



Supplementary Materials

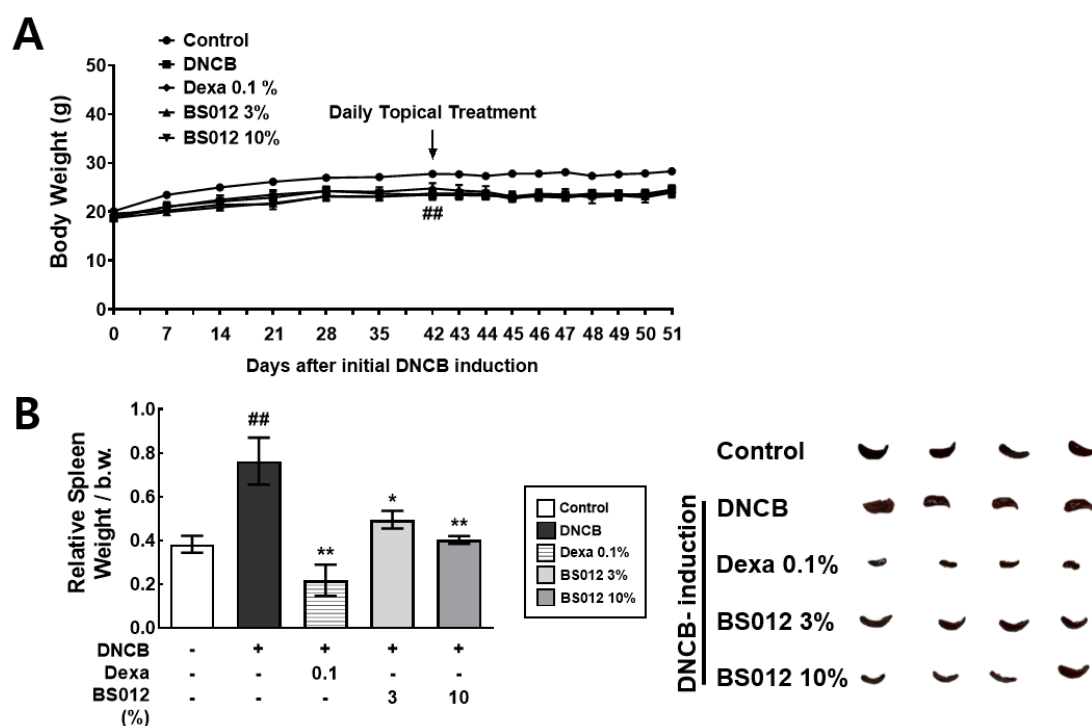


Figure S1. Relative spleen weight. The graphs represent mean \pm SEM, $n = 6$; ## $p < 0.01$ vs. control group; * $p < 0.05$, ** $p < 0.01$ vs. DNCB group, Dexa: Dexamethasone.

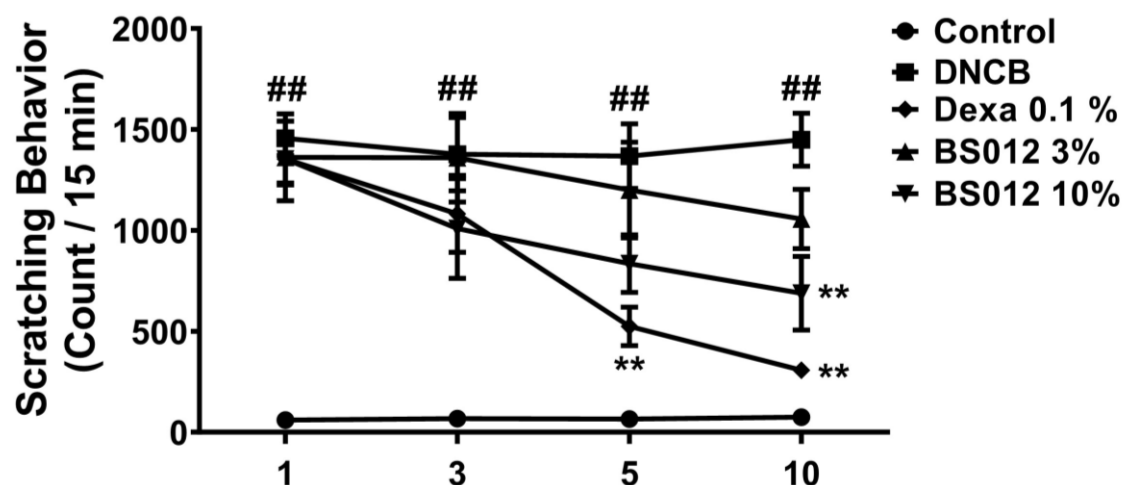


Figure S2. Effect of BS012 on the scratching behavior in NC/Nga mice (A) Total scratching behavior. Scratching behavior was quantified on days 1, 3, 5, and 10 post-treatment. The graphs represent mean \pm SEM, $n = 6$; ## $p < 0.01$ vs. control group; ** $p < 0.01$ vs. DNCB group, Dexa: Dexamethasone.

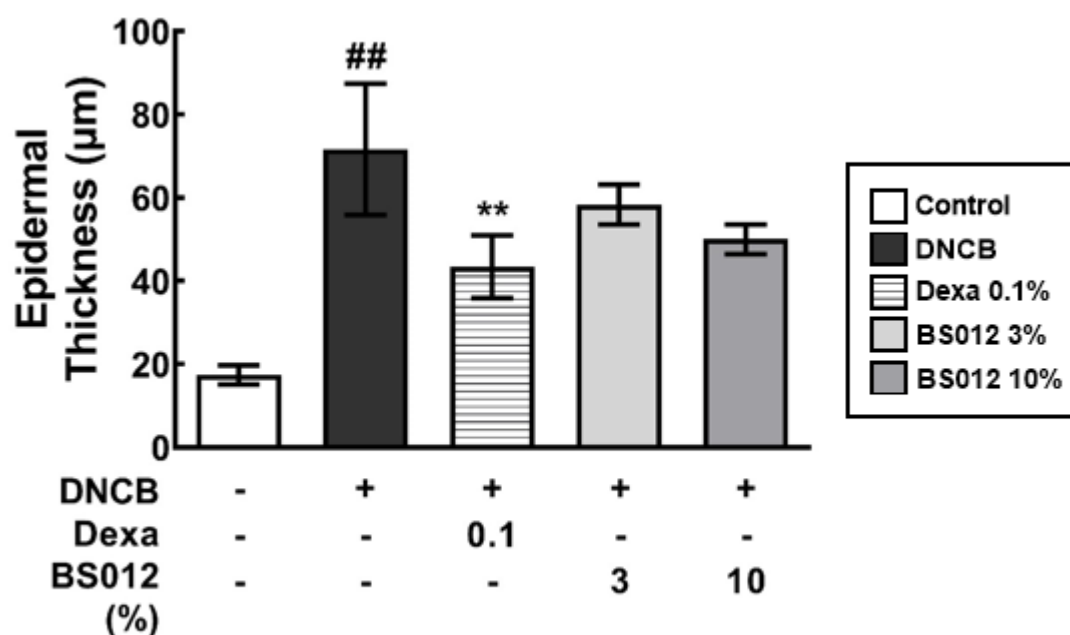


Figure S3. Effect of BS012 on the epidermal thickness in DNCB-induced NC/Nga mice. Epidermal thickness in the skin were evaluated using H&E staining. Data in the graphs are presented as mean \pm SEM. ^{##} $p < 0.01$ vs. control group; ^{**} $p < 0.01$ vs. DNCB-induced group, Dexa: Dexamethasone.

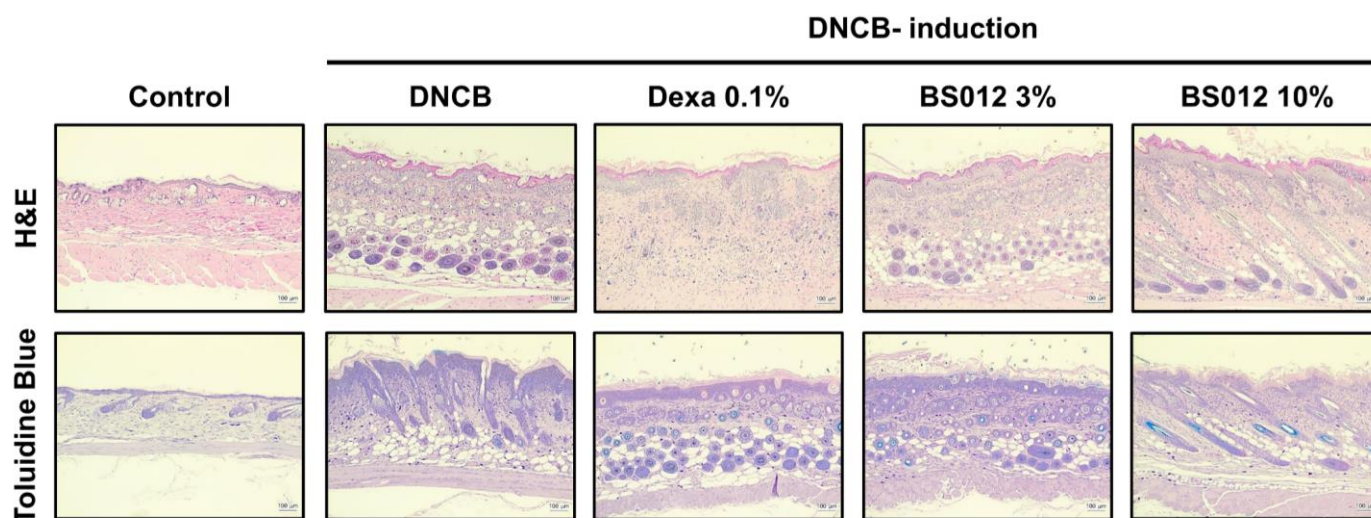


Figure S4. High-resolution original image of histological analysis. Examinations were conducted using H&E and toluidine blue staining at a magnification of 100 \times .

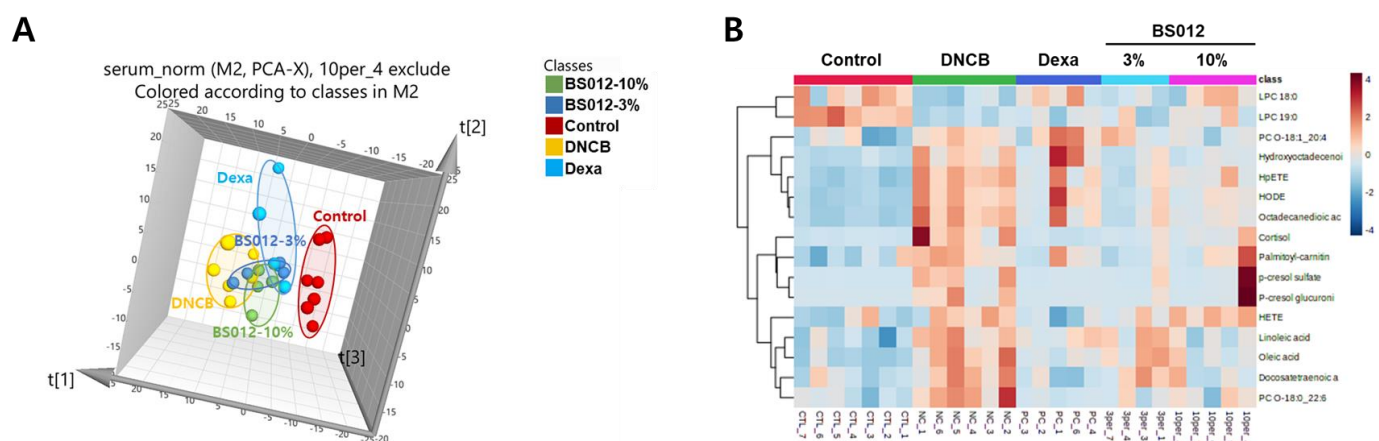


Figure S5. Effects of BS012 topical application on serum metabolome in DNCB-induced mice. (A) The principal component analysis (PCA) score plot derived from serum metabolomics. (B) Heatmap analysis using the 16 identified metabolites. The mean-centered relative abundances divided by the standard deviation of each variable. Dexa: Dexamethasone 0.1%.

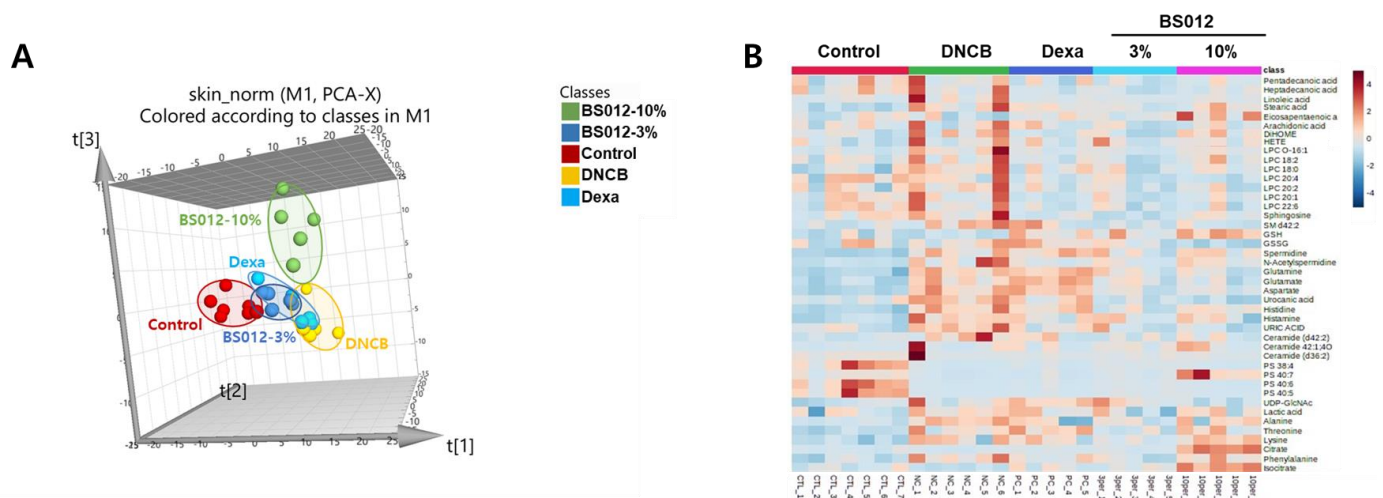


Figure S6. Effects of BS012 topical application on skin metabolome in DNCB-induced mice. (A) The principal component analysis (PCA) score plot derived from serum metabolomics. (B) Heatmap analysis using the 43 identified metabolites. The mean-centered relative abundances divided by the standard deviation of each variable. Dexa: Dexamethasone.

Table S1. List of identified metabolites significantly altered in serum from DNCB-induced mice with topical BS012 treatment.

Related metabolism	Identified metabolite	(ionization mode) Observed m/z	Mean value ^a					Trend ^b Control vs. DNCB ^d	Chage trend ^c		
			Control	DNCB	BS012 (%)		Dexa		DNCB vs. treated group		
					BS012 (%)				Dexa		
					3	10					
Fatty acid metabolism	Linoleic acid	(-) 279.231	0.183	0.312	0.296	0.224	0.259	↑**	↓	↓*	↓
	Oleic acid	(-) 281.247	0.061	0.174	0.156	0.108	0.100	↑**	↓	↓*	↓**
	HODE	(-) 295.226	0.036	0.224	0.109	0.133	0.197	↑**	↓	↓**	↓
	Hydroxyoctadecenoic acid	(-) 297.241	0.025	0.121	0.057	0.053	0.138	↑**	↓*	↓**	↑
	Octadecanedioic acid	(-) 313.236	0.029	0.163	0.077	0.087	0.114	↑**	↓*	↓**	↓
	HETE	(-) 319.226	2.640	3.251	2.976	3.433	2.520	↑*	↓	↓	↓*
	Docosatetraenoic acid	(-) 331.262	0.023	0.041	0.036	0.027	0.019	↑**	↓	↓	↓**
	HpETE	(-) 381.222	0.004	0.015	0.008	0.010	0.011	↑**	↓**	↓*	↓
Glycero-phospholipid metabolism	LPC 18:0	(-) 568.357	0.279	0.135	0.174	0.246	0.255	↓**	↑	↑	↑*
	LPC 19:0	(-) 582.372	0.084	0.038	0.044	0.056	0.046	↓**	↑	↑*	↑
	PC O-18:1_20:4	(-) 838.592	0.004	0.006	0.006	0.005	0.007	↑*	↓	↓*	↑

	PC O-18:0_22:6	(-) 864.608	0.006	0.010	0.006	0.007	0.006	↑**	↓*	↓	↓**
Others	Palmitoyl-carnitine	(+) 400.342	0.033	0.065	0.039	0.059	0.051	↑**	↓*	↓	↓
	p-Cresol sulfate	(-) 187.006	0.005	0.984	0.249	0.876	0.052	↑**	↓*	↓	↓*
	p-Cresol glucuronide	(+) 302.122	0.000	0.003	0.001	0.003	0.000	↑**	↓*	↑	↓*
	Cortisol	(-) 361.199	0.005	0.073	0.014	0.027	0.004	↑**	↓*	↓	↓**

^aAverage values of relative abundance normalized by internal standard in each group. ^bChange trends in the DNCB group compared to the control group. ^cChange trends in the BS012 or dexamethasone treated group compared to the DNCB group. [#] $p < 0.05$, ^{##} $p < 0.01$ vs. Control group; ^{*} $p < 0.05$, ^{**} $p < 0.01$ vs. DNCB group, Dexa: Dexamethasone.

Table S2. List of identified metabolites significantly altered in skin lesions from DNCB-induced mice with topical BS012 treatment.

Related metabolism	Identified metabolite	(ionization mode) Observed m/z	Mean value ^a					Trend ^b Control vs. DNCB ^d	Change trend ^c DNCB vs. treated group ^e		
			Control	DNCB	BS012 (%)		Dexa		BS012 (%)		Dexa
					3	10			3	10	
Fatty acid metabolism	Pentadecanoic acid	(-) 241.217	0.0180	0.0205	0.0128	0.0164	0.0149	↑	↓*	↓	↓
	Heptadecanoic acid	(-) 269.249	0.0678	0.0810	0.0334	0.0496	0.0392	↑	↓*	↓	↓
	Linoleic acid	(-) 279.232	0.0010	0.0037	0.0008	0.0011	0.0009	↑	↓*	↓	↓
	Stearic acid	(-) 283.263	0.0001	0.0005	0.0003	0.0004	0.0004	↑**	↓*	↓	↓
	Eicosapentaenoic acid	(+) 303.233	0.0086	0.0062	0.0085	0.0195	0.0098	↓	↑	↑**	↑
	Arachidonic acid	(-) 303.232	0.0084	0.0125	0.0068	0.0072	0.0094	↑	↓	↓	↓
	DiHOME	(-) 313.238	0.0028	0.0056	0.0022	0.0051	0.0027	↑	↓*	↓	↓
	HETE	(-) 319.227	0.0002	0.0005	0.0004	0.0003	0.0003	↑	↓	↓	↓
Polyamine metabolism	Spermidine	(+) 147.077	0.0019	0.0056	0.0049	0.0051	0.0060	↑**	↓	↓	↑
	N-Acetylspermidine	(+) 188.175	0.0245	0.3343	0.1219	0.1842	0.1091	↑**	↓	↓	↓
	Glutamine	(-) 145.061	0.0074	0.0191	0.0146	0.0109	0.0199	↑**	↓	↓**	↑
	Glutamate	(-) 146.046	0.0104	0.0177	0.0141	0.0113	0.0215	↑**	↓	↓**	↑
	Aspartate	(-) 132.030	0.0050	0.0129	0.0083	0.0076	0.0135	↑**	↓**	↓**	↑
Glutathione metabolism	GSH	(+) 308.091	0.0045	0.0042	0.0084	0.0133	0.0107	↓	↑*	↑**	↑**
	GSSG	(+) 613.160	0.0055	0.0065	0.0046	0.0019	0.0076	↑	↓	↓**	↑
Glycerophospholipid metabolism	LPC O-16:1	(-) 524.334	0.0006	0.0023	0.0006	0.0011	0.0007	↑**	↓*	↓	↓*
	LPC 18:2	(+) 520.340	0.0692	0.1201	0.0430	0.1135	0.0596	↑	↓*	↓	↓
	LPC 18:0	(-) 568.361	0.0119	0.0191	0.0127	0.0131	0.0147	↑**	↓*	↓*	↓
	LPC 20:4	(+) 544.342	0.0057	0.0059	0.0033	0.0033	0.0036	↑	↓	↓	↓
	LPC 20:2	(-) 592.360	0.0002	0.0003	0.0002	0.0002	0.0002	↑	↓*	↓*	↓
	LPC 20:1	(+) 550.387	0.0020	0.0029	0.0012	0.0015	0.0010	↑	↓	↓	↓*
	LPC 22:6	(-) 612.329	0.0021	0.0024	0.0015	0.0015	0.0015	↑	↓	↓*	↓
Sphingolipid metabo- lism	Sphingosine	(+) 300.291	0.0345	0.0570	0.0119	0.0300	0.0221	↑	↓**	↓	↓*
	SM d42:2	(-) 857.675	0.0015	0.0070	0.0009	0.0018	0.0053	↑*	↓**	↓*	↓
	Ceramide (d42:2)	(-) 692.619	0.0002	0.0009	0.0002	0.0005	0.0005	↑**	↓*	↓	↓
	Ceramide (d42:1)	(+) 684.652	0.0005	0.0021	0.0013	0.0018	0.0010	↑	↓	↓	↓
	Ceramide (d36:2)	(-) 622.541	0.0087	0.0208	0.0036	0.0022	0.0022	↑	↓	↓	↓
	PS 38:4	(-) 810.531	0.0486	0.0020	0.0077	0.0025	0.0040	↓**	↑**	↑	↑
	PS 40:7	(-) 832.515	0.0045	0.0009	0.0022	0.0179	0.0013	↓**	↑*	↑**	↑
	PS 40:6	(+) 836.544	0.0282	0.0007	0.0031	0.0011	0.0019	↓**	↑**	↑	↑
	PS 40:5	(-) 836.546	0.0248	0.0008	0.0033	0.0010	0.0014	↓**	↑**	↑	↑
Histidine metabolism	Urocanic acid	(-) 137.035	0.0137	0.0246	0.0119	0.0071	0.0190	↑**	↓**	↓**	↓
	Histidine	(-) 154.062	0.0030	0.0105	0.0058	0.0069	0.0092	↑**	↓*	↓**	↓
	Histamine	(+) 112.087	0.2223	0.7793	0.5591	0.6005	0.5457	↑**	↓	↓	↓
Amino acid metabolism	Alanine	(+) 90.055	0.0292	0.0594	0.0389	0.0468	0.0318	↑**	↓**	↓	↓
	Threonine	(+) 118.051	0.0470	0.1111	0.0665	0.1103	0.1391	↑**	↓	↓	↑
	Lysine	(+) 147.113	0.0064	0.0189	0.0126	0.0195	0.0161	↑*	↓	↑	↓
	Phenylalanine	(+) 120.080	0.1127	0.2429	0.1383	0.2071	0.1753	↑**	↓**	↓	↓
Others	Isocitrate	(-) 173.009	0.0433	0.0194	0.0548	0.2147	0.0562	↓	↑**	↑**	↑*
	Citrate	(+) 191.019	0.0194	0.0183	0.0265	0.2301	0.0251	↓	↑	↑**	↑
	UDP-GlcNAc	(-) 606.074	0.0037	0.0066	0.0065	0.0034	0.0077	↑	↓	↓*	↑
	Uric acid	(-) 167.021	0.0102	0.0212	0.0115	0.0075	0.0133	↑**	↓*	↓**	↓
	Lactic acid	(-) 89.0244	0.2902	0.3570	0.2752	0.3487	0.3490	↑	↓*	↓	↓

^aAverage values of relative abundance normalized by internal standard in each group. ^bChange trends in the DNCB group compared to the control group. ^cChange trends in the BS012 or dexamethasone treated group compared to the DNCB group. [#] $p < 0.05$, ^{##} $p < 0.01$ vs. Control group; ^{*} $p < 0.05$, ^{**} $p < 0.01$ vs. DNCB group, Dexa: Dexamethasone.

Table S3. List of the BS012-derived exogenous metabolites detected from skin tissue.

Classification		Compounds	Retention time (min)	Molecular formula	Measured m/z	Adduct	MS/MS fragment m/z (Relative intensity, %)
<i>Cinnamomum cassia</i>	Essential oil components	Cinnamaldehyde	10.50	C ₉ H ₈ O	115.0543	[M+H-H ₂ O] ⁺	116 (100), 107 (80), 88 (75)
		Methoxycinnamaldehyde	10.06	C ₁₀ H ₁₀ O ₂	163.0755	[M+H] ⁺	135 (100), 145 (80), 107 (60)
		Hydroxycinnamic acid	7.94	C ₉ H ₈ O ₃	147.0441	[M-H] ⁻	119 (100)
		Hydroxymethoxycinnamate	7.30	C ₁₀ H ₁₀ O ₄	177.0549	[M+H-H ₂ O] ⁺	177 (100)
		Dimethoxycinnamic acid	9.28	C ₁₁ H ₁₂ O ₄	209.0813	[M+H] ⁺	191 (100), 161 (20)
		Coumaraldehyde	6.98	C ₉ H ₈ O ₂	149.0598	[M+H] ⁺	131 (100), 121 (100), 107 (40)
		Coumaric acid	7.94	C ₉ H ₈ O ₃	163.0396	[M-H] ⁻	119 (100)
		Methylcoumarin	8.96	C ₁₀ H ₁₀ O ₃	161.0598	[M+H-H ₂ O] ⁺	161 (100)
		Coumaroyl Hexoside	7.18	C ₁₅ H ₁₈ O ₈	325.0919	[M-H] ⁻	163 (100), 119 (20)
		Dihydrocoumaroyl Hexo-side	6.98	C ₁₅ H ₂₀ O ₈	327.1078	[M-H] ⁻	165 (100)
	Flavonoids	Feruloyltyramine	8.21	C ₁₈ H ₁₉ NO ₄	312.1231	[M-H] ⁻	297 (100), 178 (90), 135 (40), 313 (30)
		Kaempferol	9.08	C ₁₅ H ₁₀ O ₆	285.0393	[M-H] ⁻	285 (100)
		Isokaempferide	8.07	C ₁₆ H ₁₂ O ₆	299.0915	[M-H] ⁻	255 (100), 271 (70), 284 (20)
		Dihydrokaempferol	7.70	C ₁₅ H ₁₂ O ₆	287.0552	[M-H] ⁻	259 (100), 243 (15)
		Kaempferol-3-O-glucoside-6"-p-coumaroyl	8.69	C ₃₀ H ₂₆ O ₁₃	593.1278	[M-H] ⁻	285 (100), 447 (10)
		Quercetin	7.98	C ₁₅ H ₁₀ O ₇	303.0506	[M+H] ⁺	285 (100), 257 (80), 229 (40)
		Dihydroquercetin	7.19	C ₁₅ H ₁₂ O ₇	303.0499	[M-H] ⁻	285 (100), 177 (5), 125 (5)
		Quercetin-3-O-pentoside	7.83	C ₂₀ H ₁₈ O ₁₁	433.0757	[M-H] ⁻	373 (100), 403 (50), 300 (30)
		Quercetin-3-O-rhamnoside	8.03	C ₂₁ H ₂₀ O ₁₁	447.0916	[M-H] ⁻	301 (100)
		Quercetin-4'-O-glucoside	7.62	C ₂₁ H ₂₀ O ₁₂	463.0865	[M-H] ⁻	301 (100)
Polyphenols	Quercetin-3-O-deoxyhexo-syl(1-2) pentoside	7.98	C ₂₆ H ₂₈ O ₁₅	579.1344	[M-H] ⁻	300 (100)	
	Syringaldehyde	7.00	C ₉ H ₁₀ O ₄	183.0654	[M+H] ⁺	155 (100), 123 (20)	
<i>Platycodon grandiflorum</i>	Triterpenoid saponins	Catechin	6.57	C ₁₅ H ₁₄ O ₆	289.0711	[M-H] ⁻	245 (100), 205 (30)
		Platycodin D	9.42	C ₅₇ H ₉₂ O ₂₈	1223.5704	[M-H] ⁻	1224 (100)
		Polygalacin D	9.50	C ₅₇ H ₉₂ O ₂₇	1207.5752	[M-H] ⁻	665 (100), 469 (65), 1117 (25)
		Platyconic acid D	9.44	C ₅₄ H ₈₄ O ₂₆	1147.5162	[M-H] ⁻	1117 (100), 937 (85), 485 (20)
		Platycosaponin A	8.81	C ₄₂ H ₆₈ O ₁₆	827.4433	[M-H] ⁻	828 (100), 665 (30)
<i>Asarum sieboldii</i>	Lignan	Sesamin	10.65	C ₂₀ H ₁₈ O ₆	337.1080	[M+H] ⁺	319 (100), 267 (45), 289 (35), 135 (20)
	N-acyl amines	N-isobutyl-dodecatetrae-namide	11.34	C ₁₆ H ₂₅ NO	248.2019	[M+H] ⁺	149 (100), 142 (60)