Enhanced antioxidant activity of okara through solid state fermentation of GRAS Fungi

Azis Boing SITANGGANG1,2,3*, Wenny Silvia Loren SINAGA4, Felicia WIE4, Franz FERNANDO4, Warawut KRUSONG3

Abstract

The enhancement of antioxidant activity of fermented okara was investigated using fungal solid-state fermentation. The initial fungal concentration and water activity were found to influence the enhanced antioxidant capacity of fermented okara. Extracellular enzymes, especially proteases and carbohydrases might facilitate the hydrolysis of okara protein that yielded antioxidative amino acids and peptides, as well as the release of isoflavones-derived components. Fermentation of okara with R. oligosporus with 1.04x10^3 g DCW/g okara, as initial concentration, water activity of 0.931 was found to yield the highest antioxidant strength. Whereas for A. Oryzae, 1.6x10^3 g DCW/g okara was optimum as the initial concentration with water activity of 0.913. With these optimum conditions, solid-state fermentation of okara could enhance the antioxidant activity up to 2.6- and 5.5-fold, respectively for R. oligosporus and A. oryzae. Besides having an enhanced antioxidant activity, the fermented okara also had higher protein content, amino nitrogen content, and protein digestibility.

Keywords: antioxidant; A. oryzae; okara; R. oligosporus; solid-state fermentation.

Practical Application: Fermentation conditions within this study can be used as a strategy to valorise okara.

1 Introduction

Okara is a by-product from soymilk or tofu production, and usually used as feed stocks, fertilizer, or directly discarded as waste (Li et al., 2012). For the latter, large quantities of okara produced per annum will lead to environmental problems (O’Toole, 1999). It is, in fact contradicted to inherent nutrients which are rich in okara. In okara, especially protein and its respective amino acids, can level up to 27% based on the dry basis (Rashad et al., 2011). Considering this, there are many strategies have been reported on the valorisation of okara for human consumption, such as increasing its digestibility through treatments with cellulolytic and proteolytic enzymes (Kasai et al., 2004), protein deamination (Chan & Ma, 1999), or liberation of its phenolic content, and other enhancement of its nutritive values through fermentations (Hur et al., 2014).

Microbial fermentation has been reported to influence the antioxidant activity in plant-based foods. During fermentation, phenolic compounds can be hydrolysed that entails on the depolymerisation of its complex, and liberation of its derivatives (Hur et al., 2014). Additionally, due to the production of extracellular proteases, and carbohydrases, both protein and carbohydrate complex within the structures of cereals or grains are hydrolyzed. Either liberation of phenolic compounds or the production of small molecular peptides will consequently lead to a higher radical scavenging activity of the fermented cereal/grains (Chang et al., 2009).

Compared to submerged fermentation, solid state fermentation offers several advantages, as it is more cost- and energy-effective, as well as environmentally friendly (Krishna, 2005). Commonly, solid state fermentation is used to enhance the nutritional value of cereal grains in which microbial proteases are expected to hydrolyze the proteins inherently found in the beans or grains (Barampama & Simard, 1995). For instance, the hydrolysis of proteins in soybean during tempe fermentation usually yields small molecular weight peptides and free amino acids which are active biologically (Nout & Kiers, 2005). Especially for okara, its solid state fermentations using various microorganisms (i.e., bacteria and fungi) have been reported (Chi & Cho, 2016; Mateos-Aparicio et al., 2010; Visessanguan et al., 2005). Most of the goals are to enhance okara’s antioxidant properties, flavour properties, protein digestibility, production of antibiotic, etc. (Guan et al., 2016; Ohno et al., 1996; Vong et al., 2018).

Within this work, a detailed comparative study between Rhizopus oligosporus- and Aspergillus oryzae-based okara solid state fermentation is reported. R. oligosporus is a common mold used in Indonesia for tempe fermentation, and is known to have a proteolytic activity. According to Wronkowska et al. (2015), solid state fermentation using R. oligosporus could enhance the antioxidant properties of raw and roasted buckwheat groats in terms of free radical scavenging activity. The study conducted by Vong et al. (2018) on the utilization of R. Oligosporus for okara solid state fermentation was not specific. The enhancement of okara properties by R. oligosporus was not singled out within...
their study, as they mainly discussed the detailed possible synergistic effects between *R. oligosporus* and *Farraria lipolytica* in producing enzymes for bioconversion some components in the okara cell wall. On the other hand, there have been many modifications using *A. oryzae* reported, and one of them was for the improvement of digestibility of the fermented grains or cereals (Teng et al., 2012). De Castro & Sato (2014) reported that culturing *A. oryzae* on different agroindustrial wastes as matrix supports yield high protease activity.

The use of individual *R. oligosporus* and *A. oryzae* are expected to produce extracellular protease(s) and also possible carbohydrides that concomitantly hydrolize the proteins in okara into bioactive peptides, and the intact complex carbohydrates to release the proteins itself. For fungal protease, its production has been reported to be influenced by the fermentation time, incubation temperature, initial mold concentration, and water activity (\(a_w\)) (Nuñez-Gaona et al., 2010). Thus, these factors must be considered while conducting solid state fermentation of okara. Additionally, besides having generally recognized as safe (GRAS) status, the two fungi are also grouped as "microbial food cultures" (MFC) based on three criteria, such as: regulatory systems and legal terms, scientific criteria, and inventory of microbial species in food fermentations (Bourdichon et al., 2012). Therefore, the utilization of these fungi to enhance the functional properties of okara intended for functional food applications will be appropriate.

### 2 Materials and methods

#### 2.1 Materials

Okara, tempe and tofu were purchased from a local market in Tangerang, Indonesia. Especially for okara, it is a solid waste of soymilk production. Pure culture of *R. oligosporus* and *A. oryzae* were purchased from MBIO Biotekindo, Bogor, Indonesia. Potato dextrose broth (PDB, P6685) was from Sigma-Aldrich, Thailand. For ascorbic acid (100468), 2,2-diphenyl-1-picrylhydrazyl solution (DPPH) was from Sigma-Aldrich. For methanol, hydrogen chloride, sodium hydroxide, sodium azide (106687) were purchased from Merck Millipore, Thailand. For methanol, hydrogen chloride, sodium hydroxide, phosphoric acid, and ethanol were analytical grade and purchased from PT. Smart Lab, Indonesia.

#### 2.2 Growth profile of *R. oligosporus* and *A. oryzae*

Initially, the lyophilized fungal cultures were thawed in a water bath at 30 °C. The thawed *R. oligosporus* and *A. oryzae* culture were looped into sterilized PDB (temperature of 121 °C, pressure of 1 atm for 1 h). The inoculated broth with individual fungal type was incubated for 3-5 d at temperature of 30 °C. A 10 µL of solution from each newly activated fungus was placed into a clean test tube where fresh PDB solution was added into, until 10 mL of total solution was obtained. Another incubation period was performed until 11 d at different temperatures (25, 30, and 37 °C). The growth profile was constructed for each type of fungus by monitoring the amount of biomass produced (expressed in gram dry cell weight, \(g_{DCW}\)).

#### 2.3 Effect of initial fungal concentration and water activity on the functional properties of okara

A 10 g of dried okara (see Table 1 for dried okara properties) was placed into petri dishes (P5606 Sigma-Aldrich), and added with 10 mL of distilled water to reach the water activity, \(a_w\), of 0.913. Different initial concentrations (\(g_{DCW}\)) of *R. oligosporus* and *A. Orzzyae* were inoculated separately into the substrates. The incubations were carried out for 5 d, by placing the petri dishes into an incubator at predetermined temperature. For investigating the influence of water activity, the dried okara (moisture content of 8.26% wb) was added with sterile water to vary the \(a_w\) level.

#### 2.4 Radical scavenging activity

A 2.0 mL of methanol extract (i.e., okara, fermented okara, tofu or tempe) or standard (i.e., ascorbic acid) at a concentration of 2.0 mg/mL was added to 1.0 mL of 0.3 mM 2,2-diphenyl-1-picrylhydrazyl solution (DPPH). The solution was mixed thoroughly and left in the dark to stand for 30 min at 30 °C. The blank for each sample was prepared by replacing DPPH with methanol in the prementioned mixture above. The control preparation was made with 1.0 mL of methanolic DPPH plus 2.0 mL of methanol (Rashad et al., 2011). The absorbance of each solution was determined at 517 nm. The radical scavenging activity, RSA (%) was calculated as follows (equation 1):

\[
RSA(\%) = \left(1 - \frac{A_s}{A_c}\right) \times 100\%
\]

where RSA (%) stands for radical scavenging activity, \(A_s\) and \(A_c\) refer to absorbance of sample and control, respectively (-).

#### 2.5 Protein content

The protein determination following Kjeldahl method (Association of Official Analytical Chemists, 2005) was started by weighing 1.0 g of okara, 7.0 g of K₂SO₄, 5.0 mg of selenium, 10 mL of H₂SO₄, and 10 mL of H₂O, into 100 mL Kjeldahl flask. The mixture was then boiled at 420 °C for 30 min. The end-point of the process was indicated by a clear solution without leaving charred materials. The solution was cooled down to 50-60 °C, followed by the addition of 50 mL of distilled water. A 50 mL of 35% NaOH was added to the mixture, and the distillation was performed to obtain approximately 100 mL of distillate.

<p>| Table 1. Characteristics of dried okara. |</p>
<table>
<thead>
<tr>
<th>Okara Characteristics</th>
<th>Values</th>
</tr>
</thead>
<tbody>
<tr>
<td>Moisture content [% wb]</td>
<td>8.26 ± 0.24</td>
</tr>
<tr>
<td>Water activity (a_w) [-]</td>
<td>0.183</td>
</tr>
<tr>
<td>Particle size [µm]</td>
<td>6.15 × 10⁻¹</td>
</tr>
<tr>
<td>Radical scavenging act. [%]</td>
<td>14.87 ± 0.44</td>
</tr>
<tr>
<td>Protein content [% db]</td>
<td>23.96 ± 0.43</td>
</tr>
<tr>
<td>Amino Nitrogen content [mg/g]</td>
<td>3.48 ± 0.12</td>
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</tbody>
</table>
This distillate was placed into a 125 mL Erlenmeyer flask that previously poured with 25 mL of 4% boric acid. Afterwards, about 2-3 drops of bromocresol green-methyl red indicator were added, and this final solution was titrated with 0.2 N HCl until faint pink gray color was obtained. The nitrogen and protein content were calculated as follows (equation 2 and equation 3):

\[
N(\%) = \left( \frac{V_{\text{HCl}} - V_{\text{blank}}}{W_{\text{sample}}} \right) \times 100\%
\]  
(2)

\[
P(\%) = N(\%) \times C_{\text{fact}}
\]  
(3)

where \( N(\%) \) = percentage of nitrogen in the sample, \( N = \) Normality (eq/L), \( V = \) volume (mL), \( W = \) weight (g), \( P(\%) = \) percentage of protein in the sample, and \( C_{\text{fact}} = \) conversion factor of nitrogen to protein, and in this study the value used was 5.71.

### 2.6 Amino nitrogen content and degree of hydrolysis

Formal titration was used to determine the amino nitrogen content of okara (Morais et al., 2013). Formaldehyde reagent was prepared by diluting 25 mL of formaldehyde with a 50% ethanol to reach a volume of 250 mL. The pH of this reagent was brought to 8.0 with 0.005 M NaOH. The okara hydrolysate solution was prepared where the pH was also adjusted to reach 8.0. About 5.0 mL of formaldehyde reagent and 5.0 mL sample solution were poured into the Erlenmeyer flask, continued by the addition of 2-3 drops of phenolphthalein. The final solution was titrated with 0.005 M NaOH until pink colored solution was obtained. The total volume of NaOH consumed was recorded, and the amino nitrogen content was calculated as follows (equation 4):

\[
Am.N\text{ (mg/g }_{\text{sample}}\text{)} = V_{\text{NaOH}} \times M_{\text{NaOH}} \times 14 \text{ mg/mmol} \times DF
\]  
(4)

where \( Am.N = \) amino nitrogen content (mg/g \(_{\text{sample}}\)), \( V = \) volume (mL), \( M = \) Molarity, and \( DF = \) dilution factor.

The degree of hydrolysis was calculated as the ratio of amino nitrogen content \((Am.N)\) from formal titration and total nitrogen \((N)\) from Kjeldahl method as following (equation 5):

\[
DH(\%) = \left( \frac{Am.N}{N} \right) \times 100\%
\]  
(5)

where \( DH = \) degree of analysis (%).

### 2.7 In vitro protein digestibility

The samples were weighed for 250 mg and transferred into 50 mL Erlenmeyer flask, whereas for the blank about 250 μL of deionized water were used. A 15 mL of 0.1 N HCl containing 1.5 mg/mL pepsin was added into each flask, and then incubated at 37 °C for 3 h. The reaction was stopped by neutralizing the solution with 7.5 mL of 0.5 N NaOH. For the pancreatic digestion, it was done overnight at 37 °C in which 10 mL of 0.2 M phosphate buffer (pH 8) containing 10 mg of pancreatin (107133, Merck-Millipore) was added to the neutralized solution. Additionally, 1.0 mL of 0.005 M azide was also added to the mixture to prevent microbial growth. Prior to centrifugation at 503 × g for 20 min, the mixture was added with 1.0 mL of 0.6M trichloroacetic acid (Akesson & Stahmann, 1964; Almeida et al., 2015). The protein content was determined based on nitrogen content using Kjeldahl method (Association of Official Analytical Chemists, 2005). The in vitro protein digestibility was calculated as follows (equation 6):

\[
Dig(\%) = \left( 1 - \frac{N_c}{N_s} \right) \times 100\%
\]  
(6)

where \( Dig = \) protein digestibility (%), \( N_s = \) nitrogen content in blank (mg/g \(_{\text{sample}}\)), \( N_c = \) nitrogen content in sample (mg/g \(_{\text{sample}}\)).

### 2.8 The amino acids analysis

The analysis of amino acids was performed by PT. Saraswanti Indo Genetech (18-5-17/MU SMM-SIG, UPLC) where the method has been also reported by Alen et al. (2017). The hydrolysis of dried okara and fermented okara was done at 110 °C for 24 h using 6 M HCL containing 0.5% (w/v) phenol. After the hydrolysis, the samples were dried using 20 mL of ethanol: triethylamine: water (2:1.2, v/v) for 20 min at room temperature. Prior to analysis, the sample was dissolved in 100 μL of equilibration buffer. The analysis of the amino acids was performed with a Waters C18 column (3.9 x 150 mm). The used mobile phase was a combination between acetonitrile and sodium acetate buffer (about 300 mL acetonitrile, 200 mL water, 0.2 mL CaEDTA). A 20 μL of aliquots was injected into the column, and the measurement was performed at 254 nm. The comparison was made between the individual peaks of the sample with the standards (Alen et al., 2017).

### 3 Results and discussion

#### 3.1 Okara characteristics

The initial characteristics of the dried okara are shown in Table 1. Prior to be used, the dried okara was milled and sieved through 60 mesh. The particle size obtained was about 6.146×10⁻¹ μm. The initial antioxidant activity of the dried okara, in terms of radical scavenging activity was 14.87±0.44%. Presumably, the presence of isoflavones was associated to this antioxidant strength. It must be noted that the okara used was a solid waste of soy milk production. Due to the application of wet milling process for the preparation of soy milk, the soybean proteins were extracted. In this study, the protein content of okara was about 24% which was far below from its initial content in soybean meal, ranging from 37-42% (D’Keefe et al., 2015). This value was comparable to that reported by Yokomizo et al. (2002) and Li et al. (2012), where the concentrations of okara protein in their studies were 25, and 25.4-28.4% db, respectively.

#### 3.2 The growth curve of R. oligosphorus and A. oryzae

Different cell concentrations were obtained as the incubation temperature was varied respective to incubation time for both R. oligosphorus and A. oryzae. The incubation temperature of 30 °C yielded the highest biomass concentration for both fungi (data for 25 and 37 °C are not shown). R. oligosphorus and A. oryzae are known to be mesophilic fungi. Therefore, both fungi are...
able to grow at moderate temperatures between 20 and 45 °C, and with optimum growth temperatures between 30 and 39 °C (Schiraldi & Rosa, 2014).

The optimum productions of an extracellular enzyme is normally associated with the optimum cell concentration obtained throughout the incubation time. According to Rauf et al. (2010), temperature of 30 °C was the optimum growth temperature for acidic protease production from *R. oligosporus* through solid state fermentation of sunflower meal. Whereas for *A. oryzae*, temperatures of 28-30 °C were found to be the suited range for production protease, amylase and glucoamylase (Puri et al., 2013; Kamath et al., 2010). Within this study, temperature of 30 °C was confirmed and selected as the incubation temperature for both fungal preparations, and okara solid-state fermentation.

At the first four-day of incubation at 30 °C, both *R. oligosporus* and *A. oryzae* had a growth rate that was relatively similar. *R. oligosporus* had a maximum biomass concentration of $22.34 \times 10^{-3}$ g DCW/mL at 6 d incubation. Whereas for *A. Oryzae*, it was obtained at 5 d incubation, where the value was lower approximately 34% compared to *R. Oligosporus* biomass concentration (Figure 1).

### 3.3 Effect of initial fungal concentration on okara properties

Within this study, three levels of initial biomass concentration for each fungus were designed to conduct solid state fermentations of okara. These levels of fungal concentration were obtained by following the growth profile of each fungus that previously described. Unfortunately, the levels of the concentration were not the same for both fungi. Definitely, such a simple dilution could be performed to obtained the same levels. However, one must carefully consider to have an identical substrate's water activity for both fungi.

With the highest initial *R. oligosporus* concentration (i.e., $2.44 \times 10^{-3}$ g DCW/g okara), the highest radical scavanging activity was obtained after for 4 d incubation (32.33%) (Figure 2a). At cell concentration of $1.95 \times 10^{-3}$ g DCW/g okara, the antioxidant activity also peaked at 4 d incubation. However, the value was smaller than that of the highest initial *R. oligosporus* concentration. The okara fermentation with the lowest initial concentration of *R. oligosporus* yielded the highest antioxidant activity of 40.1% after 5 d fermentation.

Okara fermentation with *A. oryzae* could obtain the highest antioxidant activity of 82.6% using the highest initial biomass concentration (i.e., $1.6 \times 10^{-3}$ g DCW/g okara). This was achieved for 4 d incubation period (Figure 2b). For the other two initial biomass concentrations of *A. oryzae*, the highest antioxidant activity also peaked at 4 d incubation, around 76 and 78%, respectively for cell concentration of $0.8 \times 10^{-3}$ and $1.4 \times 10^{-3}$ g DCW/g okara. It was found that solid state fermentation of okara using *R. oligosporus* and *A. oryzae* could enhance the antioxidant activity up to 2.6- and 5.5-fold, respectively.

![Figure 1](image1.png)  
**Figure 1.** The growth curve of *R. oligosphorus* and *A. oryzae* in PDB medium, incubated at 30 °C.

![Figure 2](image2.png)  
**Figure 2.** Influence of (a) *R. oligosporus* and (b) *A. oryzae* initial concentrations towards antioxidant activity of fermented okara, incubated at 30 °C, and water activity ($a_w$) of 0.913.
Enhanced antioxidant activity of okara

The increase of antioxidant activity of fermented okara has been also reported in many studies utilizing different microorganisms. Rashad et al. (2011) mentioned that the highest significant levels of antioxidant activities of fermented okara were achieved with solid-state fermentation of *K. marxianus* NRRL Y-8281. It was assumed that fungal extracellular β-glucosidase produced during fermentation could catalyse the release of aglycones from okara. Additionally, high concentrations of genistein and daidzein were reported during tempe fermentation which was caused by the activity of *R. oligosporus* β-glucosidase on isoflavones (Murakami et al., 1984).

Besides isoflavones, peptides and amino acids might also play a vital role for the increase of scavenging activity of fermented okara. Yokomizo et al., (2002) demonstrated that fungal and bacterial proteases are able to hydrolyse okara protein to have antioxidative activity-based peptides against the peroxidation of linoleic acid performed in an aqueous system at pH 7.0. Four antioxidative peptides were successfully isolated from the hydrolysate prepared with the addition of *B. subtilis* protease (Yokomizo et al., 2002). Related to this work, Sparringa & Owens (1999) reported that after 3 d fermentation, approximately eight-fold increase in amino acids and peptides content of tempe was pronounced.

Vong et al. (2018) reported the coculture of *R. oligosporus* and yeast *Yarrowia lipolytica* (ROYL) on okara. After fermentation, the flavor, digestibility as well as nutritional value of okara were enhanced. The enhancements that are specifically related to the increased antioxidant activities were 254% more free amino acids, 179% more isoflavone aglycones and 197% more free phenolic acids. Moreover, Guan et al. (2016) used *Actinomucor elegans* DCY-1 to ferment okara. They assumed that proteins were hydrolyzed into peptides during fermentation. Thus, fermented okara showed an increase in antioxidative activities as indicated by the enhanced ferrous ion-chelating activity, reducing power and scavenging effect of ABTS, DPPH and hydroxyl radicals of fermented okara. Within this study, it conclusively considered that release of isoflavones-derived compounds, small molecular weight peptides, and amino acids produced during the fermentation could enhance the antioxidant activities of fermented okara.

In Figure 3a and b, a combination of a longer fermentation time and a higher initial biomass concentration was able to yield a higher degree of protein hydrolysis. Okara fermentation that was conducted for 5 d at the highest initial cell concentration for both fungi yielded the highest degree of hydrolysis, of 19.2 and 26.3%, respectively for *R. oligosporus* and *A. oryzae*. The difference of the degree of hydrolysis between *R. oligosporus* and *A. oryzae* might be related to the amount of protease produced and its corresponding activity (Teng et al., 2012). According to Zhang et al. (2014), the formation of free amino acids was linear to amino nitrogen, and increased proportionally to the length of fermentation. A longer fermentation was considered to give sufficient time for the accumulated extracellular protease to hydrolyze protein into oligopeptides, small molecular weight peptides and even into free amino acids (Zhang et al., 2014). According to Zou et al. (2016), most of peptides that have antioxidant activity are in the range of tripeptides until oligopeptides. Adebiyi et al. (2009) mentioned that antioxidative peptides normally have molecular weights between 600 until 2,000 Da. Hence, the degree of hydrolysis might be able to explain the reduction of antioxidant activity in Figure 2a and b after a prolonged fermentation. After the highest antioxidant activity was obtained in fermented okara, such prolonged fermentation would facilitate further hydrolysis of oligopeptides and small molecular peptides into free amino acids that entailed in a less antioxidant strength.

3.4 Effect of water activity on okara properties

It is necessary to determine the optimum water activity in solid-state fermentation. A lower water activity in the fermentation system causes solubility problems of nutrients found in the substrate. On the other hand, high water activity, *i.e.*, higher

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**Figure 3.** Influence of (a) *R. oligosporus* and (b) *A. oryzae* initial concentrations towards degree of protein hydrolysis of fermented okara, incubated at 30 °C, and water activity (*a*<sub>w</sub>) of 0.913.
moisture level can lead to a reduced porosity of the substrate, and thereby limiting oxygen transfer (Prasad & Raju, 2013). In Figure 4a and b, the increase of water activity resulted in a higher antioxidant capacity and protein degree of hydrolysis of *R. oligosporus*-based okara solid state fermentation. However, for *A. oryzae*-based okara solid state fermentation, an increase in water activity yielded a lower value either for antioxidant activity or degree of hydrolysis. Higher water activity in *A. oryzae*-based okara solid state fermentation might result in tight mass that restricted the aeration, decreased porosity, and the available area for growth as well (Yu et al., 2012).

### 3.5 Amino acid profiles of fermented okara

The comparison of amino acids profile between unfermented and fermented okara can be seen in Figure 5. The respective amino acids that have been reported to have antioxidant activity are proline, histidine, tyrosine, cysteine, valine, leucine, phenylalanine, and tryptophan (Hur et al., 2014). Cysteine and tyrosine were increased in *A. oryzae*-based okara fermentation, while for *R. oligosporus*-based okara fermentation only showed an increase in cysteine. Moreover, essential amino acids such as methionine, cysteine, tryptophan, and threonine were increased.

![Figure 4](image4.png)

**Figure 4.** Influence of water activity $a_w$ on the degree of protein hydrolysis (a) and antioxidant activity (b) of fermented okara. Conditions: for *R. oligosporus*-based okara fermentation, the initial concentration was $1.04 \times 10^{-3}$ gDCW/gokara, incubated at 30 °C for 5 d, whereas for *A. oryzae*-based okara fermentation, the initial concentration was $1.6 \times 10^{-3}$ gDCW/gokara, incubated at 30 °C for 4 d.

![Figure 5](image5.png)

**Figure 5.** Amino acid profiles of unfermented and fermented okara with *R. oligosporus* and *A. oryzae.*
when okara was fermented by *A. oryzae*. The increase of essential amino acids in fermented okara by *R. oligosporus* was also identical, except for threonine. The fluctuation in the amino acid concentrations for fermented okara might be due to their utilizations for the metabolisms of both fungi (Kwon et al., 2010).

### 3.6 Functional Comparison between unfermented, fermented okara, and other soy-based products.

Solid-state fermentation of okara with *R. oligosporus* was optimum with $1.04 \times 10^2 \frac{g_{DCW}}{g_{soy}}$ as initial concentration, substrate’s water activity of 0.931, and incubated for 5 d. Whereas for *A. Oryzae*, $1.6 \times 10^2 \frac{g_{DCW}}{g_{soy}}$ was the optimum initial concentration with a substrate’s water activity of 0.913, and fermentation time of 4 d. The antioxidant, protein content, amino nitrogen content and protein digestibility of fermented okara were found higher than unfermented one (Table 2). Additionally, the fermented okara was also found to have higher antioxidant activity than other soy-based products such as (commercial) tofu and tempe.

The particle size of okara was reduced down to 6.146 × 10$^{-1}$ μm (see Table 1), whereas for tempe production, soybean is normally unmilled. Schmidt & Furlong (2012) reported that substrates with smaller particles will result in a higher surface area leading to a larger contact area between the fungus and the substrate, favoring its growth. Additionally, for the extracellular protease to work optimally a smaller particle size will enhance the access of the enzyme molecules to the protein located within the okara particles. Thereby, a higher degree of proteolysis will be pronounced that leads to the production of bioactive peptides 

<table>
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<tr>
<th>Radical scavenging</th>
<th>Protein</th>
<th>Amino nitrogen</th>
<th>Protein digestibility</th>
<th>Amino acids</th>
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<td>[% db]</td>
<td>content [mg/g]</td>
<td>[%]</td>
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### References


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