



TITLE:

# Enhancement of developmentally regulated daidzein secretion from soybean roots in field conditions as compared with hydroponic culture

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2       **Enhancement of developmentally regulated daidzein secretion from**  
3       **soybean roots in field conditions as compared with hydroponic culture**

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21 **Enhancement of developmentally regulated daidzein secretion from**  
22 **soybean roots in field conditions as compared with hydroponic culture**

23

24 Analyses of metabolite secretions by field-grown plants remain scarce. We  
25 analyzed daidzein secretion by field-grown soybean. Daidzein secretion was  
26 higher during early vegetative stages than reproductive stages, a trend that was  
27 also seen for hydroponically grown soybean. Daidzein secretion was up to  
28 10,000-fold higher under field conditions than hydroponic conditions, leading to  
29 a more accurate simulation of rhizosphere daidzein content.

30

31 **Keywords: Daidzein; Rhizosphere; Simulation; Soybean**

32

33 Plant specialized metabolites (PSMs) play important roles in the rhizosphere for  
34 modulation of symbiotic interactions (e.g., repelling pests and pathogens and shaping

35 microbiota), thereby promoting plant growth and improving crop production [1–3].

36 Flavonoids are a group of PSMs and consist of more than 8,000 compounds [4]. These

37 molecules function as regulators of auxin transport and reactive oxygen species and

38 protect against damage caused by ultraviolet (UV) light exposure. In legumes,

39 flavonoids are secreted from the roots to exert functions in rhizosphere plant-microbe

40 interactions, such as those necessary for defense and symbiosis [5–7].

41       Isoflavones are a subfamily of flavonoids and are found mainly in legumes [8].

42 In the rhizosphere, isoflavones such as daidzein and genistein of soybean (*Glycine max*)

43 and formononetin-7-*O*-(6"-*O*-malonylglycoside) of alfalfa (*Medicago sativa*) induce

44 *nod* genes for initiation of the nodulation process [9, 10]. In particular, daidzein was

45 recently shown to be involved in the modulation of rhizosphere bacterial communities

46 in soybean, where this compound increased the relative abundance of the

47 Comamonadaceae family of bacteria [11].

48           The secretion of metabolites from roots is a crucial process influencing  
49 interactions in the rhizosphere. Daidzein is the major isoflavone secreted into  
50 hydroponic media, the concentration of which is higher during the soybean vegetative  
51 stage than the reproductive stage [12]. Daidzein is relatively stable in soil, with a half-  
52 life of about seven days, enabling the estimation of daidzein contents in the rhizosphere  
53 based on the amount secreted in hydroponic cultures [13]. Whereas sorption filters or  
54 glass beads have been used to collect and analyze various metabolites in the rhizosphere  
55 [14, 15], the direct measurement of secreted metabolites is technically challenging,  
56 especially for field-grown plants [16]. In this study, we used the cellulose acetate  
57 membrane method utilized in hydroponic culturing to analyze flavonoid secretion for  
58 direct measurement of the amount of daidzein secreted by field-grown soybean plants  
59 during the stages of growth.

60           All chemicals used in this study were obtained from either Wako Pure  
61 Chemical Industries (Osaka, Japan) or Nacalai Tesque (Kyoto, Japan) unless otherwise

62 stated.

63           The field experiments were conducted at Kyoto University of Advanced  
64 Science, Kameoka, Kyoto, Japan (coordinates: 34°99'38"N, 135°55'14"E). Soybean  
65 seeds (“Tambaguro”) were sown on May 31, 2019. The plants were irrigated as needed,  
66 and emerging weeds were manually removed weekly. No apparent symptoms of  
67 pathogen infection were observed, and pesticides were not used. Root samples and root  
68 exudates were collected on June 14 (V1 stage), July 3 (V5 stage), July 22 (V9 stage),  
69 August 14 (R2 stage), September 4 (R4 stage), and October 2 (R6 stage) of 2019 [17].  
70 The soil around the lateral roots was partially removed with a shovel. The lateral roots  
71 were rinsed with tap water and pinched between a cellulose acetate filter (Advantec,  
72 Tokyo, Japan) using a hairpin (Fig. 1A) and then covered with soil. Additionally,  
73 cellulose acetate filters were placed in the bulk soil as a control. The cellulose acetate  
74 filters were held in the soil for 2 h, and then the filters and root tissues were collected.  
75 All samples were transferred to the laboratory in a cool container (0–10°C) within 2 h of

76 collection. The root samples were stored in pure water prior to fresh weight  
77 measurement. The roots and fully expanded leaves were taken from 2-week-old soybean  
78 seedlings (VE stage) for the quantification of isoflavones as described previously [13].  
79 Rhizosphere soil was obtained from seven plants using sterile brushes and combined  
80 into one sample as described previously [18]. The samples were immediately frozen in  
81 dry ice and transferred to the laboratory for storage at  $-80^{\circ}\text{C}$ . The bulk soil was  
82 sampled at least 20 cm away from the plant.

83           The extraction of daidzein was performed as previously described [13, 19]. The  
84 cellulose acetate filters were rinsed with tap water, and the compounds were extracted  
85 twice using 1 ml methanol with shaking on a Labo shaker BC-730 (Bio craft, Tokyo,  
86 Japan) for 5 min each time. The combined supernatant from each sample was dried  
87 under a nitrogen stream at  $50^{\circ}\text{C}$ , dissolved in 150  $\mu\text{l}$  of methanol, and filtered through a  
88 Minisart RC4 syringe filter (Sartorius, Gottingen, Germany) for LC-MS (Liquid  
89 chromatography–mass spectrometry) analysis. The exudates were analyzed by UPLC-



90 MS on an ACQUITY UPLC system (Waters Corporation) coupled with Xevo TQD.

91 The LC was performed by injecting a 2  $\mu$ l sample onto an ACQUITY UPLC BEH C18

92 column (2.1 mm  $\times$  50 mm, 1.7  $\mu$ m; Waters Corporation) at 40  $^{\circ}$ C. The LC mobile phase

93 consisted of (A) water containing 0.1% (v/v) formic acid and (B) acetonitrile. The

94 gradient program was linear over the range of 10%–35% B, 0–1 min; linear 35%–85%

95 B, 1–11 min; isocratic 85% B, 11–11.1 min; isocratic at 100% B, 11.1–15.5 min; and

96 isocratic at 10% B, 15.5–20 min. The flow rate was 0.2 ml min<sup>-1</sup>. Isoflavones were

97 detected at 260 nm. The contents of daidzein were estimated from the peak areas in

98 comparison with calibration curves constructed using known concentrations of the

99 authentic compound.

100 The extraction of isoflavones was performed as described [13]. The frozen

101 tissues were pulverized in liquid nitrogen using a mortar and pestle and then freeze-

102 dried. The tissues were extracted in 80% methanol at 60 $^{\circ}$ C for 1 h, followed by

103 centrifugation at 12,000 $\times$ g for 5 min to remove debris. The supernatant was filtered

104 through a Minisart RC4 syringe filter (Sartorius). Soil samples (1 g) were extracted in  
105 500  $\mu$ l of methanol at 50°C three times (10 min each) and centrifuged at 4,800 rpm for 5  
106 min. The combined supernatant from each sample was dried under a nitrogen stream at  
107 50°C and redissolved in 150  $\mu$ l methanol. Isoflavones were analyzed by LC-MS/MS as  
108 described [20].

109           The movement of daidzein secreted by a single cylindrical root was simulated  
110 using a two-dimensional asymmetric system. The equations, model domains, and  
111 relevant initial/boundary conditions were previously described [11]. The daidzein  
112 secretion rate at the root surface was assumed to be constant (1.06 nmol m<sup>2</sup> s<sup>-1</sup>), based  
113 on the daidzein extraction for the roots sampled on June 14 (V1 stage). The simulation  
114 period was set at 14 days with a 0.1-day time interval. The parameters used in this study  
115 were summarized in Table S1. A cylinder of soil with a diameter of 20 cm and a depth  
116 of 20 cm with a single root of diameter 2 mm and length 10 cm in the center was set as  
117 a model domain for the simulation. Root length and diameter were assumed to be

118 constant for 14 days.

119           The isoflavones secreted from field-grown soybean were analyzed at three  
120 vegetative growth stages V2, V5, and V8, corresponding to 2, 5, and 8 weeks after  
121 sowing, respectively. Moreover, samples from three reproductive growth stages R2, R4,  
122 and R6, respectively corresponding to 12, 15, and 19 weeks after sowing, were  
123 analyzed. Of all the detected isoflavones collected using cellulose acetate membranes  
124 that adsorb flavonoid aglycones [13,19], only daidzein was identified at each growth  
125 stage (Fig. 1B). The amount of secreted daidzein changed over the growth stages and  
126 peaked at V5, whereas it was constant over the reproductive stages. The trend of  
127 daidzein secretion was similar to that of hydroponically grown soybean. In contrast,  
128 field-grown soybean secreted up to a 10,000-fold higher amount of daidzein than  
129 hydroponically grown soybean (about 36 fmol mg FW<sup>-1</sup> day<sup>-1</sup> at V3) [12]. While the  
130 secretion from whole roots was analyzed in the hydroponic culture media, the secretion  
131 from the field-grown soybean was analyzed using a 3 cm root-tip section. The

132 possibility that partial soil removal induced the isoflavone biosynthesis and increased  
133 the isoflavone levels within 2 h is probably small because it is suggested to take more  
134 than 3 h for the roots to accumulate isoflavones after gene induction in *Arabidopsis*  
135 *thaliana* and soybean [20, 21]. The difference in the magnitude of secretion is,  
136 therefore, presumably attributable to environmental conditions, i.e., sterile hydroponics  
137 vs. non-sterile field environments. The contents of isoflavones in the root tissue at  
138 steady-state under the field-grown conditions were similar to those under hydroponic  
139 conditions [12, 13], suggesting that both the isoflavone synthesis and secretions are  
140 remarkably enhanced in the rhizosphere, probably due to the presence of various  
141 microorganisms.

142       The spatiotemporal distribution of metabolites in the rhizosphere is of particular  
143 importance for deciphering their functions in inter-organismal interactions such as  
144 chemotaxis response and *nod* gene induction, which are concentration-dependent [9, 22,  
145 23]; however, the distribution of PSMs remains largely unknown [24]. In our previous

146 study, we simulated the spatiotemporal distribution of daidzein in field soil based on the  
147 advection–diffusion equation [11], and we showed that daidzein distribution was limited  
148 to within a few millimeters from the root surface [11]. To further refine the simulation  
149 of daidzein distribution in the field, we applied the secreted rate of daidzein under field  
150 conditions. The distribution of daidzein was also limited to within a few millimeters  
151 from the root surface, similar to findings from the previous simulation [11]. Limited  
152 daidzein distribution withn a few millimeters is likely due to the adsorption of daidzein  
153 by the soil. In this simulation, the average daidzein content within 1 or 3 mm soils from  
154 root surface was around 0.8 and 0.5 nmol g soil<sup>-1</sup>, respectively (Fig. 2A). This  
155 concentration was within the range to induce *nod* genes in *Bradyrhizobium japonicum*,  
156 which is reported to be more than 0.1 μM [9, 25]. The isoflavone contents in the  
157 rhizosphere and plant tissues were measured in 2-week-old soybean seedlings to  
158 validate the results of this simulation. Malonylgenistin was the most predominant  
159 isoflavone in the leaves at this stage, while malonyldaidzin and daidzein were

160 accumulated in the roots (Fig. 2B). Rhizosphere soil was sampled from less than 3 mm  
161 layer from the root surface. In the rhizosphere soil, daidzein was the most abundant  
162 isoflavone, and the content was about 5 nmol g soil<sup>-1</sup> (Fig. 2C). Collectively,  
163 rhizosphere modeling based on the amount secreted by field-grown soybean led to a  
164 more accurate simulation of daidzein distribution than our previous simulation, i.e.  
165 daidzein distribution at physiologically relevant concentrations is limited to within a  
166 few millimeters from root surface.

167         Despite the importance of PSMs in the rhizosphere, our current knowledge of  
168 the dynamics in the rhizosphere of field-grown plants is still preliminary. The dynamics  
169 between proteins, metabolites, and ions in the rhizosphere have been analyzed mostly  
170 using the rhizobox [23], but they should be examined in field-grown plants as well. In  
171 this study, we showed that the secretion of daidzein by field-grown soybean followed  
172 the same trends in terms of developmental regulation, but the amount was much higher  
173 than in hydroponic condition, leading to the accurate estimation of daidzein distribution

174 in the rhizosphere. The rhizosphere microbiome affects the secretion of metabolites  
175 from roots [26]; therefore, we presume that the rhizosphere microbiome enhanced  
176 daidzein secretion in the field, in addition to the effects of other both biotic and abiotic  
177 stresses under field conditions. It is of particular importance to analyze the secretion of  
178 PSMs in the rhizosphere of field-grown plants under various conditions and to integrate  
179 the distribution of PSMs and the structure and functions of the microbiota in future  
180 studies.

181

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185

186 Author Contributions: A.S. conceived and designed the research; H.T., K.Y., and A.S.  
187 supervised the experiments; M.T., F.O., and M.N. conducted plant sampling and LC-  
188 MS/MS analysis; S.H. conducted the simulation; M.T. and A.S. wrote the article with

189 contributions of all authors; A.S. agrees to serve as the author responsible for contact  
190 and ensuring communication.

191

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193

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197

## 198 **Figure Legends**

199 Fig. 1 (A) Cellulose acetate membrane used to collect root exudates. Tips of lateral  
200 roots were washed with pure water and pinched in a cellulose acetate membrane, which  
201 was then covered with soil. The site of analysis was marked with a piece of white paper.  
202 (B) Root exudation of daidzein throughout soybean growth stages. Amount per root  
203 fresh weight of daidzein in root exudates ( $n \geq 9$ ). Significant differences ( $P < 0.05$ ;  
204 Tukey–Kramer test) are indicated with various letters. Root samples and root exudates  
205 were collected at three vegetative stages (V) and three reproductive stages (R).



206

207 Fig. 2 Simulation of daidzein distribution in soil and isoflavone contents in the  
208 rhizosphere. (A) Simulated daidzein distribution from 0 to 14 days in soil. The rate of  
209 daidzein secretion from roots was assumed to be constant at each depth, and the  
210 distribution at the middle of root at a depth of 5 mm was displayed in radial direction. It  
211 is noted that vertical distribution of daidzein was not obtained in this simulation. (B)  
212 Contents of isoflavones in leaves and roots (n = 3). (C) Contents of isoflavones in bulk  
213 and rhizosphere soils at VE stage (n = 3).

214

## 215 **Supplementary Material**

216 Supplementary Table 1. Parameters used in this study

217

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