

## Supplementary Information

**Table S1.** Transcriptomic database, chemical compounds, and medical uses of *P. lactiflora*, *P. suffruticosa*, *P. unicagranatum*, *R. husradicans* and *Coriarianepalensis*.

Species	Name	Samples	Family	Compounds	Chinese material	Reference
					medicine	
<i>Paeonia lactiflora</i>	HTIP	Flower buds	Paeoniaceae	Paeoniflorin, oxypaeoniflorin, benzoylpaeoniflorin, benzoyloxy paeoniflorin, oxybenzoylpaeoniflorin, albiflorin, lactiflorin, paeoniflorigenone, lactinolid, Paeonilactinone, gallpaeoninorin, albilforin, Paeonivayin, benzoieaeid, stearieacid, oleanolieae, hederagenin, betulinic acid, Gallic acid, <i>et al.</i>	Bai Shao	[12]
<i>Paeonia suffruticosa</i>	JI	Whole	Paeoniaceae	Paeonolum, Paeoniflorin, Paeonol, Paeonolide, Apiopaeonoside, Benzoylpaeoniflorin, Oxypaeoniflorin, Oleanolic Acid, Hydroxycoumarin, Gallic acid, <i>et al.</i>	Dan Pi	[13]

**Table S1.** Cont.

Species	Name	Samples	Family	Compounds	Chinese material	Reference medicine
<i>Punica granatum</i>	JROW	young leaves	Punicaceae	Gallic acid, Humarain, brevifolin, Pomegranatate, strictinin, Punicafolin, Tellimagrandin, Pedunculagin, Punicalin, 2-O-galloylpunicali, granatin, Casuarinin, 5-O-galloylpunicacortein, Punigluconin, Punicanolic acid, Friedelin, <i>et al.</i>	Shi Liu Pi	[14]
<i>Rhus radicans</i>	YUOM	Leaves stems	Anacardiaceae	Urushiol, catechol, tannic acid, gallic acid, fisetin, mangiferolic acid, anacardic acid, fustin, sulfuretin, myricetin, skikimic acid, glucoside, <i>et al.</i>	--	[15]
<i>Coriaria nepalensis</i>	NNGU	leaves	Coriariaceae	Braylin, norbraylin, dihydrocoriamyrtin, coriamyrtin, tutin, coriatin, apotutin, hydroxycoriatin, gallic acid, <i>et al.</i>	--	[16]

**Table S2.** Gene-specific primer sequences for gene identification and expression level assessment of qRT-PCR.

Gene	Forward primer (5'-3')	Reverse primer (5'-3')
HTIP-2056491	TGGGTGTGAAAGGTGCGAG	AATGGCAGACATCAACGACAG
HTIP-2000861	GGCTGGAGGGACTATGACTG	TAAACCAAATGATGGCTGC
HTIP-2007108	ACTGAGGCTGGTGAAGGTGAG	CATCCCGGTGCAGGACA
HTIP-2056592	CTTCCTTGCACCAAACCCCT	ATCATCTTCCTCCGTGGCTGTA
HTIP-2003301	ACTCGTCTACCTTGCTGCTACT	TCTTGCCAAGTGTACCCCT
HTIP-2003675	AGGGAGTTACCAAACAAATCGG	CGGTCTGGGAAACGGCG
HTIP-2057315	TGGACCCACCATAACCACG	AATGACAGCGGACACTCCTTAC
HTIP-2011414	CGGGAGGCAGAGGGAAACC	TTGATTAGCAAGAGCCACGACAG
HTIP-2056545	TGGAAGCGGTCTATCTGCG	CAGGAGGAAACTCGTGTCTTGAT
HTIP-2056640	TGGGTGGAAAGTAACTCGTGG	AAGGTCAGGCTTGTAGAAGTCATAG
HTIP-2010562	TGCCCAGCACCAACGGAC	CAGAACAAAGCCAGGGAGAATAAC
HTIP-2057123	CGCACGAATGCTTGCTTTAC	GATGGTAATCCTCCTGGCTTG

**Table S2.** Cont.

<b>Gene</b>	<b>Forward primer (5'-3')</b>	<b>Reverse primer (5'-3')</b>
HTIP-2009308	TGGATTGTTCGGAAGGAGG	TCGGATGTTAGGCGTGC
HTIP-2008205	TGAGAACAGAGTGAGCCTTGAA	CTCCGTCCATCCTGCCAC
HTIP-2056266	TGAGATGGCGAGGGTGC	TGAACTGAAGAAATGGCTGATACAAC
HTIP-2008416	ATGGACTTGGTAGGAAATGGGT	AGCAGTTGGCGATGGCAC
HTIP-2044815	GATTCATCCGCCATTCCAG	CCGTAACAACCTCCGCACTC
HTIP-2049497	GTCCCAGATTCTGTGGTG	TGCTGGGTTGGCTTCTTCC
HTIP-2007063	AGCGTCCCTTGGCTACATC	ACCGCCTTCCCACCTCG
HTIP-2057051	AGGGTGGGCAAAGGAGAC	CAAGTACAAGTGGAGTTGGAG
HTIP-2010607	CCCTCCTCCCACTCTCCTT	CTTGAGCCGAACTCGCACAT
HTIP-2054028	TGCTTCATTGCGTTGTGGAT	CAAGTTCATAACCTGCCTCGTC
HTIP-2055407	TTTCACCCAAATCTCACATCG	ACAACATTATCATTCTCGTCCACT
HTIP-2006951	TGGCAACCTCAATGACACC	CTGGTTCAAATGTGACTTTCTT
actin	GCAGTGTCCCCAGTATT	TCTTTCCATGTCATCCC

**Table S3.** HPLC conditions.

<b>T (min)</b>	<b>A (%)</b>	<b>B (%)</b>
0	95	5
25	93	7
35	88	12
65	86.5	13.5

Note: T, Retention time; A, mobile phase deionized water- formic acid (98:2, v/v); B, mobile phase acetonitrile.

**Table S4.** Chemical formulas of active compounds in *Paeonia*.

<b>Compounds</b>	<b>Formula</b>	<b>Linearity correlation (<i>R</i>)</b>
Peony lactone glycosides	$y = 2E + 06x - 19052$	0.9995
Paeoniflorin	$y = 1E + 06x + 18652$	0.9995
Benzoic acid	$y = 6E + 06x - 4934$	0.9995
Benzoyl paeoniflorin	$y = 6E + 06x + 41796$	0.9995
Paeonol	$y = 4E + 06x + 15731$	0.9995

Note: Calibration plots of four standards were constructed on the basis of peak areas (*y*) using seven different concentration solutions (*x*). All plots were linear in the examined ranges, and linear ranges had been shown as the concentration of the standard compounds (1 g mL<sup>-1</sup>). The *R* referred to the correlation coefficient of the equation. The standard compounds were purchased from National institutes for food and drug control, China.

**Table S5.** Orthologous genes related to the biosynthesis of paeoniflorin and gallic acid in *P. suffruticosa*, *Punicagranatum*, *Rhusradicans*, and *Coriarianepalensis*.

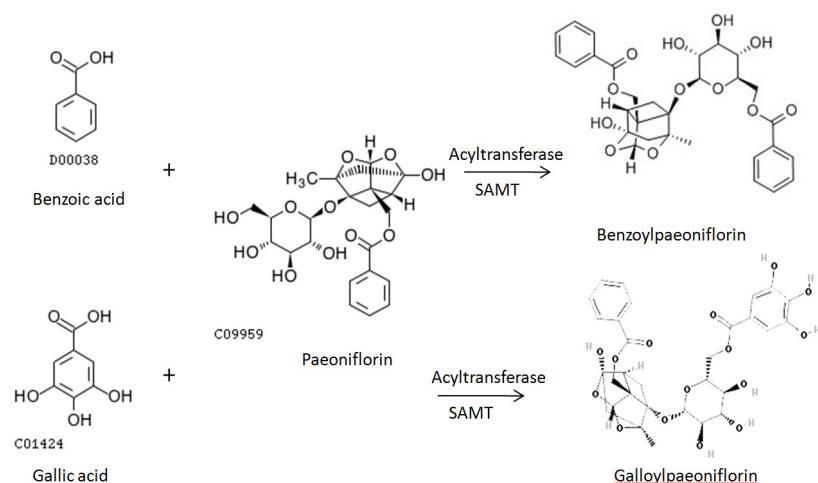
Enzyme	<i>Paeonia suffruticosa</i>	<i>Punica granatum</i>	<i>Rhus radicans</i>	<i>Coriaria nepalensis</i>
	Contig No.	Contig No.	Contig No.	Contig No.
2.5.1.54	JI449682	JROW-2002529	YUOM-2000104	NNGU-2014951
	JI444900		YUOM-2001495	NNGU-2094439
	JI444901		YUOM-2010426	
4.2.3.4	JI457522	JROW-2064854	YUOM-2006774	NNGU-2001397
				NNGU-2001398
				NNGU-2001400
4.2.1.10/1.1.1.25	JI456716	JROW-2001105	YUOM-2000314	NNGU-2094496
	JI448700	JROW-2001106	YUOM-2000315	
	JI454716	JROW-2009754	YUOM-2011625	
		JROW-2002635	YUOM-2038955	
			YUOM-2012212	
2.3.3.10	JI446744	JROW-2009760	YUOM-2011469	NNGU-2023038
	JI448691			NNGU-2023039
				NNGU-2025722
1.1.1.34	JI454933	JROW-2064932	YUOM-2004227	NNGU-2008975
			YUOM-2039378	NNGU-2008976
				NNGU-2008977
2.7.1.36	JI451094	JROW-2006872	YUOM-2004663	NNGU-2001935
				NNGU-2005819
				NNGU-2005820
2.7.4.2	JI455091	JROW-2028708	YUOM-2039067	NNGU-2091117
				NNGU-2092888
4.1.1.33	JI445250	JROW-2003445	YUOM-2038811	NNGU-2094300
		JROW-2052506		
2.3.1.9	JI447640	JROW-2060754	YUOM-2001553	NNGU-2094093
	JI451723		YUOM-2001554	NNGU-2094232
	JI447641		YUOM-2037851	
2.2.1.7	JI447268	JROW-2001269	YUOM-2010080	NNGU-2025412
	JI452657	JROW-2009533	YUOM-2039542	NNGU-2026482
	JI447269			NNGU-2094641
	JI447267			NNGU-2024936
	JI447843			
1.1.1.267	JI447843	JROW-2006718	YUOM-2005209	NNGU-2024936
2.7.7.60	JI449471	-	YUOM-2036678	NNGU-2014133
2.7.1.148	JI456924	JROW-2008008	YUOM-2037480	NNGU-2006912
				NNGU-2006914
4.6.1.12	JI453204	JROW-2064030	YUOM-2037098	NNGU-2000688
				NNGU-2000689
				NNGU-2000690

**Table S5.** Cont.

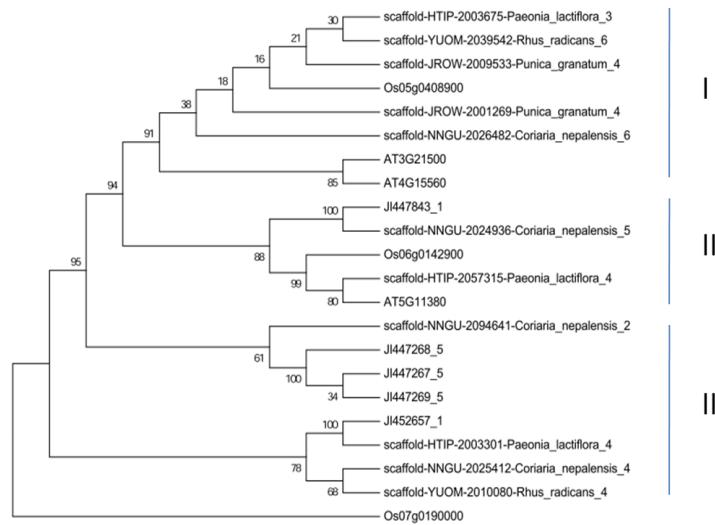
Enzyme	<i>Paeonia suffruticosa</i>	<i>Punica granatum</i>	<i>Rhus radicans</i>	<i>Coriaria nepalensis</i>
	Contig No.	Contig No.	Contig No.	Contig No.
1.17.7.1	JI448239	JROW-2005509	YUOM-2003220	NNGU-2013350
		JROW-2063357	YUOM-2003221	NNGU-2013351
		JROW-2063526		NNGU-2013743
				NNGU-2013744
1.17.7.2	JI447937	JROW-2005797	YUOM-2038387	NNGU-2026367
5.3.3.2	JI448125	JROW-2000471	YUOM-2008052	NNGU-2015100
				NNGU-2017126

**Table S6.** Correlation of active compounds content in roots of *Paeonia lactiflora*.

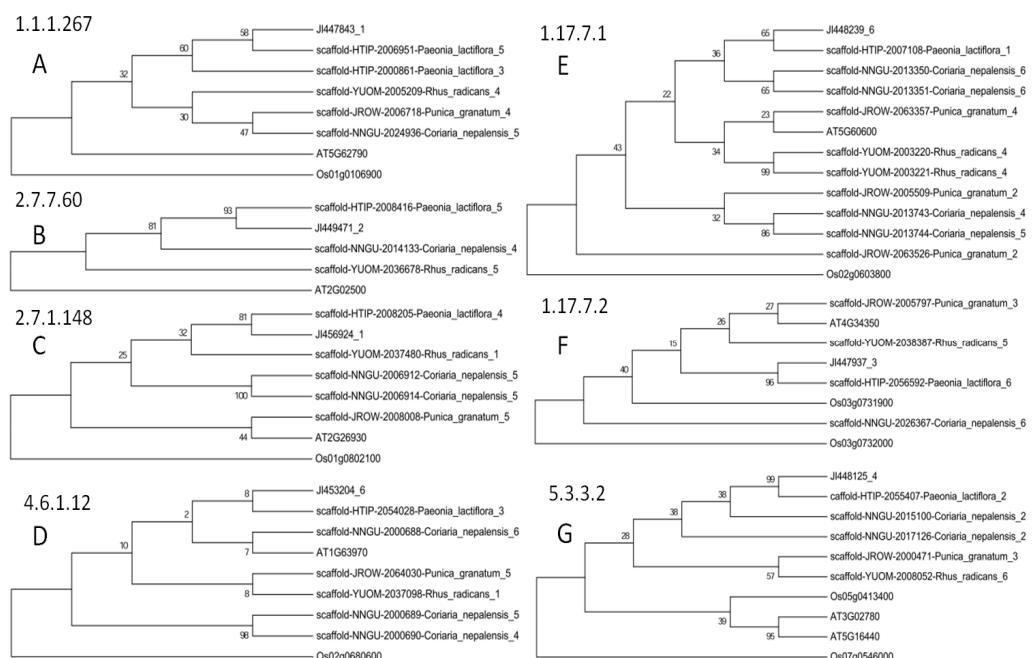
Correlation (R)	Peony lactone glycosides	Paeoniflorin	Benzoic acid	Benzoyl paeoniflorin	Paeonol
Peony lactone glycosides	1.00				
Paeoniflorin	-0.01	1.00			
Benzoic acid	-0.34	0.72	1.00		
Benzoyl paeoniflorin	-0.34	0.78	0.99	1.00	
Paeonol	0.67	0.55	0.46	0.46	1.00

**Figure S1.** Compounds in *P. lactiflora* roots.

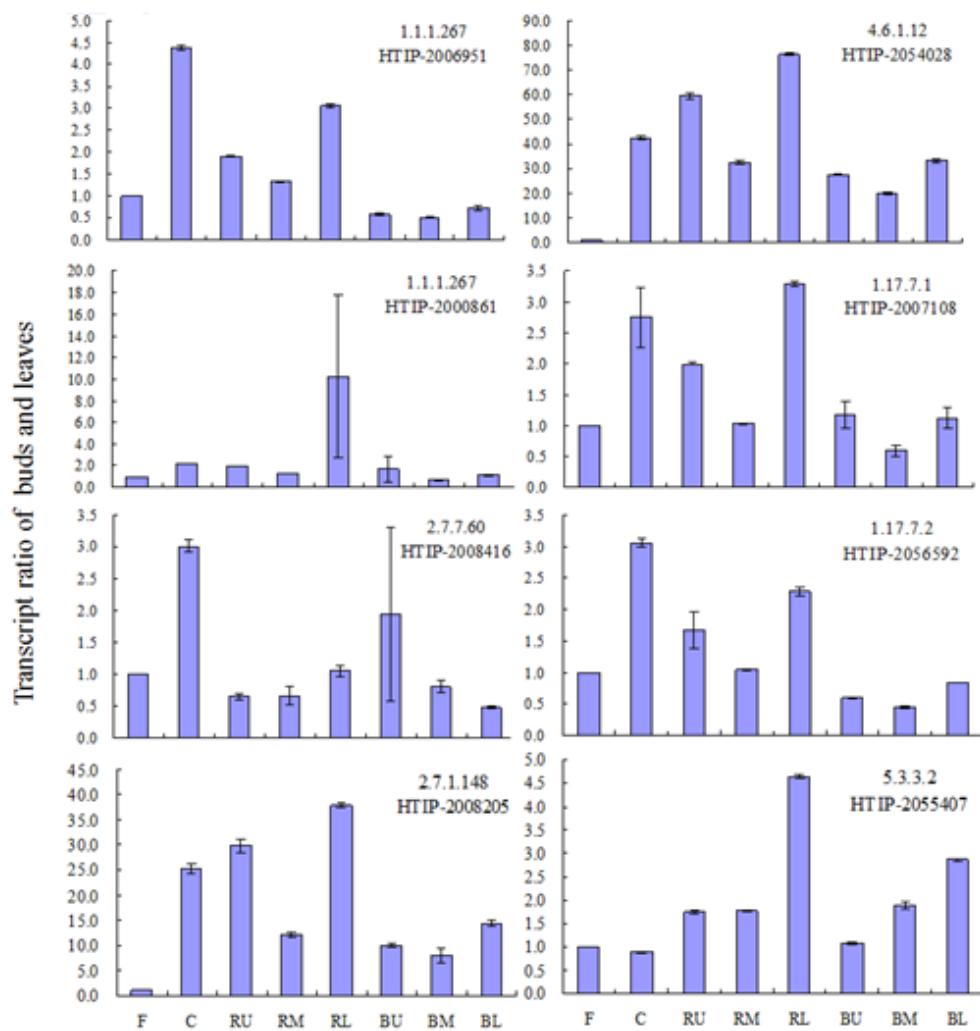
**Figure S2.** Phylogenetic relationship of plant DXPSs (E2.2.1.7). The rooted neighbor-joining tree was constructed with ClustalW [28] and all of genes were divided into three groups (I, II, and III). HTIP, *P. lactiflora*; JI, *P. suffruticosa*; JROW, *Punica granatum*; YUOM, *Rhus radicans*; NNGU, *Coriaria nepalensis*; AT, *Arabidopsis thaliana*; Os, *Oryza sativa*.



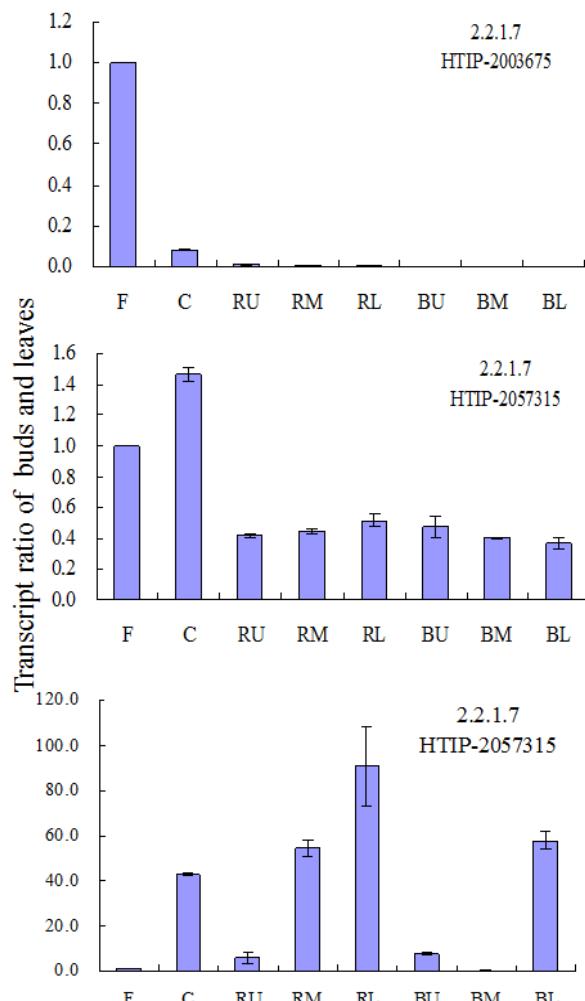
**Figure S3.** Phylogenetic relationship of plant gene families (A, DXR, E1.1.1.267; B, IspD, E2.7.7.60; C, CMK, E2.7.1.148; D, IspF, E4.6.1.12; E, HDS, E1.17.7.1; F, HDR, E1.17.7.2; and G, IDI, E5.3.3.2). The rooted neighbor-joining tree was constructed with ClustalW. HTIP, *P. lactiflora*; JI, *P. suffruticosa*; JROW, *Punica granatum*; YUOM, *Rhus radicans*; NNGU, *Coriaria nepalensis*; and AT, *Arabidopsis thaliana*.



**Figure S4.** The transcription levels of gene families (A, DXR, E1.1.1.267; B, IspD, E2.7.7.60; C, CMK, E2.7.1.148; D, IspF, E4.6.1.12; E, HDS, E1.17.7.1; F, HDR, E1.17.7.2; and G, IDI, E5.3.3.2) in different tissues and flower of *P. lactiflora*. F, flower; C, carpel; RU, RM, and RL, the upper, middle, and lower portions of the bark-free root; BU, BM, and BL, the upper, middle, and lower portions of the root bark.



**Figure S5.** Transcription levels of the DXPS (E2.2.1.7) gene family in different tissues and organs of *P. lactiflora*. F, flower; C, carpel; RU, RM, and RL, the upper, middle, and lower portions of the bark-free root; BU, BM, and BL, the upper, middle, and lower portions of the root bark.



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