

Supplementary Material

Development of a Reverse-Phase High-Performance Liquid Chromatography and Liquid Chromatography Tandem Mass Spectrometry Methods for Quality Control of Daegunjoong-Tang

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Table S1. Information on the composition of DGJT.

Herbal Medicine	Scientific Name	English name	Family	Used part	Origin	Amount (g)
Zingiberis Rhizoma	<i>Zingiber officinale</i> Rosc.	Ginger	Zingiberaceae	Rhizome	Seosan, Korea	2500.0
Ginseng Radix	<i>Panax ginseng</i> C.A.Mey.	Ginseng	Araliaceae	Root	Geumsan, Korea	1250.0
Zanthoxyli Pericarpium	<i>Zanthoxylum schinifolium</i> Sieb. et Zucc.	Zanthoxylum Peel	Rutaceae	Pericarp	Yeongyang, Korea	1250.0
Total (g)						5000.00

Table S2. HPLC parameters for quantification of DGJT.

Parameter for HPLC Analysis		
System	Shimadzu Prominence LC-20A series (Kyoto, Japan)	
Detector	Photo diode array	
Column	Waters SunFire C18 (4.6 × 250 mm, particle size 5 µm, Milford, MI, USA)	
Column oven Temperature (°C)	40.0	
Flow rate (mL/min)	1.0	
Injection volume (µL)	10.0	
Mobile phase	A: 0.1% (v/v) Aqueous formic acid B: 0.1% (v/v) Formic acid in acetonitrile	
Gradient elution	Time (min)	A (%)
	0	90
	30	40
	40	0
	45	0
	50	90
	60	90

Table S3. MRM parameters for LC–MS/MS analysis of 4 marker compounds in DGJT.

Compound	Mode	MW (<i>m/z</i>)	Transition (Q1→Q3, <i>m/z</i>)	Collision Energy (eV)	Cone Voltage (V)
Hyperoside	Negative	464.10	463.2→300.0	25	40
Quercitrin	Negative	448.10	447.0→301.0	25	20
Ginsenoside Rg1	Negative	800.49	799.2→637.0	20	50
6-Gingerol	Positive	294.18	295.3→177.1	10	13

Table S4. System suitability for the four marker analytes in the developed HPLC method.

Compound	<i>k'</i>	<i>α</i>	<i>N</i>	<i>Rs</i>	<i>Tf</i>
Hyperoside	3.94	1.17	501122	6.94	1.13
Quercitrin	4.60	1.14	557071	7.23	1.13
Ginsenoside Rg1	5.25	1.14	800846	7.23	1.11
6-Gingerol	9.91	1.89	1122666	40.42	1.07

Retention factor (*k'*), relative retention (*α*), theoretical plate number (*N*), resolution (*Rs*), and symmetry factor (*S*).

Table S5. Repeatability for retention time and peak areas of the four marker analytes using HPLC (*n* = 6).

No.	Retention Time (min)				Peak Area			
	Hyperoside	Quercitrin	Ginsenoside Rg1	6-Gingerol	Hyperoside	Quercitrin	Ginsenoside Rg1	6-Gingerol
1	14.152	16.043	17.880	31.244	1587117	356796	1013372	3787576
2	14.158	16.043	17.879	31.245	1594064	358826	1015568	3803418
3	14.151	16.038	17.869	31.233	1599084	359382	1021961	3818628
4	14.155	16.039	17.871	31.235	1606396	361835	1026638	3832737
5	14.158	16.044	17.878	31.241	1601466	361019	1024103	3826101
6	14.151	16.039	17.872	31.236	1592469	358462	1018940	3801593
Mean	14.15	16.04	17.87	31.24	1596766.00	359386.67	1020097.00	3811675.50
SD ($\times 10^{-1}$)	0.03	0.03	0.05	0.05	69147.41	18187.79	50869.11	170331.92
RSD (%)	0.02	0.02	0.03	0.02	0.43	0.51	0.50	0.45

Table S6. The linear range, regression equation, *r*², LODs, and LOQs of the analytes from YPS using LC–MS/MS (*n* = 3).

Compound	Retention Time (min)	Linear Range (ng/mL)	Regression Equation (<i>y</i> = <i>ax</i> + <i>b</i>) ^a	<i>r</i> ²	LOD (μg/mL) ^b	LOQ (μg/mL) ^c
Hyperoside	1.63	6.25–100.00	<i>y</i> = 29066.1 <i>x</i> – 155575.0	0.9953	0.002	0.006
Quercitrin	2.03	0.63–10.00	<i>y</i> = 9346.2 <i>x</i> + 5140.3	0.9956	0.001	0.003
Ginsenoside Rg1	2.71	6.25–100.00	<i>y</i> = 19.6 <i>x</i> – 108.5	0.9952	1.586	4.757
6-Gingerol	6.33	0.63–10.00	<i>y</i> = 29284.3 <i>x</i> + 10841.7	0.9975	0.002	0.005

^a*y* and *x* are peak area and concentration of compound, respectively ^bLOD = 3.3 × S/N ^cLOQ = 10 × S/N.

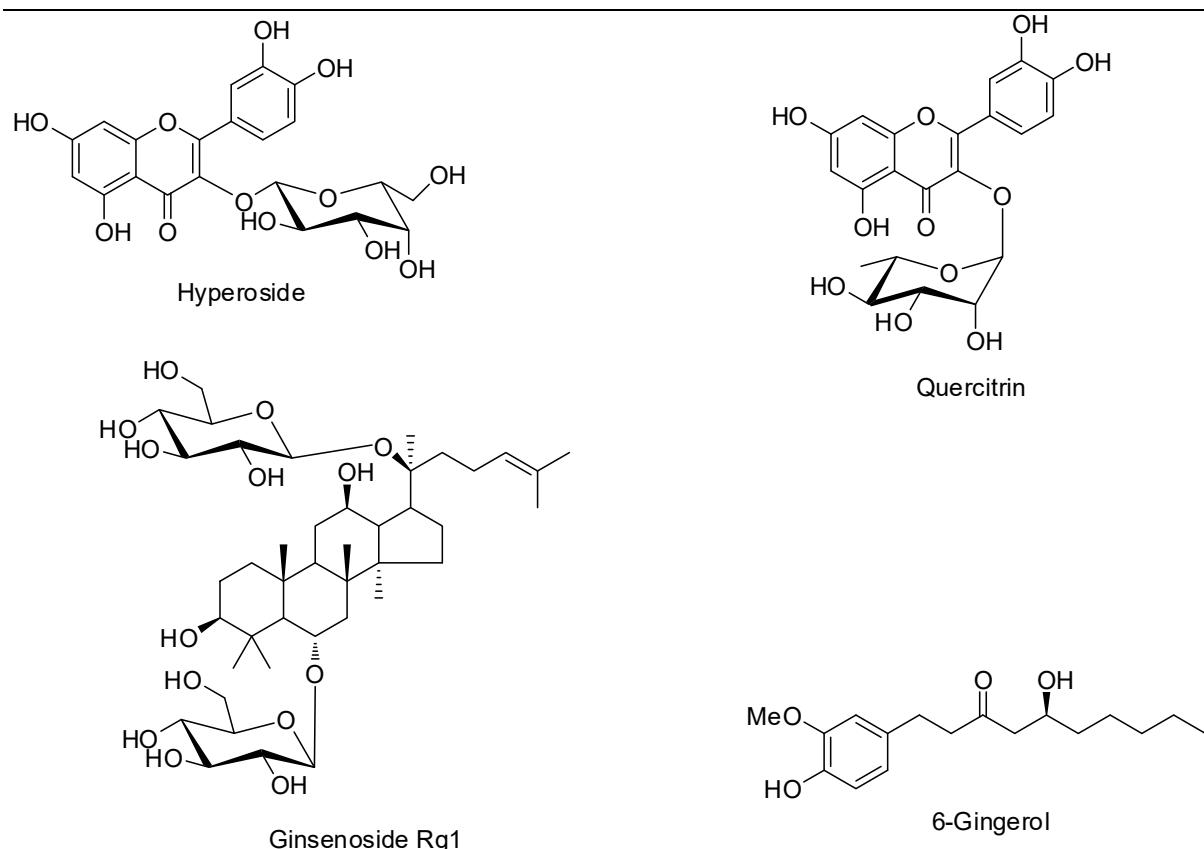
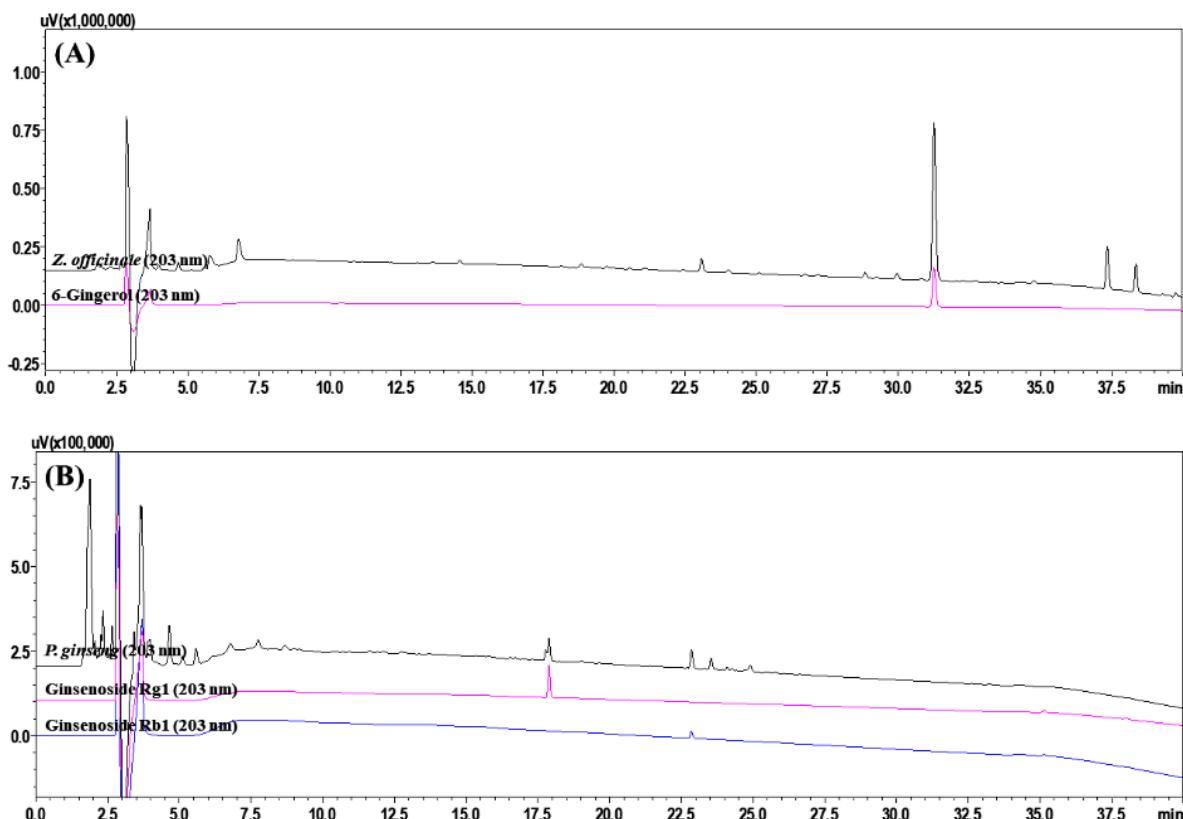
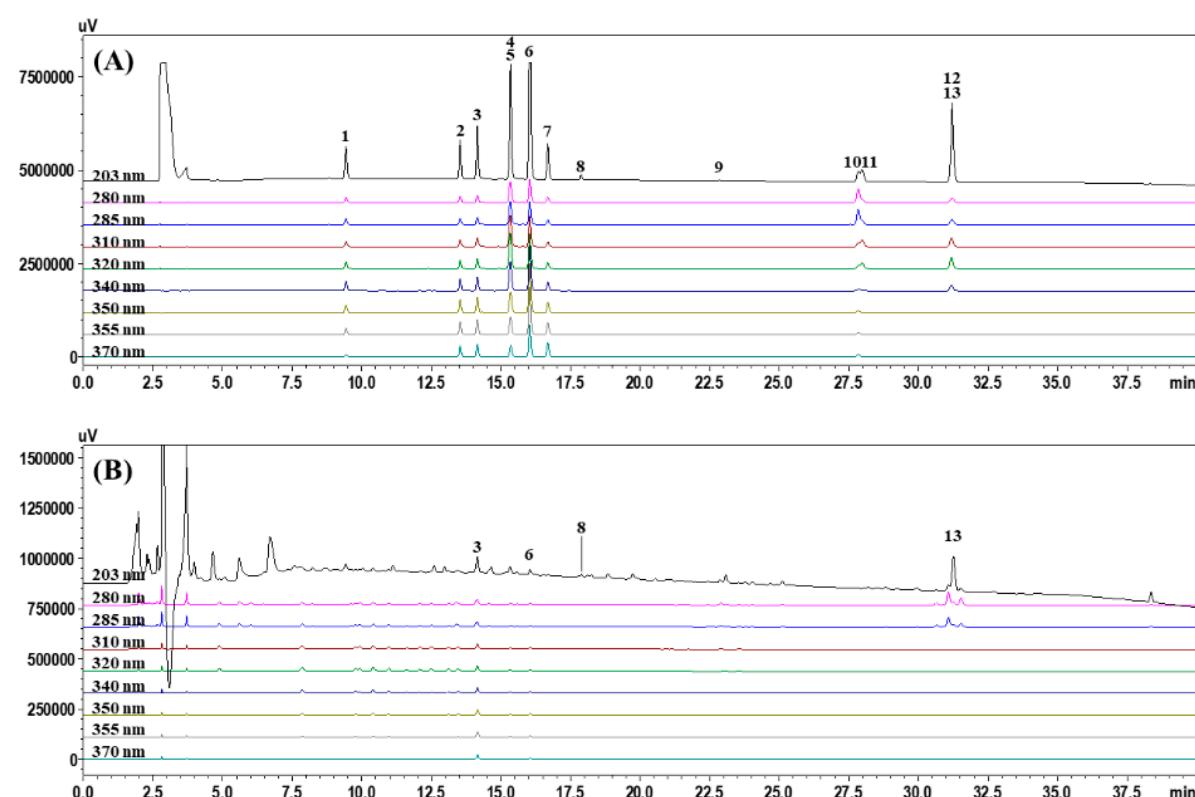
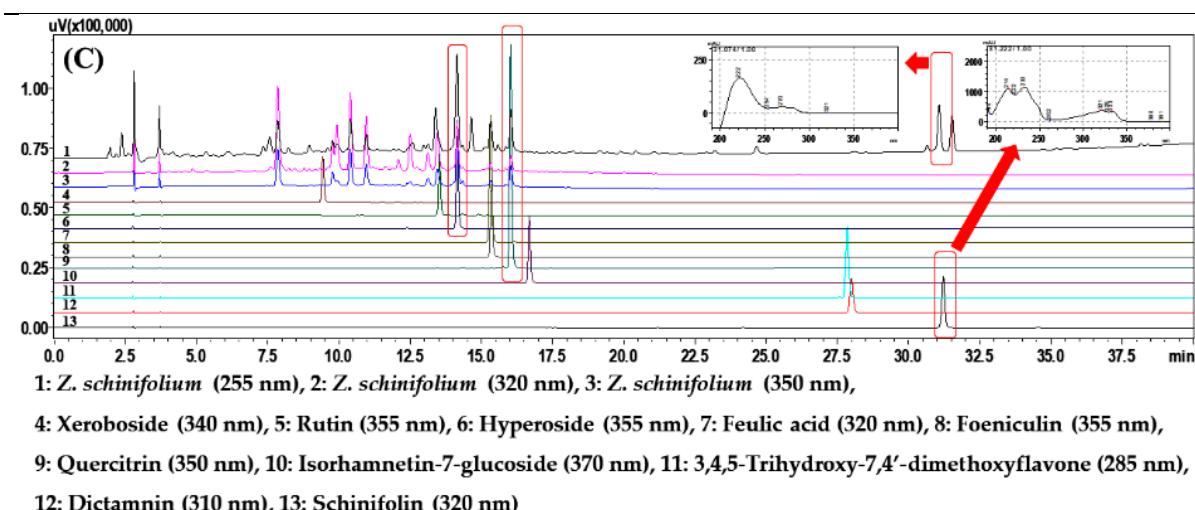
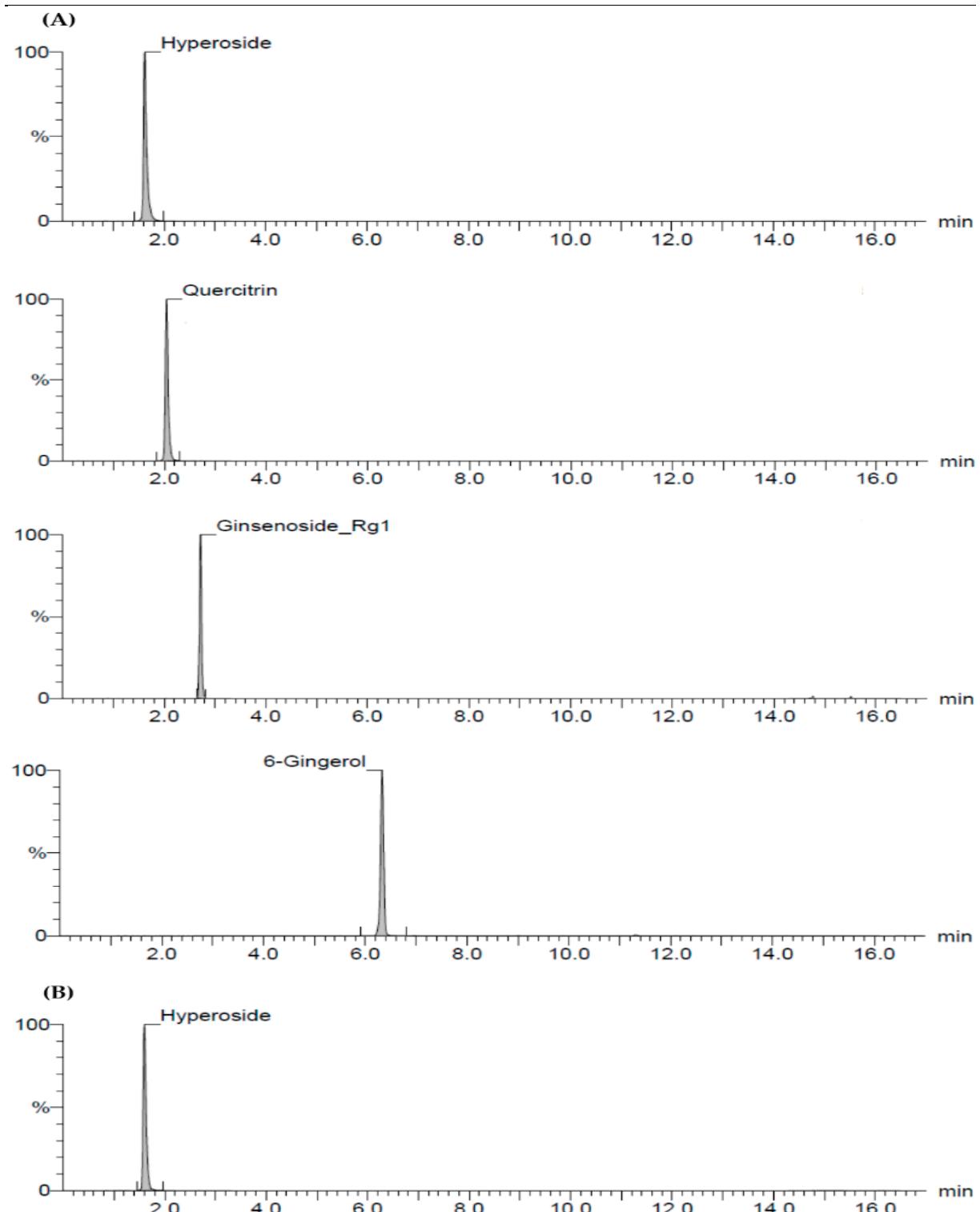


Figure S1. Chemical structures of the four marker analytes used in the quantitative analysis of DGJT.







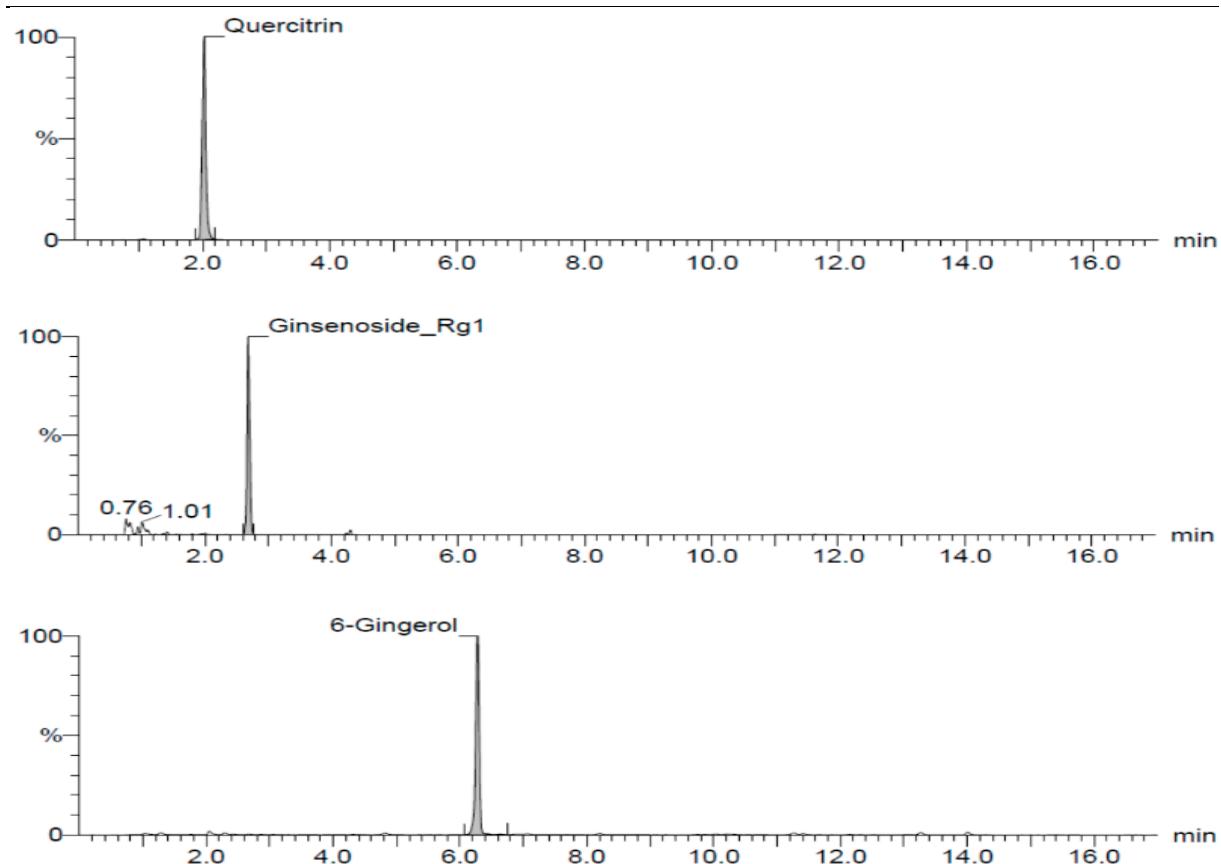


Figure S4. Extracted ion chromatograms of the reference standard (A) and DGJT sample (B) by LC-MS/MS MRM mode.