

179	G06001	China	Cultivated ginseng	Fusong, Jilin, China	Ginseng Division, Department of Herbal Crop Research, Rural Development Administration, Korea
180	G06003	China	Cultivated ginseng	Fusong, Jilin, China	Ginseng Division, Department of Herbal Crop Research, Rural Development Administration, Korea
181	G06010	China	Cultivated ginseng	Tonghua, Jilin, China	Ginseng Division, Department of Herbal Crop Research, Rural Development Administration, Korea
182	G08082	China	Cultivated ginseng	Huanren, Liaoning, China	Ginseng Division, Department of Herbal Crop Research, Rural Development Administration, Korea
183	Gaolishen	China	Cultivated ginseng	Dünhuà, Jilin, China	School of Pharmaceutical Sciences, Changchun University of Chinese Medicine, China / Zhao <i>et al.</i> 2015
184	Jilin	China	Cultivated ginseng	Changchun, Jilin, China	College of Life Science, Jilin Agricultural University, China / Wang <i>et al.</i> 2018
185	JYH-2016473	China	Cultivated ginseng	Unknown, China	Key Laboratory for Plant Diversity and Biogeography of East Asia, Kunming Institute of Botany, China / https://www.ncbi.nlm.nih.gov/nuccore/MK408938
186	Yeshanshen	China	Wild ginseng	Dünhuà, Jilin, China	School of Pharmaceutical Sciences, Changchun University of Chinese Medicine, China / Zhao <i>et al.</i> 2015
187	J-farm 1	Japan	Cultivated ginseng	Azumino, Nagano, Japan	Institute of Chinese Materia Medica, China Academy of Chinese Medical Sciences, China
188	J-farm 2	Japan	Cultivated ginseng	Azumino, Nagano, Japan	Institute of Chinese Materia Medica, China Academy of Chinese Medical Sciences, China
189	J-farm 3	Japan	Cultivated ginseng	Azumino, Nagano, Japan	Institute of Chinese Materia Medica, China Academy of Chinese Medical Sciences, China
190	J-farm 4	Japan	Cultivated ginseng	Azumino, Nagano, Japan	Institute of Chinese Materia Medica, China Academy of Chinese Medical Sciences, China
191	J-farm 5	Japan	Cultivated ginseng	Azumino, Nagano, Japan	Institute of Chinese Materia Medica, China Academy of Chinese Medical Sciences, China
192	Mimaki 1	Japan	Cultivated ginseng	Suwon-si, Gyeonggi-do, Korea	Lab. of Functional Plants, Seoul National University, Korea / Japanese cultivar
193	Mimaki 2	Japan	Cultivated ginseng	Suwon-si, Gyeonggi-do, Korea	Lab. of Functional Plants, Seoul National University, Korea / Japanese cultivar
194	Mimaki 3	Japan	Cultivated ginseng	Suwon-si, Gyeonggi-do, Korea	Lab. of Functional Plants, Seoul National University, Korea / Japanese cultivar
195	Russia 0	Russia	Wild ginseng	Vladivostok, Primorsky Krai, Russia	Lab. of Functional Plants, Seoul National University, Korea
196	Russia 1	Russia	Wild ginseng	Vladivostok, Primorsky Krai, Russia	Lab. of Functional Plants, Seoul National University, Korea
197	Russia 2	Russia	Wild ginseng	Vladivostok, Primorsky Krai, Russia	Lab. of Functional Plants, Seoul National University, Korea
198	Russia 3	Russia	Wild ginseng	Chuguevskoe, Primorsky Krai, Russia	Lab. of Functional Plants, Seoul National University, Korea
199	Russia 4	Russia	Wild ginseng	Chuguevskoe, Primorsky Krai, Russia	Lab. of Functional Plants, Seoul National University, Korea
200	Russia 5	Russia	Wild ginseng	Chuguevskoe, Primorsky Krai, Russia	Lab. of Functional Plants, Seoul National University, Korea
201	Russia 6	Russia	Wild ginseng	Chuguevskoe, Primorsky Krai, Russia	Lab. of Functional Plants, Seoul National University, Korea
202	Russia 7	Russia	Wild ginseng	Chuguevskoe, Primorsky Krai, Russia	Lab. of Functional Plants, Seoul National University, Korea
203	Russia 8	Russia	Wild ginseng	Chuguevskoe, Primorsky Krai, Russia	Lab. of Functional Plants, Seoul National University, Korea

Table S2. Plastome sequence assembly statistics of the 44 ginseng germplasms used in this study

No.	Name	Abbreviation	Origin	Type	Plastid genome				GenBank No.	Reference
					LSC	SSC	IR	Entire		
0	Sobek ²⁾	SB		Wild collection	86,106	18,070	26,071	156,318	AY582139	Kim <i>et al.</i> 2004
1	Cheongsun	CS			86,129	18,077	26,075	156,356	KM067386	
2	Chunpoong	CP			86,128	18,084	26,018	156,248	KM088019	
3	Gopoong	GO			86,128	18,077	26,075	156,355	KM067387	
4	Gumpoong	GP			86,129	18,077	26,075	156,356	KM067388	
5	Sunhyang	SH		Cultivar	86,200	18,077	26,074	156,425	KM067393	
6	Sunwon	SW			86,128	18,077	26,075	156,355	KM067390	Kim <i>et al.</i> 2015b
7	Sunpoong	SP			86,128	18,077	26,075	156,355	KM067391	
8	Sunun	SU			86,128	18,077	26,075	156,355	KM067392	
9	Yunpoong	YP			86,128	18,077	26,075	156,355	KM088020	
10	Hwangsook	HS		Landrace	86,128	18,077	26,018	156,241	KM067394	
11	Jakyung	JK			86,128	18,077	26,075	156,355	KM067389	
12	K-farm 1	KF1	Korea		86,128	18,077	26,075	156,355	MT416073	
13	K-farm 2	KF2			86,128	18,084	26,018	156,248	MT416074	
14	K-farm 3	KF3		Farm collection	86,128	18,084	26,018	156,248	MT416075	
15	K-farm 4	KF4			86,129	18,077	26,075	156,356	MT416076	
16	K-farm 5	KF5			86,128	18,077	26,018	156,241	MT416077	
17	Cheongsong 2	ChS2			86,128	18,077	26,075	156,355	MK415693	
18	Daejeon	DJ			86,128	18,077	26,075	156,355	MK415694	This study
19	Gangwon 2	G2			86,129	18,077	26,075	156,356	MK415695	
20	Gangwon 8	G8			86,128	18,077	26,075	156,355	MK415696	
21	Gangwon 13	G13	Wild collection	86,128	18,077	26,075	156,355	MK415697		
22	Gangwon 15	G15		86,129	18,077	26,075	156,356	MK415698		
23	Gangwon 16	G16			86,128	18,077	26,075	156,355	MK415699	
24	Gangwon 17	G17			86,128	18,077	26,075	156,355	MK415700	
25	Hamyang 1	HY1			86,128	18,077	26,074	156,353	MN864139	
26	C-farm 1	CF1			86,128	18,077	26,075	156,355	MT416063	
27	C-farm 2	CF2			86,128	18,077	26,075	156,355	MT416064	
28	C-farm 3	CF3		Farm collection	86,128	18,077	26,075	156,355	MT416065	This study
29	C-farm 4	CF4			86,128	18,077	26,075	156,355	MT416066	
30	C-farm 5	CF5			86,128	18,077	26,075	156,355	MT416067	
31	Jilin ³⁾	JL	China		86,128	18,077	26,018	156,241	MH049735	Wang <i>et al.</i> 2018
32	Damaya	DMY		Landrace	86,129	18,077	26,074	156,354	KC686331	
33	Gaolishen	GLS			86,129	18,077	26,074	156,354	KC686333	Zhao <i>et al.</i> 2015
34	Ermaya	EMY			86,129	18,077	26,074	156,354	KC686332	
35	Yeshanshen	YSS		Wild collection	86,130	18,077	26,074	156,355	KF431956	
36	JYH-2016473	JYH		-	86,128	18,077	26,018	156,241	MK408938	-
37	Mimaki 1	M1		Cultivar	86,128	18,077	26,075	156,355	MK415701	
38	J-farm 1	JF1			86,128	18,077	26,075	156,355	MT416068	
39	J-farm 2	JF2	Japan		86,127	18,077	26,075	156,354	MT416069	This study
40	J-farm 3	JF3		Farm collection	86,128	18,077	26,075	156,355	MT416070	
41	J-farm 4	JF4			86,128	18,077	26,075	156,355	MT416071	
42	J-farm 5	JF5			86,128	18,077	26,075	156,355	MT416072	
43	Russia 1	R1		Russia	Wild collection	86,128	18,077	26,075	156,355	
44	Russia 2	R2			86,129	18,077	26,075	156,356	MK415703	

²⁾ The sequence of this germplasm was not used for comparative analysis due to its low sequence accuracy. ³⁾ The reported length for each section of this germplasm was readjusted by comparison with other germplasms.

IR, Inverted repeat; LSC, Large single copy; SSC, Small single copy.

Table S3. Variation information of the *P. quinquefolius* plastomes

No.	Variant type	<i>P. quinquefolius</i> 1 (NC027456, 156,088 bp)		<i>P. quinquefolius</i> 2 (MK408923, 156,070 bp)		Region
		Genotype	Position	Genotype	Position	
1	InDel	(A) ₁₃	4,835	(A) ₁₂	4,834 4,835	
2	SNP	T	24,192	C	24,191	
3	InDel	(C) ₁₃	38,149 ~ 38,151	(C) ₁₀	38,147 38,148	
4	SNP	C	56,945	A	56,941	
5	SNP	C	59,389	T	59,385	
6	SNP	T	60,441	G	60,437	
7	SNP	C	60,992	A	60,988	
8	InDel	TATTAATTATTATAA	61,322 ~ 61,337	-	61,317 61,318	LSC
9	SNP	C	64,601	T	64,581	
10	InDel	(A) ₈	73,253 73,254	(A) ₉	73,234	
11	InDel	(A) ₈	73,919 73,920	(A) ₉	73,901	
12	SNP	A	78,705	T	78,687	
13	SNP	A	78,706	T	78,688	
14	SNP	T	82,075	C	82,057	
15	SNP	T	95,250 / 146,934	A	95,232 / 146,916	
16	SNP	G	95,251 / 146,933	T	95,233 / 146,915	
17	SNP	A	95,252 / 146,932	C	95,234 / 146,914	
18	SNP	T	95,253 / 146,931	A	95,235 / 146,913	IR
19	SNP	T	99,395 / 142,789	A	99,377 / 142,771	
20	SNP	C	100,912 / 141,272	T	100,894 / 141,254	

IR, Inverted repeat; LSC, Large single copy

Table S4. Sequence information for the 10 KASP markers developed based on ginseng plastome sequences

Variation No ²⁾	Marker name	Nucleotide position ³⁾	Allele	Allele-specific primer sequence (5'→3')	Common primer sequence (5'→3')
S1	pgcpKASP006	7,159	FAM	G CGTTGTCACATAAATAGAACGAAATCG	GGAAATGCTTAGTTAACTTAATATGTAAT
			HEX	T AACGTTGTCACATAAATAGAACGAAATCT	
S2	pgcpKASP001	21,344	FAM	T AAAAGATTGTGGCACCGTCCGA	CCCATTTCGGGACTCACAGAAATA
			HEX	C AAAGATTGTGGCACCGTCCGA	
S3	pgcpKASP005	22,287	FAM	T GAAATATGACCAACAGTAGTTCGAATATAT	GGGCAATATCTAATAGTAAGAAGTATAAAA
			HEX	G AAATATGACCAACAGTAGTTCGAATATAG	
S4	pgcpKASP009	23,946	FAM	A TTAAAATTCTAGAACCCCCCCCA	TGGGAACCTTGAGTAAGGAGTAGATCTTTT
			HEX	C TAAAATTCTAGAACCCCCCCCA	
S5	pgcpKASP010	40,011	FAM	C GAGAATAGTAATTCACATATTCTTAACAATAGG	AACCCAGAGCAATAGCATGATGAACTAA
			HEX	T GAGAATAGTAATTCACATATTCTTAACAATAG	
S6	pgcpKASP003	44,895	FAM	G GACATGAGTTTTAACTGAAATCTTAAATTTG	CTGAAGGTGGGAGAAAAGATAAAACCAA
			HEX	T GACATGAGTTTTAACTGAAATCTTAAATTTT	
S7	pgcpKASP004	90,858 / 151,519	FAM	G ACATCACTTCTGCTTCTATTGTAATAAAG	CGGGCCTTTTCCACATAAAAAAGGGAA
			HEX	A CACATCACTTCTGCTTCTATTGTAATAAAG	
S9	pgcpKASP002	117,376	FAM	G GCTAAGAATTATTACAGGTCCC	GACACGATAACTCCAATAATCCAAGTGT
			HEX	A CTGCTAAGAATTATTACAGGTCCC	
S10	pgcpKASP007	118,525	FAM	T AACATCTCCGGCCAACGTAATTTA	CCATATGAGATACAGAGGAATAGGCTATT
			HEX	A AACATCTCCGGCCAACGTAATTTT	
S11	pgcpKASP008	127,069	FAM	A TTCTTCTTGATTTTCGATTCTTTAATTCAAAAA	GCAAAAATGGAATTTTTTTAGACCATCAAA
			HEX	T TTCTTCTTGATTTTCGATTCTTTAATTCAAAAA	

Bold and underlined nucleotides in the primer sequences represent allele-specific positions. ²⁾ Variation number shown in Fig. 1. ³⁾ Variation positions are based on the plastome sequence of *P. ginseng* cv. 'Chunpoong' (KM088019).

FAM, Fluorescein amidite; HEX, Hexachloro-fluorescein; KASP, Kompetitive allele-specific PCR.

Table S5. Sequence information for the four InDel markers developed based on ginseng plastome sequences

Variation No ^{z)}	Marker name	Nucleotide position ^{y)}	Primer sequence (5'→3')	TR unit length (bp)	CNV	Product size (bp)
ID2	pgcp097f2*r	7,189	Forward TGGAAAGGCTGTTGTCACTG	13	1 - 2	378 / 391
			Reverse TCAGCAACGGGAGATATTCA			
ID3	pgcp137	32,850	Forward TCCTGAACCACTAGACGATGG	59	1 - 2	457 / 516
			Reverse TTTCGATAACTTCTTGATCCCTCT			
ID6	pgycf1	111,304 / 130,897	Forward GGTATTAGTCTGGATACGGCAAA	57	2 - 5	615 / 672 / 729 / 786
			Reverse TCGAAAAGAAGGGTCACAAGA			
ID7	pgcp139f*r2	115,833	Forward TGTGCGACAAACAAATAAGTCA	7	2 - 3	150 / 157
			Reverse CGAAGCGAGTTCCATTTTCAT			

All InDel markers were developed in Kim *et al.* 2015b. ^{z)} Variation number shown in Fig. 1. ^{y)} Variation positions are based on the plastome sequence of *P. ginseng* cv. 'Chunpoong' (KM088019).

CNV, Copy number variation; TR, Tandem repeat.

Table S6. Genotype and group information for 203 individual ginseng plants

Origin	Germ-plasm type	No.	Variation No. ²⁾ (Marker) Name	S1	S2	S3	S4	S5	S6	S7	S9	S10	S11	ID2	ID3	ID6	ID7	SNP combination	InDel combination	Haplotype
				pgcp KASP 006	pgcp KASP 001	pgcp KASP 005	pgcp KASP 009	pgcp KASP 010	pgcp KASP 003	pgcp KASP 004	pgcp KASP 002	pgcp KASP 007	pgcp KASP 008	pgcp 097f2*r ^{y)}	pgcp 137 ^{y)}	pg ycf1 ^{y)}	pgcp 139f*r2 ^{y)}			
		1	Cheonryang	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
		2	CP	G	C	T	A	C	G	A	A	T	A	1	1	3	3	GCTACGAATA	1133	I
		3	CS	G	T	G	A	C	G	A	G	T	A	1	1	4	2	GTGACGAGTA	1142	B'
		4	G03001	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
		5	G03002	G	T	G	A	C	G	A	A	T	A	1	1	4	2	GTGACGAATA	1142	B
		6	G03003	G	T	G	A	C	G	A	A	T	A	1	1	4	2	GTGACGAATA	1142	B
		7	G03004	G	T	G	A	C	G	A	A	T	A	1	1	4	2	GTGACGAATA	1142	B
		8	G03005	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
		9	G03006	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
		10	G03007	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
		11	G03008	G	C	T	A	C	G	A	A	T	A	1	1	3	3	GCTACGAATA	1133	I
		12	G03009	G	C	T	A	C	G	A	A	T	A	1	2	5	2	GCTACGAATA	1252	I
		13	G03010	G	T	G	A	C	G	A	A	T	A	1	1	4	2	GTGACGAATA	1142	B
		14	G03011	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
		15	G03012	G	C	G	A	C	G	A	A	T	A	1	1	2	2	GCGACGAATA	1122	A
		16	G03013	G	C	G	C	C	G	A	A	T	A	1	1	4	2	GCGCCGAATA	1142	J
		17	G03015	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
		18	G03016	G	T	G	A	C	G	A	A	T	A	1	1	4	2	GTGACGAATA	1142	B
		19	G03017	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
		20	G03018	G	T	G	A	C	G	A	A	T	A	1	1	4	2	GTGACGAATA	1142	B
		21	G03019	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
		22	G03020	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
		23	G03021	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
		24	G03022	G	C	G	A	C	T	A	A	T	A	2	2	5	2	GCGACTAATA	2252	C
		25	G03024	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
		26	G03025	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
		27	G03026	G	T	G	A	C	G	A	A	T	A	1	1	4	2	GTGACGAATA	1142	B
		28	G03027	G	C	T	C	C	G	A	A	T	A	1	2	3	2	GCTCCGAATA	1232	IJ
		29	G03028	G	C	T	C	C	G	A	A	T	A	1	2	3	2	GCTCCGAATA	1232	IJ
		30	G03029	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
		31	G03030	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
		32	G03031	G	C	G	A	C	G	A	A	A	A	1	1	4	2	GCGACGAAAA	1142	D
		33	G03034	G	T	G	A	C	G	A	G	T	A	1	1	4	2	GTGACGAGTA	1142	B'
		34	G03035	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
		35	G03036	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
		36	G03037	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
		37	G03038	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
		38	G03039	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
		39	G03040	G	T	G	A	C	G	A	A	T	A	1	1	4	2	GTGACGAATA	1142	B

40	G03041	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
41	G03042	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
42	G03043	G	T	G	A	C	G	A	G	T	A	1	1	4	2	GTGACGAGTA	1142	B'
43	G03044	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
44	G03045	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
45	G03046	G	C	G	A	C	G	A	A	T	A	1	1	3	2	GCGACGAATA	1132	A
46	G03047	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
47	G03048	G	T	G	A	C	G	A	A	T	A	1	1	4	2	GTGACGAATA	1142	B
48	G03049	G	C	T	A	C	G	A	A	T	A	1	1	3	3	GCTACGAATA	1133	I
49	G03050	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
50	G03051	G	C	G	A	C	G	A	A	T	A	1	2	4	2	GCGACGAATA	1242	A
51	G03052	G	C	G	A	C	T	A	A	T	A	1	1	4	2	GCGACTAATA	1142	C
52	G03053	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
53	G03054	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
54	G03055	G	T	G	A	C	G	A	A	T	A	1	1	4	2	GTGACGAATA	1142	B
55	G03056	G	T	G	A	C	G	A	A	T	A	1	1	4	2	GTGACGAATA	1142	B
56	G03057	G	T	G	A	C	G	A	A	T	A	1	1	4	2	GTGACGAATA	1142	B
57	G03058	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
58	G03059	G	C	G	A	C	G	A	A	T	A	1	2	4	2	GCGACGAATA	1242	A
59	G03060	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
60	G03061	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
61	G03062	G	C	G	A	C	T	A	A	T	A	1	1	4	2	GCGACTAATA	1142	C
62	G03063	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
63	G03064	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
64	G03065	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
65	G03067	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
66	G03068	G	T	G	A	C	G	A	A	T	A	1	1	4	2	GTGACGAATA	1142	B
67	G03069	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
68	G03070	G	C	G	A	C	G	A	A	A	A	1	1	4	2	GCGACGAAAA	1142	D
69	G03071	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
70	G03072	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
71	G03075	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
72	G03076	G	T	G	A	C	G	A	A	T	A	1	1	4	2	GTGACGAATA	1142	B
73	G03077	G	T	G	A	C	G	A	A	T	A	1	1	4	2	GTGACGAATA	1142	B
74	G03078	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
75	G03079	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
76	G03080	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
77	G03081	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
78	G03083	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
79	G03084	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
80	G03086	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
81	G03089	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
82	G03092	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
83	G03093	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
84	G03096	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
85	G03105	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A

C

Korea

86	G03107	G	C	G	A	C	G	A	A	T	A	1	1	3	2	GCGACGAATA	1132	A
87	G03109	G	C	G	A	C	G	G	A	T	A	1	1	4	2	GCGACGGATA	1142	E
88	G03110	G	T	G	A	C	G	A	G	T	A	1	1	4	2	GTGACGAGTA	1142	B'
89	G03111	G	T	G	A	C	G	A	A	T	A	1	1	4	2	GTGACGAATA	1142	B
90	G03112	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
91	G03114	G	T	G	A	C	G	A	A	T	A	1	1	4	2	GTGACGAATA	1142	B
92	G03118	G	T	G	A	C	G	A	A	T	A	1	1	4	2	GTGACGAATA	1142	B
93	G03119	G	C	G	A	C	G	G	A	T	A	1	1	4	2	GCGACGGATA	1142	E
94	G03140	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
95	G03180	G	C	G	A	C	G	A	A	A	A	1	1	4	2	GCGACGAAAA	1142	D
96	G04051	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
97	G04054	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
98	G04076	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
99	G05021	G	T	G	A	C	G	A	A	T	A	1	1	4	2	GTGACGAATA	1142	B
100	G05032	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
101	G05097	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
102	G08024	G	T	G	A	C	G	A	G	T	A	1	1	4	2	GTGACGAGTA	1142	B'
103	G08034	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
104	GO	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
105	GO	G	T	G	A	C	G	A	A	T	A	1	1	4	2	GTGACGAATA	1142	B
106	Gowon	G	T	G	A	C	G	A	A	T	A	1	1	4	2	GTGACGAATA	1142	B
107	GP	G	T	G	A	C	G	A	G	T	A	1	1	4	2	GTGACGAGTA	1142	B'
108	Gumjin	T	C	G	A	C	G	A	A	T	A	2	2	4	2	TCGACGAATA	2242	G
109	Gumsun	G	T	G	A	C	G	A	A	T	A	1	1	4	2	GTGACGAATA	1142	B
110	HS	G	C	G	A	C	G	A	A	T	T	1	1	3	2	GCGACGAATT	1132	F
111	JK	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
112	K-1	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
113	KF1	G	C	G	A	T	G	A	A	T	A	1	1	4	2	GCGATGAATA	1142	H
114	KF2	G	C	T	A	C	G	A	A	T	A	1	1	3	2	GCTACGAATA	1132	I
115	KF3	G	C	T	A	C	G	A	A	T	A	1	1	3	2	GCTACGAATA	1132	I
116	KF4	G	C	G	A	C	T	A	A	T	A	1	1	4	2	GCGACTAATA	1142	C
117	KF5	G	C	G	A	C	G	A	A	T	T	1	1	3	2	GCGACGAATT	1132	F
118	M05057	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
119	SH	T	C	G	A	C	G	A	A	T	A	2	2	4	2	TCGACGAATA	2242	G
120	SP	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
121	SU	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
122	SW	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
123	YP	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
124	Cheongsong 1	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
125	ChS2	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
126	Cheongsong 3	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
127	DJ	G	C	G	A	C	G	A	A	A	A	1	1	4	2	GCGACGAAAA	1142	D
128	Gangwon 1	G	C	G	C	C	G	A	A	T	A	1	1	4	2	GCGCCGAATA	1142	J
129	G2	G	T	G	A	C	G	A	G	T	A	1	1	4	2	GTGACGAGTA	1142	B'
130	Gangwon 3	G	T	G	A	C	G	A	G	T	A	1	1	4	2	GTGACGAGTA	1142	B'
131	Gangwon 4	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A

	178	EMY	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A	
	179	G06001	G	T	G	A	C	G	A	A	T	A	1	1	4	2	GTGACGAATA	1142	B	
	180	G06003	G	C	G	A	C	G	A	A	T	A	1	2	4	2	GCGACGAATA	1242	A	
	181	G06010	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A	
	182	G08082	G	T	G	A	C	G	A	A	T	A	1	1	4	2	GTGACGAATA	1142	B	
	183	GLS	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A	
	184	JL	G	C	G	A	C	G	A	A	T	T	1	1	3	2	GCGACGAATT	1132	F	
	185	JYH	G	C	G	A	C	G	A	A	T	A	1	1	3	2	GCGACGAATA	1132	A	
W	186	YSS	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A	
	187	JF1	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A	
	188	JF2	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A	
	189	JF3	G	C	G	C	C	G	A	A	T	A	1	1	4	2	GCGCCGAATA	1142	J	
Japan	C	190	JF4	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
		191	JF5	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
		192	M 1	G	C	G	A	C	G	G	A	T	A	1	1	4	2	GCGACGGATA	1142	E
		193	Mimaki 2	G	C	G	A	C	G	G	A	T	A	1	1	4	2	GCGACGGATA	1142	E
		194	Mimaki 3	G	C	T	A	C	G	A	A	T	A	1	1	3	3	GCTACGAATA	1133	I
		195	Russia 0	G	T	G	A	C	G	A	A	T	A	1	1	4	2	GTGACGAATA	1142	B
		196	R1	G	C	G	A	C	G	A	A	T	A	1	1	4	2	GCGACGAATA	1142	A
		197	R2	G	T	G	A	C	G	A	A	T	A	1	1	4	2	GTGACGAATA	1142	B
		198	Russia 3	G	T	G	A	C	G	A	A	T	A	1	1	4	2	GTGACGAATA	1142	B
Russia	W	199	Russia 4	G	T	G	A	C	G	A	A	T	A	1	1	4	2	GTGACGAATA	1142	B
		200	Russia 5	G	T	G	A	C	G	A	A	T	A	1	1	4	2	GTGACGAATA	1142	B
		201	Russia 6	G	T	G	A	C	G	A	A	T	A	1	1	4	2	GTGACGAATA	1142	B
		202	Russia 7	G	T	G	A	C	G	A	A	T	A	1	1	4	2	GTGACGAATA	1142	B
		203	Russia 8	G	T	G	A	C	G	A	A	T	A	1	1	4	2	GTGACGAATA	1142	B

The germplasms in bold letters represent the sequenced samples.²⁾ Variation number shown in Fig. 1.³⁾ InDel markers were developed in Kim *et al.* 2015b and the genotypes were represented using the copy numbers of each TR units.

C, Cultivated ginseng; TR, Tandem repeat; W, Wild ginseng.