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## Original Article

# Characterization of metabolite profiles from the leaves of green perilla (*Perilla frutescens*) by ultra high performance liquid chromatography coupled with electrospray ionization quadrupole time-of-flight mass spectrometry and screening for their antioxidant properties



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## ARTICLE INFO

## Article history:

Received 30 June 2016

Received in revised form

20 September 2016

Accepted 29 September 2016

Available online 5 November 2016

## Keywords:

antioxidant activity

green perilla leaves

metabolite

rosmarinic acid

UPLC-ESI-Q-TOF-MS/MS

## ABSTRACT

The objective of this research was to access the determination of metabolite profiles and antioxidant properties in the leaves of green perilla (*Perilla frutescens*), where these are considered functional and nutraceutical substances in Korea. A total of 25 compositions were confirmed as six phenolic acids, two triterpenoids, eight flavonoids, seven fatty acids, and two glucosides using an ultra high performance liquid chromatography coupled with electrospray ionization quadrupole time-of-flight mass spectrometry (UPLC-ESI-Q-TOF-MS/MS) technique from the methanol extract of this species. The individual and total compositions exhibited significant differences, especially rosmarinic acid (10), and linolenic acids (22 and 23) were detected as the predominant metabolites. Interestingly, rosmarinic acid (10) was observed to have considerable differences with various concentrations in three samples (Doryong, 6.38 µg/g; Sinseong, 317.60 µg/g; Bongmyeong, 903.53 µg/g) by UPLC analysis at 330 nm. The scavenging properties against 2,2-diphenyl-1-picrylhydrazyl (DPPH) and 2,2'-azino-bis(3-ethylbenzthiazoline-6-sulphonic acid) (ABTS) radicals also showed potent effects with remarkable differences at a concentration of 100 µg/mL, and their abilities were as follows: Sinseong (DPPH, 86%; ABTS, 90%) > Bongmyeong (71% and 84%, respectively) > Doryong (63% and 73%, respectively). Our results

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<http://dx.doi.org/10.1016/j.jfda.2016.09.003>

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suggest that the antioxidant activities of green perilla leaves are correlated with metabolite contents, especially the five major compositions 10 and 22–25. Moreover, this study may be useful in evaluating the relationship between metabolite composition and antioxidant activity.

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## 1. Introduction

In recent years, metabolites including phenolic compounds and triterpenoids have been of great interest in food and medical industries because of their beneficial effects on human health [1–5]. Phenolic compounds are distributed in crops, fruits, vegetables, and edible natural plants [6–8] and are associated with a wide range of health beneficial effects including antioxidant, antidiabetic, anti-inflammatory, and anticancer agents [1,3,7,9]. Triterpenoids have also been reported to play essential roles in preventing human diseases because of their anticancer, antioxidant, antibacterial, and antiatherosclerotic properties [2,4,10]. Moreover, many studies have reported that fatty acids have beneficial properties on lipoprotein profile, blood cholesterol level, and basal metabolism of humans [11,12]. For these reasons, several researchers have focused on natural sources with high phytochemical contents for the manufacture of supplements with preventive and therapeutic capacities. In our continuing survey of bioactive substances, an investigation of the metabolite profile in the leaves of green perilla was carried out.

Perilla [*Perilla frutescens* (L.) Britt], which belongs to the Labiatae family, is a widely cultivated edible and medicinal plant in Asian countries such as China, Japan, India, and South Korea [13]. Moreover, this plant has long been used as an important traditional medicine for treating diseases such as tumor, cough, allergy, and intoxication [14,15]. Numerous researches have described that the health beneficial capacities of perilla are related to its metabolite contents (phenolic acids, monoterpenes, flavonoids, and triterpenoids) [2,6,16,17]. In particular, the leaves, seeds, and stems of this species have shown to have antipyretic and antibiotic effects for treating intestinal disorders [18]. Perilla leaves are used as food and medicinal materials for their antimicrobial, antioxidant, anticancer, antidiabetic, antitumor, and antiallergic effects owing to the presence of phenolic compounds, monoterpenes, and triterpenoids [18–21]. Perilla seeds are an important source of fatty acids ( $\alpha$ -linolenic acid, linoleic acid, oleic acid, and palmitic acid) and possess health benefits, such as lowering risk of colon cancer as well as plasma lipid levels, reducing the cholesterol level, and reducing the triglyceride level [22–24]. Commonly, human health benefits are associated with many antioxidant metabolites. Among these various biological benefits, perilla plant has demonstrated potent antioxidant properties by *in vitro* method. For example, studies on purple (red) perilla have revealed that the high antioxidant activities are attributable to several anthocyanins [25]. Also, green perilla is known to be associated with phenolic compounds except anthocyanins [26]. We recently

reported the optimal harvest time for the phytochemical contents in the leaves of purple perilla [27]. Moreover, we evaluated information concerning the antioxidant effects and inhibitory activities against  $\alpha$ -glucosidase as well as aldose reductase of perilla seeds [28]. Several studies have also established the metabolite contents and their biological activities in perilla plant [18–24]. Even though many reports have evaluated the beneficial health effects of perilla regarding metabolites, the exact chemical components in the leaves of green perilla have still not been fully characterized. In addition, the antioxidant capacities of natural plants are considered to have higher synergistic activities in metabolite extracts compared to the effects of a single phytochemical in recent years [1,3,6]. For these reasons, in order to evaluate the antioxidant abilities of perilla leaves, the 2,2-diphenyl-1-picrylhydrazyl (DPPH) and 2,2'-azino-bis(3-ethylbenzthiazoline-6-sulphonic acid) (ABTS) radical methods, based on an electron transfer and involving the reduction of a colored oxidant (DPPH: purple; ABTS: blue/green), have been measured by the spectrophotometric assay. Therefore, our work was designed to investigate the metabolite profiles and antioxidant capacities from the methanol extract of a widely used green perilla leaves.

The purpose of the present research was to characterize the metabolite profiles in the leaves of Korean green perilla using ultra high performance liquid chromatography coupled with electrospray ionization quadrupole time-of-flight mass spectrometry (UPLC-ESI-Q-TOF-MS/MS) system. Furthermore, this study is the first to investigate the changes in metabolite compositions from three methanol extracts of this species. We also determined the antioxidant properties of the scavenging abilities on DPPH and ABTS radicals in view of simplicity, easy control, and cost-effectiveness aspects.

## 2. Materials and methods

### 2.1. Plant material and chemicals

The leaves of green perilla were obtained from local markets (Daejeon) in Korea. Three samples (Doryong, Sinseong, Bongmyeong) of perilla leaves were air-dried for 3 days at room temperature to remove the moisture. The collected perilla leaves were immediately freeze-dried at  $-40^{\circ}\text{C}$  until analysis. Caffeic acid and rosmarinic acid standards were isolated by chromatographic techniques using silica gel column chromatography (230–400 mesh silica gel, kieselgel 60; Merck, Darmstadt, Germany) based on the data reported in a previous study [6]. Silica gel column chromatography was

carried out using 230–400 mesh silica gel (kieselgal 60; Merck) and DPPH and ABTS were purchased from Sigma (St. Louis, MO, USA). Analytical-grade methanol, formic acid, and water were purchased from J.T. Baker (Phillipsburg, NJ, USA). Moreover, other chemicals, reagents, and solvents used in the present research were of analytical grade.

## 2.2. Instruments

The isolated phenolic compounds, caffeic acid and rosmarinic acid, were elucidated with a Bruker AM 500 [ $^1\text{H}$  nuclear magnetic resonance (NMR) at 500 MHz,  $^{13}\text{C}$  NMR at 125 MHz] spectrometer (Bruker, Karlsruhe, Germany) using  $\text{DMSO}-d_6$  with tetramethylsilane. Two phenolic contents were performed using an Agilent 1200 UPLC series (Agilent Technologies, Inc., Wilmington, DE, USA) including a quaternary gradient pump and an Agilent 1200 series photodiode array detector. For analysis of multiple compositions in perilla leaves, the UPLC system (Agilent 1200 series) was coupled with triple quadrupole time-of-flight tandem mass spectrometry (LC/MS-Triple TOF 5600<sup>+</sup>; AB SCIEX, Foster City, CA, USA) equipped with an electrospray ionization (ESI) interface. To evaluate the antioxidant capacities concerning the scavenging effects against DPPH and ABTS radicals, UV–Vis absorption spectra were measured on a Beckman DU650 spectrophotometer (Beckman Coulter, Fullerton, CA, USA).

## 2.3. Sample preparation and calibration curves

The dried leaves of green perilla were pulverized (60 mesh) using an HR 2860 coffee grinder (Philips, Drachten, Netherlands) for 3 minutes. The powdered leaves (1.0 g) were extracted with methanol (10 mL) in a shaking incubator for 8 hours at room temperature. The supernatant was centrifuged at 3000g for 3 minutes using VS-6000CFN (VISION, Seoul, Korea), and then filtered through a 0.45- $\mu\text{m}$  syringe filter (Whatman Inc., Maidstone, UK) prior to UPLC analysis. For the calibration curve, the peak area of the identified phenolic compounds (rosmarinic acid and caffeic acid) was integrated with the UPLC chromatogram at 330 nm and plotted against the concentration to create a linear curve. The stock solutions of standard material were prepared with methanol to obtain a 1 mg/mL concentration, and calibration curves were made by dilution of each stock solution in methanol to six concentrations (1  $\mu\text{g/mL}$ , 5  $\mu\text{g/mL}$ , 10  $\mu\text{g/mL}$ , 20  $\mu\text{g/mL}$ , 50  $\mu\text{g/mL}$ , and 100  $\mu\text{g/mL}$ ). The correlation coefficients ( $r^2$ ) of two phenolic standards were detected, higher than 0.999.

## 2.4. UPLC conditions

The UPLC analysis was carried out using an Agilent Technologies 1290 Infinity series instrument coupled to a binary pump, a photodiode array detector, an autosampler, and a column compartment (kept at 25°C). A 5- $\mu\text{L}$  sample of the methanol extract in perilla leaves was injected into an analytical reverse phase column (Poroshell 120 EC-C18, 150  $\times$  2.1 mm, I.D., 4  $\mu\text{m}$ ). Separation was performed with 0.1% formic acid in water (eluent A) and methanol (eluent B). The gradient elution program was follows: 0–5 minutes, 20% B; 20 minutes, 35% B; 40 minutes, 75% B; and then held for

10 minutes prior to returning to the initial conditions. The total running time was 20 minutes at a flow rate of 0.4 mL/min, and the wavelength of detection was recorded at 330 nm.

## 2.5. UPLC-ESI-Q-TOF-MS/MS conditions

The metabolite profile in the methanol extract of perilla leaves was analyzed on a triple TOF 5600<sup>+</sup> system coupled to the UPLC system. The mass spectrometer was operated in the negative ESI mode with Duo-Spray source, and the mass scan range was set at  $m/z$  100–1000 for both TOF-MS and TOF-MS/MS scan using a resolution of 2700. The following parameter conditions were used: ion spray voltage, –4500 V; ion source heater, 500°C; curtain gas, 25 psi; nebulizer gas 50°C (GS1 50 and GS2 50); collision energy, –10 eV; declustering potential –100. The analyst TF software (version 1.7) combined with the information-dependent acquisition packing was used to acquire the MS/MS data. The mobile phase was composed of 0.1% formic acid in water (elution A) and methanol (elution B) using a gradient elution of 30% elution B (0–5 minutes), from 30% to 50% of elution B (5–20 minutes), from 50% to 90% elution B (20–40 minutes), and from 90% to 100% of elution B (40–45 minutes).

## 2.6. Antioxidant activity

In order to measure antioxidant abilities, the DPPH and ABTS radical scavenging methods were used in this research. The DPPH radical scavenging effect in the methanol extract of perilla leaves was determined as described by Lee et al [6]. The grounded leaves (1.0 g, 60 mesh) were extracted with methanol (10 mL) for 6 hours at room temperature. The crude extract was filtered through Whatman No. 42 filter paper, and the supernatants were measured. BHT (positive control) or sample (0.1 mL) of various concentrations (300  $\mu\text{g/mL}$ , 200  $\mu\text{g/mL}$ , 100  $\mu\text{g/mL}$ , 50  $\mu\text{g/mL}$ , 20  $\mu\text{g/mL}$ , 10  $\mu\text{g/mL}$ , and 5  $\mu\text{g/mL}$ ) were mixed with 0.49 mL of methanol and 0.39 mL methanolic solution of 1mM DPPH. The reaction mixture was vortexed and left to stand for 30 minutes at room temperature in darkness. The scavenging activity was reported as a percentage using the following formula at 517 nm: DPPH radical scavenging effect (%) =  $(1 - \text{absorbance of sample} / \text{absorbance of control}) \times 100$ .

The scavenging capacity of the ABTS radical was defined as the ability of different substances to scavenge the  $\text{ABTS}^{*+}$  radical cation, in comparison with a positive control (Trolox) [6]. This radical cation was generated by reacting the 7mM  $\text{ABTS}^{*+}$  stock solution with 2.45mM potassium persulfate, and the mixture was maintained for 10 hours in the dark. The  $\text{ABTS}^{*+}$  stock solution was melted with ethanol to an absorbance of 0.70 at 734 nm. The  $\text{ABTS}^{*+}$  solution (0.9 mL) was mixed with 0.1 mL sample (300  $\mu\text{g/mL}$ , 200  $\mu\text{g/mL}$ , 100  $\mu\text{g/mL}$ , 50  $\mu\text{g/mL}$ , 20  $\mu\text{g/mL}$ , 10  $\mu\text{g/mL}$ , and 5  $\mu\text{g/mL}$ ) and then measured using a spectrophotometer. This effect was determined as a percentage using the following formula:  $\text{ABTS}^{*+}$  scavenging capacity (%) =  $[(\text{absorbance of control} - \text{absorbance of sample}) / \text{absorbance of control}] \times 100$ .

## 2.7. Statistical analysis

All determinations were based on three replicate samples, and the results for the caffeic acid and rosmarinic acid contents are shown as mean values. The results were determined using Microsoft Excel (Microsoft 2013; Microsoft Corporation, Roseville, IL, USA).

## 3. Results and discussion

### 3.1. Determination of two phenolic acid contents in the leaves of green perilla

Perilla contains remarkably high amounts of phytochemical contents for the manufacture of supplements with

therapeutic and preventive properties for human beneficial health effects, such as antioxidant, anticancer, and antidiabetic properties [18–21,29]. Among the various phytochemicals, caffeic acid and rosmarinic acid are of great interest in food and crop industries because of their attractive biological activities [6,19,28]. We evaluated the main phenolic acids in the methanol extract of perilla leaves using UPLC analysis. A representative UPLC chromatogram of two phenolic acids was produced within 25 minutes at a wavelength of 330 nm (Figure 1A). The identification of each peak in the sample was confirmed using the retention time in the chromatogram, and their retention times are as follows: peak 1 (caffeic acid),  $t_R = 9.8$  minutes and peak 2 (rosmarinic acid),  $t_R = 23.7$  minutes. To quantitatively analyze phenolic acids, calibration curves were constructed in the range of 100–1  $\mu\text{g/mL}$ , and their equations were calculated as  $y = 1352.35x +$

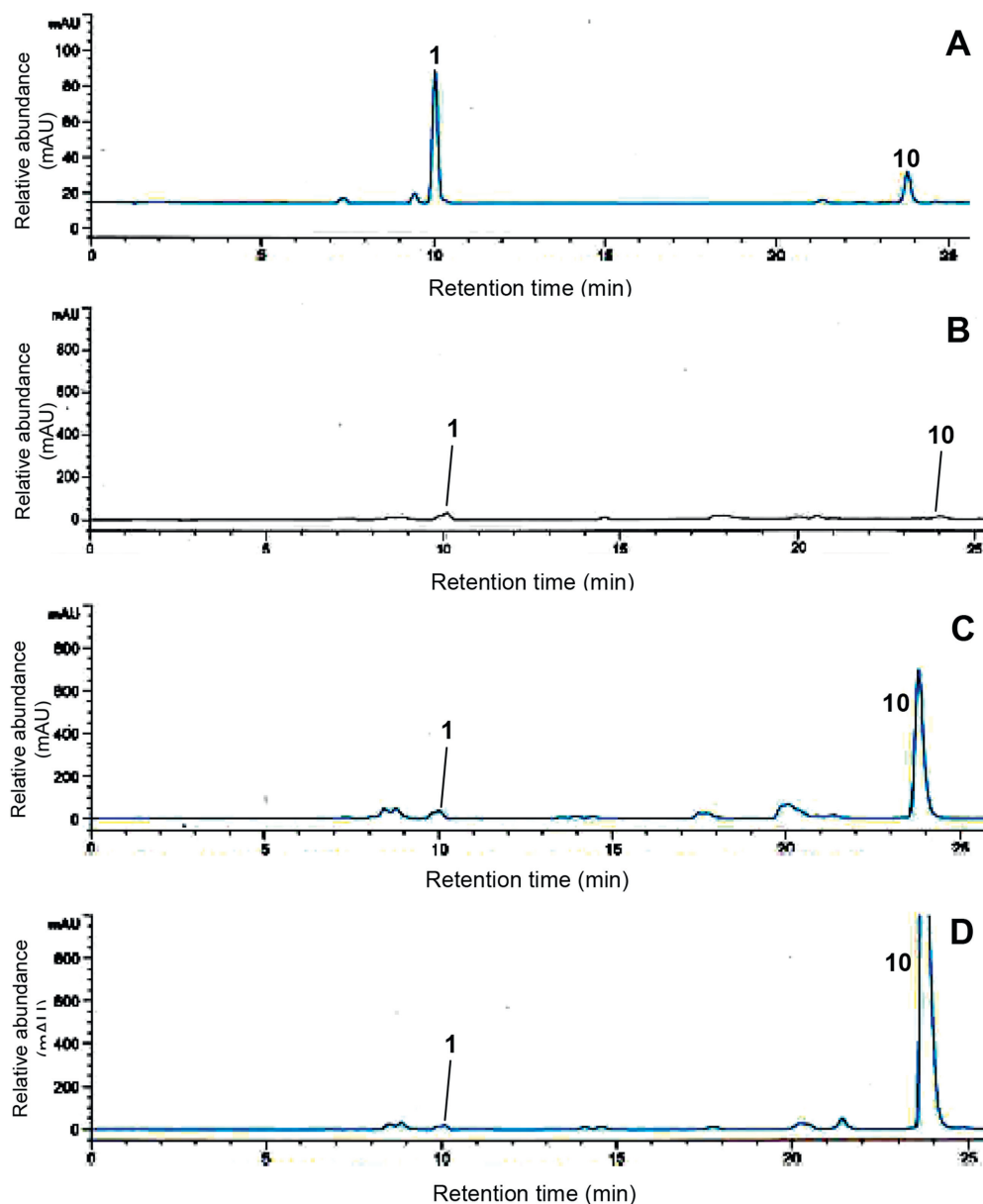


Figure 1 – UPLC chromatograms concerning caffeic acid (1) and rosmarinic acid (10) in the methanol extracts of green perilla leaves. (A) The isolated phenolic acid standard mixture. (B) Doryong sample. (C) Sinseong sample. (D) Bongmyeong sample.

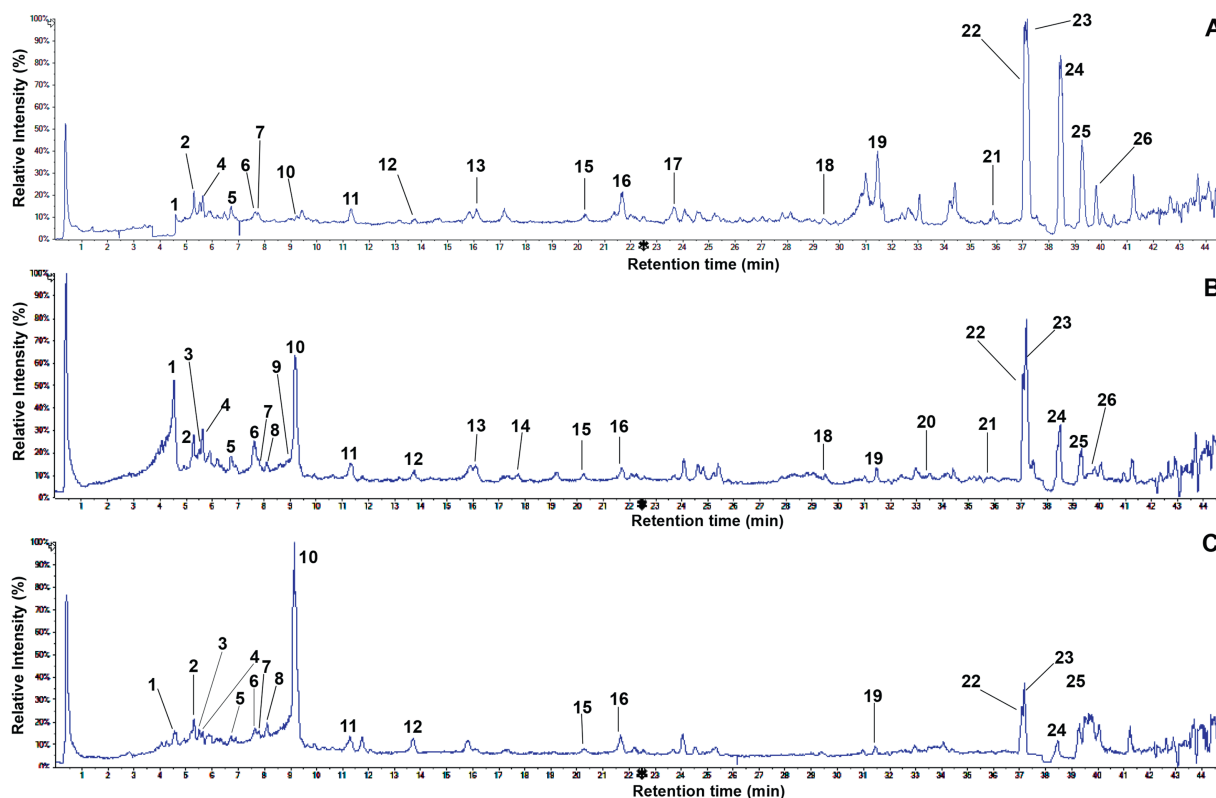


105.13 (caffeic acid,  $r^2 = 0.9998$ ) and  $y = 760.68x + 25.27$  (rosmarinic acid,  $r^2 = 0.9997$ ). We have chosen methanol for extraction of two phenolic acids in perilla leaves because it is an appropriate solvent for the maximum extraction of phenolic compounds [30,31]. The typical UPLC chromatograms of three perilla leaves are shown in Figures 1B–1D. As can be seen, the metabolite profiles exhibited similar patterns except one peak. Briefly, target phenolic acids, caffeic acid (1) and rosmarinic acid (10), presented in all samples and peak 10 exhibited significant differences. The highest rosmarinic acid content was 903.53  $\mu\text{g/g}$  in Bongmyeong (Figure 1D), whereas the lowest was 6.38  $\mu\text{g/g}$  in the Doryong sample (Figure 1B). Moreover, Sinseong showed the second highest content with 317.60  $\mu\text{g/g}$  (Figure 1C). Caffeic acid (1) exhibited little differences with 6.69  $\mu\text{g/g}$  (Doryong), 8.54  $\mu\text{g/g}$  (Sinseong), and 2.54  $\mu\text{g/g}$  (Bongmyeong), respectively. Although numerous studies have revealed that the major metabolites of perilla leaves are caffeic acid and rosmarinic acid, this study observed one major phenolic acid (rosmarinic acid) with various contents. These results suggest that the distribution of phenolic acids in natural plants may be influenced by various factors including environmental stress (light, temperature, region, and moisture) and agronomic conditions [1,7,32]. Their contents may also be affected by cultivars, sowing periods, years, and genetics, as previously reported data [6,7]. Our work agreed with earlier studies reporting the distribution of phenolics in many foods and crops [27,32,33].

### 3.2. Characterization of metabolites using UPLC-ESI-Q-TOF-MS/MS analysis

It is commonly demonstrated that the UPLC system coupled with MS is an excellent analytical method for simultaneously identifying complicated metabolites owing to the product ions produced from the fragmentation of a selected precursor ion [32,34]. Therefore, metabolite profiles in the leaves of three perilla samples were tentatively investigated by UPLC-ESI-Q-TOF-MS/MS through the full scan negative ESI mode. In the current research, a complete chromatographic separation of various metabolites, including major and minor peaks, was reached within 45 minutes (Figure 2). Their compositions were observed to have significant differences in samples. Table 1 shows the retention times, elementary compositions, and mass spectral data (molecular and fragment ions) of the identified metabolites by comparison with those of previous studies. First, we identified 21 peaks from the Doryong sample. Among the various peaks, four major compositions (peak 22,  $t_R = 37.1$  minutes; peak 23,  $t_R = 37.2$  minutes; peak 24,  $t_R = 38.5$  minutes; and peak 25,  $t_R = 39.3$  minutes) were detected with identical molecular ions  $[\text{M}-\text{H}]^-$  at  $m/z$  277.2165,  $m/z$  277.2174,  $m/z$  279.2332, and  $m/z$  255.2336, representing approximately 80% of the total peak area (Figure 2A).

The product ion mass spectrum of peak 1 ( $t_R = 4.5$  minutes) showed as deprotonated ion  $[\text{M}-\text{H}]^-$  at  $m/z$  179.0346. In the MS/MS spectrum, the distinctive product ion at  $m/z$  135.0446



**Figure 2** – Representative total ion chromatograms of the methanol extracts of green perilla leaves obtained by UPLC-ESI-Q-TOF/MS in negative ion mode: (A) Doryong sample. (B) Sinseong sample. (C) Bongmyeong sample. UPLC-ESI-Q-TOF/MS = ultra high performance liquid chromatography coupled with electrospray ionization quadrupole time-of-flight mass spectrometry.

Table 1 – Metabolite profile in methanol extract of green perilla leaves by UPLC-ESI-Q-TOF-MS/MS analysis.

Peak	$t_R$ (min)	Formula	Calculated mass [M] (m/z)	Calculated mass [M-H] <sup>−</sup> (m/z)	Observed molecular ion in MS [M-H] <sup>−</sup> (m/z)	Observed fragment ions in MS and MS/MS (m/z)	Identification	Reference
1	4.5	C <sub>9</sub> H <sub>8</sub> O <sub>4</sub>	180.0423	179.0344	179.0346	179.0336, 161.0265, 135.0446	Caffeic acid	Lee et al [6], Zhou et al [35], Chen et al [41]
2	5.3	C <sub>18</sub> H <sub>28</sub> O <sub>9</sub>	388.1733	387.1655	387.1647	387.1649, 207.1022	5'-Gluco-pyranosyxyjasmanic acid	Fujita et al [36]
3	5.9	C <sub>17</sub> H <sub>26</sub> O <sub>10</sub>	390.1526	389.1448	389.1451	167.0358	Loganin	Song et al [51], Ye et al [52]
4	6.2	C <sub>24</sub> H <sub>36</sub> O <sub>13</sub>	522.1373	521.1295	521.1847	521.1847, 359.0757, 285.0391	Rosmarinic acid-3-O- glucoside	Lee et al [6], Zhou et al [35]
5	6.7	C <sub>27</sub> H <sub>26</sub> O <sub>17</sub>	622.1170	621.1092	621.1109	621.1109, 351.0582, 269.0451, 113.0225	Apiogenin-7-O-digluconide	Kaufmann et al [38]
6	7.6	C <sub>21</sub> H <sub>18</sub> O <sub>12</sub>	462.0798	461.0720	461.0723	461.0723, 285.0408	Scutellarein-7-O-glucuronide	Yamazaki et al [39], Kaufmann et al [38]
7	7.8	C <sub>21</sub> H <sub>20</sub> O <sub>11</sub>	448.1006	447.0927	447.0907	447.0926, 285.0394	Luteolin-7-O- glucoside	Lee et al [6], Chen et al [41]
8	8.1	C <sub>36</sub> H <sub>32</sub> O <sub>16</sub>	720.1690	719.1612	719.1601	719.1601, 359.0763, 197.0417, 161.0224	Sagerinic acid	Lu and Foo [53]
9	9.0	C <sub>21</sub> H <sub>18</sub> O <sub>11</sub>	446.0849	445.0771	445.0779	445.0779, 269.0460, 176.0344, 161.0253	Apiogenin-7-O-glucuronide	Kaufmann et al [38]
10	9.2	C <sub>18</sub> H <sub>16</sub> O <sub>8</sub>	360.0845	359.0773	359.0772	359.0772, 197.0455, 179.0360, 161.0243	Rosmarinic acid	Lee et al [6], Zhou et al [35]
11	11.3	C <sub>19</sub> H <sub>18</sub> O <sub>8</sub>	374.1002	373.0923	373.1871	373.1880, 161.0758, 148.0529	Rosmarinic acid methyl ester	Zhou et al [35]
12	13.8	C <sub>20</sub> H <sub>36</sub> O <sub>12</sub>	468.2207	467.2129	467.2119	467.2119, 289.1660, 161.0462	N-Octanoylsucrose	Tirupati et al [43]
13	16.1	C <sub>19</sub> H <sub>24</sub> O <sub>4</sub>	316.1675	315.1596	315.0875	164.9840, 119.0514	trans-p-Menth-8-en-yl caffeate	Tada et al [44]
14	17.2	C <sub>15</sub> H <sub>10</sub> O <sub>5</sub>	270.0528	269.0450	269.0459	269.0445, 117.0359	Apigenin	Lee et al [6], Zhou et al [35]
15	20.2	C <sub>15</sub> H <sub>12</sub> O <sub>4</sub>	256.0736	255.0657	255.0658	255.0674, 213.0565, 145.0652	Liquiritigenin	Liu et al [45]
16	21.8	C <sub>18</sub> H <sub>32</sub> O <sub>5</sub>	328.2250	327.2171	327.2174	233.1152, 229.1437	Not identified	—
17	23.7	C <sub>16</sub> H <sub>12</sub> O <sub>6</sub>	300.0634	299.0556	299.0921	299.0921, 284.0715, 269.0450, 241.0508	Chrysoeriol	Lee et al [6]
18	29.5	C <sub>30</sub> H <sub>48</sub> O <sub>5</sub>	488.3502	487.3423	487.3422	487.3432, 469.3319, 249.1881	Tomentic acid	Chaturvedula and Prakash [48]
19	31.5	C <sub>30</sub> H <sub>48</sub> O <sub>4</sub>	472.3553	471.3474	471.3477	471.3397, 361.1995, 315.1947, 293.2123, 275.2013	Corosolic acid	Chaturvedula and Prakash [48]
20	33.0	C <sub>34</sub> H <sub>58</sub> O <sub>16</sub>	722.3725	721.3647	721.3660	721.3660, 470.3346, 451.3192, 277.2169, 235.0860	Palmitoleic–linolenic glucoside	Pierson et al [50]
21	35.4	C <sub>32</sub> H <sub>60</sub> O <sub>16</sub>	700.3881	699.3803	699.3812	397.2595	Palmitic–oleic glucoside	Pierson et al [50]
22	37.1	C <sub>18</sub> H <sub>30</sub> O <sub>2</sub>	278.2246	277.2168	277.2165	277.2164, 127.0762	α-Linolenic acid	Pierson et al [50]
23	37.2	C <sub>18</sub> H <sub>30</sub> O <sub>2</sub>	278.2246	277.2168	277.2174	277.2170	Linolenic acid	Pierson et al [50]
24	38.5	C <sub>18</sub> H <sub>32</sub> O <sub>2</sub>	280.2402	279.2324	279.2332	279.2334	Linoleic acid	Pierson et al [50]
25	39.3	C <sub>16</sub> H <sub>32</sub> O <sub>2</sub>	256.2402	255.2324	255.2329	255.2336	Palmitic acid	Pierson et al [50]
26	39.8	C <sub>18</sub> H <sub>34</sub> O <sub>2</sub>	282.2559	281.2481	281.2480	281.2490	Oleic acid	Pierson et al [50]

$t_R$  = retention time; UPLC-ESI-Q-TOF-MS/MS = ultra high performance liquid chromatography coupled with electrospray ionization quadrupole time-of-flight mass spectrometry.

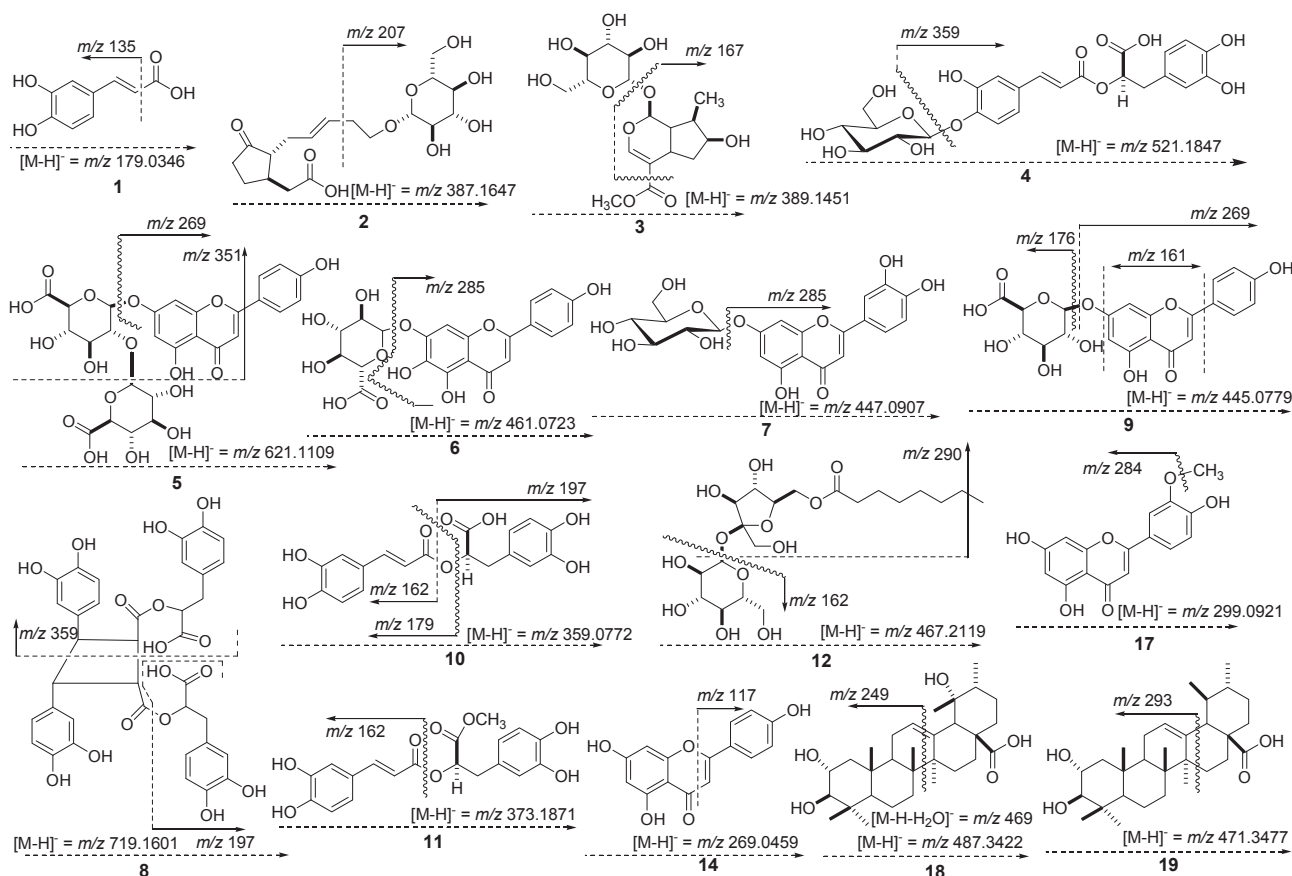
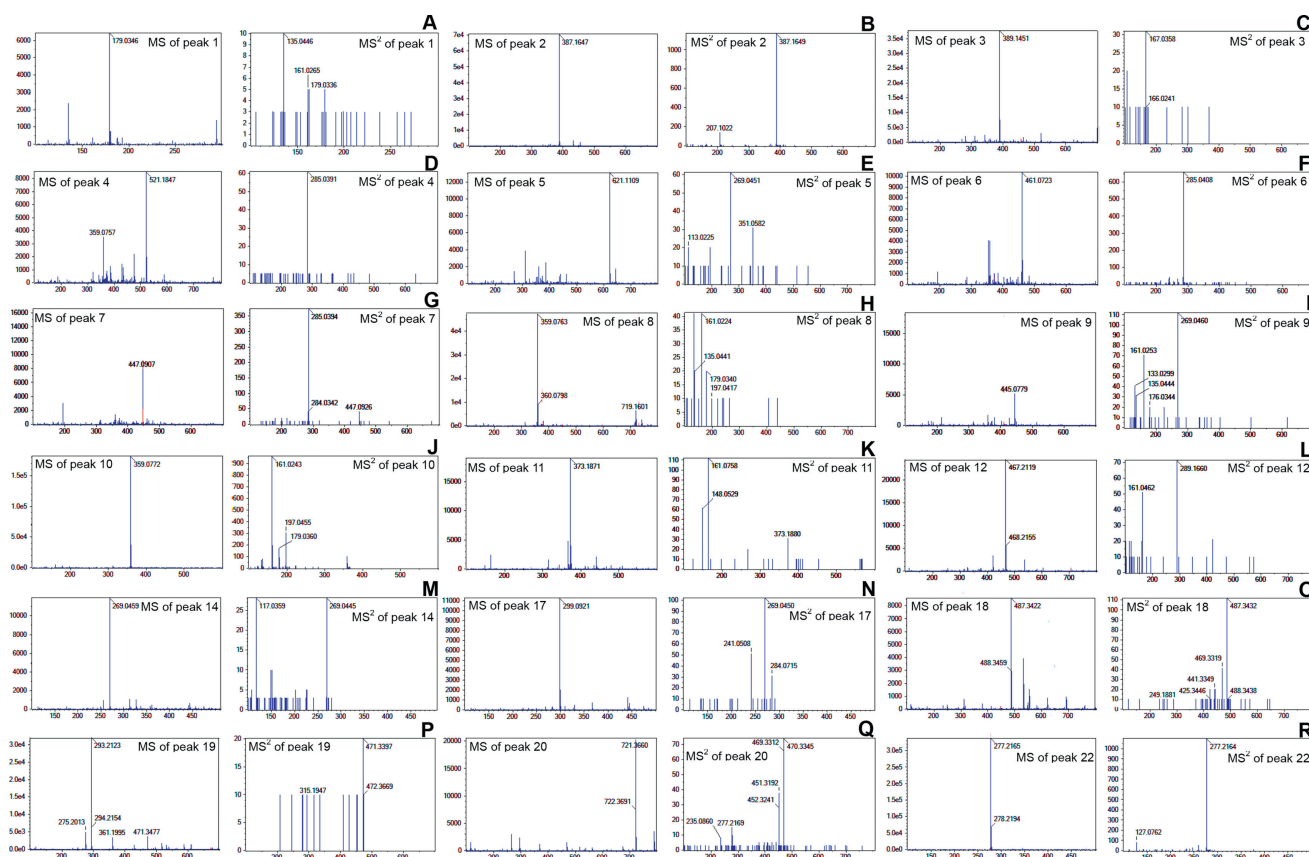


Figure 3 – Mass fragmentation patterns of metabolite compositions in the leaves of green perilla.

( $[M-H]^- - 44$  amu) corresponded to the loss of a carboxylic acid ( $-\text{COOH}$ ) moiety (Figures 3 and 4A). According to the fragment interpretation and the published data, peak 1 was confirmed to be caffeic acid (1) with the elemental formula  $\text{C}_9\text{H}_8\text{O}_4$  (Figures 4A and 5) [19,35]. The TOF-MS data of peak 2 ( $t_R = 5.3$  minutes) was detected with the precursor ions at  $m/z$  387.1647 ( $[M-H]^-$ ), and the fragmentation patterns of the MS/MS spectrum were detected with signals at  $m/z$  387.1649 and  $m/z$  207.1022 ( $[M-H]^- - 180$  amu, loss of glucopyranosy moiety) (Figures 3 and 4B). This peak was identified as 5'-glucopyranosyoxymasmanic acid (2) ( $\text{C}_{18}\text{H}_{28}\text{O}_9$ ) by comparison with jasmonoid glucoside of the earlier data [36]. Subsequently, peak 4 ( $t_R = 6.2$  minutes) presented a deprotonated molecular ion  $[M-H]^-$  at  $m/z$  521.1847 and one fragment ion at  $m/z$  359.0757 in negative ESI-TOF-MS analysis (Figure 4D). The fragmentation at  $m/z$  359.0757 was determined to be characteristic of rosmarinic acid by the loss of a glucose residue (162 amu) (Figure 3) [35]. Therefore, peak 4 was assumed to be rosmarinic acid-3-O-glucoside (4) with the formula  $\text{C}_{24}\text{H}_{26}\text{O}_{13}$  (Figure 5) [6,35]. Full TOF-MS scan analysis of peak 5 ( $t_R = 6.7$  minutes) showed an identical molecular ion  $[M-H]^-$  ion at  $m/z$  621.1109 (Figure 4E). In addition, the MS/MS spectrum possessed two peaks including one major fragment ion at  $m/z$  269.0451 and one minor ion at  $m/z$  351.0582 (Figure 4E). The fragment ion ( $m/z$  269.0451) was characteristic of apigenin aglycone through the loss of the deprotonated glucuronide residues (C-7 site,  $2 \times 176$  amu) (Figure 3) from the molecular

ion  $[M-H]^-$  at  $m/z$  621.1109 [35]. The remaining MS/MS fragment ion ( $m/z$  351.0582) was observed as the loss of the glucuronide moiety (176 amu) and 2-phenyl group (93 amu) in flavone skeleton (Figure 3) [37]. Based on these observations, peak 5 was identified as apigenin-7-O-diglucuronide (5) ( $\text{C}_{27}\text{H}_{26}\text{O}_{17}$ ) (Figure 5) [38]. The ESI-TOF-MS in the negative ion mode of peak 6 ( $t_R = 7.6$  minutes) exhibited the deprotonated precursor  $[M-H]^-$  ion at  $m/z$  461.0723 (Figure 4A). The fragment ion in the MS/MS spectrum exhibited one major peak at  $m/z$  285.0408 (Figure 4F). This fragment ion ( $m/z$  285.0408) was formed by the loss of  $m/z$  176 fragment ( $m/z$  461  $\rightarrow$   $m/z$  285;  $[M-H]^- - 176$  amu) (Figure 3) [37], indicating the characteristic of a glucuronide moiety [37,38]. Furthermore, the major fragmentation at  $m/z$  285.0428 ( $-176$  amu) was characterized as luteolinidin moiety (Figure 3), in agreement with a previous study [39]. On the basis of these observations, this compound was tentatively identified as scutellarein-7-O-glucuronide (6) with the elemental formula of  $\text{C}_{21}\text{H}_{18}\text{O}_{12}$  [39,40]. In the TOF/MS and MS/MS spectra, peak 7 ( $t_R = 7.8$  minutes) was detected as a deprotonated ion  $[M-H]^-$  at  $m/z$  447.0907 and one major fragment ion at  $m/z$  285.0394 ( $[M-H]^- - 162$  amu) (Figures 3 and 4G). The main fragmentation at  $m/z$  285.0394 resulted from the loss of a glucose moiety, which corresponding to luteolin aglycone ( $m/z$  285) [6]. On the basis of the fragment interpretation, peak 7 has already been identified as luteolin-7-O-glucoside (7) with the formula  $\text{C}_{21}\text{H}_{20}\text{O}_{11}$  (Figure 3). Also, this peak was confirmed by comparison of data reported in



**Figure 4** – The negative ESI-TOF/MS and MS/MS spectra of metabolites in the leaves of green perilla. (A) caffeic acid (1), (B) 5'-gluco-pyranosyoxyjasmonic acid (2), (C) loganin (3), (D) rosmarinic acid-3-O-glucoside (4), (E) apigenin-7-O-diglucuronide (5), (F) scutellarein-7-O-glucuronide (6), (G) luteolin-7-O-glucoside (7), (H) sagerinic acid (8), (I) apigenin-7-O-glucuronide (9), (J) rosmarinic acid (10), (K) rosmarinic acid methyl ester (11), (L) *N*-octanoylsucrose (12), (M) apigenin (14), (N) chrysoeriol (17), (O) tomentonic acid (18), (P) corosolic acid (19), (Q) palmitoleic–linolenic glucoside (20), and (R)  $\alpha$ -linolenic acid (22).

ESI = electrospray ionization; MS = mass spectrometry; TOF = time of flight.

previous studies [6,41]. The TOF mass scan analysis of peak 10 ( $t_R = 9.2$  minutes) exhibited a deprotonated molecular ion at  $m/z$  359.0772. The MS/MS fragmentation patterns possessed three fragment ions at  $m/z$  197.0455,  $m/z$  179.0360, and  $m/z$  161.0243 (Figure 4J). A fragment ion at  $m/z$  197.0455, which was formed by the loss of  $m/z$  162 fragment ( $m/z$  359  $\rightarrow$   $m/z$  197;  $[M-H]^- - 162$  amu) (Figure 3). This ion peak was determined to be the characteristic of a caffeoyl moiety [42]. In addition, the fragment ion at  $m/z$  161.0243 corresponded to the deprotonated caffeoyl residue, and the fragmentation at  $m/z$  179.0360 was in agreement with the pattern of peak 1 (caffeic acid). As a result, peak 10 was tentatively assigned as rosmarinic acid (10) ( $C_{18}H_{16}O_8$ ) (Figure 3) [6,19,35]. The ESI-TOF-MS and MS/MS in the negative ion mode of peak 11 ( $t_R = 11.3$  minutes) presented the molecular ion at  $m/z$  373.1871 and two fragment ions at  $m/z$  161.0758 and  $m/z$  148.0529 (Figure 4K). The fragmentation at  $m/z$  161.0758 resulted from the deprotonated caffeoyl residue [42] as the ion pattern of peak 10 (Figure 3). Moreover, the pseudomolecular ion at  $m/z$  373.1871 was in agreement with the rosmarinic acid derivative of published data [35]. As discussed previously, this peak was elucidated as rosmarinic acid methyl ester (11) with the elemental composition of  $C_{19}H_{18}O_8$  (Figure 3) [6,35]. The product ion mass spectrum of peak 12

( $t_R = 13.8$  minutes) appeared as deprotonated ion  $[M-H]^-$  at  $m/z$  467.2119. In addition, three fragment ions at  $m/z$  289.1660 and  $m/z$  161.0462 were observed in the MS/MS spectrum (Figure 4L). A fragment ion was observed at  $m/z$  289.1660, which was formed by the loss of  $m/z$  178 fragment ( $m/z$  467  $\rightarrow$   $m/z$  289;  $\{[M-H]^- - (\text{glucose} + \text{methyl})\}$ ) (Figure 3). In brief, this ion peak ( $m/z$  289.1660) may correspond to the deprotonated fructose lipid residue, and the fragmentation ion at  $m/z$  161.0462 was characterized as the deprotonated glucose moiety [6,27]. Based on the described fragmentation ions, peak 12 was assumed to be *N*-octanoylsucrose (12) ( $C_{20}H_{36}O_{12}$ ). Although this composition was previously identified by *Synecocystis* sp. PCC [43], we have documented for the first time its presence in perilla plant. The negative ion mode of peak 13 possessed the molecular ion at  $m/z$  315.0875, and its chemical structure was identified as *trans-p*-menth-8-en-yl caffeate (13) ( $C_{19}H_{24}O_4$ ) based on the earlier literature [44]. Peak 15 was observed with a retention time at 20.2 minutes in UPLC-ESI-Q-TOF-MS chromatogram, and its mass measurement was observed at  $m/z$  255.0658 ( $[M-H]^-$ ). The chemical structure was tentatively assigned as liquiritigenin ( $C_{15}H_{12}O_4$ ) in reference to a previously reported study [45]. As far as we know, it is reported for the first time in perilla plant. Peak 16 was



detected as a molecular ion  $[M-H]^-$  at  $m/z$  327.2174 with the formula  $C_{18}H_{32}O_5$  and two fragment ions ( $m/z$  233.1152 and  $m/z$  229.1437). The above fragmentation ions may be formed by the losses of  $m/z$  94 and  $m/z$  98. However, with this peak, it was not possible to completely characterize the chemical structure. The TOF-MS and MS/MS analyses of peak 17 ( $t_R = 23.7$  minutes) presented the molecular ion  $[M-H]^-$  at  $m/z$  299.0921 with three fragmentations at  $m/z$  284.0715,  $m/z$  269.0450, and  $m/z$  241.0508 (Figure 4N). In particular, a fragment ion was found at  $m/z$  284.0715, which was formed by the loss of the  $m/z$  15 fragment (methyl moiety) (Figure 3) [6,46]. According to the molecular weight in published data [6], the structure of peak 17 was determined as chrysoeriol (17) ( $C_{16}H_{12}O_6$ ). The deprotonated molecular ion  $[M-H]^-$  at  $m/z$  487.3422 was found in the UPLC ESI-TOF-MS spectrum of peak 18 (Figure 4O). Furthermore, two fragment ions at  $m/z$  469.3319 and  $m/z$  249.1881 were produced from the parent ion at  $m/z$  487.3422 in MS/MS data (Figure 4O). The precursor ion at  $m/z$  469.3319 resulted from the loss of a  $H_2O$  ( $-18$  amu), and the fragmentation pattern of the ion at  $m/z$  249.1881 would correspond to the molecular ion of diterpenoid moiety regarding B/C ring retro-Diels-Alder cleavage (Figure 3) [47]. According to the above summarized fragmentation, peak 18 was tentatively characterized as tomentonic acid (18) ( $C_{30}H_{48}O_5$ ) [48]. The full mass scan analysis of peak 19 ( $t_R = 31.5$  minutes) was observed with  $[M-H]^-$  at  $m/z$  471.3477 and three fragment ions at  $m/z$  361.1995,  $m/z$  293.2123, and  $m/z$  275.2013 (Figure 4P). The major fragmentation at  $m/z$  293.2123

( $[M-H]^- - 178$  amu) resulted from the cleavage of the D-ring, which is in accordance with an earlier study [49]. Therefore, the fragmentation characteristics were attributed to the corosolic acid (19) ( $C_{30}H_{48}O_4$ ) (Figure 3) [48]. The ESI-TOF-MS spectrum of peak 21 ( $t_R = 35.4$  minutes) displayed a  $[M-H]^-$  ion at  $m/z$  699.3812, and its characteristic fragment ion was observed at  $m/z$  397.2595 ( $[M-H]^- - 302$  amu) in MS/MS analysis. Based on these lines of evidence, this composition was identified as palmitic–oleic glucoside (21), with the formula  $C_{32}H_{60}O_{16}$  [50]. The ESI-TOF-MS in negative ion mode of peaks 22–26 exhibited the molecular ions at  $m/z$  277.2165 (Figure 4R),  $m/z$  277.2174,  $m/z$  279.2332,  $m/z$  255.2329, and  $m/z$  281.2480. Their molecular weights and retention times were coincident with those of mango fatty acids in a previously reported study [50]. Therefore, five peaks were confirmed as  $\alpha$ -linolenic acid (22), linolenic acid (23), linoleic acid (24), palmitic acid (25), and oleic acid (26) (Figure 5).

Second, we examined the metabolite profile from the methanol extract of Sinseong sample to gather more information in the leaves of Korean green perilla (Figure 2B). Twenty-five compositions were characterized by the molecular weights of mass spectral data in previously reported studies (Figure 5). As illustrated in Figure 2B, metabolite peaks exhibited important differences by comparison with those of Doryong. In particular, caffeic acid (1), rosmarinic acid (10),  $\alpha$ -linolenic acid (22), and linolenic acid (23) were the predominant metabolites, representing approximately 80% of the total peak area (Figure 2B). Moreover, five compositions were newly

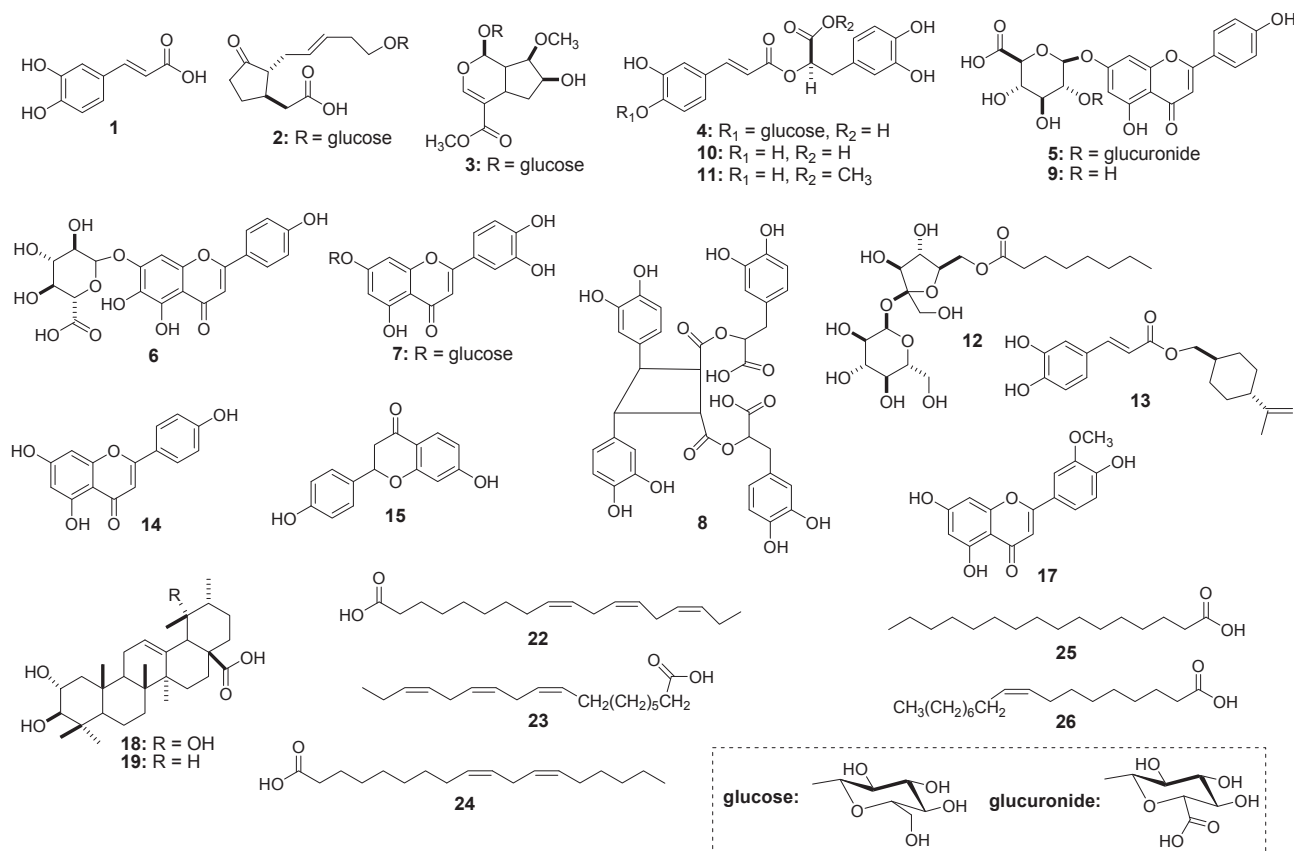


Figure 5 – Chemical structures of metabolites found in the methanol extract of green perilla leaves.

observed—loganin (3), sagerinic acid (8), apigenin-7-O-glucuronide (9), apigenin (14), and palmitoleic–linolenic glucoside (20)—with identical molecular ions  $[M-H]^-$  at  $m/z$  389.1451,  $m/z$  719.1601,  $m/z$  445.0779,  $m/z$  269.0459, and  $m/z$  721.3660 (Figure 4). In this study, we report the fragment patterns of UPLC-ESI-TOF-MS/MS data, as detailed below.

Peak 3 ( $t_R = 5.9$  minutes) possessed an identical molecular ion  $[M-H]^-$  at  $m/z$  389.1451 and the fragmentation pattern at  $m/z$  167.0358 (Figure 4C). The fragment ion ( $m/z$  167.0358,  $-222$  amu) resulted from the losses of glucose ( $-162$  amu) and ester ( $-60$  amu,  $-COOCH_3$ ) groups (Figure 3). By comparing its mass values with data from earlier reports [51,52], this peak was tentatively evaluated as loganin (3) ( $C_{17}H_{26}O_{10}$ ). To the best of our knowledge, this composition is the first reported in perilla plant. The product ion mass spectrum of peak 8 ( $t_R = 8.1$  minutes) presented the molecular ion  $[M-H]^-$  at  $m/z$  719.1601 and a fragment ion at  $m/z$  359.0763, which was formed by the loss of the  $m/z$  360 fragment (Figure 4H). The fragment ion resulted from the loss of a rosmarinic acid moiety in dimer skeleton (Figure 3) by comparison with an earlier report [53]. A fragment ion at  $m/z$  197.0417 in the MS/MS spectrum, which was formed by the loss of  $m/z$  161.0224 fragment ( $m/z$  359  $\rightarrow$   $m/z$  197;  $[M-H]^- - 162$  amu) (Figures 3 and 4H), was related with the characteristic of a caffeoyl group [6]. Briefly, peak 8 has already been identified as rosmarinic acid dimer, where dimerization had occurred by olefinic moieties. According to these observations, peak 8 was tentatively assigned as sagerinic acid (8) ( $C_{36}H_{32}O_{16}$ ). We identified for the first time that peak 8 was present in perilla plant. Using UPLC-ESI-TOF/MS, peak 9 was detected with the precursor ion of  $m/z$  445.0779 ( $[M-H]^-$ ) at the retention time of 9.0 minutes (Figure 4I). Moreover, two major fragment ions at  $m/z$  269.0460 and  $m/z$  161.0253 as well as one minor peak at  $m/z$  176.0344 were observed in the TOP/MS/MS spectrum (Figure 4I). The fragmentation patterns of this peak were similar to those of peak 5; in particular, the fragmentation at  $m/z$  269.0460 ( $-176$  amu) was characterized as apigenin aglycone. A minor ion at  $m/z$  176.0344 ( $[M-H]^- - 269$ ), which would correspond to the loss of an apigenin moiety, and a major ion at  $m/z$  161.0253 resulted from the losses of a glucuronide ( $m/z$  176) group as well as 2-phenyl ( $m/z$  93) and hydroxyl ( $m/z$  17) moieties of apigenin aglycone. Thus, peak 9 was assumed to be apigenin-7-O-glucuronide (9) with the formula  $C_{21}H_{18}O_{11}$  (Figure 5) [38]. Peak 14 ( $t_R = 17.2$  minutes) possessed an identical molecular ion  $[M-H]^-$  at  $m/z$  269.0459 as the fragmentation patterns of peaks 5 and 9 (Figure 4M). This ion was characteristic of apigenin aglycone by comparison with published data [6]. In addition, a fragment ion at  $m/z$  117.0359 of the MS/MS spectrum may be attributed to the result from the cleavage of C–ring bonds in deprotonated flavonoids (Figures 3 and 4M) [54]. This structure was tentatively assigned as apigenin (14) (Figure 5) [6,35]. The ESI TOF/MS profile of peak 20 ( $t_R = 33.0$  minutes) that exhibited an identical molecular ion  $[M-H]^-$  at  $m/z$  721.3660 was characterized by an ion signal at  $m/z$  721.3660, and the MS/MS fragment ions were observed with four values, including  $m/z$  470.3346,  $m/z$  451.3192,  $m/z$  277.2169, and  $m/z$  235.0860 (Figure 4Q). On the basis of the lines of evidence presented, this structure was coincident with those of palmitoleic–linolenic glucoside (20) in mango fatty acids [50]. The crude methanol extract of the Bongmyeong

sample was also screened for metabolite profile on Korean green perilla. This material was observed to have various minor peaks in total ion chromatogram (Figure 2C), and the individual component was not fully characterized. We revealed 18 metabolites, including the identified compositions of Doryong and Sinseong samples (Figure 5). Interestingly, only one main phenolic acid (rosmarinic acid 10) was found in the chromatography, representing approximately 55% of the total peak area (Figure 2C). Consequently, metabolite compositions and their contents in the leaves of Korean green perilla showed remarkable differences in each sample. This phenomenon suggests that metabolites may be affected by multiple factors, such as environmental stresses, cultivars, and genetics [5–7,27,39]. In addition, our result may be an important factor for the determination of the quality of green perilla.

### 3.3. Antioxidant effects against DPPH and ABTS radicals from the leaves of green perilla

The evaluation of antioxidant capacity is commonly used as a screening method for crops, foods, fruits, vegetables, and edible plants [6,19,45]. Specifically, the scavenging activities against radical sources have been used in the screening of antioxidants with regard to phenolic compounds, flavonoids, carotenoids, and anthocyanins for beneficial human health properties [3,6,9,12,27]. Among various radical sources, DPPH and ABTS radical assays are of great interest in food and crop industries owing to their simple quality control and reproducibility [6,9,35]. For these reasons, we investigated antioxidant properties including DPPH and ABTS radicals in the present research. The scavenging effects of green perilla leaves were determined by comparing the percentage inhibition through the formation of DPPH and ABTS radicals by the methanol extract of each sample, BHT (positive control), and Trolox (positive control). The scavenging abilities against DPPH and ABTS radicals of the samples and positive controls increased with increasing concentrations (5  $\mu\text{g/mL}$ , 10  $\mu\text{g/mL}$ , 20  $\mu\text{g/mL}$ , 50  $\mu\text{g/mL}$ , 100  $\mu\text{g/mL}$ , 200  $\mu\text{g/mL}$ , and 300  $\mu\text{g/mL}$ ). Although 100% scavenging capacity was observed at 200  $\mu\text{g/mL}$  and 300  $\mu\text{g/mL}$ , the antioxidant properties on the radical scavengers were carried out at 100  $\mu\text{g/mL}$  because of the dose-dependent changes and comparison of the inhibition rates in samples. In the current study, the rank order of the scavenging effects against DPPH radical was consistent with the results of the ABTS assay. The highest scavenging effects were observed in the Sinseong sample with 86% (DPPH) and 90% (ABTS) at a concentration of 100  $\mu\text{g/mL}$ . These values were slightly higher than BHT (82%) and Trolox (86%) of positive controls (Table 2).

In more detail, the DPPH and ABTS radical scavenging abilities of different perilla leaves were, in increasing order, as follows: Sinseong (86 and 90%) > Bongmyeong (71 and 84%) > Doryong (63 and 73%) (Table 2). These results reveal that the identified metabolite contents in the methanol extract of perilla leaves may be the major portion of antioxidant properties through DPPH and ABTS radical scavenging capacities, as described in previous studies [6,19,44]. The unidentified peaks may also be potential factors in determining the two radical scavenging properties [55]. Interestingly, three samples showed higher ABTS radical capacities than the results on

**Table 2 – Antioxidant effects against DPPH and ABTS radicals from the leaves of three green perilla samples.**

Sample	Radical scavenging activity (%) <sup>a</sup>	
	DPPH radical	ABTS radical
Doryong	63 <sup>b</sup>	73
Sinseong	86	90
Bongmyeong	71	84
Positive control (DPPH: BHT) (ABTS: Trolox)	82	86

ABTS = 2,2'-azino-bis(3-ethylbenzthiazoline-6-sulphonic acid); BHT = butylated hydroxytoluene; DPPH = 2,2-diphenyl-1-picrylhydrazyl.

<sup>a</sup> All values of samples and positive controls are expressed as the mean of triplicate determinations.

<sup>b</sup> Radical scavenging effects of the methanol extracts of samples and positive controls were carried out at 100 µg/mL.

DPPH radical. This phenomenon suggests that DPPH radical may be attributed to the scavenging activities regarding hydrogen donating compositions of various metabolites, whereas ABTS radical was related to the scavenging abilities of chain breaking and hydrogen donating metabolites, which was shown in previous studies [6,7]. Additionally, it is well documented that perilla contains many metabolites, namely, phenolic acids, flavonoids, triterpenoids, and fatty acids [2,6,17,19,27]. Even though all of the identified metabolites do not indicate the antioxidant effects, phenolic acid, flavonoids, and triterpenoid derivatives are responsible for the beneficial health properties such as antioxidant effects against radicals in previously published data [35,56–59]. Therefore, the metabolite contents may also be attributed to the scavenging activities on DPPH and ABTS radicals in the methanol extract of perilla leaves. As a result, the leaves of green perilla were observed to have potential antioxidant properties against DPPH and ABTS radicals because of their high metabolite contents. Moreover, this species may be considered a radical scavenger for food and nutraceutical uses.

#### 4. Conclusion

We have characterized metabolite profiles from the leaves of green perilla using UPLC-ESI-Q-TOF-MS/MS analysis. Twenty-five metabolites were demonstrated as six phenolic acids, two triterpenoids, eight flavonoids, seven fatty acids, and two glucosides. Among them, loganin (3), sagerinic acid (8), N-octanoylsucrose (12), and liquiritigenin (15) were reported for the first time in this species. The metabolite compositions exhibited remarkable differences in three samples, especially Doryong, which exhibited the highest fatty acid contents. Moreover,  $\alpha$ -linolenic acid (22), linolenic acid (23), and linoleic acid (24) showed high contents in Doryong and Sinseong samples. The well-known phenolic acid, rosmarinic acid (10), was also observed to have significant differences in samples as follows: Doryong, 6.38 µg/g; Sinseong, 317.60 µg/g; and Bongmyeong, 903.53 µg/g. In addition, the antioxidant properties such as DPPH and ABTS radical scavenging effects differed significantly according to the sample (100 µg/mL):

Doryong, 63% and 73%; Sinseong, 86% and 90%; and Bongmyeong, 71% and 84%. We believe that metabolites may be an important factor for the quality and antioxidant property of green perilla leaves. To the development of functional food and nutraceutical sources, future researches are needed to evaluate the human health beneficial properties such as anticancer, antidiabetic, antioxidant, anti-inflammatory, and antiatherogenic effects of various metabolites in green perilla.

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