Germination changes the isoflavone profile and increases the antioxidant potential of soybean

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Abstract

Isoflavones, which exist in their conjugated or aglycone forms, are well recognized for their potential health benefits. However, isoflavones as aglycone have been regarded as the most bioactive form. In the present study, the profile of isoflavones and their scavenging activity as affected by germination were investigated in several soybean fractions, namely cotyledons, epicotyls, radicles, and hypocotyls. Only aglycones were detected in the radicles from 144 h until 168 h of germination, which makes this component a potential feedstock for studies aiming at isolation, especially of daidzein, which was present in higher concentrations. In terms of total yield and contribution to the total weight of the germinated soybeans, the cotyledons are the best source of aglycones, which was achieved at 144 h of germination. The higher scavenging activity of high-aglycone components from germinated soybeans supports the use of germination to obtain functional foods and/or ingredients with potentially superior bioactivities.

Keywords: Germinated soybeans; Phenolic compounds; β-glycoside isoflavone; Malonylglucoside isoflavones; Isoflavone aglycones; Antiradical activity.

1. Introduction

There are 12 different isoflavones, three conjugated groups [β-glycosides (e.g., daidzin, genistin and glycitin), acetylglucosides (e.g., acetyldaidzin, acetylenistin and acetylglycitin) and malonylglucosides compounds (e.g., malonyldaidzin, malonylgenistin and malonylglycitin)] present in soybeans; the latter representing the aglycones (e.g., daidzein, genistein and glycine) which do not have any glucose moiety linked to the flavonoid structure (Falcão et al., 2018). During soybean seed development in the field, aglycones are synthesized by the phenylpropanoid metabolic pathway and are further stored in vacuoles as β-glucosides and malonylglucosides (Graham, 1991; Kudou et al., 1991). The concentration of isoflavones in soybean seeds may vary because their synthesis and accumulation pathways are affected by multiple environmental variations and genetic factors (Hoeck et al., 2000). Regardless, conjugated isoflavones present in soybeans make a higher contribution when compared to their aglycone
counterparts (Kudou et al., 1991). In a study by Quinhone Júnior and Ida (2015), the content of conjugated isoflavones (β-glucoside plus malonylglycoside form) accounted for 99% of the isoflavones present, while the remaining 1% as aglycones made a minor contribution. Furthermore, isoflavones in the acetylglucoside form are generally absent in soybeans or are present at very low concentrations (Paucar-Menacho et al., 2010). Isoflavones have been found to be absent or make an insignificant contribution to the phenolic content of the soybean seed coat (Kudou et al., 1991; Ribeiro et al., 2006; Yuan et al., 2009), thus demonstrating that the distribution and the identities of phenolics are tissue specific.

Germination may increase the nutritional value of the seed while reducing the amount of undesirable substances such as flatulence-inducing components and antinutrients (Kim et al., 2013; Martin-Cabrejas et al., 2008; Paucar-Menacho et al., 2010). Bioactive compounds such as phytosterols, tocopherols, and flavonoids have also been found to be positively affected by germination (Lee et al., 2007; Paucar-Menacho et al., 2010; Shi et al., 2010; Yuan et al., 2009). Furthermore, soybean germination activates β-glucosidases thus causing changes in the concentration as well as in the structural pattern of isoflavones (Yoshiara et al. 2018; Ribeiro, et al., 2006). Soybean sprouts have already been investigated as potential feedstocks in the development of soymilk and tofu (Murugkar, 2015).

Although several studies have supported the conversion of conjugated isoflavones into their aglycone counterparts during soybean germination (Quinhone Júnior and Ida, 2015), there are still some apparent gaps in the literature, especially when it comes to the effects of such a process in the potential bioactivity of soybean sprouts and, in turn, their industrial products, which may enhance the knowledge of the soybean industry in developing functional foods. Furthermore, most studies focusing on germination have investigated as potential feedstocks in the development of soymilk and tofu (Murugkar, 2015).

2. Material and methods

2.1. Materials and chemicals

Soybean seeds (Glycine max (L.) Merrill), cv. BRS 257, were kindly provided by Embrapa Soja, Londrina, Paraná state, Brazil. All reagents were of analytical or HPLC grade and were used as received.

2.2. Germination process of soybean

The germination process was conducted as described elsewhere (Yoshiara et al., 2011). At each germination time (72, 96, 120, 144 and 168 h), radicles, hypocotyls, cotyledons and epicotyls were manually separated and freeze dried while the coats were discard. The radicle and epicotyls began to germinate after 72 h. All germinated soybean components were ground and stored at −22 °C for further analysis. For comparison purposes, non-germinated soybeans, which were lyophilized, milled, and defatted, were used.

2.3. Extraction of phenolic compounds

Phenolic extraction was conducted with 70% ethanol containing 0.1% acetic acid. In brief, defatted soybeans (100 mg) were extracted with acidified ethanol (4.0 mL) in a RT 500 gyratory shaker (Marconi, Piracicaba, Brazil) for 20 min. After centrifugation at 2000 × g (Cientec-Piracicaba, Brazil), the upper layer was collected and extraction was repeated twice. The combined supernatant was evaporated to remove the organic solvent and then stored at −26 °C until used for further analyses.

2.4. Identification and quantification of isoflavones by high performance liquid chromatography (HPLC)

The HPLC instrument (Shimadzu, LC-10AT VP, Kyoto, Japan) coupled to a diode array detector (SPD-M10A VP) set at 260 nm was employed for identification and quantification purposes. The oven (CTO-10AS VP) temperature was kept constant (25 °C). Phenolic extracts were filtered through nylon membranes (0.20 µm) and injected (20 µL) in a reversed-phase ODS (M) C18 column (15 cm × 4 mm × 5 µm). Acidified water, prepared with glacial acetic acid (pH 3.0), was the mobile phase A and the mobile phase B consisted of acetonitrile. The elution gradient used was as follows: 0 min, 100% A; 25 min, 45% A; and 27 min, 100% B. The initial condition was re-established at 40 min. The total run took 40 min and the flow rate was 1 mL min⁻¹. The identification and quantification of the β-glycosides and aglycone isoflavones were performed by using external standard curves for daidzin, glycitin, genistin, daidzein, glycine and genistein. The quantification of malonylglycoside isoflavones (e.g., malonyldaidzin, malonylglycitin and malonylgenistin) was made based on the standard curves of the corresponding β-glycosides isoflavones (i.e., daidzin, glycitin and genistin, respectively) using the similarity of the molnar extinction coefficients as described by Coward et al. (1993). The concentrations are expressed as mg of isoflavone 100g⁻¹ dry weight of sample.

2.5. Total phenolic content (TPC)

The Folin-Ciocalteau method (Swain and Hillis, 1959), described by de Camargo et al. (2015), was slightly modified to estimate the TPC. Phenolic extracts (0.50 mL) of a diluted solution of phenol reagent (10%) and 2.0 mL of sodium carbonate saturated solution (7.5%, m/v) were mixed thoroughly and incubated in a water bath at 50 °C for 5 min. The absorbance was read at 760 nm using a spectrophotometer (Biochrom Libra S22, Cambridge, England). The results were expressed as mg gallic acid equivalents 100 g⁻¹ dry weight of defatted sample.

2.6. DPPH radical scavenging activity

The ability of the phenolic extracts in scavenging DPPH radical was evaluated according to Brand-Williams et al. (1995). The ex-
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Tracts (50 μL) were mixed with 1 mL of acetate buffer (0.1 M, pH 5.5), 1 mL of ethanol and 0.5 mL of DPPH solution (250 μM). After 30 min standing in the dark, the absorbance was read 517 nm using a spectrophotometer (Biochrom Libra S22, Cambridge, England). The DPPH radical scavenging activity of the extracts was calculated using the following equation.

\[
\text{DPPH radical scavenging activity (\%) = \left( \frac{\text{Abs}_{\text{control}} - \text{Abs}_{\text{sample}}}{\text{Abs}_{\text{control}}} \right) \times 100}
\]

where \(\text{Abs}_{\text{control}}\) is the absorbance of DPPH radical + ethanol; and \(\text{Abs}_{\text{sample}}\) the absorbance of DPPH radical + phenolic extract or trolox. The results were expressed as \(\mu\)mol of trolox equivalents g\(^{-1}\) dry weight of sample.

### 2.7. ABTS radical cation scavenging activity

The ABTS assay (Re et al., 1999) was performed with minor changes in the method described by de Camargo et al (2015). The ABTS radical cation, prepared in 100 mM phosphate buffer saline (PBS) solution (pH7.4, 0.15 M sodium chloride), was generated by oxidation with potassium persulphate. The stock solution consisted of potassium persulphate (2.45 mM) and ABTS (7 mM) in PBS. The oxidation was conducted in the dark for 16 h. A working solution of ABTS radical cation was prepared by diluting the stock solution in PBS to reach an absorbance value of 0.700 ± 0.020 at 730 nm. Phenolic extracts (10 μL) were added to 4 mL of ABTS radical cation solution and the absorbance was read at 730 nm after 6 min using a spectrophotometer (Biochrom Libra S22, Cambridge, England). ABTS radical scavenging activity was calculated using the following equation.

\[
\text{ABTS radical scavenging activity (\%) = \left( \frac{\text{Abs}_{\text{control}} - \text{Abs}_{\text{sample}}}{\text{Abs}_{\text{control}}} \right) \times 100}
\]

where \(\text{Abs}_{\text{control}}\) is the absorbance of ABTS radical cation + ethanol; and \(\text{Abs}_{\text{sample}}\) the absorbance of ABTS radical cation + phenolic extract or trolox. The results were expressed as \(\mu\)mol of trolox equivalents g\(^{-1}\) dry weight of sample.

### 2.8. Experimental design and statistical analysis

A randomized complete block design with three replications was used. The means were compared by Tukey’s test \((p < 0.05)\) and Statistica 8.0 software (StatSoft Inc., Tulsa, OK, USA).

### 3. Results and discussion

#### 3.1. Germination-induced changes of isoflavones

The amounts of different isoflavone forms from cotyledons, epicotyls, radicles and hypocotyls of germinated soybeans are shown in Table 1. Irrespective of the germination period, acetylglycoside isoflavones were absent in all components of the test material.
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Table 2. Total isoflavone concentration and contribution of conjugated and aglycone isoflavones of different germinated soybean components

| GT (h) | Radicles | Hypocotyls | Epicotyls | Cotyledons |
|--------|----------|------------|-----------|------------|
| 0      | Total (mg 100 g⁻¹ dw) | * | 2330.58 ± 135.59A⁴ | * | 356.17 ± 2.78B⁵ |
|        | Conjugated (%) | * | 98.36 | * | 99.37 |
|        | Aglycones (%) | * | 1.64 | * | 0.63 |
| 72     | Total (mg 100 g⁻¹ dw) | 557.36 ± 7.92A² | 406.20 ± 8.72C⁶ | nd | 436.25 ± 2.54B⁶ |
|        | Conjugated (%) | 90.09 | 62.26 | nd | 91.78 |
|        | Aglycones (%) | 9.91 | 37.74 | nd | 8.22 |
| 96     | Total (mg 100 g⁻¹ dw) | 519.51 ± 7.87A⁵ | 114.03 ± 0.04C⁷ | nd | 388.65 ± 0.59B⁴ |
|        | Conjugated (%) | 90.10 | 25.75 | nd | 81.92 |
|        | Aglycones (%) | 9.90 | 74.25 | nd | 18.08 |
| 120    | Total (mg 100 g⁻¹ dw) | 559.00 ± 7.87A⁵ | 58.82 ± 0.52C⁸ | 34.75 ± 0.20D⁹ | 402.37 ± 0.13B⁵ |
|        | Conjugated (%) | 91.48 | 52.00 | 0.00 | 77.78 |
|        | Aglycones (%) | 8.52 | 48.00 | 100.00 | 22.22 |
| 144    | Total (mg 100 g⁻¹ dw) | 41.49 ± 0.13C⁹ | 42.80 ± 0.85C⁹ | 211.7 ± 5.77A⁰ | 164.31 ± 2.48B¹ |
|        | Conjugated (%) | 0.00 | 0.00 | 83.84 | 8.95 |
|        | Aglycones (%) | 100.00 | 100.00 | 16.16 | 91.05 |
| 168    | Total (mg 100 g⁻¹ dw) | 40.47 ± 0.04D⁹ | 75.32 ± 1.97C⁹ | 253.35 ± 3.80B⁸ | 613.59 ± 0.82A² |
|        | Conjugated (%) | 0.00 | 26.82 | 62.24 | 88.95 |
|        | Aglycones (%) | 100.00 | 73.18 | 37.76 | 11.05 |

*The radicles and epicotyls were absent at time zero. Total isoflavone represent mean values ± standard deviation (n = 3). Values followed by different capital letters within a row show differences among different germinated soybean components (p < 0.05). Values followed by different lowercase letters within a column show differences among different germination times (p < 0.05). Abbreviations: GT, germination time; and nd, not detected.

which was expected since acetylglycoside isoflavones are generated upon heating treatment (Kudou et al., 1991). This is also in agreement with a previous study (Falcão et al., 2018) in which these isoflavone forms were not detected. The concentration of total isoflavones in hypocotyls of non-germinated soybeans was higher than that of cotyledons, while the remaining soybean components, namely epicotyls and radicles, only appeared during the germination process. Regardless of the germination time, cotyledons made the highest contribution to the total weight of the germinated soybeans (Yoshiara et al., 2011). According to the aforementioned study, at the latest stages of germination, the contribution of each fraction was in the order cotyledons > hypocotyls > radicle > epicotyl, and the latter one made only a minor contribution, ranging from 0.4 to 2.3%. The variation in the concentration of different isoflavone forms is helpful to better understand the biochemistry involved in the germination process. However, in terms of application in functional foods and/or functional ingredients, the changes taking place in the cotyledon are most important. The concentration of aglycones in the cotyledon increased up to 144 h of germination (Table 1), after which a decrease was noted. Because of the higher number of hydroxyl groups, the aglycones may act better as antioxidants. Therefore, such decreasing trend may be related to the oxidation of these phenolic antioxidants. In general, hypocotyls from germinated soybeans also showed higher amounts of aglycones compared to their non-germinated counterparts. The radicles and epicotyls started to develop later and, at 72 h of germination, 9.9% of the isoflavones present in the radicles were in the aglycone form (Table 2). However, at the end of the germination period, only aglycones were detected in this part of the plant material.

In general, the higher yield of isoflavones in the aglycone form was detected at 144 h of germination: 100% for radicles and hypocotyls and 91.05% for cotyledon, except the epicotyls, that presented only 16.2% of aglycones. However, as mentioned earlier, the epicotyls make a minor contribution to the weight of the germinated soybean (Yoshiara et al., 2011). Therefore, 144 h of germination appears to be the best time to obtain high-aglycone soybean feedstocks (Table 2), for which a total characterization of the isoflavones was undertaken (Table 3). Malonylgenistin and daidzin were the predominant malonylglycoside and β-glycosides isoflavones in non-germinated soybeans, respectively, while no significant differences were found between both aglycones (daidzein and genistein) detected. The concentration of both was minor (0.70–0.90 mg 100g⁻¹) compared to the total isoflavones (274.0 ± 0.10 mg 100g⁻¹). Furthermore, glycitein was not detected in non-germinated soybeans, which is in accordance with a recent report by Falcão et al. (2018). Soybean components subjected to 144 h of germination (Table 3) showed the following levels of total isoflavones: epicotyl (211.7 ± 4.08 mg 100g⁻¹), cotyledon (164.31 ± 1.76 mg 100g⁻¹), hypocotyl (42.80 ± 0.60 mg 100g⁻¹) and radicle (41.49 ± 0.09 mg 100g⁻¹) and any presence of conjugated isoflavones was detected in hypocotyl and radicle. Furthermore, different from hypocotyls, the amount of total isoflavones was quite constant in the radicle, which showed 100% aglycones until the end of the germination period (Table 2), thus making this component the best feedstock for studies aiming at isolation, especially of daidzein, which was present in higher concentrations.

Germinated soybean epicotyl was the only component that contained all three aglycone forms. Glycitin, which was present in non-germinated soybeans, was not detected in the germinated samples. In contrast, the aglycone glycitin was detected only...
in epicotyls of germinated soybeans. According to Falcão et al. (2018), there is a lack of evidence exists for hydrolysis of conjugates of glycitin upon ultrasound treatment (up to 25 min). A low specificity of the endogenous β-glucosidase towards glycinin and malonylglycitin has been reported (Ismael and Hayes, 2005). However, the decrease in the concentration of conjugated isoflavones (glycitin and malonylglycitin) with a parallel increase in their corresponding free aglycones has been reported (Yoshiara et al., 2018). Enzyme activity induced changes upon germination have also been reported by other authors (Wanasundara et al., 1999). Furthermore, the presence of insoluble-bound glycitin has also been reported for soybeans (Peng et al., 2017). According to Yeo and Shahidi (2015), the changes induced by germination are not limited to soluble phenolics as those linked to the cell wall of plant material (insoluble-bound phenolics) have also been affected.

Isoflavones have been associated with disease risk reduction or prevention of various ailments, such as breast cancer and prostate cancer (Liggins et al., 2000), osteoporosis, menopause symptoms (Levis et al., 2010), cardiovascular disease (Liu et al., 2017; Rimbach et al., 2008), memory improvement (Lephart et al., 2002), as well as estrogenic and antioxidant activity (Ma et al., 2010). Isoflavones may also inhibit adipogenesis and lipogenesis via interaction with several transcription factors and upstream signaling molecules (Wang et al., 2017), therefore, may render potential anti-obesity effects. Furthermore, according to a meta-analysis, consumption of dietary flavonoids is associated with a reduced risk of type 2 diabetes (Liu et al., 2014). Soybean aglycones are the most biologically active forms under physiological conditions and have been regarded as the best form for absorption in the human body (Falcão et al., 2018). This contribution supports the use of germination as an alternate process to promote the bioconversion of conjugated isoflavones into their corresponding aglycones. Finally, the results presented here (Table 3) demonstrate that germinated soybean cotyledons constitute a rich source of aglycones as their concentration of 149.6 mg 100g−1 is much higher than that present in the hypocotyls (42.8 mg 100g−1) and radicles (4.19 mg 100g−1) thus demonstrating that different parts of the plant should be evaluated when germination is carried out.

### 3.2. Total phenolic content (TPC)

The TPC of germinated and non-germinated soybeans is presented in Figure 1. The various components of germinated and non-germinated soybean differed from each other (p < 0.05). The cotyledon (570.88 ± 28.80 GAE 100 g−1 dry weight of sample) showed the highest TPC, which was followed by radicle (398.43 ± 18.77 GAE 100 g−1 dry weight of sample), hypocotyl (227.93 ± 1.00 GAE 100 g−1 dry weight of sample) and epicotyl (308.93 ± 3.77 GAE 100 g−1 dry weight of sample). A recent report has discussed the use of colorimetric methods in the screening of different sources of food phenolic antioxidants (Granato et al., 2018). A significant positive correlation existed between TPC and daidzein (r = 0.9030, p = 0.036) in the present study. However, no correlation existed between TPC and both total isoflavones and genistein (p > 0.05) as evaluated

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**Table 3. The contents of isoflavones (mg 100 g−1 dw) of non-germinated soybean and samples germinated for 144 h at 35 °C in the presence of light**

| Compound     | NGS  | Germinated soybean components | Cotyledons | Epicotyls | Hypocotyls | Radicles |
|--------------|------|-------------------------------|------------|-----------|------------|----------|
|              |      |                               |            |           |            |          |
| Daidzein     | 0.90 ± 0.03 f | 89.32 ± 0.89 a | 15.56 ± 0.12 c | 33.11 ± 0.41 a | 35.71 ± 0.06 a |
| Glycitein    | nd   | nd                            | 12.94 ± 0.44 d | nd         | nd         |
| Genistein    | 0.72 ± 0.1 f  | 60.28 ± 0.56 b | 5.73 ± 0.01 f  | 9.69 ± 0.20 b | 5.78 ± 0.03 b |
| Daidzin      | 27.04 ± 0.32 c | nd                    | 8.59 ± 0.01 e  | nd         | nd         |
| Glycitin     | 8.61 ± 0.17 a | nd                    | nd         | nd         | nd         |
| Genistin     | 21.32 ± 0.22 d | 14.70 ± 0.31 c | 7.83 ± 1.93 f  | nd         | nd         |
| Malonyldaidzin| 77.04 ± 0.85 b | nd                    | 69.69 ± 0.00 b | nd         | nd         |
| Malonylglycitin| 21.88 ± 0.14 d | nd                    | 2.79 ± 0.16 e  | nd         | nd         |
| Malonylgenistin| 116.5 ± 0.03 b | nd                    | 88.59 ± 2.31 a | nd         | nd         |
| Total        | 274.1 ± 0.10 | 164.3 ± 1.76 | 211.7 ± 4.08 | 42.80 ± 0.60 | 41.49 ± 0.09 |

Data represent mean values ± standard deviation (n = 3). Values followed by the same letters within a column are not significantly different (p > 0.05). Abbreviations: NGS, non-germinated soybeans; and nd, not detected. Acetyldaidzin, acetylglycitin, and acetylgenistin were absent in all samples.
by HPLC. This suggests that the reaction taking place during the Folin-Ciocalteu assay is dependent on the structure of individual compounds. Furthermore, all germinated soybean components had a higher TPC compared to non-germinated soybeans (204.73 ± 5.08 mg GAE 100 g⁻¹ dry weight of sample). According to the data reported for several food legumes (Xu and Chang, 2007), the TPC of non-germinated soybeans was lower than that of lentils (244.00 mg GAE 100 g⁻¹) and black beans (320.00 mg GAE 100 g⁻¹) but it was higher than that of green peas (134.00 mg GAE 100 g⁻¹) and chickpeas (154.00 mg GAE 100 g⁻¹). However, except for black beans, all components of germinated soybeans showed higher TPC than the aforementioned legume seeds. Therefore, the results of this study suggest that the germination process may improve the TPC thus serving better as a source of phenolic compounds.

### 3.3. Scavenging activity as affected by germination

Free radicals may induce changes in DNA, oxidize LDL-cholesterol, inactivate enzymes, and cause cell damage, which are important biomarkers related to higher incidences of several chronic ailments (Amarowicz and Pegg, 2017; Fernandes et al., 2011; Peng et al., 2015; Perera et al., 2018). In food systems, oxidation caused by free radicals may induce sensory changes, decrease nutritional quality, including depletion of vitamins (e.g. vitamins C and E) and affect polyunsaturated fatty acids (Ramírez-Cahero and Valdivia-López, 2018; Sartori et al., 2018; Shahidi and de Camargo, 2016). Bueno-Borges et al. (2017) carried out several antioxidant assays (e.g. ABTS, DPPH, ORAC, and Rancimat) and have suggested superior antioxidant properties of soybean isoflavones compared to that of tocopherols.

The antioxidant properties of food phenolics may be explained by single electron transfer (SET) or hydrogen atom transfer (HAT) (de Camargo et al., 2017; Leopoldini et al., 2004). While both electron and hydrogen transfer may be contemplated in DPPH assay, the ABTS assay is based on electron transfer reactions. Furthermore, ABTS is used to evaluate radical scavenging activity of both hydrophilic and lipophilic compounds. Hence, in the current study, both methods were used to evaluate the effects of germination in different soybeans components.

In general, germinated soybean components showed higher antioxidant activity compared to non-germinated samples. In accordance to TPC results, the cotyledons exhibited the greatest ability in scavenging both radicals (Table 4). While cotyledons had 2.4-fold higher scavenging capacity towards ABTS radical cation, a lower difference was noted against DPPH radical (12% higher) which may due to different mechanisms that are operative in each assay/reaction. A high positive correlation existed between the concentration of genistein (r = 0.9088, p = 0.033) and the ability of phenolic extracts in neutralizing the ABTS radical cation, which was not detected between total isoflavones and daidzein (p > 0.05) as evaluated by HPLC. These findings suggest the critical role of genistein as an antioxidant compound. In addition, no correlations were found between these aglycones and DPPH results. Therefore, considering the great difference for both assays and the lack of correlation between any aglycone and DPPH radical scavenging activity, it is possible to suggest that genistein may act through SET rather than via HAT.

### 4. Conclusions

Germination positively influenced the isoflavone profile of soybean. The cotyledons made the highest contribution to the total weight of the germinated soybean cotyledons. Hence, after germination for 144 h, the cotyledons rendered a higher yield of aglycones (genistein and daidzein). Furthermore, only aglycones were detected in the radicles from 144 h until 168 h of germination, thus making this component a potential feedstock in developing functional ingredients and/or processing products as well as in future studies aiming at isolation, especially of daidzein, which was present in higher concentrations. Finally, this contribution lends support on the use of germination to improve the antioxidant potential of soybeans.

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### Conflict of interest

The authors declare no conflict of interest.

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