

Supplementary Information, Table S2. Datasets for construction of Genus *Panax* cDNA database

Dataset ^a	Plant	Organ	Year-old	Sampling location	Sequence method	Bases/reads	References
SRS006698	<i>Panax quinquefolius</i>	root	4	Beijing	454	113.4M	[1]
SRX017444	<i>Panax notoginseng</i>	root	4	Yunnan province	454	99.9M	[2]
SRS024769	<i>Panax ginseng</i>	root	11	Liaoning province	454	116.9M	[3]
ERS093423 ^b	<i>Panax ginseng</i>	root	5	Ji Lin province	HiSeq 2000	3.6G	/
ERS088918 ^b	<i>Panax ginseng</i>	stem	5	Ji Lin province	HiSeq 2000	3.5G	/
ERS088917 ^b	<i>Panax ginseng</i>	leaf	5	Ji Lin province	HiSeq 2000	3.5G	/
LIBEST_018294	<i>Panax ginseng</i>	leaf	/	/	Sanger	3160 reads	[4]
LIBEST_026760	<i>Panax ginseng</i>	root	/	/	Sanger	4226 reads	[5]
LIBEST_015794	<i>Panax ginseng</i>	MeJA treated hairy root	/	/	Sanger	3135 reads	[6]

^a accession number for SRA (Sequence Read Archive) or number for cDNA library in NCBI (<http://www.ncbi.nlm.nih.gov/>).

^b These datasets were available from DNA Data Bank of Japan(<http://www.ddbj.nig.ac.jp/searches-e.html>)

References

- 1 Sun C, Li Y, Wu Q *et al.* De novo sequencing and analysis of the American ginseng root transcriptome using a GS FLX Titanium platform to discover putative genes involved in ginsenoside biosynthesis. *BMC Genomics* 2010; **11**:262.
- 2 Luo H, Sun C, Sun Y *et al.* Analysis of the transcriptome of *Panax notoginseng* root uncovers putative triterpene saponin-biosynthetic genes and genetic markers. *BMC Genomics* 2011; **12**:S5.
- 3 Chen S, Luo H, Li Y *et al.* 454 EST analysis detects genes putatively involved in ginsenoside biosynthesis in *Panax ginseng*. *Plant Cell Rep* 2011; **30**:1593-1601.
- 4 Kim MK, Lee B-S, In J-G, Sun H, Yoon J-H, Yang D-C. Comparative analysis of expressed sequence tags (ESTs) of ginseng leaf. *Plant Cell Rep* 2006; **25**:599-606.
- 5 Han J-Y, Kim H-J, Kwon Y-S, Choi Y-E. The Cyt P450 enzyme CYP716A47 catalyzes the formation of protopanaxadiol from dammarenediol-II during ginsenoside biosynthesis in *Panax ginseng*. *Plant Cell Physiol* 2011; **52**:2062-2073.
- 6 Choi D-W, Jung J, Im Ha Y *et al.* Analysis of transcripts in methyl jasmonate-treated ginseng hairy roots to identify genes involved in the biosynthesis of ginsenosides and other secondary metabolites. *Plant Cell Rep* 2005; **23**:557-566.