

## Supplementary Materials

### Quantitative and Differential Analysis between *Bupleurum chinense* DC. and *Bupleurum scorzonerifolium* Willd. Using High-Performance Liquid Chromatography-Mass Spectrometry and Gas Chromatography-Mass Spectrometry Coupled with Multivariate Statistical Analysis

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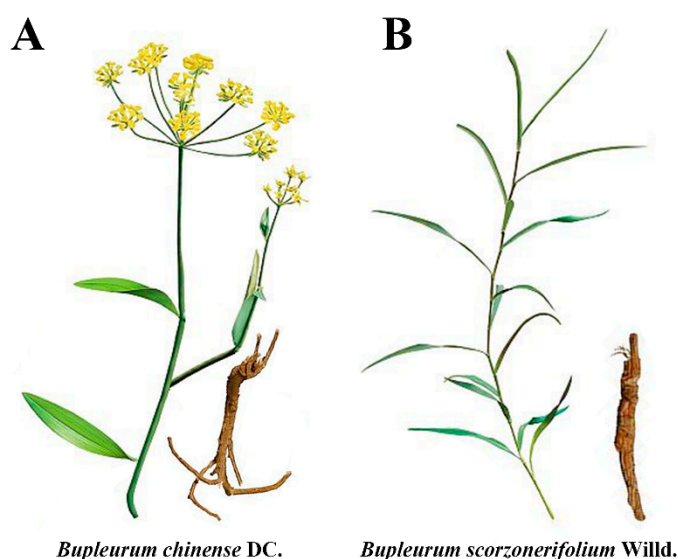
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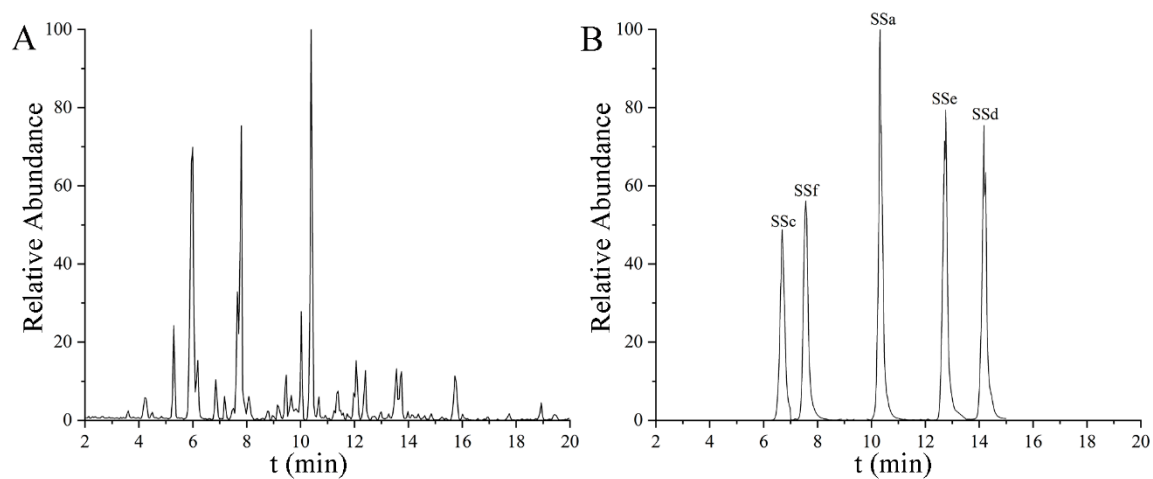
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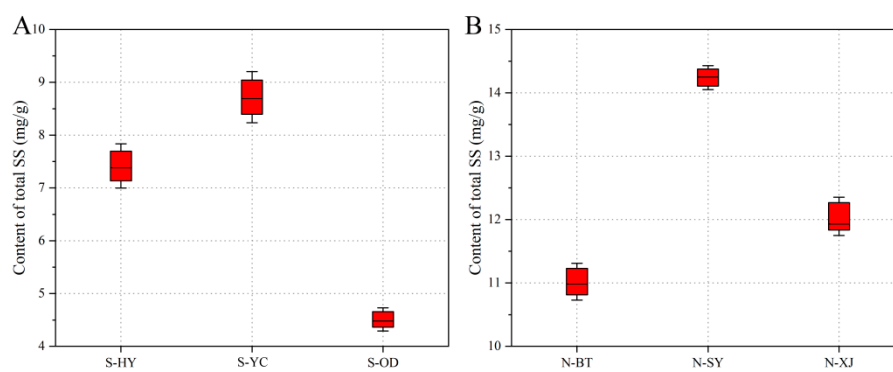
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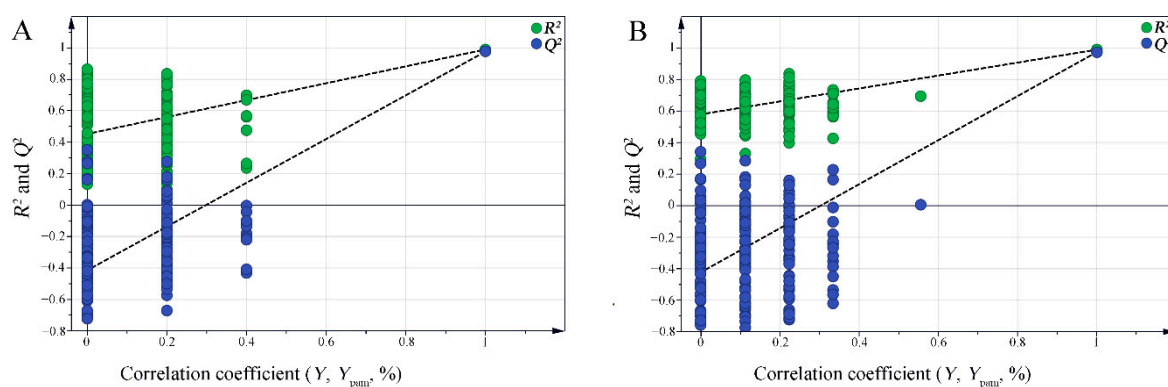
**Figure S1.** The pictures of *Bupleurum chinense* DC. (A) and *Bupleurum scorzonerifolium* Willd. (B).



**Figure S2.** Total ion chromatograms of extracted saponins detected by HPLC-MS in full scan (A) and MRM mode (B).



**Figure S3.** Boxplots of the total saikosaponin contents in the South (A) and North (B) RB samples from different growing regions.



**Figure S4.** Permutation tests of PLS-DA models derived from the HPLC-MS data sets of saikosaponins (A) and the GC-MS data sets of volatile oils (B) in the Radix Bupleuri samples.

**Table S1.** The absolute contents of the five and total saikosaponins in the North and South RB samples.

No.	Content of SS <sup>(1)</sup> (μg/mg)					
	SSa	SSc	SSd	SSe	SSf	Total SS
N-BT-1 <sup>(2)</sup>	3.523±0.008	0.820±0.006	5.853±0.007	0.048±0.004	0.334±0.011	10.578±0.047
N-BT-2	3.572±0.023	0.829±0.002	6.070±0.029	0.036±0.013	0.398±0.009	10.904±0.050
N-BT-3	3.632±0.012	0.834±0.010	6.233±0.009	0.055±0.007	0.308±0.006	11.063±0.016
N-BT-4	3.568±0.009	0.821±0.031	5.877±0.004	0.057±0.017	0.328±0.007	10.652±0.023
N-BT-5	3.710±0.012	0.815±0.043	6.314±0.023	0.033±0.008	0.349±0.009	11.221±0.011
N-BT-6	3.714±0.012	0.845±0.022	6.730±0.004	0.042±0.002	0.378±0.012	11.710±0.021
N-SY-1	4.564±0.051	0.935±0.066	8.076±0.046	0.095±0.003	0.619±0.007	14.290±0.056
N-SY-2	4.639±0.064	0.962±0.005	7.873±0.034	0.107±0.009	0.628±0.006	14.209±0.087
N-SY-3	4.378±0.039	0.955±0.005	8.470±0.035	0.094±0.011	0.638±0.009	14.535±0.077
N-SY-4	4.589±0.054	0.983±0.009	8.164±0.067	0.141±0.002	0.651±0.023	14.528±0.022
N-SY-5	4.219±0.006	0.968±0.015	7.971±0.008	0.111±0.013	0.613±0.029	13.882±0.052
N-SY-6	4.182±0.051	0.951±0.013	8.097±0.005	0.105±0.002	0.664±0.019	13.998±0.067
N-XJ-1	3.816±0.028	0.892±0.011	6.514±0.006	0.079±0.007	0.522±0.022	11.823±0.039
N-XJ-2	3.670±0.039	0.891±0.009	6.888±0.041	0.055±0.009	0.503±0.002	12.037±0.022
N-XJ-3	3.822±0.077	0.911±0.024	6.469±0.019	0.057±0.006	0.515±0.008	11.775±0.091
N-XJ-4	3.956±0.008	0.922±0.003	7.452±0.014	0.062±0.008	0.469±0.009	12.866±0.068
N-XJ-5	3.914±0.009	0.894±0.005	6.739±0.091	0.091±0.001	0.478±0.007	12.116±0.011
N-XJ-6	3.855±0.042	0.860±0.029	6.418±0.006	0.066±0.007	0.494±0.004	11.694±0.041
S-HY-1 <sup>(3)</sup>	2.499±0.049	0.611±0.016	3.851±0.009	0.024±0.002	0.146±0.014	7.118±0.087
S-HY-2	2.533±0.023	0.614±0.007	3.993±0.047	0.020±0.002	0.174±0.007	7.327±0.059
S-HY-3	2.576±0.025	0.611±0.003	4.101±0.079	0.019±0.003	0.135±0.004	7.438±0.073
S-HY-4	2.530±0.024	0.605±0.005	3.867±0.038	0.024±0.001	0.143±0.002	7.169±0.079
S-HY-5	2.632±0.028	0.623±0.007	4.154±0.047	-	0.152±0.001	7.573±0.091
S-HY-6	2.634±0.025	0.631±0.005	4.428±0.068	0.019±0.003	0.165±0.001	7.873±0.089
S-YC-1	2.888±0.042	0.705±0.007	4.573±0.072	0.027±0.002	0.182±0.003	8.375±0.075
S-YC-2	2.928±0.027	0.715±0.014	4.742±0.074	0.024±0.004	0.216±0.002	8.625±0.086
S-YC-3	2.977±0.072	0.721±0.026	4.870±0.086	0.019±0.001	0.168±0.003	8.754±0.087

S-YC-4	2.924±0.043	0.708±0.007	4.592±0.087	0.022±0.002	0.178±0.004	8.419±0.091
S-YC-5	3.041±0.045	0.714±0.015	4.933±0.064	0.021±0.002	0.189±0.001	8.898±0.091
S-YC-6	3.044±0.036	0.721±0.007	5.258±0.075	-	0.206±0.007	9.239±0.098
S-OD-1	1.367±0.017	0.586±0.007	2.314±0.035	0.010±0.002	0.065±0.007	4.342±0.058
S-OD-2	1.406±0.035	0.546±0.015	2.400±0.053	-	0.078±0.014	4.436±0.058
S-OD-3	1.410±0.046	0.570±0.016	2.464±0.062	0.019±0.002	0.070±0.005	4.527±0.078
S-OD-4	1.445±0.017	0.570±0.005	2.323±0.033	0.021±0.017	0.064±0.018	4.415±0.059
S-OD-5	1.461±0.035	0.575±0.015	2.496±0.046	0.020±0.001	0.061±0.007	4.599±0.076
S-OD-6	1.452±0.014	0.553±0.013	2.660±0.035	0.018±0.003	0.074±0.012	4.748±0.085

- <sup>(1)</sup> SSa, SSc, SSd, SSe, SSf, and SS are the abbreviations of saikosaponin a, saikosaponin c, saikosaponin d, saikosaponin e, saikosaponin f, and saikosaponins, respectively.
- <sup>(2)</sup> N-BT, N-SY, and N-XJ represent *Bupleurum chinense* DC. grown in Baotou city, Sanyuan county, and Xinjiang county, respectively.
- <sup>(3)</sup> S-HY, S-YC, and S-OD represent *Bupleurum scorzonerifolium* Willd. grown in Heyang county, Yuncheng city, and Ordos city, respectively.

**Table S2.** ANOVA of the effects of variety and growing region on the contents of the five saikosaponins.

Factor	SSa	SSc	SSd	SSe	SSf
Variety	*	*	*	*	*
Growing region	*	*	*	*	*

\* significant difference at the level of  $p < 0.05$ .

**Table S3.** Comparison of current and reported work

	Present study	Reference 6*
Plants	18 batches of two-year-old North RB and 18 batches of two-year-old South RB, collected from Shaanxi Province, Shanxi Province, and Inner Mongolia Autonomous Region, China.	12 wild North RB and 12 wild South RB, at least one-year-old, collected from the suburban mountainous areas of Beijing, China,
Parts of the plants	Root	Root, stem, leaf, and flower
Extraction method	Sonication in 15% ammonia-methanol	Immersion in 70% aqueous methanol and

	solution and steam distillation	HS-SPME
LC-MS conditions	HPLC-QqQ/MS in MRM and fullscan mode, C18 column, mobile phase: H <sub>2</sub> O with 0.1% formic acid (A) and acetonitrile (B)	HPLC-QqQ-LIT/MS in MRM and fullscan mode, C18 column, mobile phase: H <sub>2</sub> O (A) and acetonitrile with 0.1% acetic acid (B)
GC-MS conditions	GC-QqQ/MS in SIM and fullscan mode, EI at 70 eV, DB-5MS capillary column, helium at 1.2 mL/min in split mode	EI at 70 eV, DB-5MS capillary column, helium at 1.2 mL/min in splitless mode
Quantitative analysis	Absolute saikosaponin contents	Relative metabolite contents
Qualitative analysis	Tandem MS analysis, retrieval of references and NIST 14 database	Self-compiled database
Statistical analysis	HCA, PCA, PLS-DA, ANOVA, T-test	HCA, PCA, OPLS-DA, Venn diagrams, pathway annotation and enrichment analysis
Results and conclusions	<ol style="list-style-type: none"> <li>1. The absolute content of 5 saikosaponins in Radix Bupleuri from different varieties and regions were obtained.</li> <li>2. Twenty-one saponins and fifty-two volatile oils were identified as the differential compounds.</li> <li>3. The content of saikosaponins was significantly higher in North RB than in South RB.</li> <li>4. There were significant differences in saikosaponins and volatile oils between the two varieties, whether grown in the same region or not.</li> <li>5. The identified differential compounds may be the main active components responsible for the differences in their clinical efficacy.</li> </ol>	<ol style="list-style-type: none"> <li>1. There were significant differences in metabolites between different tissues.</li> <li>2. 144 common differential metabolites were identified between different tissues.</li> <li>3. The content of active components was higher in the root of North RB than in South RB.</li> <li>4. Differential metabolites of the aboveground parts mainly concentrated in monoterpene biosynthesis, while those of the root mainly concentrated in sesquiterpene and triterpene biosynthesis.</li> <li>5. The difference in metabolites and metabolic networks between the two varieties resulted in the difference in the clinical efficacy of Radix Bupleuri.</li> </ol>

\* Qu, X.; Hu, S.; Li, T.; Zhang, J.; Wang, B.; Liu, C. Metabolomics Analysis Reveals the Differences Between *Bupleurum chinense* DC. and *Bupleurum scorzonifolium* Willd. *Front. Plant. Sci.* **2022**, *13*, 933849.