



Full Length Article

Breeding of Selenium Rich Red Glutinous Rice, Protein Extraction and Analysis of the Distribution of Selenium in Grain

Yuanke Liang^{1†}, Muhammad Umer Farooq^{1†}, Rui Zeng^{1†}, Zhicheng Tang¹, Yujie Zhang¹, Tengda Zheng¹, Hla Hla Ei¹, Xiaoying Ye¹, Xiaomei Jia¹ and Jianqing Zhu^{1*†}

¹Rice research institute, Sichuan Agricultural University, Chengdu, Sichuan 611130, China

*For correspondence: zhujianqing1963@163.com

†These authors contributed equally to this work

Abstract

Selenium acts as a chemo preventive agent for specific cancers, which can be useful as protective agent against stress. The objective of this study was to breed healthy selenium rich red glutinous rice serving as diet and search for convenient methods to extract protein from grain and measure the protein and selenium concentration. Furthermore, determine the interaction of selenium and protein in different parts of red glutinous rice seeds. After progress of five years breeding, the red glutinous rice achieved the standard (≥ 40 ng/g) selenium contents of 121.75 ± 3.01 ng/g. Total selenium concentrations in the rice fractions varied in the following order: husk < polished rice < whole grain < bran. Through the analysis of the protein content in different red glutinous rice parts, compared with the selenium concentration, we have made a conclusion that more than 80% selenium in the grain is organic selenium (in selenoprotein form) and more than half of the total selenium is in the endosperm of the red glutinous rice. Overall, the selenium rich red glutinous rice might be used as a natural healthy food, which could be applied as preventive measures against cancer and supplements in medical treatments near future. © 2018 Friends Science Publishers

Keywords: Selenium protein interaction; Selenium accumulation; Anticancer; Life-protecting agent; *Heterosis*

Introduction

Cancer is generally considered as one of the biggest threat to human health. In previous study, the most commonly diagnosed forms of cancers were lung (1.82 million), breast (1.67 million), and colorectal (1.36 million); while the most lethal were lung cancer (1.6 million deaths), liver cancer (745,000 deaths), and stomach cancer (723,000 deaths) all over the world in 2012 (Ferlay *et al.*, 2015). However, the majority of epidemiological studies provided evidence for selenium as a chemo preventive agent for specific cancers: lung (Knekt *et al.*, 1998), breast (Cann *et al.*, 2000), colorectal (Ghadirian *et al.*, 2000), liver (Yu *et al.*, 1997), stomach (Scieszka *et al.*, 1997) and other multiple cancers (Patrick, 2004; Wallace *et al.*, 2009). Cancer is a major public health problem across the world. Besides, previous studies have reported that more than 30% of human cancers could be prevented by an alternative strategy of appropriate dietary modification (Liu, 2004). So, supplement of selenium in food products could be important to the human health.

Selenium (Se), an essential micronutrient for many organisms, is important for preventing disease, improving health, and preventing aging and has been considered a life-protecting agent (Zhao *et al.*, 2017), with a recommended

dietary allowance (RDA) for healthy adults of 55 $\mu\text{g/d}$ (Institute of Medicine (US) Panel on Dietary Antioxidants and Related Compounds, 2000). But people in many areas of the world are selenium deficient, with the consequence that they are unable to express their selenoproteins fully (Xia *et al.*, 2005). For humans, the boundary between inorganic Se deficiency and toxicity is narrow (Augilar *et al.*, 2009). So it has a great potential risk to regard inorganic selenium as selenium supplement directly. Organic Se, selenomethylcysteine (SeMeSeCys) and selenomethionine (SeMet) species are better assimilated by the natural body than inorganic Se and also serves as effective anticarcinogens, leading by SeMeSeCys (Carey *et al.*, 2012). Nevertheless, for most people worldwide, as well as for livestock, plants are the main source of dietary organic selenium; thus, plant selenium metabolism is of great importance for the selenium nutrition of humans and animals (Zhu *et al.*, 2009).

At present, food health is a burning issue in the world. Avoiding selenium deficiency and toxicity, it is important to monitor and optimize crop selenium quality and concentrations, through it can vary greatly between different crops and regions (Zhu *et al.*, 2009). Rice (*Oryza sativa* L) is one of the most principal food crops in the world, serving as the staple food source for more than half of the world's

population (Gealy *et al.*, 2003), which contributes 40% to the total calories intake of people. Increased healthy rice production can play a vital role to address this issue successfully.

Since long ago, Chinese used *Monascus purpureus* to ferment red glutinous rice for cuisine and medicinal food owing to its especial function such as promoting blood circulation (Chairote *et al.*, 2009), reducing cholesterol (Heber *et al.*, 1999) and antioxidant activities (Que *et al.*, 2006). Pigmented rice (*Oryza sativa* L.) has been consumed for a long time in Asia, especially China, Japan, Korea and many countries in Southeast Asia (Kanitha and Wanida, 2010). Besides, pigmented rice genotypes have high nutritive value compared to white rice genotypes (Faiz *et al.*, 2015). Several varieties of pigmented rice, particularly red rice, have been cultivated in many countries. Many studies have demonstrated antioxidant activity and radical scavenging ability of the pigmented rice or its extract in both *in vitro* and *in vivo* models (Nam *et al.*, 2006; Bae *et al.*, 2014). Glutinous rice (*Oryza sativa* L.) is popular rice type due to its superior qualities of fineness, aroma, taste and protein content (Loypimai *et al.*, 2017). It also contains very high quantities of bioactive components which are beneficial for health such as tocopherols, tocotrienols, γ -oryzanol and phenolic compounds especially in the outer pericarp and aleurone layers covering the grains (Loypimai *et al.*, 2009). These scientific reports assist us to do research on selenium accumulation and protein contents deviation in red selenium rich glutinous rice than other traditional rice. Hence, breeding of selenium rich red glutinous rice has very high application value.

The beneficial health-related effects of selenium rich red glutinous rice are of great importance to consumers, breeders and the rice industry. The objective of this study was to determine the interaction of selenium and protein in different parts of red glutinous rice seeds. Using *Heterosis* (Huang *et al.*, 2016), we cross-fertilized different rice species for five years, then we obtained the natural selenium rich red glutinous rice Z5097A (sterile lines) and Z5097B (maintainer line). The selenium contents in them are high enough to meet the daily selenium intake. Moreover, analysis of successive abrasive brown rice milling fraction has shown that nutrients aren't uniformly distributed in brown rice. In addition, we proved that the selenium and protein are in direct relationship in red glutinous rice, the more protein concentration the more will be selenium contents in different parts of rice. More than 80% selenium in the grain is organic selenium (in selenoprotein form) and more than half of the total selenium is in the endosperm of the red glutinous rice.

Materials and Methods

Plant Material

The rice germplasm (2045B red rice, maintainer line, as

female parent) used in this study while local glutinous rice and D62 A (Sterile line) were of Chinese origin. Then conventional hybridization was conducted between local glutinous rice and 2045B. The progenies since F3 hybrids were selfed and then test crossed with D62A. Continuously backcrossed till the material entered into most advanced F7 generation. The red rice sterile and maintainer line materials (numbered 5097A and B) observed thoroughly with identical characters in both parents were preferably selected for backcross, selection and preservation in the Winter of 2013. After years of generations of breeding, being named as Z5097A and Z5097B (Fig. 1).

Preparation of Red Glutinous Rice Powder

The fresh red glutinous rice samples (cleaned with distilled water) were put into the drying oven (DHG-9240B, Shen Xian Co., Ltd, Shanghai, P. R. China) 80°C for 3 h. Then the brown rice machine (JLG-II, Da Ji Co., Ltd, Hangzhou, P. R. China) was used to separate hulls and brown rice (with episperm, embryo, aleurone layer and endosperm). Then, tweezers and dissecting needles were used to fetch the episperm (red surface), embryo of brown rice and residues with white and a little yellow color. Moreover, we used the rice milling machine (JNM-III, CHINA GRAIN RESERVES CORPORATION) for 45 sec to separate the aleurone layer and endosperm, referred as Lamberts *et al.* (2007), Singh *et al.* (2000). All parts of material were then put into pulverizer (F160, Zhong Xing Co., Ltd, Beijing, P. R. China) to make powder, filtered with 100-grade sifter to obtain the tested flour for different studies.

Protein Extraction

The total protein extracted from red glutinous rice according to the method of Daiana *et al.* (2016) with some modifications. Weighted red glutinous rice powder (filtered with 100-grade sifter) 100 g, added into NaOH (0.05 mol/L, 1500 mL), DTT (DL-Dithiothreitol) 2.5 g and sodium hyposulfite 0.5 g. Then put them inside the ultrasonic cleaners (WD-9415B, LiuYi Co., Ltd, Beijing, P.R. China) with frequency of 100 Hz at 40°C for 4 h.

Afterwards, the suspension was centrifuged (5804R, Eppendorf Co., Ltd, Germany) at 3000×g for 20 min. According to Ju *et al.* (2001), the isoelectric points of albumin (pH 4.1), globulin (pH 4.3 and pH 7.9), and glutelin (pH 4.8), proteins were precipitated from the protein extract with HCl at pH 5.0. Herein after centrifugation at 3000×g for 20 min, precipitate referred to as the isoelectric precipitation protein concentrate (IPPC). Washed with deionized water and packed into dialysis bag for dislodging salt and micro molecular impurities. Then put the protein into ultra-low temperature refrigerator (MDF-U3386S, Panasonic Co., Ltd, Japan) at -80°C for 6 h. In the end, after vacuum refrigeration protein flour can be obtained.

Determination of Protein Content

The content of protein determined by Kjeldahl method (Ayalew *et al.*, 2017) and National food safety standard determination of protein in food (GB 5009.5—2010), with a little modification. First, weigh 1.00 g powder samples, put in glass digestive tube (250 mL), add into CuSO₄ (0.2 g), K₂SO₄ (6 g) and H₂SO₄ (10 mL). After gently shaking, put a small funnel on the top of every tube. The mixture was then heated with graphite digestion apparatus (SH220, Hanon Instruments Co., Ltd, Jinan, P. R. China) until the liquid turned into blue-green and clear. Afterwards, we used the *Kjeldahl* nitrogen determination apparatus (K9860, Hanon Instruments Co., Ltd, Jinan, P.R. China) to measure the protein content. Made the blank control group at the same time. The formula for calculating total protein is as follows:

$$\text{protein content (\%)} = \frac{(V_1 - V_2) \times c \times 0.0140}{m \times V_3 / 100} \times F \times 100$$

Where, V_1 is the volume (mL) of samples' liquid H₂SO₄ consumption standard titration solution; V_2 is the volume (mL) of control group (without sample); V_3 is the volume (mL) of absorbed digestive solution; c is the concentration (mol/L) of H₂SO₄ standard titration solution; 0.0140 is the mass of nitrogen of 1.0 mL H₂SO₄; m is the mass of samples (g); F means the coefficient of nitrogen conversion for protein (rice is 5.95).

Determination of Se Content

The content of Se determined by the National food safety standard determination of selenium in foods (GB 5009.93—2010), Wu *et al.* (2007) with a little modification. Primarily weigh 0.1g flour samples into glass digestive tube, added into HNO₃ (9 mL) and HClO₄ (1 mL), mixed up then covered. After ultrasonic cleaners (WD-9415B, LiuYi Co., Ltd, Beijing, P.R. China) at 20°C for 4 h with frequency 100 Hz, then heated using electric hot plate (EH20A Plus, Labtech, USA), HNO₃ added occasionally. When the solution's volume left less than 2 mL and turned into colorless and became clear with blank smoke, made it cools down. Then added HCl (5.0 mL, 6 mol /L), and heated again until the solution turned into colorless and became clear with blank smoke. Cooled it down, poured into volumetric flask, made the blank control group at the same time. After that it was necessary to make the standard curve by adding 0.00 mL, 0.10 mL, 0.20 mL, 0.30 mL, 0.40 mL and 0.50 mL selenium standard solution in centrifugal tube (15 mL), added deionized water to the volume of total 10 mL, HCL (2 mL, guarantee reagent) and potassium ferricyanide (1 mL, 100 g/L). Afterwards, we used the atomic fluorescence spectrophotometer (RGF-6800, Bo Hui Co., Ltd, Beijing, P. R. China) to determine the Se content and set the parameter as follows: Negative high voltage: 340 V; Lamp current: 100 mA; The atomization temperature: 800°C; High furnace: 8 mm; The carrier gas flow rate: 500

mL/min; Shielding gas flow rate: 1000 mL/min; Measurement methods: standard curve; Reading: peak area; Delay time: 1 s; Reading time: 15 s; Charging time: 8 s; Sample size: 2 mL. Se content (mg/kg) was calculated with the following formula:

$$\text{Se content} = \frac{(C - C_0) \times V \times 1000}{m \times 1000 \times 1000}$$

Where, C is the sample measured concentration of digestive solution (ng/mL); C_0 is concentration of blank control group (ng/mL); m is mass of samples; V is the total volume of digestive solution.

Statistical Analysis

The experiment was replicated thrice under the same conditions. Data were expressed as the mean \pm standard error (SEM). One-way ANOVA was carried out with multiple comparisons using Duncan's test to compare the means of different treatments at $p \leq 0.01$. All statistical analyzes were performed using the SPSS 21.0 statistical package (SPSS Inc., Chicago, IL, USA).

Results

Protein Content

Results of protein content determined by *Kjeldahl* method showed that different parts of red glutinous rice have different protein concentrations. Brown red glutinous rice has more total protein than polished red glutinous rice (Fig. 2A). Among all the parts of red glutinous rice, there are significant different protein contents (Fig. 2B). The content of protein is in the following descending order: embryo > aleurone layer > glume > episperm. The protein extracts with protein content $81.65 \pm 1.75\%$ is very high compared with others (Fig. 2C). But the result of the protein extract residue illustrated that there is still a little protein remains in the residue after protein extraction.

Se Content

As can be seen from the Fig. 3A, a noteworthy difference ($p \leq 0.01$) was detected in the tested material; the selenium content of brown red glutinous rice is higher than polished red glutinous rice. Brown red glutinous rice have episperm, endosperm, embryo and aleurone layer, while polished red glutinous rice has endosperm and little embryo. So, we divided different parts of rice to measure the selenium content Fig. 3B. Compared with protein content, we observed relatively the similar trend in selenium content of different parts of rice. The embryo has the highest selenium content than other parts, while the episperm with the minimum protein content has the minimum selenium content. Fig. 3C showed that the selenium content in the red glutinous rice protein extract flour is much higher in comparison to the residue.

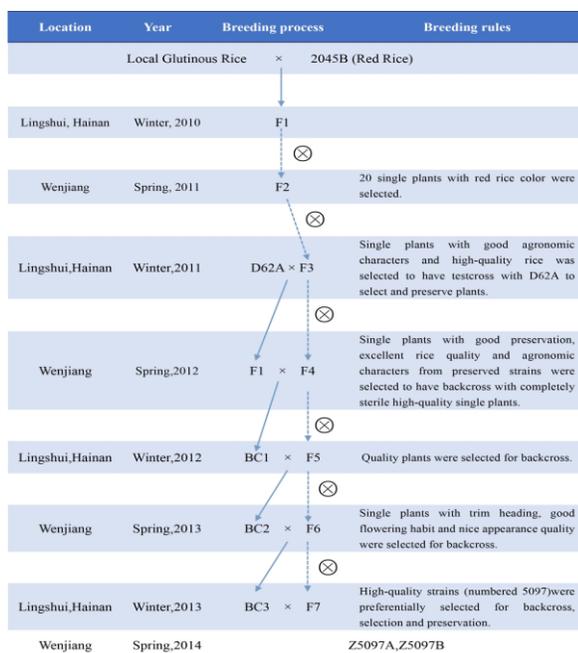


Fig. 1: Breeding process of sterile lines Z5097A and maintainer line Z5097B

Wenjiang, a district in Chengdu, China. Lingshui, Hainan, China. × means crossing; ⊗ means selfing; A means sterile lines; B means maintainer line; F means crossing generation; BC means backcross generation

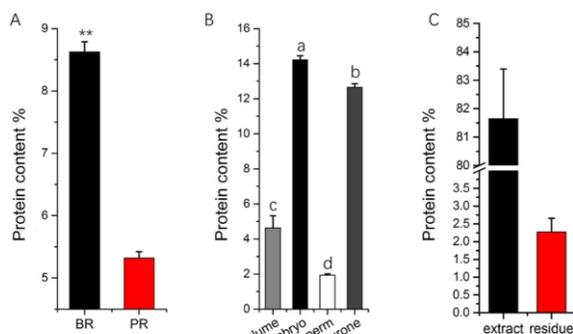


Fig. 2: The protein content from different parts in red glutinous rice

(A) The protein content of brown red glutinous rice and polished red glutinous rice. BR means brown red glutinous rice; PR is polished red glutinous rice. Error bars show the SEM (n = 3). Different letters on the top of column indicate significant differences at $p \leq 0.01$

(B) The protein content in different parts of red rice grain. Error bars show the SEM (n = 3). Lowercase letters on the top of column indicate the statistical significance between different parts of rice according to the Duncan's test ($p \leq 0.01$)

(C) The protein content of protein extracts and residue. Error bars show the SEM (n = 3)

Se Distribution

In red glutinous rice grain, there are five parts making up the whole seed: glume, episperm, endosperm, aleurone layer and embryo (Fig. 4). In a grain of red glutinous rice, the weight of glume, accounts for 17.84%,

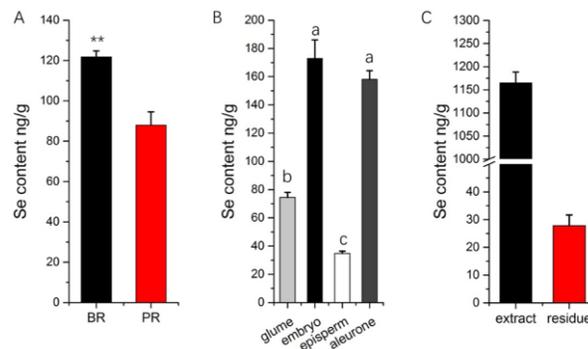


Fig. 3: The Se content from different parts in rice the red glutinous rice

(A) The Se content of brown red glutinous rice and polished red glutinous rice. BR means glutinous rice; PR is polished red glutinous rice. Error bars show the SEM (n = 6). Different letters on the top of column indicate significant differences at $p \leq 0.01$

(B) The Se content in different parts of red rice grain. Error bars show the SEM (n = 6). Lowercase letters on the top of column indicate the statistical significance between different parts of rice according to the Duncan's test ($p \leq 0.01$)

(C) The Se content of protein extracts and residue. Error bars show the SEM (n = 6)

while endosperm occupies 60.66% weight. Thus, in one-gram red glutinous rice seed, all of the five parts contain Se while endosperm significantly has the most of it (Fig. 5A). Besides, the episperm is the only part that has the red color (Fig. 4), which accounts for 7.35% in a grain of red glutinous rice while it holds Se just 2.76%.

Discussion

Breeding of Red Glutinous Rice

In China, most of the selenium rich rice is produced from inorganic selenium fertilizer by spraying on the leaves such as Na_2SeO_3 (Chen *et al.*, 2002; Yong *et al.*, 2009). But it is difficult to ensure the selenium content in rice is within safety boundary. As previously reported, excessive inorganic selenium compounds are toxic to human health (Spallholz, 1994). Uptake of Selenium by plant roots depends greatly on rhizospheric conditions of crops such as competing anions, pH, type of clay minerals and hydrous oxides of iron while the form and concentration of Se in the soil also affects the absorption rate of it (Dhillon and Dhillon, 2003). Thus, to ensure healthy food, it is necessary to monitor and optimize crop selenium concentrations (Zhu *et al.*, 2009) and electing the rice that has strong ability to uptake selenium from soil is a good way to avoid the selenium deficiency or toxicity problem. After five years breeding, the sterile lines Z5097A and maintainer line Z5097B holds selenium contents as 121.74 ± 3.01 ng/g (National standard of selenium rich rice is ≥ 40 ng/g). In addition, the total Se concentrations is more than most other rice in China, Japan, Egypt, France and other countries compared with Williams *et al.* (2009) reports.

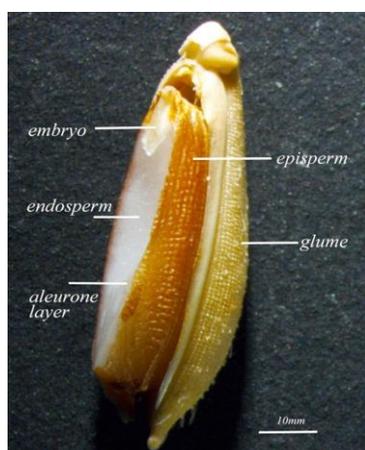


Fig. 4: The endoscopic observation of red glutinous rice
The scale bar represents 10 mm

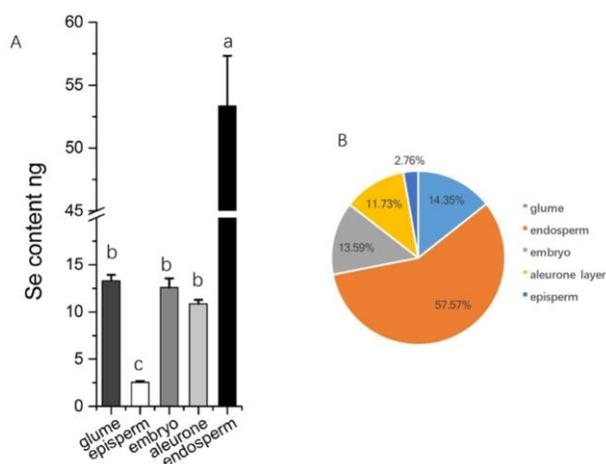


Fig. 5: Se distributes in different part of red glutinous rice grain

(A) The content of Se in different parts of one-gram red glutinous rice seed. Error bars show the SEM ($n = 6$). Lowercase letters on the top of column indicate the statistical significance between different parts of rice according to the Duncan's test ($p \leq 0.01$)

(B) Proportion of Se in a grain of red glutinous rice seed

So the range of selenium in red glutinous rice is not too toxic or deficient. Without inorganic Se fertilizer, the natural Se rich red glutinous rice has a high application value.

Protein Concentrations

Chaiyakul *et al.* (2009) believed that rice has relatively low protein content ranged between 6–8%. While Xie *et al.* (2014) found a range of 6.7–13.8% for the protein content in 335 samples of milled indica rice grown in China, which is a little different from ours. While the range of protein contents in polished rice is determined by the different degree of milling (Chen and Siebenmorgen, 1997; Lamberts *et al.*, 2007; Wang *et al.*, 2007). According to our results, we proved that the high protein concentration in glutinous

rice different parts is located mostly in the outermost layers. Some previous studies reported that the brown rice accumulates more nutrient components such as protein than polished rice which supports this study (Chen *et al.*, 1998; Ohtsubo *et al.*, 2005).

In one-gram red glutinous rice seed, the distribution from outermost to innermost layers was in the following order: glume, episperm, aleurone layer, embryo and endosperm (Fig. 4). It is easily observed that only episperm (the surface of brown rice) contains pigments in natural pigmented rice (Zhu *et al.*, 2017). Lamberts *et al.* (2007) believed that approximately 80% of the kernel proteins were located in the starchy endosperm (the degree of milling > 12%). In this research, there is a trend that the protein content gradually starts declining from embryo to episperm (Fig. 2A and B), which is in accordance with Itani *et al.* (2002) and Resurrection *et al.* (1979). The protein contents found maximum in the embryo is due to the reason that the embryo is the rich source of nutrients and is the next generation plant (Jia *et al.*, 2017). So it is obvious that the maximum protein contents in embryo are needed for plants proper growth and development. Rice protein, with a high nutrition value and good digestion, has a strong ability in antioxidant and antiatherosclerotic (Burriss *et al.*, 2010). Thus, researching for a fast and simply method to extract red glutinous rice protein can improve the application value in the food industry. Compared with Daiana *et al.* (2016), the extract protein content is approximate equality, but the residue with $2.27 \pm 0.38\%$ protein is a little more than $1.30 \pm 0.03\%$ as reported (Fig. 2C). So, how to enhance the protein concentration in extract and decline protein in residue deeply needs to be studied in the future.

Se Concentrations and Distribution

As Lamberts *et al.* (2007) reported, minerals were more abundant in the outer bran layers, losses of minerals reached up to 84.7%, during the milling process for brown rice to polished rice. Promuthai *et al.* (2007) believed that milling process resulted in 25–84% iron loss from different brown rice cultivars. Like other nutrients, selenium is lost inevitably from brown rice to polished rice during milling process (Fig. 3A). Embryo and aleurone layer have more Se than polished rice (endosperm). Fig. 3A, B, which coincides with Sun *et al.* (2010) stated that the Se concentration in bran is 1.9 times higher than corresponding polished rice. The Se content of the protein extract has the highest concentration as 1164.86 ± 23.67 ng/g (Fig. 3C) and it demonstrated that Se prevails in the rice grain is organic Se (selenoprotein), which are in settlement with the findings of Sun *et al.* (2010). Above all the results showed that there is positive correlation between selenium and protein contents, the more protein concentration the more selenium content, while the protein content of embryo is nearly 7.3 times to episperm but the Se concentration of embryo is five times to

episperm. This revealed that availability of Se in the grain is not only in selenoprotein form; plant also accumulated other forms of Se such as Se-polysaccharide, selenium nucleic acid (Stadtman, 1983; Wang et al., 2013).

Total Se concentrations in the rice fractions varied in the following order: husk < polished rice < whole grain < bran. The Se distribution for the one gram of red glutinous rice seed (Fig. 5A and B) showed that organic Se deposited into the embryo, aleurone layer and ultimately in the endosperm. More than half of the total selenium is found in the endosperm of the red glutinous rice as disclosed by Carey et al. (2012). Williams et al. (2009) demonstrated that mature rice grain's endosperm predominantly has organic form of Se while limited to the bran layer was inorganic Se. Kadasi et al. (2010) indicated that the dispersal of S and Se was coincident, evenly distributed throughout the endosperm's aleurone layer, while specifically deposited around the starchy endosperm cells (starch granules) a rich proteinaceous area. Only 5% of the Se was present as inorganic Se in the endosperm. On the contrary, we found more than 80% selenium in the grain as organic Se (selenoprotein).

Conclusion

Based on the results observed in the present study and the aforementioned discussion, it is concluded that the selenium and protein are in direct relationship in red glutinous rice, the more the protein the more will be selenium contents in different parts of rice. Moreover, more than 80% selenium in the grain is organic selenium (in selenoprotein form) and more than half of the total selenium is in the endosperm of the red glutinous rice. Besides, there is still need to identify, what molecular mechanisms or pathways involved in the uptake of selenium, protein and aids their accumulation within specific parts. Meanwhile, the bio-fortified red glutinous rice obtained through genetic improvement by breeding is a natural source of selenium and can be used for further application uses, as natural alternate source to fulfill body selenium demands.

Acknowledgments

This work was supported by the International Cooperation and Exchanges Research Program of the Department of S&T of Sichuan Province (2015HH0028), Technology R&D Program of Sichuan Province (2016NZ0106).

References

- Augilar, F., U.R. Charrandiere, B. Dusemund and P. Galtier, 2009. L-selenomethionine as a source of selenium added for nutritional purposes to food supplements. *J. Eur. Food Saf. Authority*, 13: 1–39
- Ayalew, Y., N. Retta, G. Desse, A. Mohammed and A. Mellese, 2017. Amino acid profile and protein quality in tuber and leaf of *Coccoloba abyssinica* (Lam.) (Cogn.) accessions of Ethiopia. *Food Sci. Nutr.*, 5: 722–729
- Bae, H.J., C.W. Rico, N.R. Su and Y.K. Mi, 2014. Hypolipidemic, hypoglycemic, and antioxidative effects of a new pigmented rice cultivar “superjami” in high fat-fed mice. *J. Kor. Soc. Appl. Biol. Chem.*, 57: 685–691
- Burris, R.L., C.P. Xie, X. Wu, S.B. Melnyk and S. Nagarajan, 2010. Dietary rice protein isolate attenuates atherosclerosis in apoE-deficient mice by upregulating antioxidant enzymes. *Atherosclerosis*, 212: 107–115
- Cann, S.A., J.P. van Netten and N.C. Van, 2000. Hypothesis: iodine, selenium and the development of breast cancer. *Cancer Causes Cont.*, 11: 121
- Carey, A.M., K.G. Scheckel, E. Lombi, M. Newville, Y. Choi and G.J. Norton, 2012. Grain accumulation of selenium species in rice (*Oryza sativa* L.). *Environ. Sci. Technol.*, 46: 5557–5564
- Chairote, E.O., G. Chairote and S. Lumyong, 2009. Red yeast rice prepared from Thai glutinous rice and the antioxidant activities. *Chiang Mai J. Sci.*, 36: 42–49
- Chaiyakul, S., K. Jangchud, A. Jangchud, P. Wuttijumng and R. Winger, 2009. Effect of extrusion conditions on physical and chemical properties of high protein glutinous rice-based snack. *LWT - Food Sci. Technol.*, 42: 781–787
- Chen, H. and T.J. Siebenmorgen, 1997. Effect of rice kernel thickness on degree of milling and associated optical measurements. *Cereal Chem.*, 74: 821–825
- Chen, H., T.J. Siebenmorgen and K. Griffin, 1998. Quality characteristics of long-grain rice milled in two commercial systems. *Cereal Chem.*, 75: 560–565
- Chen, L., F. Yang, J. Xu, Y. Hu, Q. Hu, Y. Zhang and G. Pan, 2002. Determination of selenium concentration of rice in china and effect of fertilization of selenite and selenate on selenium content of rice. *J. Agric. Food Chem.*, 50: 5128–5130
- Daiana, D.S., A.F. Sbardelotto, D.R. Ziegler, L.D. Marczak and I.C. Tessaro, 2016. Characterization of rice starch and protein obtained by a fast alkaline extraction method. *Food Chem.*, 191: 36–44
- Dhillon, K.S. and S.K. Dhillon, 2003. Distribution and management of seleniferous soils. *Adv. Agron.*, 79: 119–184
- Faiz, A., M.M. Hanafi, M.A. Hakim, M.Y. Raffi and S.N.A. Abdullah, 2015. Micronutrients, Antioxidant Activity, and Tocochromanol Contents of Selected Pigmented Upland Rice Genotypes. *Int. J. Agric. Biol.*, 17: 741–747
- Ferlay, J., I. Soerjomataram, R. Dikshit, S. Eser, C. Mathers, M. Rebelo, D.M. Parkin, D. Forman and F. Bray, 2015. Cancer incidence and mortality worldwide: sources, methods and major patterns in globocan 2012. *Int. J. Cancer*, 136: 359
- Gealy, D.R., D.H. Mitten and J.N. Rutger, 2003. Gene flow between red rice (*Oryza sativa*) and herbicide-resistant rice (*O. sativa*): implications for weed management I. *Weed Technol.*, 17: 627–645
- Ghadirian, P., P. Maisonneuve, C. Perret, G. Kennedy, P. Boyle and D. Krewski, 2000. A case-control study of toenail selenium and cancer of the breast, colon, and prostate. *Cancer Detect. Prevent.*, 24: 305–313
- Heber, D., I.J. Yip, D. Elashoff, R. Elashoff and V. Go, 1999. Cholesterol-lowering effects of a proprietary Chinese red-yeast-rice dietary supplement. *Amer. J. Clin. Nutr.*, 69: 231–236
- Huang, X., S. Yang, J. Gong, Z. Qiang, F. Qi and Q. Zhan, 2016. Genomic architecture of heterosis for yield traits in rice. *Nature*, 537: 629
- Institute of Medicine (US) Panel on Dietary Antioxidants and Related Compounds, 2000. *Dietary Reference Intakes for Vitamin C, Vitamin E, Selenium, and Carotenoids*. National Academies Press
- Itani, T., M. Tamaki, E. Arai and T. Horino, 2002. Distribution of amylose, nitrogen, and minerals in rice kernels with various characters. *J. Agric. Food Chem.*, 50: 5326–5332
- Jia, W., J.M. Baskin, C.C. Baskin, G. Liu, X. Yang and Z. Huang, 2017. Seed dormancy and germination of the medicinal holoparasitic plant *Cistanche deserticola*, from the cold desert of northwest china. *Plant Physiol. Biochem.*, 115: 279–285
- Ju, Z.Y., N.S. Hettiarachchy and N. Rath, 2001. Extraction, denaturation and hydrophobic properties of rice flour proteins. *J. Food Sci.*, 66: 229–232
- Kadasi, L.M., W.C. Dent and A.M. Malek, 2010. Nanosims analysis of arsenic and selenium in cereal grain. *New Phytol.*, 185: 434–445

- Kanitha, T. and T. Wanida, 2010. Extraction and application of antioxidants from black glutinous rice. *LWT - Food Sci. Technol.*, 43: 476–481
- Knekt, P., J. Marniemi, L. Teppo, M. Heliövaara and A. Aromaa, 1998. Is low selenium status a risk factor for lung cancer. *Amer. J. Epidemiol.*, 148: 975
- Lamberts, L., E.D. Bie, G.E. Vandeputte, W.S. Veraverbeke, V. Derycke and W.D. Man, 2007. Effect of milling on colour and nutritional properties of rice. *Food Chem.*, 100: 1496–1503
- Liu, R.H., 2004. Potential synergy of phytochemicals in cancer prevention: mechanism of action. *J. Nutr.*, 134: 3479–3485
- Loypimai, P., K. Sittisuanjik, A. Moongarm and W. Pimthong, 2017. Influence of sodium chloride and vacuum impregnation on the quality and bioactive compounds of parboiled glutinous rice. *J. Food Sci. Technol.*, 54: 1–9
- Loypimai, P., A. Moongarm and P. Chottanom, 2009. Effects of ohmic heating on lipase activity, bioactive compounds and antioxidant activity of rice bran. *Aust. J. Basic Appl. Sci.*, 4: 3642–3652
- Nam, S.H., S.P. Choi, M.Y. Kang, H.J. Koh, N. Kozukue and M. Friedman, 2006. Antioxidative activities of bran extracts from twenty-one pigmented rice cultivars. *Food Chem.*, 94: 613–620
- Ohtsubo, K., K. Suzuki, Y. Yasui and T. Kasumi, 2005. Bio-functional components in the processed pre-germinated brown rice by a twin-screw extruder. *J. Food Compos. Anal.*, 18: 303–316
- Patrick, L., 2004. Selenium biochemistry and cancer: a review of the literature. *Altern. Med. Rev.*, 9: 239–258
- Promuthai, C., C. Sanchai, B. Rerkasem, S. Jamjod, S. Fukai and I.D. Godwin, 2007. Effect of grain morphology on degree of milling and iron loss in rice. *Cereal Chem.*, 84: 384–388
- Que, F., L. Mao and X. Pan, 2006. Antioxidant activities of five Chinese rice wines and the involvement of phenolic compounds. *Food Res. Int.*, 39: 581–587
- Resurrection, A.P., B.O. Juliano and Y. Tanaka, 1979. Nutrient content and distribution in milling fractions of rice grain. *J. Sci. Food Agric.*, 30: 475
- Scieszka, M., A. Danch, M. Machalski and M. Drózd, 1997. Plasma selenium concentration in patients with stomach and colon cancer in the upper Silesia. *Neoplasma*, 44: 395–397
- Singh, N., H. Singh, K. Kaur and M.S. Bakshi, 2000. Relationship between the degree of milling, ash distribution pattern and conductivity in brown rice. *Food Chem.*, 69: 147–151
- Spallholz, J.E., 1994. On the nature of selenium toxicity and carcinostatic activity. *Free Radical Biol. Med.*, 17: 45–64
- Stadtman, T.C., 1983. New biologic functions—selenium-dependent nucleic acids and proteins. *Fundament. Appl. Toxicol.*, 3: 420
- Sun, G.X., X. Liu, P.N. Williams and Y.G. Zhu, 2010. Distribution and translocation of selenium from soil to grain and its speciation in paddy rice (*Oryza sativa* L.). *Environ. Sci. Technol.*, 44: 6706–6711
- Wallace, K., K.T. Kelsey, A. Schned, J.S. Morris, A.S. Andrew and M.R. Karagas, 2009. Selenium and risk of bladder cancer: a population-based case-control study. *Cancer Prevent. Res.*, 2: 70–73
- Wang, Y.J., L. Wang, D. Shephard, F. Wang and J. Patindol, 2007. Properties and structures of flours and starches from whole, broken, and yellowed rice kernels in a model study. *Cereal Chem.*, 79: 383
- Wang, Y., J. Chen, D. Zhang, Y. Zhang, Y. Wen, L. Li and L. Zheng, 2013. Tumoricidal effects of a selenium (Se)-polysaccharide from Ziyang green tea on human osteosarcoma U-2 OS cells. *Carbohydr. Polym.*, 98: 1186–1190
- Williams, P.N., E. Lombi, G.X. Sun, K. Scheckel, Y.G. Zhu and X.B. Feng, 2009. Selenium characterization in the global rice supply chain. *Environ. Sci. Technol.*, 43: 6024–6030
- Wu, H., Y. Jin, Y. Shi and S. Bi, 2007. On-line organoselenium interference removal for inorganic selenium species by flow injection coprecipitation preconcentration coupled with hydride generation atomic fluorescence spectrometry. *Talanta*, 71: 1762–1768
- Xia, Y., K.E. Hill, D.W. Byrne, J. Xu and R.F. Burk, 2005. Effectiveness of selenium supplements in a low-selenium area of china. *Amer. J. Clin. Nutr.*, 81: 829–834
- Xie, L.H., S.Q. Tang, N. Chen, J. Luo, G.A. Jiao and G.N. Shao, 2014. Optimisation of near-infrared reflectance model in measuring protein and amylose content of rice flour. *Food Chem.*, 142: 92–100
- Yong, F., Y.F. Zhang, B. Catron, Q.L. Chan, Q.H. Hu and J.A. Caruso, 2009. Identification of selenium compounds using HPLC-ICPMS and nano-ESI-MS in selenium-enriched rice via foliar application. *J. Anal. Atom. Spectr.*, 24: 1657–1664
- Yu, S.Y., Y.J. Zhu and W.G. Li, 1997. Protective role of selenium against hepatitis B virus and primary liver cancer in Qidong. *Biol. Trace Elem. Res.*, 56: 117–124
- Zhao, W., W. Xu, Y. Chai, X. Zhou, M. Zhang and W. Xie, 2017. Differences in selenium uptake, distribution and expression of selenium metabolism genes in tomatoes. *Int. J. Agric. Biol.*, 19: 528–534
- Zhu, Q., S. Yu, D. Zeng, H. Liu, H. Wang, Z. Yang and X. Zhao, 2017. Development of “Purple Endosperm Rice” by Engineering Anthocyanin Biosynthesis in the Endosperm with a High-Efficiency Transgene Stacking System. *Mol. Plant*, 10: 918–929
- Zhu, Y.G., E.A.H. Pilonismit, F.J. Zhao, P.N. Williams and A.A. Meharg, 2009. Selenium in higher plants: understanding mechanisms for biofortification and phytoremediation. *Trends Plant Sci.*, 14: 436

(Received 04 October 2017; Accepted 02 January 2018)