

Comparative Analyses of Flavonoids for *nod* Gene Induction in *Bradyrhizobium japonicum* USDA110

RYU, JI-YOUNG¹ AND HOR-GIL HUR^{1*}

¹International Environmental Research Center, UNU-GIST Joint Program on Science and Technology for Sustainability, and Department of Environmental Science and Engineering, Gwangju Institute of Science and Technology, Gwangju, Republic of Korea

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Abstract Using the *nodY::lacZ* fusion system in *Bradyrhizobium japonicum* USDA110, 22 flavonoids, which have structurally different features, were tested to define the role of the substituted functional groups as an inducer or inhibitor for the *nod* gene expression. A functional group of 4'-OH on the B-ring and the double bond between 2-C and 3-C on the C ring were required to induce the *nod* gene expression in *B. japonicum* USDA110. In the case of isoflavones, the 4'-methoxyl group, which blocks the open 4'-OH functional group, did not significantly lower inducing activity, as compared with isoflavones with 4'-OH. However, all flavonols tested, which have a 3-OH functional group on the C-ring, did not induce, but inhibited the *nod* gene expression. Flavone, 7-hydroxyflavone, and kaempferol (5,7,4'-trihydroxyflavonol) at 1 μ M concentration significantly inhibited the *nod* gene expression induced by 7,4'-dihydroxyflavone. However, 7-hydroxy-4'-methoxyflavone at 1 μ M concentration showed a synergistic effect with genistein and 7,4'-dihydroxyflavone on the induction activity.

Key words: Nodulation, *Bradyrhizobium*, isoflavonoid, flavonoid, induction

Rhizobia can make nitrogen-fixing nodules on the host plants in the sequential consequences of chemical communication between both partners [7]. The mutual interactions between the symbiotic partners determine the early stage of nodulation processes prior to infection, which are fairly specific [3, 4, 13, 18, 23]. The chemical compounds responsible for the early stage signals to rhizobia have been known to be flavonoids [14, 18, 19] which are synthesized through the phenylpropanoid pathway present in the host plants.

Once released into the root zone, flavonoids serve as chemoattractants and regulate expression of nodulation (*nod*) genes of symbiotic rhizobia at the transcriptional level [14, 16, 21]. For the expression of *nod* genes, regulatory protein NodD, which is constitutively expressed in the rhizobia [13], is necessary to control the expression of the inducible *nod* genes via interaction with the *cis*-acting promoter element *nod* box. When NodD recognizes appropriate flavonoids on its C-terminal residue, deformation of the DNA helix in the vicinity of the *nod* box occurs, and transcription of adjacent *nod* genes starts [5, 9, 16]. As consequences of the expression of *nod* genes, vast biochemical reactions, including synthesis of Nod factors, are carried out for the formation of nodules on the host plants [9, 11, 17].

Since Redmond *et al.* [18] elucidated that flavones induce the expression of nodulation genes in *Rhizobium*, an extensive variety of additional compounds such as anthocyanin derivatives, betaines, trigonelline, and stachydrine have been revealed as inducers [1, 8, 15]. In addition, isoflavones, genistein, daidzein, and their glycoside derivatives have been found to work as *nod* gene inducers in *Bradyrhizobium japonicum* [1, 9, 22]. Using a *nodY::lacZ* fusion system in *B. japonicum*, Cunningham *et al.* [2] tested over 1,000 compounds for their induction or inhibition activity to regulate *nod* gene expression. The compounds tested included representatives of coumarines, benzoic acids, benzophenones, chalcones, aurones, xanthenes, isoflavones, and flavones. Although extensive studies revealed that a rather broad range of derivatives of plant exudates influence induction or inhibition of *nod* gene expression in the corresponding host bacteria, there is little information available about the role of a chemical functional group(s) in the flavonoids for the *nod* gene induction. In this study, 22 representative flavonoids, which contain a hydroxyl or methoxyl group(s) differently located on the carbon skeleton

*Corresponding author
Phone: 82-62-970-2437; Fax: 82-62-970-2434;
E-mail: hghur@gist.ac.kr

of the compounds, were selected to test their induction activity, using a *nodY::lacZ* fusion system in *B. japonicum* USDA110.

MATERIALS AND METHODS

The bacterial strain was *B. japonicum* USDA110 (pZB32) [1], which has the *nod*-box and *nodY* gene fused with the *lacZ* gene on the plasmid. All the flavonoids were purchased from Indofine Chemical Co. (Somerville, NJ, U.S.A.). The *nod* gene induction activity of 22 flavonoids (Table 1) were determined by β -galactosidase assay, which has previously been described [13]. The flavonoid stock solutions were dissolved in methanol at a concentration of 1 mM. *B. japonicum* USDA110 (pZB32) was grown on AG medium [20] containing 30 μ g/ml tetracycline. For the assay, the initial turbidity of the culture was adjusted to O.D.₆₀₀=0.05 in 5 ml of AG medium containing 2 μ M flavonoids or solvents as controls. The bacterial cultures were incubated for 24 h at 27°C with shaking at 200 rpm. Assays were performed at 27°C, and replicated more than twice. To examine the synergistic or inhibitory effect of flavonoids, 1 μ M each of genistein, 7,4'-dihydroxyflavone, flavone, 7-hydroxyflavone, 7,4'-dihydroxyflavonol, apigenin (5,7,4'-trihydroxyflavone), kaempferol (5,7,4'-trihydroxyflavonol), and 7-hydroxy-4'-methoxyflavone was added to the culture medium of *B. japonicum* USDA110 (pZB32) containing either 1 μ M of genistein or 7,4'-dihydroxyflavone.

RESULTS AND DISCUSSION

Current study was undertaken to define the important features of the functional groups necessary for the flavonoids to be an active inducer or inhibitor to express the *nod* gene expression in *B. japonicum* USDA110. Table 1 shows the effect of 22 representative flavonoids at 2 μ M concentration on the induction of *nod* gene expression. Isoflavones with a 4'-OH functional group on the B-ring, genistein and daidzein, showed strong induction activity, as previously reported [2]. Comparing these two isoflavones, genistein with an additional 5-OH functional group on the A-ring induced the *nod* gene expression more significantly than daidzein with its additional 7-OH functional group on the A-ring. Formononetin and biochanin A, in which the 4'-OH functional group is blocked with a methyl functional group, showed relatively weak induction activity, compared with those of genistein and daidzein. Flavones with both 4'-OH functional groups on the B-ring induced *nod* gene expression. 4'-Hydroxyflavone showed strong induction activity, comparable to genistein. Interestingly, the hydroxyl functional group on the A-ring of the flavones had an effect to lower the induction activity. In addition, multiple hydroxyl

functional groups on the A-ring of apigenin (5,7,4'-trihydroxyflavone) significantly lowered the induction activity, compared with 5,4'-, 6,4'-, and 7,4'-dihydroxyflavones. Surprisingly, flavonols 7,4'-dihydroxyflavonol, kaempferol (5,7,4'-trihydroxyflavonol), fisetin (7,3',4'-trihydroxyflavonol), and quercetin (5,7,3',4'-tetrahydroxyflavonol), which have both 4'-OH and 3-OH functional groups on the rings, did not show the *nod* gene induction activity. Blocking of the 4'-OH functional group of flavones with the methyl group, such as 7-hydroxy-4'-methoxyflavone, acacetin (5,7-dihydroxy-4'-methoxyflavone), and kaempferide (5,7-dihydroxy-4'-methoxyflavonol), dramatically lowered the *nod* gene induction activity. Flavones with no 4'-OH functional group, such as flavone, 5-hydroxyflavone, and 7-hydroxyflavone, did not induce *nod* gene expression. The hydroxyl group located on 2'- or 3'-C of the flavone on the B-ring, instead of 4'-C, could not induce *nod* gene expression. In addition, naringenin (5,7,4'-trihydroxyflavanone), which has a single bond between 2-C and 3-C on the C-ring, did not induce *nod* gene expression, although it contains a 4'-OH functional group.

Taken together, a general rule for the flavonoids to act as an effective inducing compound to express the *nod* genes in *B. japonicum* USDA110 could be summarized as follows: (1) the double bond between 2-C and 3-C on the C-ring of (iso)flavones should be present; (2) the 4'-OH functional group is required; (3) the 3-OH functional group should not be present; and (4) in the case of flavones, the 4'-OH functional group should not be blocked with the methyl group (Fig. 1). However, this contention is not applied for the induction of the *nodA* promoter of *Rhizobium leguminosarum* Sym plasmid pRL1J1 [23]. As the (iso)flavones induced *nod* genes in *B. japonicum* USDA110, some of them could induce the *nodA* promoter. The known (iso)flavones able to induce the *nodA* promoter are flavanones (naringenin and eriodictyol), flavones (apigenin, luteolin, and 7-hydroxyflavone), and flavonol (kaempferol). However, isoflavone genistein did not act as an inducing compound for the *nodA* promoter. The general structural features required to induce the *nodA* promoter of *Rhizobium leguminosarum*

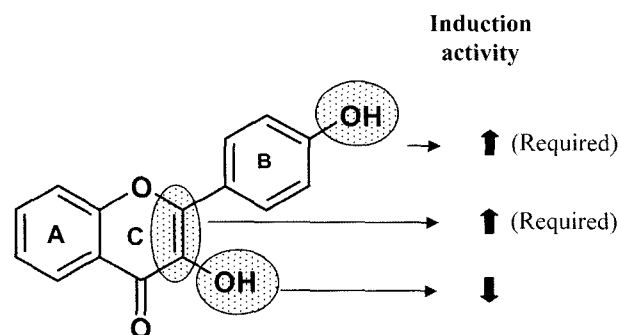


Fig. 1. Effect of the functional group of flavonoids on the induction of *nod* genes in *Bradyrhizobium japonicum* USDA110.

Table 1. β -Galactosidase activity of the *nodY::lacZ* fusion gene induced by (iso)flavones, flavonols, and flavanone at 2 μ M concentration.

Flavonoids ^a	Chemical structure	Unit of β -galactosidase activity ^b
1) <i>No Flavonoids</i>	—	8.6 \pm 0.5
2) <i>Isoflavones with 4'-OH</i>		
Daidzein (7,4'-dihydroxyisoflavone)		121.0 \pm 3.0
Genistein (5,7,4'-trihydroxyisoflavone)		256.8 \pm 39.0
3) <i>Isoflavones with 4'-OCH₃</i>		
Formononetin (7-hydroxy-4'-methoxyisoflavone)		110.7 \pm 24.8
Biochanin A (5,7-dihydroxy-4'-methoxyisoflavone)		196.4 \pm 14.4
4) <i>Flavones with 4'-OH</i>		
4'-Hydroxyflavone		200.1 \pm 10.9
5,4'-Dihydroxyflavone		151.3 \pm 32.9
6,4'-Dihydroxyflavone		160.0 \pm 26.9
7,4'-Dihydroxyflavone		168.0 \pm 35.5
Apigenin (5,7,4'-trihydroxyflavone)		86.3 \pm 17.5
5) <i>Flavonol with 4'-OH</i>		
7,4'-Trihydroxyflavonol		9.2 \pm 1.4
Kaempferol (5,7,4'-trihydroxyflavone)		12.9 \pm 5.3
Fisetin (3,7,3',4'-trihydroxyflavonol)		11.6 \pm 3.0
Quercetin (5,7,3',4'-Tetrahydroxyflavonol)		8.7 \pm 0.4

Table 1. Continued.

Flavonoids ^a	Chemical structure	Unit of β -galactosidase activity ^b
6) Flavones and flavonol with 4'-OCH ₃		
7-Hydroxy-4'-methoxyflavone		8.9±1.4
Acacetin (5,7-dihydroxy-4'-methoxyflavone)		8.5±0.2
Kaempferide (5,7-dihydroxy-4'-methoxyflavonol)		15.9±2.0
7) Flavones with no 4'-OH		
5-Hydroxyflavone		3.5±0.4
7-Hydroxyflavone		8.4±0.4
7,2'-Dihydroxyflavone		5.2±0.7
7,3'-Dihydroxyflavone		6.6±0.7
8) Flavone		
Flavone		10.2±3.2
9) Flavanone		
(R) and (S)-Naringenin mixture (5,7,4'-Trihydroxyflavanone)	 and 	12.4±1.4

^aFinal concentration of each chemical was 2 μ M.

^bThe activity is expressed as the Miller unit. Values are means and standard deviations of duplicated or more assays.

are likely to have flavone or flavanone structures with the 4'-OH functional group.

In order to examine the role of flavones as an inhibitor of *nod* gene expression, certain flavones, which did not increase *nod* gene expression (Table 1), were added at 1 μ M concentration to *B. japonicum* USDA110 broth culture

containing a relatively strong inducer, 1 μ M genistein. The *nod* gene expression was likely to be saturated at 1 μ M genistein concentration, because the level of β -galactosidase activity was almost the same at both 1 and 2 μ M concentration (data not shown). Smit *et al.* [22] reported the same pattern of induction in which β -galactosidase activity was

Table 2. β -Galactosidase activity of the *nodY::lacZ* fusion gene induced by combinations of 7,4'-dihydroxyflavone and selected flavones or flavonols.

7,4'-Dihydroxyflavone+ Selected flavones or flavonols ^a	Unit of β -galactosidase activity ^b
No Flavones	8.5 \pm 1.1
7,4'-Dihydroxyflavone	85.6 \pm 42.0 ^c 168.0 \pm 35.5 ^d
Flavone	11.5 \pm 2.1
7-Hydroxyflavone	22.5 \pm 15.7
7,4'-Dihydroxyflavonol	81.7 \pm 18.5
Apigenin (5,7,4'-trihydroxyflavone)	125.3 \pm 17.5
Kaempferol (5,7,4'-trihydroxyflavonol)	40.6 \pm 12.7
7-Hydroxy-4'-methoxyflavone	103.9 \pm 44.3

^a7,4'-Dihydroxyflavone and selected flavone were added to media at a concentration of 1 μ M, respectively. The combined final concentration of 7,4'-dihydroxyflavone and selected flavone was 2 μ M.

^bThe activity is expressed as the Miller unit. Values are means and standard deviations of duplicated or more assays.

^{c,d} β -Galactosidase activities at the concentration of 1 μ M and 2 μ M of 7,4'-dihydroxyflavone, respectively.

saturated at around 1 μ M of genistein, using the *nodY::lacZ* system in *B. japonicum* USDA135. Flavone and kaempferol (5,7,4'-trihydroxyflavonol), which did not induce *nod* gene expression, weakly decreased the level of the *nod* gene expression induced by 1 μ M genistein. However, non-inducing compounds, such as 7-hydroxyflavone and 7,4'-dihydroxyflavonol, did not affect the induction activity of genistein. Interestingly, 7-hydroxy-4'-methoxyflavone showed a slightly synergistic effect with genistein to increase the induction level of *nod* gene expression, although it itself did not increase *nod* gene expression (data not shown).

Table 2 shows the result of the inhibitory effect of flavones on *nod* gene expression, when 7,4'-dihydroxyflavone was used instead of genistein. 7,4'-Dihydroxyflavone induced stoichiometric expression of the *nod* gene as the concentration of the compound increased from 1 μ M to 2 μ M. However, flavone, 7-hydroxyflavone, and kaempferol (3,5,7,4'-tetrahydroxyflavone) significantly inhibited the induction effect of 7,4'-dihydroxyflavone on *nod* gene expression by 8-, 4-, and 2-fold, respectively. 3,7,4'-Trihydroxyflavone did not affect the *nod* gene expression induced by 7,4'-dihydroxyflavone. Induction activity by 7,4'-dihydroxyflavone on *nod* gene expression slightly increased with the addition of 7,3'-dihydroxyflavone, apigenin (5,7,4'-trihydroxyflavone), and 7-hydroxy-4'-methoxyflavone. With 1 μ M genistein, there was no significant influence of other (iso)flavones tested on the *nod* gene expression in *B. japonicum* USDA110. This is likely to be due to a competitive effect of genistein against the (iso)flavones tested in binding the *nodD* gene product or an uptake system [3, 4, 21]. However, the inhibitory effect of other selected flavonoids on the *nod* gene induction activity was observed with 7,4'-dihydroxyflavone at 1 μ M

concentration. The inhibitory activity decreased in the following order: flavone>7-hydroxyflavone>kaempferol (5,7,4'-trihydroxyflavonol)>7,4'-trihydroxyflavonol. In general, the inhibitory activity was shown with flavones that have no 4'-OH functional group and/or have the 3-OH functional group. A similar result was also reported by Cunningham *et al.* [4], who suggested inhibitory activity with flavonol (3-OH), flavones (no 4'-OH), and flavanones (no double bond in 2-C and 3-C) in decreasing order.

In conclusion, using the *nodY::lacZ* expression system in *B. japonicum*, we were able to define certain functional features for the flavonoids to act as inducing or inhibitory compounds. This information could contribute to understanding rhizobia-soybean interaction and strain-strain competition as a way to increase legume production [4, 9].

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