
			Iron-chelating DPPH	3.81	8.28	μmol CE/L	
			Iron-chelating DPPH	3.81	7.86	μmol CE/L	
			Iron-chelating DPPH	3.81	7.97	μmol CE/L	
			Iron-chelating DPPH	3.81	8.31	μmol CE/L	
			Iron-chelating DPPH	1.00	0.81 to 1.20	mg AAE/100 mL	
Soymilk	<i>Lb. acidophilus</i> CCRC 14079	Methanol	RP	1.00	0.58 to 1.10	mg AAE/100 mL	(Zhao and Shah 2014)
			RP	1.00	0.58 to 1.10	mg AAE/100 mL	
			RP	1.00	0.58 to 1.10	mg AAE/100 mL	
			RP	1.00	0.58 to 1.10	mg AAE/100 mL	
			RP	1.00	0.58 to 1.10	mg AAE/100 mL	
			RP	1.00	0.58 to 1.10	mg AAE/100 mL	

(Continued)

Table 5—Continued.

Edible seeds and their products	Inoculated microbes	Extracts	Antioxidant assay	Antioxidant effect		Unit	References
				Nonfermented samples	Fermented samples		
	<i>Lb. zeae</i> ASCC 15820	80% Methanol + Hexane	RP	1.00	0.64 to 1.21	mg AAE/100 mL	
	<i>Lb. rhamnosus</i> WQ2	80% Methanol + Hexane	RP	1.00	0.74 to 1.60	mg AAE/100 mL	
	<i>Lb. plantarum</i> CCFM8610	Unknown	RP	25.0	117	umol CE/L	(Zhai and others 2015)
	<i>Lb. bulgaricus</i> CCFM8004	Unknown	RP	25.0	81.5	umol CE/L	
Soy germ	<i>A. niger</i> M46	70% Methanol	HRSP	164 ^a	0.8 ^a	ug/mL	(Sheih and others 2014)
Soy whey	<i>Lb. plantarum</i> B1-6	80% Ethanol	ABTS	0.72 ^a	0.19 ^a	mg/mL	(Xiao and others 2015b)
			RP	1.36 ^a	1.16 ^a		
			HRSP	0.12 ^a	0.03 ^a		
			SASP	2.71 ^a	2.30 ^a		
			ABTS	22.9	15.7	μmol TE/g DW	(Malgorzata and others 2015)
Buckwheat groats (<i>Fagopyrum esculentum</i> Moench)	<i>R. oligosporus</i> NRRL 2710	80% methanol	ACL PCL ACW PCL	17.3 3.97	11.2 4.57	μmol TE/g DW μmol TE/g DW	
			DPPH	9.46	10.8	μmol TE/g DW	(Starzynska-Jamiszewska and others 2016)
Buckwheat groats (<i>Fagopyrum esculentum</i>)	<i>R. oligosporus</i> ATCC 64063	50% Acetone	ABTS	27.2	42.7	μmol TE/g DW	
		PBS (0.1 M, pH 7.4)					

AAE, ascorbic acid equivalent; ABTS, ABTS free radical scavenging value; CVA, Cyclic voltammetry assay; DPPH, DPPH free radical scavenging value; ORAC, oxygen radical absorbance capacity; RP, reducing power; HRSP, hydroxyl radical scavenging capacity; SASP, superoxide anion scavenging capacity; CBA, β-carotene bleaching assay; ACL PCL, lipophilic-based superoxide anion radicals scavenging value; ACW PCL, hydrophilic-based superoxide anion radicals scavenging value; CE, cysteine equivalent; TE, trolox equivalent; VCE/AAE, vitamin C/Ascorbic acid equivalent; DW, dry weight; N.M., not measured. A., *Aspergillus*; B., *Bacillus*; Bb, *Bifidobacterium*; C., *Corylopsis*; G., *Critifol*; Gd, *Conoderma*; Lb., *Lactobacillus*; R., *Rhizopus*; S., *Saccharomyces*; Sc., *Streptococcus*; T., *Thamnidium*. ^a EC₅₀ value.

Table 6—Anti-hypertensive effect of fermented edible seeds and their products.

Edible seeds and their products	Inoculated microbes	Anti-hypertensive effect	Main anti-hypertensive components	References
Chickpea (<i>Cicer arietinum</i> L.)	<i>C. militaris</i> SN-18	Inhibit ACE activity <i>in vitro</i>	Potential ACE-inhibitory peptides	(Xiao and others 2015c)
Lentil (<i>Lens culinaris</i> L.)	<i>Lb. plantarum</i> CECT 748 ^T	Inhibit ACE activity <i>in vitro</i>	Potential ACE-inhibitory peptides	(Torino and others 2013)
Mung bean (<i>Vigna radiata</i>)	<i>Lb. plantarum</i> B1-6	Inhibit ACE activity <i>in vitro</i>	Potential ACE-inhibitory peptides	(Wu and others 2015)
Pea seeds (<i>Pisum sativum</i> var. Bajka)	<i>Lb. plantarum</i> 299v.	Inhibit ACE activity <i>in vitro</i>	ACE-inhibitory peptides	(Jakubczyk and others 2013)
Soybean (<i>Glycine max</i>)	<i>M. pilosus</i> KFRI-1140	Inhibit ACE activity <i>in vitro</i>	Potential ACE-inhibitory peptides	(Pyo and Lee 2007)
Soy protein	<i>Lb. casei</i> spp. pseudoplantarum	Inhibit ACE activity <i>in vitro</i>	ACE-inhibitory peptides	(Vallabha and Tiku 2014)
Black soybean (<i>Glycine max</i>)	<i>B. subtilis</i> BCRC 14715 and <i>Bacillus</i> sp. CN11	Inhibit ACE activity <i>in vitro</i>	Potential ACE-inhibitory peptides, nicotiamide, isoflavone aglycones and other flavonoids	(Juan and others 2010)
A mixture of black beans, soybeans, and wheat bran (1:1:1, w/w/w)	<i>B. subtilis</i> B060	Lower blood pressure in hypertensive rats	GABA and nattokinase	(Suwanmanon and Hsieh 2014)
Navy bean (<i>Phaseolus vulgaris</i>) milk	Different LAB	Inhibit ACE activity <i>in vitro</i>	ACE-inhibitory peptides	(Rui and others 2015)
Soymilk	A mixture of LAB	Inhibit ACE activity <i>in vitro</i> and Lower blood pressure in hypertensive rats	ACE-inhibitory peptides and GABA	(Tsai and others 2006)
	Different LAB	Inhibit ACE activity <i>in vitro</i>	N.M.	(Ewe and others 2011)
	<i>Lb. plantarum</i> TWK10	Inhibit ACE activity and lower blood pressure in hypertensive rats	N.M.	(Liu and others 2016)
Soy yogurt	Different LAB	Inhibit ACE activity <i>in vitro</i>	Potential ACE-inhibitory peptides	(Donkor and others 2005)
Soy flour sourdoughs	Different LAB	Inhibit ACE activity <i>in vitro</i>	Potential ACE-inhibitory peptides	(Omedi and others 2016)
Tofuyo (based on soybean)	Unknown	Inhibit ACE activity <i>in vitro</i>	ACE-inhibitory peptides	(Kuba and others 2003)
Casein miso paste (based on rice and soybean)	<i>A. oryzae</i>	Lower blood pressure in hypertensive rats	ACE-inhibitory peptides	(Inoue and others 2009)
A <i>dosa</i> made of wheat and rice flour	<i>Lb. plantarum</i> MNZ	Inhibit ACE activity <i>in vitro</i> and Lower blood pressure in hypertensive rats	GABA and potential bioactive peptides	(Zareian and others 2015)
White wheat, wholemeal wheat and rye	Different LAB	Inhibit ACE activity <i>in vitro</i>	ACE-inhibitory peptides	(Rizzello and others 2008)

N.M., not mentioned; ACE, angiotensin converting-1-enzyme; GABA, γ -aminobutyric acid; LAB, lactic acid bacteria; *Lb.*, *Lactobacillus*; *A.*, *Aspergillus*; *B.*, *Bacillus*; *C.*, *Cordyceps*; *M.*, *Monascus*.

renin–angiotensin–aldosterone system in humans. As a central component of the system, ACE functions to convert angiotensin I into angiotensin II, which is able to increase blood pressure by directly causing blood vessels to constrict (Skeggs and others 1956). As a result, ACE plays a central role in the control of blood pressure, and natural or synthetic compounds with ACE-inhibitory activity have been reported to lower blood pressure in experimental animals and humans (Fang and others 2008).

Recent studies demonstrate that some fermented edible seeds and their products exhibit anti-hypertensive effects *in vitro* and *in vivo*. A variety of fermented edible seeds and their products, mainly edible bean products, such as fermented chickpea, lentil, mung bean, pea, soybean, soymilk, and navy bean milk, have been reported to possess ACE-inhibitory activity *in vitro* or lower blood pressure in animal models (Table 6). However, the anti-hypertensive effect of fermented grains has been less investigated. Their bioactive components, such as GABA, ACE-inhibitory peptides, nattokinase, vitamins, and various antioxidant phenolics, can be responsible for the anti-hypertensive effect (Kim and others 2008; Lee and Pan 2012; Huang and others 2013). Especially, GABA and ACE-inhibitory peptides are found in many fermented edible seeds and their products (Table 2 and 4), which have been reported to be predominantly responsible for their anti-hypertensive effect (Table 6). As a result, fermented edible seeds and their products can be an important dietary component consumed by people to prevent hypertension. In addition, in light of the importance of

cereal grains as staple foods for humans, more studies are needed to investigate the potential anti-hypertensive effect of fermented cereal grains in the future.

Anti-cancer effect

The anti-cancer effect has been investigated in different fermented edible seeds and their products, such as fermented soybean, black bean, green bean, soymilk, wheat germ, and rice (Table 7). LAB and fungi are the main inoculated microbes to produce anti-cancer products. Various bioactive compounds, such as natural phenolics, peptides, amino acids, benzoquinones, GABA, polysaccharides, monacolin K, and vitamin E, have been suggested to be potential anti-cancer components. Below, we highlight the anti-cancer effect of *S. cerevisiae*-fermented wheat germ and *A. oryzae*-fermented brown rice and rice bran, which have been extensively investigated for their anti-cancer effect.

S. cerevisiae-fermented wheat germ was first produced by Dr. Mate Hidvegi in the early 1990s with anti-cancer effect in animal models and cancer patients, and subsequently developed into a commercial product, with the brand name Avemar (Demidov and others 2008; Rizzello and others 2013). This fermented wheat germ has been reported to exhibit anti-cancer effect on a variety of cancer cells, such as leukemia, lymphoid, pediatric, colon, skin melanoma, ovarian, and liver cancer/tumor cells *in vitro* and *in vivo* (Table 7). Two benzoquinones, 2-methoxy benzoquinone (2-MBQ) and 2,6-dimethoxy-benzoquinone (2,6-DMBQ), have

Table 7—Anti-cancer effect of fermented edible seeds and their products.

Edible seeds and their products	Inoculated microbes	Possible effective components	Actions and potential mechanisms	References
Soybean, black bean, and green bean mixture	<i>Lb. paracasei</i> and <i>S. cerevisiae</i>	Isoflavones	Induce cancer cell apoptosis <i>in vitro</i> and inhibit tumor xenografts <i>in vivo</i>	(Chia and others 2013)
Italian legumes	<i>Lb. plantarum</i> C48 and <i>Lb. brevis</i> AM17	Lunasin-like polypeptides	Inhibit cancer cell proliferation <i>in vitro</i>	(Rizzello and others 2015)
Soybean and mung bean	<i>Rhizopus</i> sp. 5351	GABA, free amino acids and phenolics	Induce cytotoxicity by blocking G0/G1 cell cycle phase and inducing apoptosis <i>in vitro</i>	(Ali and others 2016)
Soymilk	<i>Sc. thermophilus</i> 14085 and <i>Bb. infantis</i> 14603	Phenolics	Inhibit cancer cell proliferation <i>in vitro</i>	(Lai and others 2013)
Wheat germ	<i>S. cerevisiae</i>	Benzoquinones	Inhibit proliferation and induce apoptosis in Jurkat T-cell leukemia tumor cells <i>in vitro</i>	(Comin-Anduix and others 2002)
	<i>S. cerevisiae</i>	2,6-DMBQ	Induce apoptosis and reduce MHC class I expression in lymphoid tumor cells <i>in vitro</i>	(Fajka-Boja and others 2002)
	<i>S. cerevisiae</i>	N.M.	Reduce chemotherapy-induced febrile neutropenia in pediatric cancer patients	(Carami and others 2004)
	<i>S. cerevisiae</i>	N.M.	Inhibit proliferation, and induced both necrosis and apoptosis in human HT-29 colon cancer cells <i>in vitro</i>	(Illmer and others 2005)
	<i>S. cerevisiae</i>	N.M.	Induce apoptosis and inhibit ribonucleotide reductase in human HL-60 promyelocytic leukemia cells <i>in vitro</i>	(Saiko and others 2007)
	<i>S. cerevisiae</i>	N.M.	Improve the survival of high-risk skin melanoma patients	(Demidov and others 2008)
	<i>S. cerevisiae</i>	N.M.	Attenuate the growth and induce apoptosis in human H9 lymphoma cells <i>in vitro</i>	(Saiko and others 2009)
	<i>S. cerevisiae</i>	2-MBQ and 2,6-DMBQ	Induce apoptosis in a variety of cancer cells <i>in vitro</i>	(Mueller and others 2011)
	<i>S. cerevisiae</i>	2,6-DMBQ	Inhibit proliferation and potentiate cisplatin-induced apoptosis human ovarian cancer cells <i>in vitro</i>	(Judson and others 2012)
	<i>Lb. plantarum</i> LB1 and <i>Lb. rossiae</i> LB5	2-MBQ and 2,6-DMBQ	Inhibit cancer cell proliferation <i>in vitro</i>	(Rizzello and others 2013)
	<i>S. cerevisiae</i>	N.M.	Induce cell death and enhance cytotoxicity of cisplatin and 5-fluorouracil on human liver cancer cells <i>in vitro</i>	(Tai and others 2013)
	<i>S. cerevisiae</i>	N.M.	Inhibit proliferation and induce apoptosis in human ovarian cancer cells <i>in vitro</i>	(Wang and others 2015)
	<i>Lb. plantarum</i> dy-1 (LFWGE)	2,6-DMBQ	Inhibit proliferation and induce apoptosis in human HT-29 colon cancer cells <i>in vitro</i>	(Zhang and others 2015a)
	<i>Lb. plantarum</i> dy-1 (LFWGE)	N.M.	Induce colon cancer cell apoptosis in xenograft mouse model	(Zhang and others 2015b)
Brown rice and rice bran	<i>S. cerevisiae</i>	2,6-DMBQ	Inhibit proliferation <i>in vitro</i> , probably associated with the induction of autophagy	(Otto and others 2016)
	<i>A. oryzae</i>	N.M.	Inhibit the formation of AOM-induced aberrant crypt foci and tumors in the colon of F344 rats	(Katayama and others 2002)
	<i>A. oryzae</i>	N.M.	Decrease the incidence and multiplicity of hepatocellular carcinoma in F344 rats	(Katayama and others 2003)
	<i>A. oryzae</i>	N.M.	inhibit NMBA-induced esophageal tumor development in rats possibly through inhibition of cell proliferation	(Kuno and others 2004)
	<i>A. oryzae</i>	N.M.	Inhibit chemical-induced urinary bladder carcinogenesis in mice through anti-proliferative mechanisms	(Kuno and others 2006)
	<i>A. oryzae</i>	N.M.	Suppress the development of 4-NQO-induced oral carcinogenesis in rats via inhibiting proliferation and scavenging free radicals	(Long and others 2007)
	<i>A. oryzae</i>	N.M.	Inhibit MNNG-induced development of gastric tumors in rats	(Tomita and others 2008)
	<i>A. oryzae</i>	N.M.	Inhibit NNK-induced pulmonary tumorigenesis in A/J mice via induction of Cyp2a5 in the lung and the reduced proliferation rate of tumor cells	(Phutthaphadoong and others 2009)
	<i>A. oryzae</i>	N.M.	Inhibit inflammation-related colorectal carcinogenesis in Apc(Min/+) mice	(Phutthaphadoong and others 2010)
	<i>A. oryzae</i>	N.M.	Inhibit N-nitrosobis (2-oxopropyl) amine-induced pancreatic tumorigenesis in male hamsters	(Kuno and others 2015)
	<i>A. oryzae</i>	Vitamin E and phenolic acids	Decrease the incidence and progression of prostate carcinogenesis in rats	(Kuno and others 2016)
Rice bran	<i>A. oryzae</i>	N.M.	Induce apoptosis of human acute lymphoblastic leukemia cells <i>in vitro</i>	(Horie and others 2016)
Brown rice	<i>L. edodes</i>	Polysaccharides	Inhibit the growth of B16/B16 melanoma xenograft in mice via enhancing natural killer cell activity	(Kim and others 2007)
Rice	<i>A. oryzae</i>	N.M.	Induce apoptosis of human HCT116 colorectal cancer cells <i>in vitro</i> by activating mitochondrial pathway	(Itoh and others 2012)
	<i>M. purpureus</i> NTU 803	Monacolin K and phenolics	Induce apoptosis and cell cycle arrest at G2/M phase in human MCF-7 breast cancer cells <i>in vitro</i>	(Lee and others 2013)
Black rice bran	<i>B. subtilis</i> KU3	Phenolics	Inhibit proliferation <i>in vitro</i>	(Yoon and others 2015)

A., *Aspergillus*; B., *Bacillus*; Bb., *Bifidobacterium*; L., *Lentinus*; Lb., *Lactobacillus*; M., *Monascus*; S., *Saccharomyces*; Sc., *Streptococcus*; 2-MBQ, 2-methoxybenzoquinone; 2,6-DMBQ, 2,6-dimethoxy-1,4-benzoquinone; 4-NQO, 4-nitroquinoline 1-oxide; AOM, azoxymethane; GABA, γ -aminobutyric acid; MHC, major histocompatibility complex; MNNG, N-nitrosomethyl-N'-nitro-N-nitrosoguanidine; NMBA, N-nitrosomethylbenzylamine; NNK, 4-(methylnitrosamino)-1-(3-pyridyl)-1-butanone; N.M., not mentioned.

Table 8—Other bioactivities of fermented edible seeds and their products.

Bioactivities	Edible seeds and their products	Inoculated microbes	Possible effective components	Actions and potential mechanisms	References
Anti-depressant effect Anti-diabetic effect	Black soybean milk Soybean and rice bran	<i>Lb. brevis</i> FPA 3709 <i>Bacillus</i> spp.	GABA N.M.	Reduce the duration of immobility in rats Reduce serum glucose, HbA1c and triglyceride in diabetic mice and stimulate glucose uptake via activation of PI3K/Akt signaling in C2C12 cells <i>in vitro</i>	(Ko and others 2013) (Lim and Lee 2010)
	Mung bean	<i>Rhizopus</i> sp. strain 5351	GABA and free amino acids	Reduce blood sugar and lipids, and improve insulin secretion and antioxidant level in diabetic mice	(Yeap and others 2012)
	Soymilk	<i>Lb. rhamnosus</i> CRL981	Isoflavone aglycones	Ameliorate hyperglycemia, lipid profiles and increase antioxidant enzyme activities in diabetic mice	(Marazza and others 2013)
	Bambara groundnut, African locust bean, and soybeans	Natural fermentation	Bioactive polypeptides and flavonoids	Ameliorate hyperlipidemia and inhibit ACE activity in streptozotocin-induced diabetic rats	(Ademiluyi and Oboh 2015)
	Wheat	<i>Gd. neo japonicum</i> (KUM61076) mutant <i>M. purpureus</i> 254	N.M.	Possess insulin-like potential <i>in vitro</i>	(Subramaniam and others 2015)
	Rice	<i>Lb. plantarum</i> LB1 and <i>Lb. rossiae</i> LB5 N.M.	N.M.	Lower blood glucose and alter lipid profiles in diabetic rats	(Rajasekaran and Kalaivani 2015)
Anti-fungal effect	Wheat germ		Organic acids and peptides	Inhibit various fungi isolated from bakeries	(Rizzello and others 2011)
Anti-inflammatory effect	Wheat germ		N.M.	Alleviate severe rheumatoid arthritis in patients	(Balint and others 2006)
	Brown rice	<i>A. oryzae</i>	N.M.	Inhibit colonic inflammation in rat colon by decreasing the ulcer and erosion area and myeloperoxidase activity in the colonic mucosa	(Kataoka and others 2008)
	Rice bran	<i>I. orientalis</i> MFST1	N.M.	Suppress allergic and inflammatory reactions through inhibition of degranulation, histamine release, and pro-inflammatory cytokine production from mast cells	(Fan and others 2010)
Anti-obesity effect	Brown rice and rice bran	<i>A. oryzae</i>	N.M.	Inhibit inflammatory cell infiltration in mice	(Onuma and others 2015)
	Rice bran	<i>I. orientalis</i> MFST1	Phenolics	Ameliorate oxidative stress-induced insulin resistance in 3T3-L1 adipocytes <i>in vitro</i>	(Kim and Han 2011)
	Soymilk	<i>Lb. paracasei</i> subsp <i>paracasei</i> NTU 101	N.M.	Reduce body weight and lipogenesis in obese rats	(Cheng and others 2015)
	Soybean	<i>B. subtilis</i> MORI	Isoflavones	Inhibit the differentiation of 3T3-L1 preadipocytes and facilitate its glucose utilization	(Hwang and others 2015)
Anti-stress and fatigue effects	Rice bran	<i>S. cerevisiae</i> IFO 2346	N.M.	Inhibit major changes in the weight of the adrenal, thymus, spleen and thyroid and prolong the swimming time in rats and mice	(Kim and others 2001, Kim and others 2002)
Anti-trypanosomal activity	Wheat germ	<i>S. cerevisiae</i>	Glycoside, alkaloids and saponins	Decrease in the proliferation of <i>T. brucei</i> , and extend the surviving days of <i>T. brucei</i> -infected rats	(Yusuf and Ekanem 2010)
Cardiovascular protective effect	Red rice	<i>M. ruber</i> IFO32318	N.M.	Induce NO-mediated endothelium-dependent relaxation of rat thoracic aorta	(Rhyu and others 2000)
	Rice	<i>Monascus</i> spp.	N.M.	Suppress hypertriglyceridemia and hyperlipidemia in rats fed with high-fructose diet	(Wang and others 2000)
	A mixture of coffee bean, soybean and rice bran	<i>B. subtilis</i> Natto	N.M.	Inhibit the TBARS formation and atherosclerotic lesions in the aorta of rabbits	(Naito and others 2003)
	Rice	<i>M. purpureus</i>	N.M.	Reduce serum TC, TG, LDL-C and ameliorate the severity of atherosclerosis in cholesterol fed rabbits	(Wei and others 2003)
	Brown rice	<i>Lb. acidophilus</i> KCTC 2182	N.M.	Lower plasma and hepatic TG, TC, LDL-C and VLDL-C, and increase HDL-C in cholesterol-fed rats	(Baek and others 2005)
	Rice	<i>Basidiomycola</i> (sangwhang) and <i>M. ruber</i> N.M.	N.M.	Improve the lipid metabolism and reduce oxidative stress in high-cholesterol-fed rats via upregulating the hepatic antioxidant enzymes	(Jang and others 2007)
	Wheat germ		N.M.	Attenuate chronic hypertension, diabetes or metabolic syndrome-induced cardiovascular symptoms along with metabolic abnormalities	(Iyer and Brown 2011)

(Continued)

Table 8—Continued.

Bioactivities	Edible seeds and their products	Inoculated microbes	Possible effective components	Actions and potential mechanisms	References
	Rice	<i>M. purpureus</i>	Naturally occurring statin	Inhibit TNF α -induced upregulation of MMP2 and 9 and intracellular ROS in human aortic smooth muscle cells <i>in vitro</i>	(Lin and others 2011)
	Rice bran and soybean	<i>Bacillus</i> spp.	N.M.	Decrease total lipids and triglyceride in the serum and liver of rats fed with high fat diet	(Lim and others 2011)
	Indian rice	<i>M. purpureus</i> MTCC 1090	N.M.	Ameliorate lipid profiles and oxidative stress markers in high cholesterol diet fed rats	(Rajasekaran and Kalaivani 2011)
	Rice	<i>M. purpureus</i>	Antioxidants and amino acids	Reduce the cholesterol level enhance the antioxidant level in hypercholesterolemia mice	(Yeap and others 2014)
	Wheat powder	<i>Lactobacilli</i> and natural yeast strains	Phenolics, lipic acid, tocopherols and polyunsaturated fatty acids	Improve blood lipids and oxidative status in healthy rabbits	(Pozzo and others 2015)
	Mung bean	<i>Rhizopus</i> sp. strain 5351	GABA and phenolics	Increase antioxidant activity, reduce blood lipids and regulate atherosclerosis related gene expression in hypercholesterolemia mice	(Yeap and others 2015)
Gastrointestinal protective effect	Brown rice	<i>A. oryzae</i>	Nucleobases	Improve the incidence rate of diarrhea, increase the protein content in small intestinal mucosa, and improve the survival rate in methotrexate-treated rats	(Ochiai and others 2013)
Hepatoprotective effect	Waxy brown rice	<i>Cr. cinereus</i>	N.M.	Inhibit carbon tetrachloride induced hepatotoxicity in rats	(Lee and others 2004)
	Brown rice and rice bran	<i>A. oryzae</i>	N.M.	Inhibit acute hepatitis in Long-Evans Cinnamon rats via protecting the liver against free radicals induced by copper accumulation in the liver	(Shibata and others 2006)
	Waxy brown rice	<i>Ac. cylindracea</i>	N.M.	Attenuate carbon tetrachloride-induced hepatotoxicity in rats	(Lee and others 2006)
	Rice	<i>M. purpureus</i> NTU 568	Monacolin K, GABA, dimeric acid and various pigments	Attenuate oxidative stress, inflammatory response, and steatosis in mice with alcoholic liver disease	(Cheng and Pan 2011)
	Soymilk	<i>Sc. thermophilus</i> grx02	Phenolics	Ameliorate alcohol-induced liver damage in mice by lowering ALT, AST and enhancing ADH and SOD activity as well as GSH content	(Xu and others 2012)
	Mung bean	<i>Rhizopus</i> sp. strain 5351	GABA and amino acids	Improve antioxidant activity, serum markers (ALT, AST, TG and TC) and NO level in ethanol-induced liver damage of mice	(Ali and others 2013)
	Rice bran	<i>Bacillus</i> species	N.M.	Ameliorate CCl4-induced hepatic fibrosis in mice	(Park and others 2014)
	Thai glutinous rice	<i>M. purpureus</i> CMU 002U	N.M.	Reduce blood and hepatic cholesterol and hepatic steatosis in hypercholesterolemic rats	(Bunnoy and others 2015)
Laxative effect	Rice	<i>S. cerevisiae</i> and <i>W. paramesenteroides</i>	N.M.	Exhibit a laxative effect without causing diarrhea in rats with loperamide-induced constipation	(Choi and others 2014a)
	Rice	N.M.	N.M.	Ameliorate loperamide-induced constipation in rats	(Choi and others 2014b)
Immunomodulatory effect	Rice bran	<i>S. cerevisiae</i>	N.M.	Activate macrophage and enhance secretion of hematopoietic growth factors from Peyer's patch cells of mice	(Koh and others 2003)
	Rice bran	<i>L. edodes</i>	Arabinoxylan	Increase IFN- γ production without causing adverse effects in health people	(Choi and others 2014c)
	Wheat germ	N.M.	N.M.	Induce a better immune response compared to pigs exposed to T-2 toxin	(Arpad and others 2009)
	Wheat germ	N.M.	N.M.	Enhance the blastogenic response of lymphocytes to nonspecific mitogens, phagocytic activity and phagocytic index in pigs	(Rafai and others 2011)
	nonsalty soybean powder	N.M.	Phenolics	Stimulate the cellular immune response, but suppress the acquired humoral immune response in C3H/HeN mice	(Karasawa and others 2013)

(Continued)

Table 8—Continued.

Bioactivities	Edible seeds and their products	Inoculated microbes	Possible effective components	Actions and potential mechanisms	References
Neuroprotective effect	Soybean	<i>Lb. paracasei</i> , <i>Lb. bulgaricus</i> and <i>S. cerevisiae</i>	Isoflavones	Induce T-cell proliferation, enhance the function of Th1 cells and activity of NK cells in mice	(Chin and others 2015)
	Mung bean and soybean	<i>Rhizopus</i> sp. 5351	GABA, free amino acids and phenolics	Induce splenocyte proliferation and enhance the levels of serum IL-2 and IFN- γ <i>in vitro</i>	(Ali and others 2016)
	Chickpea milk	<i>Lb. plantarum</i> M-6	GABA	Inhibit MnCl ₂ -induced PC12 cell death partially by retaining the integrity of cell membrane	(Li and others 2016)
Heavy metal protective effect DNA protective effect	Soymilk Rice	<i>Lb. plantarum</i> TWK10 <i>M. purpureus</i> NTU 568	N.M. DFC, DMA	Improve learning and memory in rats	(Liu and others 2016)
	Soymilk	<i>Lb. plantarum</i> CCFM810	Isoflavone aglycones	Ameliorate 6-hydroxydopamine-induced neurotoxicity in SH-SY5Y cells and the rat model of Parkinson's disease	(Tseng and others 2016a)
	Chickpea, mung bean, oats (<i>Avena sativa</i> L.)	<i>C. militaris</i> SN-18	Phenolics	Increase fecal Cd excretion, reduce tissue Cd burden, alleviate tissue oxidative stress in mice	(Zhai and others 2015)
Regulation of gut microbiota	Soy whey	<i>Lb. plantarum</i> B1–6	Phenolics, especially isoflavone aglycones	Inhibit hydroxyl radical-induced supercoiled DNA strand scission <i>in vitro</i>	(Xiao and others 2014, Xiao and others 2015a, Xiao and others 2015d)
	Brown rice	<i>A. oryzae</i>	N.M.	Inhibit hydroxyl radical-induced supercoiled DNA strand increase the number of lactobacilli species already resident in the rat intestine	(Xiao and others 2015b)
Regulation of sleep	Rice germ	N.M.	GABA	Counter the sleep disturbance induced by caffeine in mice	(Kataoka and others 2007)
Reproductive protective effect	Rice	<i>M. purpureus</i>	N.M.	Prevent Zn deficiency-induced testis and sperms injury in rats	(Lee and others 2011)
Skin protective effect	Rice bran	<i>Lb. rhamnosus</i> and <i>S. cerevisiae</i>	N.M.	Inhibit α -MSH-induced Melanogenesis in B16F1 Melanoma via downregulating MITF-Expression	(Chung and others 2009)
	Rice bran	<i>Lb. rhamnosus</i> and <i>S. cerevisiae</i>	N.M.	Increase the synthesis of type I collagen, decrease the expression of MMP-1, and inhibit the production of IL-1 α in UV-B irradiated human fibroblasts <i>in vitro</i>	(Seo and others 2010)
	Soymilk	<i>Lb. plantarum</i> TWK10	Isoflavone aglycones	Inhibit mushroom tyrosinase activity and melanin production in B16F0 melanocytes	(Chen and others 2013)

A., *Aspergillus*; *Ac.*, *Agroclype*; *Bb.*, *Bifidobacterium*; *C.*, *Coryzeus*; *Cr.*, *Coryzeus*; *Gd.*, *Ganoderma*; *I.*, *Issatchia*; *Lenkig.*, *L.*, *Lentinus*; *M.*, *Monascus*; *S.*, *Saccharomyces*; *Sc.*, *Streptococcus*; *T.*, *Trypanosoma*; *W.*, *Weissella*; *ACE*, angiotensin converting I-enzyme; *ADH*, alcohol dehydrogenase; *ALT*, alanine transaminase; *AST*, aspartate transaminase; *Cd*, cadmium; *DFC*, dimeric acid; *DMA*, dimeric acid; *DMBG*, 2,6-dimethoxy-1,4-benzoquinone; *GABA*, γ -aminobutyric acid; *GSH*, glutathione; *MDL-C*, high-density lipoprotein cholesterol; *IFN- γ* , interferon γ ; *IL-2*, interleukin-2; *LDL-C*, low-density lipoprotein cholesterol; *MMP*, matrix metalloproteinase; *MHC*, histocompatibility complex class II; *MIF*, microphthalmia-associated transcription factor; *MnCl₂*, manganese(II) chloride; *NK* cells, natural killer cells; *NO*, nitric oxide; *SOD*, superoxide dismutase; *ROS*, reactive oxygen species; *TBARS*, thiobarbituric acid reactive substances; *Tc*, total cholesterol; *TG*, triglycerides; *Th1* cells, T helper type 1 cells; *VLDL-C*, very low-density lipoprotein cholesterol; *N.M.*, not mentioned.

been reported as the main anti-cancer compounds in it (Mueller and others 2011; Rizzello and others 2013). *S. cerevisiae*-fermented wheat germ exhibits different actions on cancer cells (Table 7). It can inhibit cancer cell proliferation, induce cancer cell necrosis and apoptosis, potentiate the efficiency of chemotherapy, as well as improve the survival of cancer patients. For the anti-cancer mechanism, it can mainly induce cell cycle arrest and activate apoptotic signaling. Studies found that it inhibited the G1 phase cell-cycle progression in HT-29 cells (Illmer and others 2005), blocked cell cycle from G2-M to G0-G1 phase in human HL-60 promyelocytic leukemia cells (Saiko and others 2007), and activated caspase-3 and caspase-7-dependent apoptotic signaling in human ovarian cancer cells (Wang and others 2015). In addition, LAB-fermented wheat germ has also been reported to exhibit anti-cancer effect (Rizzello and others 2013; Zhang and others 2015a,b).

A. oryzae-fermented brown rice and rice bran is another extensively investigated food with anti-cancer effect. It has also been reported to inhibit various cancer cells *in vitro* and *in vivo* (Table 7), mainly by inhibiting cancer cell proliferation and inducing cancer cell apoptosis. Kuno and others (2016) reported that it controlled the growth of prostate tumor *in vivo* by activating AMP-activated protein kinase (AMPK) signaling. Another study suggested that it induced the apoptosis of human acute lymphoblastic leukemia cells probably through the death receptor-mediated apoptotic pathway (Horie and others 2016). However, its anti-cancer components have not been clarified, with vitamin E and phenolic acids having been suggested to be responsible for its anti-cancer effect (Kuno and others 2016). In addition, fermented common rice, black rice, and their brans have also been reported to possess anti-cancer effects on different cancer cells (Table 7).

Overall, fermented wheat and rice, especially their germ and bran, can be excellent anti-cancer foods consumed by humans for the prevention and treatment of cancer. On the other hand, the anti-cancer effect of other fermented edible beans, grains, and their products has been less investigated, and future studies are required to investigate their anti-cancer effect and anti-cancer components.

Other bioactivities

Fermented edible seeds and their products have also been reported to possess many other bioactivities (Table 8), such as anti-depressant, anti-diabetic, anti-fungal, anti-inflammatory, anti-obesity, anti-stress and -fatigue, anti-trypanosomal, cardiovascular protective, gastrointestinal protective, hepatoprotective, neuroprotective, reproductive protective, skin protective, DNA-protective, heavy metal protective, laxative, immunomodulatory, gut microbiota regulatory, and sleep regulatory effects. Likewise, bioactive components, including GABA, phenolics, bioactive peptides, naturally occurring statins, nucleobases, amino acids, tocopherols, polyunsaturated fatty acids, monacolin K, dimeric acid, and arabinoxylan, have been proposed to be associated with the bioactivities (Table 8).

In general, fermented edible seeds and their products possess versatile bioactivities, implying that they should have comprehensive health benefits. Therefore, it is recommended to consume fermented edible seeds and their products as a part of the diet to prevent chronic diseases such as cancer and cardiovascular diseases.

Potential problems of food safety

Although fermented edible seeds and their products have been demonstrated to contain a variety of bioactive components and to confer a great number of health benefits, their potential safety

problem cannot be ignored. However, it should be pointed out that there are few studies focused on the potential safety problems of fermented products. On one hand, fermented seed products without sterilization may contain pathogenic microbes or those with potential pathogenic risk, especially naturally fermented products. For example, naturally fermented and LAB-fermented soybeans and lentils have been reported to contain substantial numbers of aerobic mesophilic bacteria (Fernandez-Orozco and others 2007; Torino and others 2013), which may include pathogenic bacteria and increase the risk of food poisoning. On the other hand, microbes may release toxic or harmful substances during fermentation, which may be harmful for health and cannot be excluded without toxicological analysis. Recent emerging food technologies, such as ultraviolet radiation, pulsed electric field, and high-pressure processing, have been reported to inactivate *Bacillus* spores (Soni and others 2016), and these technologies may also be employed to treat fermented products. Therefore, from a food safety viewpoint, fermented products may be processed to kill microbes in them and/or toxicological analysis may be undertaken prior to further applications. However, it should also be pointed out that some fermented products, such as yogurt, known as safe and requiring a live culture (such as LAB) to provide the full health benefits, on the contrary, require maximization of the viability of their live cultures.

Conclusions

Fermented edible seeds and their products generally contain increased bioactive components, such as some vitamins, GABA, natural phenolics, and bioactive peptides, compared to unfermented materials. More importantly, they exhibit various bioactivities, such as antioxidant, anti-hypertensive, and anti-cancer effects, suggesting that they may have potential and comprehensive health benefits. Therefore, it is highlighted that fermented edible seeds and their products are excellent natural sources of bioactive components and can be recommended for consumption as a part of dietary components and developed into functional foods to prevent chronic diseases.

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Author Contributions

R.Y.G and H.C initially constructed and prepared the manuscript draft, and H.B.L, A.G, Z.Q.S, and H.C edited and revised the final manuscript.

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