



Nutritional Significance and Antioxidant-Mediated Antiaging Effects of Finger Millet: Molecular Insights and Prospects

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Kumar A, Rani M, Mani S, Shah P, Singh DB, Kudapa H and Varshney RK (2021) Nutritional Significance and Antioxidant-Mediated Antiaging Effects of Finger Millet: Molecular Insights and Prospects. Front. Sustain. Food Syst. 5:684318. doi: 10.3389/fsufs.2021.684318 Aging is a multifaceted process that is associated with progressive, lethal, and unalterable changes like damage to different molecules (DNA, proteins, and lipids), cells, tissues, and organs. It is an inevitable process but can be delayed by both genetic and dietary interventions. Besides aging, premature death and age-associated diseases can be dealt with diet regulation and the use of compounds that inhibit the stress responsiveness or promote the damage repair signaling pathways. Natural compounds offer a repertoire of highly diverse structural scaffolds that can offer hopeful candidate chemical entities with antiaging potential. One such source of natural compounds is millets, which are minor cereals with an abundance of high fiber, methionine, calcium, iron, polyphenols, and secondary metabolites, responsible for numerous potential health benefits. The present review article elucidates the nature and significance of different phytochemicals derived from millets with a major focus on finger millet and highlights all the important studies supporting their health benefits with special emphasis on the antiaging effect of these compounds. The present article also proposes the possible mechanisms through which millets can play a significant role in the suppression of aging processes and aging-related diseases by influencing genetic repair, protein glycation, and stress-responsive pathways. We further discuss well-established natural compounds for their use as antiaging drugs and recommend raising awareness for designing novel formulations/combinations from them so that their maximum antiaging potential can be harnessed for the benefit of mankind.

Keywords: aging, antioxidant, bioactive compounds, millets, natural products, phytochemicals

INTRODUCTION

In the past few decades, there is a tremendous growth in research work connoting that certain major component in diet influence aging and associated age-related diseases. These dietary components include a myriad of constituents like fiber for the glycemic index in diabetes, vegetables, and fruits for cardiovascular disease, certain fats (saturated, polyunsaturated, and trans fatty acids) for

cardiovascular disease, and vitamin D and calcium for osteoporosis and bone fracture (Everitt et al., 2006) and many more. With increasing public awareness for the wellness of health considering nutrition as a major role-playing factor, phytochemicals such as polyphenols and dietary fiber have gained popularity for their innumerable health beneficial properties. Several crops like millets are known for their numerous health benefits attributed to their dietary fiber and polyphenol content (Taylor, 2017).

Aging is often a major risk factor associated with noncommunicable chronic diseases (NCDs) like cardiovascular disease, type II diabetes, certain cancers, obstructive pulmonary disease, osteoporosis, dementia, etc. (Shlisky et al., 2017). It is a complex multifaceted process, and many theories have been proposed to explain it. These theories include the free radical theory of aging which was later modified to the mitochondrial theory of aging by Harman (1972), the telomere shortening theory as proposed by Lu and Liu (2010), and the protein translational modification theory as proposed by Santos and Lindner (2017). Dietary components such as antioxidants and protein-modifying compounds influence different aspects as proposed in these theories and help delay and in cases overcome aging. Ribarič (2012) have described the epigenetic modifications by dietary factors which include direct effect on gene expression by influencing DNA methylation, histone modification, activation of nuclear receptors by ligands, and through modification of membrane receptor signaling cascades. Furthermore, the influence of caloric restriction on aging was reviewed in the above article with increasing clarity on molecular pathways like the mammalian target of rapamycin (mTOR), sirtuin pathway, and insulin/insulin-like growth factor signaling (IIS) cascades which are all involved in aging (Altintas et al., 2016; Chen et al., 2020; Yu et al., 2021). How these pathways are influenced by dietary components have helped us in establishing the major impact of dietary components on the aging process.

Millets are considered to have immense significance because of their nutritional and nutraceutical potential. Millets are an abundant source of minerals, dietary fibers, and phenolic compounds and offer health benefits such as antimicrobial, antidiabetic, anticancerous, antiartherosclerogenic effects, antioxidant, and antiaging properties (Yang et al., 2012; Si and Liu, 2014; Kumar et al., 2016). Among different types of millets, finger millet is known to possess almost the highest nutritional value (Taylor, 2017). In the recent past, various studies have also reported the preliminary antiaging effect of millet grains, mainly the finger millet (Hegde et al., 2002; Dykes and Rooney, 2006; Shobana et al., 2010; Zhang et al., 2011; Kakkar and Bais, 2014; Khan et al., 2015; Pei et al., 2016).

The present review covers the nutritional and phytochemical composition of finger millet and how they influence aging and age-related disorders. We further discuss wellestablished natural compounds for their use as antiaging drugs and recommend raising awareness for designing novel formulations/combinations from them so that their maximum antiaging potential can be harnessed for the benefit of mankind.

MILLET: A WONDER GRAIN

Millet includes the grains from the diverse group of forage grasses of the family Poaceae. They are considered "coarse cereals" similar to sorghum mostly due to their grain texture, which makes their processing as well as cooking inconvenient when compared with rice and wheat (Rao et al., 2017; Hassan et al., 2021). They are mainly grown in Asia and Africa and are in cultivation in East Asia for the last 10,000 years (Bhat et al., 2018). Millets produce small seed grains and are often cultivated as cereals. Millets are characterized by their remarkable abilities to survive in stress conditions, fairly resistant to pests and diseases, and are widely adaptable. Millet grains can be stored for a long duration during natural disasters such as famine (Obilana and Manyasa, 2002; Adekunle, 2012; Yang et al., 2012; Kumar et al., 2018).

India, several countries of Africa, and China produce the bulk of the world's millet crops. Several important millets grown in the Asian and African countries include barnyard millet (Echinochloa spp.), pearl millet (Pennisetum glaucum), foxtail millet (Setaria italica), finger millet (Eleusine coracana), little millet (Panicum sumatrense), kodo millet (Paspalum scrobiculatum), and proso millet (Panicum miliaceum) (Rao et al., 2017). Whereas, Tef (Eragrostic tef) and Fonio (Digitaria exilis) are grown in Ethiopia, Nigeria, Niger, Togo, Senegal, and Mali (Bhat et al., 2018). Most millets are grown in different regions of the world from east to west, which is evident from their colloquial names, foxtail millet as Italian millet, proso as French millet, barnyard as Japanese barnyard millet, indicating their growing region. Proso millet is cultivated in the USA as well as in Russia and is a popular bird feed in both countries, whereas foxtail millet is largely cultivated in China and India. Millets are predominantly used for food purposes in India as well as in other developing countries but are mainly used as feed ingredients in developed countries.

Pearl millet is the most commonly grown species with large grains and accounts for 40% of the world's total millet production (ICRISAT, 2007; Mariac et al., 2007). Except for sorghum and pearl millet, all other millets are known as small or minor millets (Obilana and Manyasa, 2002; ICRISAT, 2007; FAO, 2012) (**Figure 1**). Finger millet is mainly grown in Eastern and Southern African countries (Uganda, Kenya, the Democratic Republic of the Congo, Zimbabwe, Zambia, Sudan, Tanzania, Nigeria, and Mozambique) and Southern Asia (mainly India and Nepal).

Nutritional Composition

The nutritional content of food is an important aspect for the overall health and well-being of humans especially for better health and development to maximize the human genetic potential. Millets as food are a rich source of micronutrients and phytochemicals (Hassan et al., 2021). In comparison with main cereals like wheat, rice, sorghum, and maize, millets possess a high nutritional value (Parameswaran and Sadasivam, 1994). Being an abundant source of nutrients especially minerals, vitamins, dietary fiber, and phytochemicals, millets provide numerous health benefits. They are nutritionally important as they possess high calcium content (0.38%), roughage or dietary fiber (18%), and phenolic compounds (0.3–3%) (Rao et al., 2017).



Apart from these, proteins found in millet are rich in essential amino acids such as tryptophan, threonine, and sulfur-containing amino acids excluding lysine and threonine. They have a high-fat content in comparison with other cereals especially unsaturated fatty acids (Nithiyanantham et al., 2019). The abundance of phytochemicals and micronutrients of immense therapeutic potential signifies the importance of all varieties of millets (Mal et al., 2010; Singh et al., 2012). The average of the nutrient composition of different millets is shown in **Table 1**.

Millets possess proteins ranging from 7 to 12%, carbohydrates in the range of 65-75%, fat varying from 2 to 5%, and dietary fiber in the range of 15-20%. Of the different millet varieties, pearl millet is the richest in both protein (12-16%) and lipids (4-6%). Millet proteins contain greater amounts of healthy rich essential amino acids (Bhat et al., 2018; Shah et al., 2021). Finger millet, for example, contains a distinctive amount of sulfur-containing amino acids and is a better source of calcium and several micronutrients and pyridoxine. Furthermore, millets are rich in vitamin B complex. Due to the ample quantity of these key nutrients, millets or millet-based products are being used as food products (Subramanian and Viswanathan, 2007; Liu et al., 2012). On the other hand, millet proteins are poor in lysine similar to other cereals, and hence need to be complemented with lysine-enriched vegetables and animal proteins. The major portion of dietary fiber is made up of non-starch polysaccharides (95%), and they are present in seed bran as well as endosperm. Traditionally prepared food items from millets have been a major part of the food consumed in Central America, Africa, and the Indian subcontinent. Most traditional products are flour based, and these include pancakes made from either fermented or unfermented dough, porridge or mudde, snacks, deep-fried products, sweet or sour local liquor, some non-alcoholic beverages, and decorticated grains, prepared similar to rice by boiling (Taylor, 2017; Bhat et al., 2018). Based on their nutritional components and henceforth nutraceutical aspect, different types of millets are consumed in different seasons of the year, conventionally. Almost all of these millets are a good source of both soluble and insoluble dietary fibers, which is contributed by both seed coat as well as endosperm cell walls, unlike in rice, wherein mostly the bran layers account for its dietary fiber content. Since the dietary fiber component of the food offers several physiological benefits including the health of the gastrointestinal tract, millets could be gainfully utilized in the preparation of functional and healthy foods for the target population. Along with nutrition, bioactive compounds found in millets are also recognized for their health beneficial effects, such as antioxidant, antidiabetic, antiaging properties, etc. (Sripriya et al., 1997; Kumari and Sumathi, 2002; Shobana et al., 2007; Banerjee et al., 2012; Shahidi and Chandrasekara, 2013; Kumar et al., 2016).

Natural Compounds

Phytochemicals obtained from millets are mostly secondary metabolites with molecular weights ranging from 150 to 30,000 Da in size. Phenolic acids, flavonoids, and tannins are the major polyphenols reported in different parts of the millet grain (Rao and Muralikrishna, 2002; Sreeramulu et al., 2009). Furthermore, it also consists of terpenes, betalians, organosulfides, indoles/glucosinolates/sulfur compounds, protein inhibitors, and other organic acids, along with dietary fiber and phytates. Most millets possess colored coats and some possess colored endosperm. Carotene constitutes the major coloring pigment and hence several millet varieties are known **TABLE 1** | Proximate nutrient composition and nutritive value of various millets (g/100 g db and mg/100 g db).

Variety	Carbohydrate (g)	Protein (g)	Fat (g)	Ash (g)	Fiber (g)	Ca (mg)	Fe (mg)	Zn (mg)	Thiamin (mg)	Riboflavin (mg)	Niacin (mg)	Energy (kcal)
Sorghum	71	10.4	3.1	1.8	2.0	25	5.4	3.1	0.38	0.15	4.3	329
Finger millet	59–75	6.9–10.9	1.5	2.6	15.2	350	3.9	3.13	0.42	0.19	1.1	336
Kodo millet	72–76	6.2–13.1	3.2-4.9	3.3	5.2	35	1.7	1.9–2.4	0.15	0.09	2.0	353
Foxtail millet	55-69	11.2	4.0	3.3	9.4	31	2.8	2.92	0.59	0.11	3.2	351
Fonio millet	68–75	8.4	3.3	3.4	18.2	20	2.1	1.5	0.17	0.22	1.15	379
Little millet	76	15	4.5	5.4	2.5	17	9.3	5.25	0.30	0.09	3.2	329
Barnyard millet	74	11.0	5.2	4.5	13.6	22	18.6	3	0.33	0.10	4.2	300
Pearl millet	67–72	11.8	5.1	2.2	13.8	42	11.0	3.29	0.38	0.21	2.8	363
Proso millet	64–76	12.6	2.9–11.6	2.7	13.1	15	2.2	2.36	0.41	0.28	4.54	316

All values represent mean/range of reliable published data.

Source: Adapted from Taylor (2017), Serna-Saldivar and Espinosa-Ramírez (2018), Kumar et al. (2018), Selvi et al. (2015), Shankaramurthy and Somannavar (2019), and Renganathan et al. (2020).

to have beta-carotene. Rich polyphenol content is responsible for the antioxidant potential of millets. The gelatinization temperature of the millet starch is slightly higher compared with rice and wheat. It also differs from other cereal starches concerning the crystallinity and the amylose and amylopectin organization. Hence, millet-based diet/foods may digest slowly compared with other cereal foods.

Polyphenols

Polyphenols are naturally occurring secondary metabolites which are generally characterized by the presence of multiple phenol rings (Pandey and Rizvi, 2009; Manuja et al., 2013). Several polyphenols have been detected from different plant species, and a majority of them are derivatives of phenylalanine or another precursor shikimic acid. Polyphenols are categorized on the basis of the number of phenol rings and how these rings are bound to each other. The major classes consist of phenolic acids (derivatives of benzoic acid and cinnamic acid), flavonoids, lignans, and stilbenes (Supplementary Figures S1, S2). Polyphenols are abundant in millets and are one of the most marketed dietary supplements (Ferguson, 2001). The finger millet seed coat is abundant in polyphenols when compared with many other kinds of cereal such as wheat, rice, barley, and maize (Viswanath et al., 2009).

Phenolic Compounds

Grain phenolics can exist both as free, soluble forms as well as insoluble bound forms. The majority of the phenolics found in millet exist as glycosides, whereas ferulic acid is the major bound phenolic acid (18.60 mg/100 g) and protocatechuic acid is the major free phenolic acid (45.0 mg/100 g) (Rao and Muralikrishna, 2002). The seed coat of finger millet is known to be rich in many potent antioxidant phenolics mostly benzoic acid derivatives (85%) such as gallic acid, protocatechuic acid, *p*-hydroxybenzoic acid, vanillic acid, and syringic acid (Hegde et al., 2002; Rao and Muralikrishna, 2003; Chandrasekara and Shahidi, 2010), whereas the rest consists of either flavonoid like quercetin or cinnamic acid derivatives such as *trans*-cinnamic acid, *p*-coumaric acid, caffeic acid, and sinapic acid followed by condensed tannins

(Chethan and Malleshi, 2007; Shah et al., 2021). Apart from these bound phenolic acids such as ferulic and *p*-coumaric acid are also found in abundance in finger millet (**Table 2**).

Apart from these phenolics, direct infusion electrospray ionization mass spectrometry (ESI MS) of the extract from seed coat elucidated the presence of luteolin, naringenin, kaempferol, phloroglucinol, apigenin, catechins, malic acid, epigallocatechins, diadzein, and catechingallates (Shobana et al., 2009). Among various millets studied, finger millet had the highest content of \leq 300 µmol catechin equivalents/g of the defatted meal followed by foxtail, little, pearl, and proso millets. Furthermore, finger millet is the richest in condensed tannins (Dykes and Rooney, 2006). A recent report showed that finger millet varieties with darker color from Northern Malawi had higher polyphenols and antioxidant properties in comparison with the lighter ones. It means that the dark-colored finger millet varieties have greater potential for use as functional food ingredients as they possess higher amounts of natural antioxidants than lightcolored varieties (Xiang et al., 2019).

Micronutrients

As the name suggests, micronutrients are required in very few quantities for human health. However, micronutrient deficiency and their imbalance cause various age-related diseases in humans like obesity, diabetes, osteoporosis, etc. The minerals and vitamins are called micronutrients. Worldwide, more than 2 billion people are suffering from one or other age-related chronic micronutrient deficiencies (MNDs) whereas more than 800 million people are undernourished (Kumssa et al., 2015; Beal et al., 2017). In millets, the mineral content range is from 1.7 to 4.3 g/100 g which is a comparatively higher side than that of main cereals such as wheat (1.5%) and rice (0.6%). Minerals play a chief role in structure and rigidity, muscular contractions, formation of strong bones and teeth, oocyte activation, blood clotting, oxygen transport, regulation of heartbeat, maintenance of fluid balance, nerve signal transmission, etc. (Kulkarni et al., 2018; Kumar et al., 2018). Calcium and iron deficiencies are mainly observed in developing countries. A large number of people are suffering from osteoporosis in India. The calcium TABLE 2 | Active ingredients of finger millet and their mechanism of action for enhancing the lifespan.

Active ingredients/bioactive peptides	Experimental model	Efficacy	Mechanism	References
Dietary fiber, phytate, and total phenolics of seed coat matter of finger millet	Streptozotocin- induced diabetes model rat	Hypoglycaemic, hypocholesterolaemic, nephoprotective, and delayed cataractogenesis	Reduced urinary glucose excretion, regulation of postprandial glycemia, reduced the lipid levels and atherogenic index, reduce levels of blood and urinary creatinine and attenuated nephromegaly, anticataractogenic property, lower the levels of AGE, HbA1c, AR activity, and lesser lenticular opacity	Shobana et al., 2010
Diadzene, gallic, coumaric, synringic, and vanillic acids of finger millet	_	Antioxidant activity	Inhibition of oxidation of linoleic acid and reduction of ferrous to ferric iron	Viswanath et al. (2009)
Seed coat matter of black finger millet	Streptozotocin- induced diabetes model rat	Antioxidative and antidiabetic activity	Reduced the serum levels of alanine transaminase (ALT), aspartate transaminase (AST), and alkaline phosphatase (ALP) and elevation in catalase (CAT) and superoxide dismutase (SOD) and lowered the thiobarbituric acid reactive substances (TBARS)	Okoyomoh et al., 2013
Formulated finger millet	Alloxan-induced diabetes mellitus model male Wistar albino rats	Hypoglycaemic and hypolipidemic properties	Lower the glucose level, cholesterol, triglycerides, and low-density lipoproteins (LDL)	Nadro and Elkanah, 2017
Ferulic and cinnamic acids of kodo millet	-	Antioxidant activity	Free radical scavenging activity	Sharma S. et al. (2017)
Whole grain and bran of finger millet	High-fat-diet-fed LACA model mice	Antioxidant and anti-inflammatory properties	Prevention of obesity by regulation of obesity-related genes, improved lipid profile and anti-inflammatory status, suppresses oxidative stress	Murtaza et al., 2014
Polyphenol of finger millet	_	Antioxidant activity	Inhibits superoxide, hydroxyl, and nitric oxide radicals	Bindu and Malleshi, 2003
Ethanolic extract from finger Italian millet	_	Antidiabetic and antioxidant activity	Inhibits glycation	Ofosu et al., 2020
Carotenoids and vitamin E of finger millet	-	Antioxidant activity	Free radical scavenging activity	Chethan et al., 2008
Methanolic extract of finger millet	-	Protective role against diabetes mellitus	Inhibits glycosylation of collagen	Hegde et al., 2002
Whole grain extracts of finger millet (Oshadha and Rawana), proso millet, white finger millet, kodo millet, foxtail millet	-	Antidiabetic, antialpha amylase and antiglucosidase activity	Inhibition of early glycation, middle glycation and reversal of antiglycated products	Senevirathne et al., 2021
Free and bound phenolics from finger millet varieties	_	Antioxidant activity	Free radical scavenging activity	Xiang et al., 2019
Proanthocyanidins of finger millet	-	Antioxidant activity	Free radical scavenging activity	Dykes and Rooney, 2006
Benzoic acids				
Protocatechuic acid (PCA) of finger millet seed coat	-	Antibacterial, antiviral, neurological effect, antiatherosclerotic, antifibrotic, antiaging, antiulcer, anticancer	Antiaging ability is induced as it increases the activity of glutathione peroxidase and catalase and decreases the malondialdehyde level.	Zhang et al., 2011; Kakkar and Bais, 2014; Khan et al., 2015
Gallic acid of finger millet seed coat	_	Antioxidant, anticarcinogenic, antimicrobial, anti-inflammatory, antimelanogenic, neuroprotective, hepatoprotective	Induces antiaging effect due to its ROS quenching and antiglycation activity	Panich et al., 2012; Badhani et al., 2015

(Continued)

TABLE 2 | Continued

Active ingredients/bioactive peptides	Experimental model	Efficacy	Mechanism	References
Syringic acid of finger millet seed coat	_	Antimitogenic, antidiabetic	Eradicates free radicals due to their antioxidant potential	Wei et al., 2012; Muthukumaran et al., 2013
Vanillic acid and its derivatives of finger millet seed coat	-	Antimicrobial, antiaging, hepatoprotective, antivenom	Inhibits collagenase and elastase activity	Dhananjaya et al., 2009; Itoh et al., 2009; Widowati et al., 2016
Cinnamic acid derivatives				
p-Coumaric acid of finger millet seed coat	-	Antioxidant,anti- inflammatory, antimutagenic, antiulcer, antiplatelet, and anticancer	Due to its potent antioxidant potential	Pragasam et al., 2013; Pei et al., 2016
Ferulic acid of finger millet seed coat	-	Antioxidant, antimicrobial, anti-inflammatory, antithrombosis, and anticancer activities	Inhibits the expression of some cytotoxic enzymes, such as nitric oxide synthase, caspase, and cyclooxygenase-2	Rao and Muralikrishna, 2002; Ou and Kwok, 2004; Pei et al., 2015
Caffeic acid of finger millet seed coat	-	Antioxidant, anti-ischemia reperfusion, antithrombosis, antihypertension, antifibrosis, antivirus, and antitumor	Potent antioxidant potential	Jiang et al., 2005
Sinapic acid of finger millet seed coat	-	Antioxidant, antimicrobial, anti-inflammatory, anticancer and antianxiety activities	Elastase, tyrosinase and collagenase inhibitory activities	Chen, 2016; Taofiq et al., 2017
Flavonoids				
Quercetin of finger millet seed coat	_	Antioxidant, antiaging, antiobesity, anticarcinogenic, antiviral, antibacterial, and anti-inflammatory effects	Proteasome activator with antioxidant properties that consequently influence cellular lifespan	Chethan et al., 2008; Chondrogianni et al., 2010; Wang et al., 2016
Catechin and its derivatives of finger millet seed coat (gallocatechin, epicatechin, and epigallocatechin)	-	Antioxidant, anti-inflammatory, antimutagenic effects, antiaging, as well as cardiovascular disease-preventive properties.	Eradicates free radicals, stimulates the activity of the regulatory protein SIRT1	Sripriya et al., 1996; Unno et al., 2011; Hong et al., 2013

content of finger millet (ca. 340 mg/100 g) is almost eight times higher than major cereals wheat and rice. Therefore, finger millet is the richest source of calcium to overcome age-related diseases like osteoporosis (Puranik et al., 2017; Sharma D. et al., 2017). The consumption of pearl millet and barnyard millet help in protection from anemia because they have been found as the richest source of iron. The highest content of zinc (4.1 mg/100 g) in foxtail millet as compared with other millets and stable cereals make it special to combat zinc-associated diseases. Foxtail millet is also a good source of iron (2.7 mg/100 g). The immune system is boosted by the intake of zinc and iron (Kulkarni et al., 2018; Kumar et al., 2018). Millets are also the ultimate source of β -carotene and B vitamins like riboflavin, niacin, and folic acid as compared with wheat and rice. The foxtail millet has the richest source of thiamine (0.60 mg/100 g) whereas the barnyard millet has the highest content of riboflavin (4.20 mg/100 g) (**Table 1**). The regular consumption of millets in our diets aids to curb nutritional deficiencies and age-related diseases (Chandel et al., 2017). The presence of antinutrients in millets like phytates, polyphenols, and tannins is responsible for the reduction in mineral bioavailability by chelating multivalent cations such as Ca, Zn, Fe, Mg, and K. Besides this, the rich content of protease and amylase inhibitors in millets also causes reduced digestibility of millet grains. Therefore, food processing techniques are used to enhance nutritional content, increase digestibility, and bioavailability of nutrients with a reduction in antinutrients and increase the accumulation of nutrients in milled grains (Sharma D. et al., 2017; Vinoth and Ravindhran, 2017). The applications of genetic engineering and genome-editing tools are also useful in facilitating nutrient



accumulation in grains and the prevention of the synthesis of antinutrients.

HEALTH BENEFITS OF MILLET CONSUMPTION

Millets are better alternatives to common cereal grains due to their nutraceutical properties and immense health-promoting attributes. Millets contain ample amounts of fiber and lesser amounts of simple sugars responsible for their relatively low glycemic index. Millets have been shown to aid low blood sugar levels in comparison with wheat or rice (Kumari and Sumathi, 2002) and thus possess antidiabetic properties. Millets similar to whole grains are abundant in magnesium, which acts as a cofactor for enzymes especially involved in the glucose uptake and secretion of insulin. Magnesium has been reported to reduce the effects of migraines and heart attacks. Niacin in millets is hypocholesteremic in nature. Consumption of millets also reduces the C-reactive protein and unhealthy lipids and thus may prevent cardiovascular diseases. Millets provide plenty of phosphorus which plays an important role in cell structure determination. Aside from constituting the bone mineral matrix, phosphorus is an essential component of adenosine triphosphate (ATP), the energy currency of the cell. Apart from the abovementioned health benefits, the compounds of millets are also known to possess antioxidant and antiaging properties. Previous reports have well-established the antioxidant potential of many millet varieties (Chandrasekara and Shahidi, 2010). Regulation of oxidative stress, through antioxidant administrations, is also capable of controlling different healthrelated issues such as diabetes, cancer, neurodegenerative disorders, cardiovascular problems, and age-associated diseases. Thus, in the benefit of the current review, the different properties of millets are explained in relation to aging.

BIOACTIVE COMPOUNDS: ANTIAGING AND ANTIOXIDANT PROPERTIES

Antioxidants derived from plants, such as phenolics and flavonoids have numerous biological effects. Antioxidants play a major role in the reduction of lipid peroxidation, a process that plays a key role in cancer and aging (Namikii, 1990). Antioxidants provide stable radical intermediates which prevent the oxidationinduced damage of fatty acids and oils (Lobo et al., 2010). Polyphenols like phenolic acids, flavonoids, and tannins found in the seed coat of millet seeds act as reducing agents such as free radical quenchers, metal chelators, and singlet oxygen quenchers (Banerjee et al., 2012). Potent antioxidant attributes of polyphenols arise as a result of their ability to act as hydrogen



atom donors *via* hydroxyl groups on phenol rings to electron-deficient free radicals.

Polyphenols overcome oxidative stress by the maintenance of balance between oxidants and antioxidants. Being antioxidant, these compounds may show different health benefits and primarily may act as antiaging compounds as well. Polyphenols presently have gained the status of "life span essentials" as they play a crucial role in maintaining health (Chandrasekara and Shahidi, 2010). Millet grain is entirely edible, and conventional food recipes are prepared from wholemeals throughout Asian and African countries. This indicates that the phytochemicals of millets such as polyphenols are consumable without any adverse effects on human health. The content and composition of these polyphenols vary depending on the type of millet grain and have received considerable interest because of their antioxidant, antiaging, and other nutraceutical properties (Chandrasekara and Shahidi, 2011). Additionally, some of the other health benefits such as hypoglycemic (Kumari and Sumathi, 2002), hypocholesterolemic (Hegde et al., 2002) and antiulcerative (Tovey, 1994) properties can further be contributed to a large extent to their polyphenol content.

Different types of millets are known to be rich in different polyphenols which may act as potent antioxidants which are critical for exhibiting antiaging properties. For example; different types of millet grains contain dimers and trimers of ferulates which possess comparatively higher antioxidant activity (Chandrasekara and Shahidi, 2011). Very recent findings on phenolics of millet grain varieties that are cultivated in South Korea like millet (M), Italian millet (IM), barnyard millet (BM), and finger Italian millet (FIM) have shown their potent antioxidant potential and antiglycation properties effective against diabetes mellitus which is a main age-related disorder. It has also been reported that the finger Italian millet has the highest total phenolics and total flavonoid content as compared with barnyard millet and Italian millet. Therefore, finger Italian millet is a crucial nutraceutical as it possesses high antioxidant and antidiabetic activity (Ofosu et al., 2020). Colored (brown, reddish, or red) finger millet varieties from the Northern Malawi region exhibit more significant antioxidant activity than



the white variety due to the presence of high amounts of phenolic compounds, tannins, and flavonoids in them (Xiang et al., 2019). Amadou et al. (2011) have studied the antioxidant efficiency of defatted foxtail millet (DFMB) and observed that it can scavenge free radicals and superoxide anions.

Watanabe (1999) isolated two flavones viz., luteolin and tricin, with antioxidant activity from Japanese barnyard millet. Luteolin along with its glycosides possesses health-beneficial properties such as antioxidant, anti-inflammatory, cancer preventive, and antiarrhythmic activities (Lin et al., 2008). Tricin is also known for its antitumor and antimetastatic properties (Yue et al., 2020). Finger millet showed the highest total flavonoid content in defatted meals followed by kodo and foxtail millets. Studies were carried out on the natural antioxidants in edible flours of small millets. Ferulic acid exhibits very strong antioxidant, free radical scavenging, and anti-inflammatory activity (Shahidi et al., 1992; Castelluccio et al., 1995).

Free radical production in excess and lipid peroxidation are the major reasons for chronic diseases such as diabetes, cardiovascular disease, cancer, cataract, and aging. Nonenzymatic glycosylation whereby chemical reaction occurs in between the aldehyde group of reducing sugar and the amino group of proteins plays a major role in the complications of diabetes and aging (Rowan et al., 2018). Non-enzymatic glycosylation and cross-linking of collagen are induced by free radicals, whereas free radical scavengers such as polyphenols inhibit these reactions (Fu et al., 1992). Polyphenols act as antioxidant compounds that suppress excessive oxidation responsible for cancer and aging (Namikii, 1990). Thus, any compound which can neutralize these free radicals can in



FIGURE 5 | Model of the mammalian target of rapamycin (mTOR) pathway: the mTOR pathway has two major components called TOR complex 1 (TORC1) and TOR complex 2 (mTORC2) which regulate different cellular functions. mTORC1 is sensitive to rapamycin and play role in protein synthesis, ribosome biogenesis, cell survival, proliferation, angiogenesis, invasion, migration, and metastasis whereas mTORC2 is insensitive to rapamycin and regulates cell survival, cell cycle, and actin remodeling. In response to growth nutrients, the mTOR pathway is regulated by the PI3K cascade. mTOR is negatively regulated by tuberous sclerosis proteins called TSC1 and TSC2 which exist as heterodimers. In response to insulin signaling, Akt phosphorylates TSC2 which makes it functionally inactive. Shortage of nutrients mainly amino acid starvation leads to dephosphorylation of well-known substrates of mTORC1 such as ribosomal S6 kinase (S6K) and eukaryotic translation initiation factor 4E-binding protein (4E-BP) whereas in the supply of nutrients, restoration of amino acids enables phosphorylation of S6K and 4E-BP which accelerates protein synthesis. Amino acids activate mTORC1 in two ways either via inhibition of TSC1-TSC2 or by stimulation of Rheb. AMPK is activated under low energy (high AMP/ATP ratio) and promotes fatty acid oxidation and inhibits protein synthesis. The tumor suppressor LKB1 is responsible for the activation of AMPK which leads to inhibition of mTORC1-mediated phosphorylation of S6K1 and 4E-BP1. AMPK also directly activates TSC2 via its phosphorylation and increases its activity which ultimately leads to repression of mTORC1 signaling. High-energy situation mediates upregulation of mTORC1 activity. Upon stress conditions, p53 is activated and inhibits the mTOR activity by the AMPK-TSC2 mechanism. In contrast to mTORC1, mTORC2 is insensitive to nutrients but responds to growth factors such as insulin which requires PI3K. For the activation of mTORC2, ribosomes are very crucial. mTORC2 binds to ribosomes in an aPI3K-dependent manner and regulates protein kinase B (Akt), glucocorticoid-induced protein kinase 1 (SGK1), and protein kinase C-a (PKC-a). SGK-1 is also activated by mTORC2 and controls the growth and ion transport. mTORC2 regulates the actin remodeling by the activation of PKC-a. mTORC1 regulates protein synthesis, ribosome biogenesis, cell survival, proliferation, angiogenesis, invasion, migration, and metastasis by regulation of S6K1 and 4E-BP1.

turn exhibit antiaging effect. Senevirathne et al. (2021) have very recently observed antidiabetic properties from different varieties of millets such as finger millet, proso millet, white finger millet, kodo millet, and foxtail millet cultivated in Sri Lanka. The inhibition of early glycation, middle glycation, and reversal of antiglycated products by these millets in diabetes make them crucial functional food for delaying the aging process. These millet varieties also showed anti- α -amylase activity, and suppression of this enzyme is required for better carbohydrate digestion and maintains the blood glucose levels for diabetic patients (Table 2).

Hegde et al. (2005) studied the effects of the antioxidant properties of millets on oxidative stress and glycemic status in alloxan-induced rats. The alloxan-induced rats when fed with a diet enriched with finger millet showed a significant decrease in blood glucose and cholesterol level. Tail tendon collagen glycation was reduced in the finger millet-fed rats. Diabetic groups fed on finger millet diet showed alleviated



levels of enzymatic and non-enzymatic antioxidants along with lipid peroxides. This was associated with the abundant amounts of phenolics, tannins, and phytates present in finger millets. Furthermore, the effects of methanolic extracts of finger millet and kodo millet on glycation and cross-linking of collagen were studied (Hegde et al., 2002). It is thus proposed that finger millet can have a potential therapeutic role as dietary supplements for the prevention of glycation-induced complications, as in diabetes or aging.

Free radicals are regularly generated in our body as a byproduct of normal aerobic respiration and substrate oxidation. However, the excessive accumulation of these radicals causes damage to vital biomolecules such as carbohydrates, lipids, proteins, and DNA, consequently leading to the manifestation of various age-related diseases. Polyphenols are suggested to exhibit their beneficial antioxidant effects either by scavenging free radicals, chelating metal ions, quenching singlet oxygen, or by inhibiting the reactive oxygen species (ROS) producing enzymes such as myeloperoxidase, lipoxygenase, cyclooxygenase, and xanthine oxidase. Chandrasekara and Shahidi (2011) evaluated the antioxidant properties of phenolic extracts from seven millet varieties [kodo, finger (Ravi), finger (local), proso, foxtail, little, and pearl] on HT-29 cells. All varieties used in this study showed a significant reduction in lipid peroxidation in liposomes, singlet oxygen quenching, and DNA scission inhibition to varying degrees. Furthermore, the cytoprotective effects of various types of millet phenolics (catechin, ferulic acid, vanillic acid, and resveratrol) were investigated, and it was observed that they exhibit preventive effects on protein and human erythrocyte peroxidation (Palaniswamy and Govindaswamy, 2017).

Different studies have also given exclusive attention to finger millet for its pharmacological significance. For instance, the study



conducted on early diabetic rats indicated that the administration of finger millet exhibited its beneficial effects on antioxidant status of the skin, production of nerve growth factor, and woundhealing parameters (Rajasekaran et al., 2004). Ajiboye (2017) also evaluated the antioxidant potential and free radical scavenging ability of whole-grain finger millet. The study revealed that ethanolic extract of whole finger millet effectively scavenged H_2O_2 , NO, and DPPH.

Oxidative stress, caused by ROS, is responsible for modulating several pathological conditions during aging. Thus, aging can also be defined as low energy and high oxidative stress state. As studies indicate that the administration of millet may greatly affect the level of reactive oxygen species and hence may also combat age-associated health issues. Though there is a lack of evidence showing the molecular mechanism for antiaging effect of millets, the possible molecular mechanism for antiaging may be explained using different aging theories and preliminary shreds of evidence supporting antiaging effect of millets.

ANTIAGING PROPERTIES AND MOLECULAR INSIGHTS

Natural compounds offer a repertoire of highly diverse structural scaffolds that can offer hopeful candidate chemical entities with antiaging potential. As per a large number of published pieces of literature, it is highly emphasized that consumption of millet may help in reducing aging and age-related complications. Though, its exact mechanism of action is not known. However, based upon these studies, it can be suggested that phytochemicals obtained

from millets can play a significant role in the suppression of aging processes and aging-related diseases by influencing genetic repair, protein glycation, and stress-responsive pathways. The effect of millets on all these pathways can be attributed to their antioxidant property and antiaging potential, as explained below by various aging theories.

Free Radical Theory

The free radical theory of aging and the antiaging property stated that progressive accumulation of endogenous oxygen radicals causes damage to macromolecular components of the cellrendering cells and organs to stop functioning, ultimately leading to death by senescence (Finkel and Holbrook, 2000). With the discovery of the mitochondrial genome, the theory was modified to the mitochondrial theory of aging in the year 1972. This theory stated that reactive oxygen species (ROS) generated during mitochondrial respiration damages macromolecules including mitochondrial DNA (mtDNA) (Muller et al., 2007). Such mutations in mtDNA which accumulate over time result in increased ROS generation and oxidative damage as suggested by studies conducted in a variety of experimental animal models and tissues (Muller et al., 2007). The grains of millet possess natural antioxidants which help to combat damage induced by free radicals and prolong the lifespan of organisms. This theory has been proven by different experiments as listed in Table 2.

Telomere Shortening Theory

Excessive oxidative stress leads to the accumulation of oxidative DNA damage in telomeres and is strongly related to premature aging (Lu and Liu, 2010). As per telomere shortening theory,

aging occurs due to the loss of telomeres. Cellular senescence is initiated by erosion or improper functioning of telomeres resulting in cell-cycle exit after several cell cycles (Holliday, 2014). In each subsequent cellular division, telomere length gets shorter which obstructs the chromosomal ends from being identified as double-strand breaks. DNA damage response gets triggered due to telomere shortening (Karlseder et al., 1999). Apart from telomere shortening, damage in DNA in regions other than telomere is also responsible for aging. Similarly, DNA damage theory also proposes that loss of its functionality during aging process may stem from the accumulation of unrepaired naturally occurring DNA damages like DNA breaks, cross-linking, and base modifications (Hayflick, 2007). With the accumulation of such mutations over time in aged organisms, there is an increased risk of tumor formation and other age-related disorders (Iyama and Wilson, 2013). As per literature, there are several factors inducing DNA damage which in turn are responsible for aging and associated diseases (Figure 2). Naturally occurring compounds in millets act as potent antioxidants which may reduce the level of oxidative stress and hence minimize the rate of telomere shortening.

Protein Translational Modifications

During aging, the functioning of the proteins can be disturbed due to disorganized protein synthesis and aberrant types of posttranslational modifications (PTMs) such as acetylation, phosphorylation, glycosylation, ubiquitination, sumoylation, and neddylation ultimately leading to the accretion of inactive, denatured, misfolded, and mutated proteins in cells of aged organisms (Santos and Lindner, 2017) (Figure 3). The posttranslational modifications of proteins are chiefly responsible for the activity, stability, as well as protein-protein interactions, influencing the cellular function and metabolism (Santos and Lindner, 2017). It has been observed that different compounds influence the PTMs of different proteins and help in maintaining cellular function during the aging process (Lan et al., 2016; Peleg et al., 2016). In the recent past, a good number of drugs and natural compounds including millets and millet-based products have been studied for their antiaging potential. It was reported that millet extracts inhibited the glycosylation of collagen protein and thus protects its function (Hegde et al., 2002).

NEW PARADIGM FOR DESIGNING ANTIAGING DRUGS DERIVING INFORMATION FROM MILLETS AND OTHER COMPOUNDS OF PLANT ORIGIN

Various natural products from grains, fruits, and vegetables are studied for their antiaging properties. These natural products or compounds are suggested to delay the aging process by regulating genes involved in different pathways and involved in the aging process. Though the detailed mechanism of action of bioactive compounds of millets has not been explored, preliminary investigations highlight the effect of millet extract on longevity as well as aging-associated diseases. Hence, it is imperative to have detailed investigations of bioactive compounds of millets for their mode of action as antiaging compounds. As per various studies, the premature aging and aging-associated diseases are found to be associated with the fault in the key cellular pathways such as the IIS (**Figure 4**), mTOR (**Figure 5**), and sirtuin (**Figure 6**) molecular networks. Hence, all these crucial pathways involved in the aging process need to be studied focusing on the effect of bioactive compounds of millet extracts on the key molecules of all these pathways in suitable *in vitro* model systems (**Figure 7**). Once the molecular mechanism of action of bioactive compounds obtained from millets is revealed, formulations based on these compounds can be processed and targeted to the specific targets in a nanoencapsulated form.

Millets are a rich natural source of antioxidants and thus hold potential as a nutraceutical and functional food ingredient promoting health and reducing the risk of disease. However, more validation in animal models and human subjects needs to be performed to verify their activity and health benefits. The effect of different cellular processes such as methylation, glycosylation, acetylation, phosphorylation, ubiquitination, etc. on aging has been studied by many scientists to find the mechanism by which these processes affect the extent of aging and associated diseases. Several efforts have been made to design some potential drugs that can bind with the target enzyme of cellular processes associated with aging. Some therapeutics such as nutulin-3, HDAC, and MLN4924 have shown their effectiveness in tackling the aging and associated diseases, but no success story of any drug is available in the market that can be recommended for use to prevent aging and its symptoms. The role of some pathways such as the IGF-1 signaling pathway, mTOR signaling pathway, and sirtuin-related pathways is also associated with the aging process. Modulation of these pathways by the use of some potential therapeutics can also be one of the possible approaches to curb the aging process. A comprehensive network of different processes and pathways associated with aging can be designed to find some potential targets for the treatment of aging and associated diseases. Furthermore, validation studies related to these targets can be carried out by in vitro and in vivo studies, if not validated earlier. Structure-based drug designing approach can be applied to design and screen some potential candidate drugs that can modulate the agingassociated pathways by specific binding with the drug target. The druggability of these candidate drugs can be predicted using absorption, distribution, metabolism, and toxicity (ADMT) prediction tools to avoid the chance of drug failure in clinical trials.

Many synthetic and natural compounds such as indolepropionaamide (IPAM), acetyl L-carnitine (ALC), spermidine, metformin, lithocholic acid, caffeine, reserveratrol, rapamycin, epicatechin, and quercetin have shown their potential role in the suppression of the aging process by modulating different drug targets. Mechanisms of action for these compounds are also different for different targets by which they generate the antiaging response. It is important to mention that most of such studies related to aging have been performed in *Caenorhabditis elegans*, *Drosophila*

melanogaster, mice, rat, and worms. The effect of these antiaging compounds needs to be studied in humans to explore the mechanism of action and interaction of compounds with the drug target. Natural ingredients in millets have also shown their antiaging response, but their target and mechanism of action are not explored well scientifically. More studies on the antiaging response of millets are required to validate the target and mechanism of action. Compounds with known antiaging effects can be used as a lead molecule to design more specific and potential candidate drugs for a given drug target. Optimization of lead compounds is very necessary to achieve better bioavailability, specificity, and affinity for the drug target. Computational tools can be more helpful in deciphering the chemical modification required to optimize the activity of a lead molecule. Potential candidate drugs resulted from lead optimization can be recommended for in vitro or clinical testing.

CONCLUSION

A balanced diet with definite nutritional supplements plays a key role in promoting the lifespan and health of human beings. Various natural compounds from different plants have been found to curb aging and associated diseases. Millets are a chief source of proteins, carbohydrates, and minerals and are low in fat content which validates their utilization in many healthy diets. Furthermore, millets contain high content of total phenolics, flavonoids, and other phytochemicals that have been reported to possess antiaging potential. Henceforth, millets might be utilized in the formulation of nutraceutical and functional foods that can be consumed daily due to the presence of many bioactive compounds that prevent aging and associated diseases. The prospects include the identification, isolation, and characterization of bioactive molecules present in millets that can slow down the aging process. Once characterized, further validation of their mechanism of action at the molecular level can be explored by the wet lab and *in silico* analysis. The effectiveness

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of these bioactive compounds requires evaluation in a cell or tissue-specific manner followed by the animal model system for safety concerns and to determine their therapeutic potency.

AUTHOR CONTRIBUTIONS

AK gave the manuscript concept. MR and PS wrote the manuscript. SM, DS, HK, and RKV contributed critically in revising the draft and updating the manuscript for publication. All authors contributed to the article and approved the submitted version.

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SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fsufs. 2021.684318/full#supplementary-material

Supplementary Figure S1 | Structure of major phenolic compounds present in millets.

Supplementary Figure S2 | Basic structure of flavonoids reported in millets.

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