Curcubinoyl flavonoids from wild ginseng adventitious root cultures

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Wild ginseng (Panax ginseng) adventitious root cultures were prepared by elicitation using methyl jasmonate and investigated further to find new secondary metabolites. Chromatographic fractionation of wild ginseng adventitious root cultures led to the isolation of eleven compounds. The chemical structures of isolated compounds were identified as four known flavanone derivatives (1–4), one new curcubinoyl derivative, jasmogin A (5) and six new curcubinoyl-flavanone conjugates, jasmoflagins A-F (6–11) by extensive spectroscopic analysis. Newly isolated curcubinoyl derivatives showed inhibitory activity against lipopolysaccharide-stimulated nitric oxide production in RAW 264.7 macrophages. Therefore, our present study suggested that elicitor stimulated plant cell cultures might contribute to the production of new metabolites.

Natural products contain a variety of ingredients and have long been used to prevent and treat diseases. However, securing natural products is essential in order to develop these natural products, which is sometimes not easy due to various constraints. Plant tissue culture technology is suggested as a powerful tool for obtaining natural substances1–3. This is widely used for the production of plant materials because it is less affected by weather and other external conditions than plant cultivation and relatively for a short period of time.

For maximum productivity, culture conditions such as culture medium and the incubation conditions, etc., are optimized when growing plant tissues4–6. In particular, the use of elicitors is widely used for increased productivity and useful substances. As elicitors, salicylic acid and methyl jasmonate (MJ), which control the immune of plants, are most widely used7–4. These elicitors greatly increase the content of biomass and useful metabolites4,9. Moreover, new ingredients have been reported in elicitor-stimulated plant cell culture10–11. Therefore, plant tissue culture has become an important tool not only for securing plant materials but also finding new metabolites.

Panax ginseng C.A. Meyer (Araliaceae) is commonly known as Korean ginseng. It is one of the most widely used tonic to enhance immune response and consequent health and longevity for over 2000 years in Oriental countries12. Various efficacy of P. ginseng, including anti-cancer, anti-inflammatory, anti-diabetic, anti-fatigue and neuroprotective activities have been also reported from a lot of research13–16.

Ginseng grows in wild environment or is cultivated on farm. Cultivated ginseng is systematically grown on farm under the control of growth condition and harvested after 4–6 year cultivation. The wild ginseng, also called mountain ginseng in Korea, grows without human touch in deep areas with low sunlight and temperature changes. This difference in the cultivation environment and genotypes leads to differences in the composition and efficacy of the two specimens. Wild ginseng has been reported to have enhanced host defense components and biological activities. The concentration of ginsenosides and amino acids in wild ginseng were much higher than those of cultivated ginseng17,18.

However, due to the short supply and consequent high price of wild ginseng has limited its usage despite of beneficial biological activities. Therefore, sufficient production is required for the development as products. As a preparation of wild ginseng, tissue culture system is considered as a valuable tool to achieve rapid and stable production of excellent individual. We previously established efficient adventitious root cultures of wild ginseng with fast growth and stable production9,20. In addition, we also demonstrated the increased yield and antioxidant activity of MJ-elicitated wild ginseng adventitious root cultures compared to MJ-untreated samples21. In the present study, MJ-treated wild ginseng adventitious root cultures were investigated further to find new secondary metabolites.

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Results and discussion

Isolation of compounds from MJ-treated wild ginseng adventitious root cultures. Plant cell cultures were used not only for the stable production but also useful to find new secondary metabolites for better pharmacological activity9–11. Investigation on the constituents of the adventitious root cultures of *P. ginseng* yielded eleven compounds including seven new compounds (Fig. 1). Known compounds were identified as naringenin (1), naringenin-4′-O-β-glucoside (2), naringenin-7-O-β-glucoside (3) and hesperetin 7-O-β-glucoside (4) by the analysis of their spectroscopic data and comparison with literature values22–24.

Structural determination of the new compounds. Compound 5 was obtained as a white amorphous powder. The molecular formula of 5 was determined as C_{12}H_{20}O_{4} by its HRESI-MS (m/z 251.1254 [M + Na]+, calcd. C_{12}H_{20}NaO_{4} 251.1253). In the ^1H and ^13C NMR spectrum, the presence of an olefinic moiety in cis-configuration was deduced from the signals at [δ_H 5.52 (1H, m, H-9), 5.42 (1H, dd, J = 10.4, 8.8 Hz, H-10); δ_C 128.8 (C-9), 134.0 (C-10)]. The ^1H and ^13C NMR spectrum showed the signals attributed to two hydroxymethines at [δ_H 4.15 (1H, m, H-6), 4.68 (1H, dq, J = 8.8, 6.4 Hz, H-11)], two methines at [δ_H 2.16 (1H, m, H-3), 1.46 (1H, m, H-7)] and one methyl group at [δ_H 1.23 (1H, d, J = 6.4 Hz, H-12)]. The ^1H NMR spectrum and the corresponding carbon signals in HSQC spectrum revealed the presence of four methylenes from the signals at [δ_H 2.55 (1H, dd, J = 14.4, 3.6 Hz, H-2a), 2.15 (1H, m, H-2b); δ_C 39.1 (C-2)], [δ_H 1.88 (1H, m, H-4a), 1.64 (1H, m, H-4b); δ_C 32.8 (C-4)], [δ_H 2.11 (1H, m, H-5a); 1.33 (1H, m, H-5b); δ_C 28.8 (C-5)] and [δ_H 2.24 (1H, m, H-8a); δ_C 25.5 (C-8)]. In addition, a carbonyl signal was observed at δ_C 175.9 (C-1) in the 13C NMR spectrum. In the HMBC spectrum, correlations from H-4 to C-6, 7 and from H-5 to C-3 suggested the presence of cyclopentyl moiety in 5. These NMR spectroscopic data of 5 were quite similar to those of curcurbic acid, a hydroxylated jasmonate derivative25, except for the additional hydroxyl group. The position of an additional hydroxyl group was determined by the NOESY correlations between H-6, H-7 and H-2 and between H-8 and H-11 (Fig. 2). Taken together, compound 5 was defined as shown and named jasmogin A.

Compound 6 was purified as a white amorphous powder and assigned the molecular formula as C_{33}H_{40}O_{13} by its HRESI-MS (m/z 667.2354 [M + Na]+, calcd. C_{33}H_{40}NaO_{13} 667.2331). The NMR spectroscopic clearly showed that 6 has hydroxylated curcurbic acid moiety of 5 as a partial structure. Additionally, compound 6 was supposed to be a glycoside from the anomeric signals at [δ_H 4.96 (1H, d, J = 7.4 Hz, H-1″); δ_C 99.8] together with [δ_H 3.35–3.70 (4H, m, H-2″, 3″, 4″, 5″), 4.43 (1H, dd, J = 11.7, 2.1 Hz, H-6″a), 4.21 (1H, m, H-6″b); δ_C 76.4 (C-2″), 73.2 (C-3″), 70.3 (C-4″), 74.2 (C-5″), 63.2 (C-6″)] in the ^1H and ^13C NMR spectrum. Besides aforementioned...
moieties of curcurbic acid and glucose, the presence of disubstituted and tetrasubstituted aromatic rings were
deduced from the signals at \([\delta_H 7.34 (2H, d, J = 8.0 \text{ Hz}, H-2', 6'), 6.84 (2H, d, J = 8.0 \text{ Hz}, H-3', 5')]; \delta_C 129.5 (C-1'),
128.9 (C-2', 6'), 157.7 (C-4'), 115.0 (C-3', 5'))\] and from the signals at \([\delta_H 6.20 (1H, d, J = 2.1 \text{ Hz}, H-6), 6.23 (1H, d, J = 2.1 \text{ Hz}, H-8); \delta_C 163.2 (C-5), 96.7 (C-6), 165.5 (C-7), 95.7 (C-8), 163.5 (C-9), 103.6 (C-10)],\) respectively, in
the \(^1\text{H}\) and \(^{13}\text{C}\) NMR spectrum. From these two aromatic rings and together with additional signals at \([\delta_H 5.40 (1H, dd, J = 13.0, 3.0 \text{ Hz}, H-2), \delta_C 79.3 (C-2)]; \delta_H 3.17 (1H, dd, J = 17.0, 13.0 \text{ Hz}, H-3a), 2.77 (1H, dd, J = 17.0, 3.0, H-3b); \delta_C 43.1 (C-3)]\] and carbonyl signal at \(\delta_C 197.2 (C-4),\) compound 6 was supposed as a flavanone derivative,
which was identified as naringenin (1)\(^{23}\). Taken together, 6 was suggested as a flavanone glycoside consisting of
naringenin, glucose and curcurbic acid moieties. The linkage of each unit was determined by cross peaks between
H-1" of glucose and C-7 of naringenin, and between H-6" of glucose and C-1‴ of curcurbic acid moiety in the
HMBC spectrum (Fig. 3). Collectively, compound 6 was defined as shown and named jasmoflagin A.

Compound 7 was purified as a white amorphous powder and showed an HRESI-MS ion at \(m/z\) 667.2354
([M + Na]\(^{+}\), calcld 667.2361) for \(C_{33}H_{40}NaO_{13}\). The spectroscopic data of 7 were quite similar to those of 6, which
suggested that 7 is also a curcurbinoyl derivative of naringenin glycoside. Careful comparison of \(^1\text{H}\) and \(^{13}\text{C}\) NMR
data of 7 with those of 6 showed the differences in the chemical shifts of H-11 and H-12. The hydroxymethine
and methyl signals at δ_H 4.62 (H-11) and 1.20 (H_3-12) in 6 were replaced by methylene at [δ_H 2.37 (1H, m, H-11a) and 2.25 (1H, m, H-11b); δ_C 30.3] and hydroxymethylene at δ_H 3.56 (2H, m, H-12); δ_C 61.2] in 7. The correlation between H-10‴ and C-11‴ and between H-11‴ to C-12‴ confirmed the attachment of hydroxyl group to C-12 (Fig. 3). Taken together, compound 7 was determined as shown and named jasmoflagin B.

Compound 8 was also purified as a white amorphous powder. The molecular formula was established as C_{33}H_{40}O_{13} from an HRESI-MS ion at m/z 651.2404 ([M + Na]+, calc 651.2204). The 1H NMR spectrum of 8 were quite similar to those of 6, except for the disappearance of hydroxymethylene proton at δ_H 4.05 (H-6‴) of 6. Additional carbonyl signal at δ_C 220.1 in the 13C NMR spectrum suggested that 8 is a jasmonate derivative of naringenin glycoside. Further HMBC correlation from H-4‴, H-5‴, H-7‴ to C-6‴ confirmed the presence of carbonyl moiety at C-6‴ (Fig. 3). Taken together, compound 8 was determined as shown and named jasmoflagin C.

Compound 9 was purified as a white amorphous powder and showed an HRESI-MS ion at m/z 651.2404 ([M + Na]+, calc 651.2412) for C_{33}H_{40}NaO_{12}. The spectroscopic data of 9 suggested that 9 is also a curcurbinoyl derivative of naringenin glycoside. However, on the contrary to 6 and 7, hydroxymethine signals in curcurbic acid were not observed in the 1H and 13C NMR data of 9. Further analysis demonstrated that the hydroxymethine signals at [δ_H 4.62 (1H, m, H-11); δ_C 62.5] in 6 were replaced by methylene signals at [δ_H 2.06 (2H, m, H-11); δ_C 20.1] in 9. In addition, HMBC correlation between H-10‴ and C-11‴ and between H-11‴ and C-12‴ confirmed the detachment of hydroxyl group at C-11 in 6 (Fig. 3). Taken together, compound 9 was determined as shown and named jasmoflagin D.

Compound 10 was purified as white amorphous powder and assigned the molecular formula as C_{33}H_{40}O_{13}, which is same as 6. The spectroscopic data of 10 were quite similar to those of 6 and suggested 10 is comprised of naringenin glucoside and curcurbic acid with a hydroxyl group. The correlations in the 1H and 13C NMR data of 10 from those of 6 were observed as downfield shift of CH_{3}-12‴ from δ_H 1.20 to δ_H 1.68 and upfield shift of H-8‴ from δ_H 2.04, 2.31 to δ_H 1.60, 1.70, suggesting the change in the positions of hydroxyl group and double bond in curcurbic acid moiety. The 1H-1H COSY correlations of H-8‴, H-5‴, H-7‴ to C-6‴” confirmed the presence of amine group to curcurbinoyl moiety of 10, which is also confirmed by the presence of nitrogen in 11 from MS analysis. The positions of hydroxyl and amine groups were determined to be C-9‴ and C-10‴, respectively, by the HMBC correlations from H-7‴/H-8‴ to C-9‴ and from H-8‴/12‴ to C-10‴” (Fig. 3). Taken together, compound 11 was defined as shown and named jasmoflagin E.

**NO inhibitory activity of isolated compounds.** Next, we investigated the anti-inflammatory effects of newly isolated compounds by measuring the production of NO in LPS-stimulated RAW 264.7 macrophages. As shown in Fig. 4, compounds 5, 7 and 10 dose-dependently reduced NO production stimulated by LPS without any significant cytotoxic effects at the concentration ranging from 5 to 50 μM. Compound 5, which has only curcurbinoyl moiety, inhibited NO production. However, addition of flavanone moiety to compound 5 reduced the inhibitory activity, as observed in compound 6. Interestingly, among the curcurbinoyl flavanone derivatives, compounds 7 and 10 showed stronger inhibitory activity compared to others, which suggested the importance of the position of hydroxyl group in curcurbinoyl moiety.
Conclusion
Fractionation of using various chromatographic techniques yielded eleven compounds from the MJ-treated adventitious root cultures of wild ginseng. The chemical structures of isolated compounds were identified by spectroscopic analysis and further identified seven new compounds. The newly reported compounds are curcubinoyl derivative, named jasminog A (5) and curcubinoyl-conjugated flavanone derivatives, named jasmo-
flagins A-F (6–11). Considering the structural similarity between methyl jasmonate and curcubinoyl moiety, addition of elicitor can affect not only the increase of biosynthesis of active metabolite, but MJ itself also partici-
pate in biosynthetic pathway as a substrate, which needs to be clarified by further study.

Materials and methods
General experimental procedure. IR spectra were obtained using JASCO FTIR 4100 spectrometer in
CH3OH solvent. Optical rotations were measured on a JASCO DIP-1000 polarimeter (Tokyo, Japan). HRESIMS data were measured on maXis 4G (Bruker) and LCQ Fleet (Thermoscientific), respectively. NMR spectra were recorded on a Bruker Avance 400, 500 and 800 MHz spectrometers using CD3OD as solvent. Silica gel (200–400 mesh, Merck), Sephadex LH-20 (20–100 μm, Sigma) and Diaion HP-20P (Mitsubishi Kasei, Japan) for column chromatography. TLC was performed on silica gel 60 F254 (0.2 mm, Merck) or silica gel 60 RP-18 F254S (0.2 mm, Merck), and spots were detected by a 10% vanillin-H2SO4 in EtOH spray reagent. MPLC was performed on a Biotage Isolera Prime chromatography system and a Lichroprep RP-18 column (40–63 μm). Semi-prep HPLC was performed using a Waters system (three 515 pumps and a 996 photodiode array detector) with a Phenome-

Plant material. Wild ginseng was collected at Mt. Ohdae of Korea by the government certificated dig-
ger. It was identified by the Emeritus Prof. Kee-Yoeup Paek and certificated by Korea ginseng institutes. The
permisions were obtained from concerned authorities for collection and use of sample. And all methods were performed in accordance with the relevant regulations.

Adventitious root cultures of wild ginseng (P. ginseng) were produced from a 100-year-old wild ginseng through callus culture as we described previously20. The root cultures were proliferated in a 5 L airlift balloon type bioreactor containing 4.0 L Murashige and Skoog (MS) liquid medium (3/4 strength) supplemented with 5.0 mg/L IBA, 0.1 mg/L kinetin, and 5% (w/v) sucrose for seven weeks. The stock solution of MJ was prepared in ethanol as 50 mM and MJ were added to the culture as an elicitor as final concentration of 100 μM, one week before harvest. The adventitious roots were harvested from the culture and washed three times with distilled water to remove the medium on the surface of the adventitious roots. Then, it was immediately frozen with liquid nitrogen and stored in deep-freezer at − 70 °C, and then freeze-dried before further experiments. A voucher specimen (CBNU-WGAR2014) was deposited at the Herbarium of the College of Pharmacy, Chungbuk National University, Korea.

Extraction and isolation of compounds. The dried MJ-treated wild ginseng adventitious root cultures (5.0 kg) were extracted twice with 25 L of 80% MeOH for 24 h at room temperature. The methanol extract (1.7 kg) was suspended in H2O and partitioned successively with n-hexane, CH2Cl2, EtOAc and n-BuOH to yield corresponding fractions. The EtOAc fraction (WGE, 21.9 g) was subjected to MPLC over silica gel (CH2Cl2-MeCN, 1:0 → 0:1) to afford sixteen subfractions (WGE1-WGE16). WGE5 was subjected to MPLC over reverse phase silica gel (MeOH-H2O, 10:90 → 1:0) to give five subfractions (WGE5A-WGE5E). Compounds 9 (17.1 mg) and 10 (12.6 mg) were purified from WGE5 by semi-preparative HPLC eluting with MeCN- H2O (30:70). WGE9 was subjected to MPLC over reverse phase silica gel (MeOH-H2O, 10:90 → 1:0) to give three subfractions (WGE9A- WGE9C). Compounds 3 (29.7 mg) were purified from WGE9B by semi-preparative HPLC eluting with MeCN- H2O (30:70). WGE15 was subjected to MPLC over reverse phase silica gel (MeOH-H2O, 10:90 → 1:0) to give two subfractions (WGE15A-WGE15B). Compounds 1 (19.9 mg) and 5 (13.4 mg) were purified from WGE15A by semi-preparative HPLC eluting with MeCN- H2O (30:70).

The n-BuOH fraction (WGB, 218.3 g) was subjected to HP-20 (MeOH-H2O, 0:1 → 0:1) to afford five subfractions (WGB1-WGB5). WGB5 was subjected to MPLC over reverse phase silica gel (CH2Cl2-MeCN, 1:0 → 0:1) to give nine subfractions (WGE5A- WGE5I). WGB5C was subjected to MPLC over silica gel (MeOH-H2O, 10:90 → 1:0) to give three subfractions (WGE5C1- WGE5C3). Compounds 8 (18.1 mg) and 11 (11.7 mg) were purified from WGE5C1 by semi-preparative HPLC eluting with MeCN- H2O (30:70). Compounds 2 (15.0 mg), 4 (15.3 mg), 6 (16.2 mg) and 7 (16.7 mg) were purified from WGE5C2 by semi-preparative HPLC eluting with MeCN- H2O (40:60).

Jasminogin (5) Light yellow gum; [α]25D +53.9° (c 0.01, MeOH); IR (KBr) νmax 224, 281 nm; ESIMS m/z 249 [M + H]+; HRESIMS m/z 251.1254 [M + Na]+ (calcd for C12H20NaO4 251.1253); 1H-NMR (500 MHz, CD3OD) and 13C-NMR (225 MHz, CD3OD), see Table 1.
Jasminoflagin A (6) Light yellow gum; [α]25D +79.2° (c 0.05, MeOH); IR (KBr) νmax 3537, 2915, 2337, 1662, 1362, 1052 cm−1; UV (MeOH) λmax 249 [M + Na]+; HRESIMS m/z 667.2354 [M + Na]+ (calcd for C30H25NaO12 667.2331); 1H-NMR (500 MHz, CD3OD) and 13C-NMR (175 MHz, CD3OD), see Table 2 and 3.
Jasminoflagin B (7) Light yellow gum; [α]25D +97.1° (c 0.05, MeOH); IR (KBr) νmax 3698, 3250, 2044, 1662, 1362, 802 cm−1; UV (MeOH) λmax 212, 281, 327 nm; ESIMS m/z 667 [M + Na]+; HRESIMS m/z 667.2354 [M + Na]+ (calcd for C30H25NaO12 667.2361); 1H-NMR (700 MHz, CD3OD) and 13C-NMR (175 MHz, CD3OD), see Tables 2 and 3.
Jasmoflagin C (8) Light yellow gum; [α]_D^25 ~34.2 (c 0.01, MeOH); IR (KBr) ν_max 3472, 2928, 1645 cm⁻¹; UV (MeOH) λ_max 283, 327 nm; ESIMS m/z 672 [M + Na]⁺; HRESIMS m/z 667.2360 [M + Na]⁺ (calcd for C33H40NaO13 667.2361); 1H-NMR (700 MHz, CD3OD) and 13C-NMR (175 MHz, CD3OD), see Tables 2 and 3.

Jasmoflagin D (9) Light yellow gum; [α]_D^25 ~38.5 (c 0.05, MeOH); IR (KBr) ν_max 3478, 2928, 1645 cm⁻¹; UV (MeOH) λ_max 212, 281, 389 nm; ESIMS m/z 667 [M + Na]⁺; HRESIMS m/z 651.2404 [M + Na]⁺ (calcd for C33H40NaO12 651.2412); 1H-NMR (50 MHz, CD3OD) and 13C-NMR (100 MHz, CD3OD), see Tables 2 and 3.

Table 1. NMR spectroscopic data for compound 5 (CD3OD).

| δ_H | δ_C |
|-----|-----|
| 1   | 175.9 |
| 2   | 2.55 (dd, 14.4, 3.6), 2.15 (m) | 39.1 |
| 3   | 2.16 (m) | 38.5 |
| 4   | 1.88 (m), 1.64 (m) | 32.8 |
| 5   | 2.11 (m), 1.33 (m) | 28.8 |
| 6   | 4.15 (m) | 73.4 |
| 7   | 1.46 (m) | 50.9 |
| 8   | 2.29 (m), 2.24 (m) | 25.5 |
| 9   | 5.52 (m) | 128.8 |
| 10  | 5.42 (dd, 10.4, 8.8) | 134.0 |
| 11  | 4.68 (dq, 8.8, 6.4) | 63.2 |
| 12  | 1.23 (d, 6.4) | 22.5 |

Table 2. 1H NMR spectroscopic data for compounds 6–11 (CD3OD).

| No | 6  | 7   | 8   | 9   | 10  | 11   |
|----|----|-----|-----|-----|-----|------|
| 2  | 5.40, dd (13.0, 3.0) | 5.41, dd (12.5, 3.5) | 5.41, dd (12.6, 3.6) | 5.38, dd (13.0, 2.8) | 5.42, dd (13.0, 3.0) | 5.42, dd (12.7, 2.9) |
| 3  | 3.17, dd (17.0, 13.0) | 3.17, dd (17.0, 12.5) | 3.17, dd (17.5, 12.6) | 3.14, dd (17.1, 13.0) | 3.16, dd (17.0, 13.0) | 3.18, dd (17.1, 12.7) |
| 6  | 6.20, d (2.1) | 6.20, d (2.2) | 6.20, d (2.3) | 6.21, d (2.1) | 6.21, d (2.1) |
| 8  | 6.23, d (2.1) | 6.23, d (2.2) | 6.25 d (2.2) | 6.23, d (2.3) | 6.24, d (2.1) | 6.24, d (2.1) |
| 2′/6′ | 7.34, d (8.0) | 7.34, d (8.5) | 7.34, d (8.5) | 7.33, d (8.5) | 7.35, d (8.3) | 7.35, d (8.6) |
| 3′/5′ | 6.84, d (8.0) | 6.84, d (8.5) | 6.83, d (8.5) | 6.84, d (8.5) | 6.85, d (8.3) | 6.85, d (8.6) |
| 1″ | 4.96, d (7.4) | 4.96, d (7.4) | 4.97, d (7.5) | 4.97, d (7.5) | 4.97, d (7.5) | 4.96, d (7.5) |
| 2″ | 3.48 m | 3.48 m | 3.50 m | 3.47 m | 3.49 m | 3.48 m |
| 3″ | 3.46 m | 3.47 m | 3.46 m | 3.47 m | 3.47 m | 3.46 m |
| 4″ | 3.34 m | 3.34 m | 3.33 m | 3.34 m | 3.34 m | 3.33 m |
| 5″ | 3.70 m | 3.71 m | 3.69 m | 3.72 (t, 9.8) | 3.72 m | 3.72 m |
| 6″ | 4.43, dd (11.7, 2.1) | 4.43, dd (11.9, 2.1) | 4.51, dd (11.8, 2.1) | 4.42, dd (11.8, 2.1) | 4.48 dd (11.8, 2.1) | 4.47 dd (11.8, 1.9) |
| 2″/6″ | 2.52 m | 2.54 m | 2.67 m | 2.54 m | 2.33, m | 2.35 m |
| 3″ | 2.22 m | 2.20 m | 2.36 m | 2.18 m | 2.31 m | 2.30 m |
| 4″ | 2.08 m | 2.08 m | 2.19 m | 2.06 m | 1.96 m | 1.96 m |
| 5″ | 1.99 m | 1.74 m | 1.38 m | 1.96 m | 1.81 m | 1.85 m |
| 6″ | 1.19 m | 1.53 m | 2.10 m | 1.20 m | 0.99 m | 1.52 m |
| 7″ | 1.75 m | 1.98 m | 2.30 m | 1.73 m | 1.87 m | 1.83 m |
| 8″ | 1.54 m | 1.18 m | 2.31 m | 1.55 m | 1.47 m | 1.12 m |
| 9″ | 4.05 m | 4.07 m | – | 4.07 m | 4.48 m | 4.48 m |
| 7″/8″ | 1.24 m | 1.24 m | 1.71 m | 1.23 m | 2.14 m | 2.13 m |
| 3″/4″ | 2.31 m | 2.20 m | 2.16 m | 2.12 m | 1.70 m | 1.70 m |
| 2″/3″ | 2.04 m | 2.09 m | 2.23 m | 2.12 m | 1.60 m | 1.62 m |
| 9″ | 5.43 m | 5.51 m | 5.26 m | 5.37 m | 4.15 m | 3.69 m |
| 10″ | 5.41 m | 5.41 m | 5.40 m | 5.36 m | 5.37 m | 3.24 m |
| 11″ | 4.62 m | 2.37 m, 2.25 m | 4.58 m | 2.06 m | 5.65 m | 1.40 m |
| 12″ | 1.20, d (6.4) | 3.56 m | 1.18, d (6.3) | 0.96, d (7.5) | 1.68, d (6.3) | 0.97, dd (14.1, 7.1) |
Jasmoflagin E (10) Light yellow gum; \([\alpha]_{D}^{25} -92.2 (c 0.05, \text{MeOH})\); IR (KBr) \(\text{n}\text{max} 3751, 3311, 2331, 1662, 824 \text{ cm}^{-1}\); UV (MeOH) \(\lambda\text{max} 283, 325 \text{ nm}\); ESIMS \(m/z 667 [M + \text{Na}]^{+}\); HRESIMS \(m/z 651.2334 [M + \text{Li}]^{+}\) (calcd for \(C_{33}H_{40}LiO_{13} 651.2624\); 1H-NMR (400 MHz, CD3OD) and 13C-NMR (100 MHz, CD3OD), see Tables 2 and 3.

**Measurement of LPS-induced NO production.** Inhibitory effect of compounds on lipopolysaccharide (LPS)-induced nitric oxide (NO) production was assessed using RAW264.7 macrophage cell lines. RAW264.7 cells were treated with 1 μg/ml LPS in the presence or absence of compounds. After 24 h incubation, the cell medium was mixed with Griess reagent and the amount of NO formed was determined by measuring the absorbance at 550 nm in an ELISA reader. Cell viability of the remaining cells was determined by MTT assay.

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**Author contributions**
Q.L. and M.K.L. designed and coordinated the project. Q.L., S.B.K., Y.H.J, J.H.A. and A.T. performed the extraction, isolation and structural identification of the compounds. D.E.K., B.Y.C. and S.Y.K. carried out the biological assays. C.-S.J. and S.Y.P. performed the mountain ginseng root cultures. Q.L., S.Y.P., B.Y.H. and M.K.L. analyzed the data. Q.L. and M.K.L. wrote the manuscript and all authors reviewed the manuscript.

**Competing interests**
The authors declare no competing interests.

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