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Studies on chemical profiling and pharmacokinetics of traditional Chinese medicine Formula *Kang Shuai Lao Pian*

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ABSTRACT

Kang Shuai Lao Pian (KSLP) is a traditional Chinese medicine (TCM) preparation used to delay aging. However, due to the lack of research on the chemical composition and pharmacokinetic behavior of KSLP, its material basis and *in vivo* components with high exposure remain ambiguous. The UPLC/Q-Orbitrap-MS/MS was performed to identify the preliminary chemical profile of KSLP. A total of 138 compounds, including ginsenosides, phenyl-ethanol glycosides, iridoids, alkaloids, ionones and others, were identified in accordance with their retention times, accurate masses and characteristic MS/MS fragment patterns. Moreover, considering the active components and characteristic components of KSLP, the extraction process of KSLP was optimized, and the quantitative analysis by UPLC/QQQ-MS/MS of 13 compounds in KSLP was established. The method was stable and sensitive, and could be used for the quality control of KSLP. Then, the pharmacokinetic study was carried out by further refining the components of KSLP. Besides, quantitative method for 6 compounds in rat plasma was validated and developed by UPLC/QQQ-MS/MS. The established approach was successfully applied to characterize the pharmacokinetic features of components in KSLP and it was found that the absorption and elimination of ginsenosides in KSLP was slow. Altogether, this study laid a solid foundation and provided theoretical guidance for further clarification of bioactive components of KSLP.

1. Introduction

Aging refers to the process of degenerative changes in various tissues and organs as the body ages after reaching maturity in growth and development, which is affected by genetic, lifestyle and environmental factors (Antell and Taczanowski, 1999, Lee et al., 2016, Noroozi et al., 2021). Population aging is reportedly one of the most significant trends of the 21st century. Therefore, there is a growing need for aging management, and healthy aging is the main goal of aging intervention and research. In recent years, although new drugs have been developed to improve aging, most of them are expensive and have side effects, so it is particularly important to find safe and effective drugs to delay aging (Blagosklonny, 2007, Li et al., 2013). Clinical practice has proved that

traditional Chinese medicine (TCM) has the advantages of multiple targets and few side effects, and has played a pivotal role in maintaining human healthcare (Zhao et al., 2020).

Kang Shuai Lao Pian (KSLP) is a famous TCM formulated from a court prescription of the Ming Dynasty. It is prepared from *Rehmannia radix*, *Ginseng radix et rhizoma rubra*, *Asparagi radix*, *Ophiopogonis radix*, *Lych cortex* and *Poria* in the weight ratio of 409, 167, 26, 26, 26, 77 (Gong et al., 2020a, 2020b). In China, it has been widely accepted as a health care product for delaying senescence (Gong et al., 2020a, 2020b). Obviously, the identification and detection of the main components in KSLP is the premise and key to reveal its active ingredients. The chemical constituents of each crude drug in KSLP have been reported in previous studies, but little attention has been paid to the integral

Abbreviations: KSLP, *Kang Shuai Lao Pian*; TCM, traditional Chinese medicine; MRM, multiple reaction monitoring; QC, quality control; CE, collision energy; CV, cone voltage; LOD, limit of detection; LOQ, limit of quantification; S/N, single to noise ratios; IS, internal standard solution; LLOQ, lower limit of quantification; BPI, base peak ion chromatograms; PPD, protopanaxadiol; PPT, protopanaxatriol.

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chemical composition of KSLP. In the process of drug development, pharmacokinetic research is an indispensable part of the process, which has important reference value for drug development, evaluation and clinical application (Zhang et al., 2017). However, to the best of our knowledge, no pharmacokinetic study on KSLP has been reported. Thus, the establishment of comprehensive qualitative, quantitative and pharmacokinetic methods for KSLP is urgently needed.

In recent years, with the development of analytical technology, UPLC/Q-Orbitrap-MS/MS has been widely used for the analysis of TCM due to its fast separation speed and high sensitivity (Han et al., 2015). The UPLC/QQQ-MS/MS technology is a powerful tool for high-throughput quantitative analysis of TCM owing to its high-selective simultaneous detection of multiple compounds with a multiple reaction monitoring (MRM) mode (Liu et al., 2017, Ren et al., 2022). The integration of UPLC/Q-Orbitrap-MS/MS and UPLC/QQQ-MS/MS is a potentially effective approach for in-depth chemical profiling and pharmacokinetic of KSLP.

In this study, the UPLC/Q-Orbitrap-MS/MS analysis method was established for the global characterizations of chemical composition in KSLP. Besides, 13 compounds were further quantitatively analyzed with UPLC/QQQ-MS/MS considering the representative components of KSLP, the abundance and activity of chemicals. The extraction and refining process of components in KSLP was optimized to maximize the content of compounds, and then the UPLC/QQQ-MS/MS method was applied to study the pharmacokinetics of refined components in KSLP. Through the chemical analysis of KSLP, the material basis was clarified and a reference was provided as its quality evaluation. Pharmacokinetic studies were conducted to reveal the changes of refined components in KSLP *in vivo* and provided a basis for their clinical application.

2. Materials and methods

2.1. Chemicals and materials

Acetonitrile (chromatographic purity) was purchased from Fisher company (USA), formic acid (MS grade) was obtained from ACS company (USA). Reference standards of acteoside, isoacteoside, echinacoside, jionoside A1, ginsenoside Rb1, ginsenoside Rb2, ginsenoside Rc, ginsenoside Rd, ginsenoside Re, ginsenoside Rf, ginsenoside Rg1, ginsenoside Rg2, ginsenoside Rg3 and digoxin were acquired from Sichuan Weikeqi Biochemical Co., Ltd. (Sichuan, China). KSLP (batch number: 2207001) was supplied by Chiatai Qingchunbao Medicine Co., Ltd. (Hangzhou, China).

2.2. Qualitative analysis by UPLC/Q-Orbitrap-MS/MS

2.2.1. Sample solutions preparation

0.1 g KSLP powder was added to 10 mL of 50 % ethanol, and ultrasonically extracted for 60 min. The solution was centrifuged at 15,000 *g* for 10 min, and the supernatant was used for UPLC/Q-Orbitrap-MS/MS qualitative analysis.

2.2.2. Standard solutions preparation

The standards of acteoside, isoacteoside, echinacoside, jionoside A1, ginsenoside Rb1, ginsenoside Rb2, ginsenoside Rc, ginsenoside Rd, ginsenoside Re, ginsenoside Rf, ginsenoside Rg1, ginsenoside Rg2 and ginsenoside Rg3 were weighed accurately and dissolved in 50 % ethanol to 1 mg/mL single standard solutions. 50 μ L of each standard solutions were added to 350 μ L of 50 % ethanol, and the mixed standard solution was collected after vortex mixing 1 min and the following centrifugation (12000 *g*, 10 min, 4°C).

2.2.3. Chromatographic and mass spectrographic conditions

The UPLC separation was performed on Ultimate 3000 UPLC system (Thermo Fisher Scientific, USA) with an ACQUITY UPLC HSS T3 (2.1 \times 100 mm, 1.7 μ m, Waters) maintained at 40°C. The mobile phase

was acetonitrile (A) and water with 0.1 % formic acid (B) at a flow rate of 0.3 mL/min, and injection volume of samples was 4 μ L. The gradient elution procedure was as follows: 0 ~ 2 min, 3 % (A); 2 ~ 5 min, 3 ~ 21 % (A); 5 ~ 7 min, 21 ~ 22 % (A); 7 ~ 9 min, 22 ~ 30 % (A); 9 ~ 13 min, 30 ~ 33 % (A); 13 ~ 18 min, 33 ~ 34 % (A); 18 ~ 23 min, 34 ~ 55 % (A); 23 ~ 28 min, 55 ~ 70 % (A); 28 ~ 33 min, 70 ~ 98 % (A).

The mass spectrometry analysis was completed under Q-Orbitrap MS system (Thermo Fisher Scientific, USA). The optimized MS parameters were: ESI positive and negative (+/-) mode; spray voltage, 3.5kV; capillary temperature, 350°C; aux gas heater temperature, 350 °C; sheath gas was nitrogen and its flow was 35 L/h; auxiliary gas was nitrogen and its flow was 10 L/h; full scan mode, *m/z* 100–1500.

2.3. Quantitative analysis by UPLC/QQQ-MS/MS

2.3.1. Sample solutions preparation

In this study, the effects of four single factors (extraction method, extraction solution, extraction time and solid-to-liquid ratio) on the contents of compounds were investigated. The maximum total content of the 13 target compounds was used as the criterion to determine the optimal extraction process.

The levels of the four single factors were as follows: ultrasonic and reflux for extraction method; 30 % ethanol, 50 % ethanol, 70 % ethanol and 90 % ethanol for extraction solution; 30, 60, 90 and 120 min for extraction time; 1:20, 1:50, 1:100 and 1:200 g/mL for solid-to-liquid ratio. When the influence of any single factor was investigated on the total content of 13 target compounds, 50 % ethanol, ultrasonic, 90 min and 1:100 g/mL were selected for the other three factors.

The samples were extracted according to each extraction process, diluted with the corresponding solutions, and finally 100 μ g/mL of sample solutions were obtained for UPLC/QQQ-MS/MS quantitative analysis.

2.3.2. Standard solutions preparation

The stock solutions of acteoside, isoacteoside, echinacoside, jionoside A1, ginsenoside Rb1, ginsenoside Rb2, ginsenoside Rc, ginsenoside Rd, ginsenoside Re, ginsenoside Rf, ginsenoside Rg1, ginsenoside Rg2 and ginsenoside Rg3 were prepared in 50 % ethanol. Working standard solutions containing each of the 13 compounds were prepared by diluting the stock solutions with 50 % ethanol to a series of proper concentrations. Similarly, the working solutions of quality control (QC) with low, medium and high concentrations were prepared. QC samples contained acteoside, isoacteoside, echinacoside, jionoside A1, ginsenoside Rg2 and ginsenoside Rg3 of 1, 10 and 80 ng/mL, ginsenoside Rb1, ginsenoside Rb2, ginsenoside Rc, ginsenoside Rd, ginsenoside Re, ginsenoside Rf and ginsenoside Rg1 of 10, 100 and 800 ng/mL.

2.3.3. Chromatographic and mass spectrographic conditions

The chromatographic separation was achieved on an ACQUITY UPLC Ultra Performance Liquid Chromatograph (Waters, American) with an ACQUITY UPLC HSS T3 (2.1 \times 100 mm, 1.7 μ m, Waters) maintained at 40°C. The mobile phase was acetonitrile (A) and water with 0.1 % formic acid (B) at a flow rate of 0.3 mL/min, and injection volume of samples was 4 μ L. The gradient elution was performed as follows: 0 ~ 2.5 min, 20 ~ 30 % (A); 2.5 ~ 10 min, 30 ~ 60 % (A); 10 ~ 10.5 min, 60 ~ 98 % (A).

MS detection was carried on Waters Xevo TQ-S Triple Quadrupole Mass Spectrometer (Waters, American), and the mass spectrometer was operated in a MRM mode. The optimized MS parameters: capillary voltage, 3.0 KV (positive ion mode)/ 2.0 KV (negative ion mode); solvent removal temperature, 350°C. The remaining specific parameters, such as parents ion (Parents), daughters ion (Daughters), collision energy (CE) and cone voltage (CV), were optimized and summarized in Table S1.

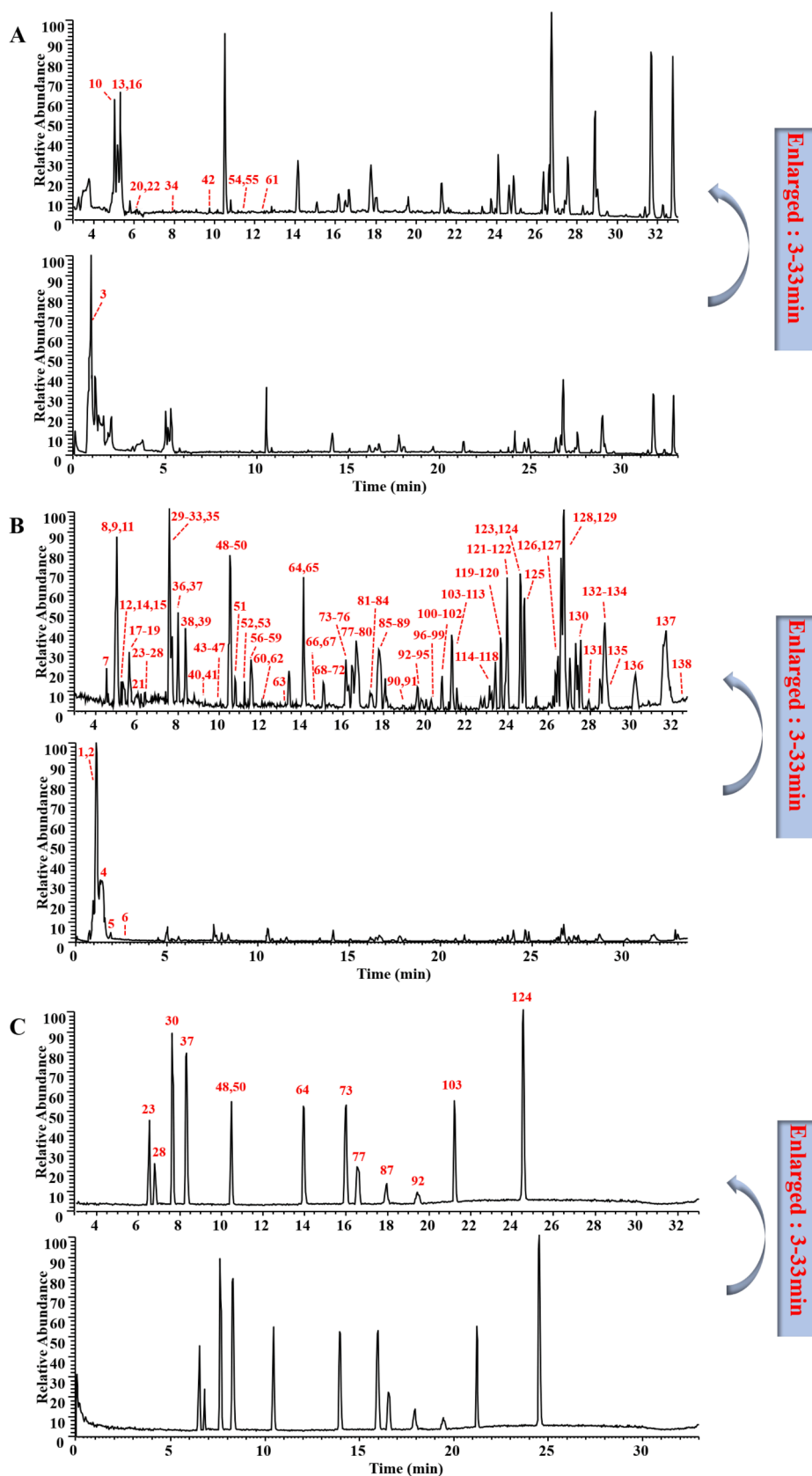


Fig. 1. The base peak ion chromatograms of KSLP samples and standards (A: the sample under positive ion mode; B: the sample under negative ion mode; C: the standards under negative ion mode).

Table 1
Characterization of chemical constituents of KSLP by UPLC/Q-Orbitrap-MS/MS.

NO.	Observed (m/z)	Formula	t _R (min)	Mass error (ppm)	Adducts	Identification	Source	ESI-MS ²
1	407.1197	C ₁₅ H ₂₂ O ₁₀	0.94	0.493	[M + COOH] ⁻	Monomelittoside	RG	361.1140 199.0605 169.0497
2	191.0191	C ₆ H ₈ O ₇	0.96	-0.626	[M-H] ⁻	Citric Acid	RG	173.0441 127.0388 87.0074 85.0282
3	268.1043	C ₁₀ H ₁₃ N ₅ O ₄	1.13	1.006	[M + H] ⁺	Adenosine	PG	136.0618 119.0353
4	191.0191	C ₆ H ₈ O ₇	1.46	-0.626	[M-H] ⁻	3-carboxy-2,3-dideoxy-1-hydroxypropan-1,2,3-tricarboxylic acid	RG	173.0083 129.0180 111.0075 87.0074 85.0282
5	361.1144	C ₁₅ H ₂₂ O ₁₀	1.83	1.052	[M-H] ⁻	Catalpol	RG	199.0604 181.0497 169.0497
6	125.0232	C ₆ H ₆ O ₃	2.86	-9.737	[M-H] ⁻	5-hydroxymethylfurfural	AC	97.0281 87.9239 81.0333 69.0332
7	731.2261	C ₂₇ H ₄₂ O ₂₀	4.38	1.305	[M + COOH] ⁻	Rehmannioside D	RG	505.1034 341.1085 179.0553
8	373.1141	C ₁₆ H ₂₂ O ₁₀	4.62	0.08	[M-H] ⁻	Geniposidic acid	RG	211.0605 193.0499 179.0554 167.0704 123.0439
9	167.0343	C ₈ H ₈ O ₄	4.91	-4.083	[M-H] ⁻	Vanillic acid	RG	152.0105 137.0230 124.0755 93.0332
10	251.1393	C ₁₃ H ₁₈ N ₂ O ₃	5.02	1.119	[M + H] ⁺	N-caffeoyl putrescine	LC	234.1125 190.4467 163.0389 135.0441
11	373.1142	C ₁₆ H ₂₂ O ₁₀	5.07	0.18	[M-H] ⁻	Gardoside	RG	211.0606 149.0597 123.0439
12	347.1349	C ₁₅ H ₂₄ O ₉	5.21	0.416	[M-H] ⁻	Ajugol	RG	317.1079 235.1352 225.1024 167.0707 123.0802
13	531.3179	C ₂₈ H ₄₂ N ₄ O ₆	5.26	0.355	[M + H] ⁺	Kukoamine A	LC	513.3179 367.2735 293.1857 222.1124 165.0547
14	461.1669	C ₂₀ H ₃₀ O ₁₂	5.31	0.977	[M-H] ⁻	Decaffeoyl verbascoside	RG	315.1082 297.0962 161.0446 135.0439
15	375.1294	C ₁₆ H ₂₄ O ₁₀	5.38	-0.72	[M-H] ⁻	Mussaenosidic acid	RG	213.0761 195.0652 179.0553 169.0860
16	531.318	C ₂₈ H ₄₂ N ₄ O ₆	5.5	0.543	[M + H] ⁺	Kukoamine B	LC	513.3069 293.1858 222.1124 165.0547
17	345.1178	C ₁₄ H ₂₀ O ₇	5.52	-3.783	[M + COOH] ⁻	Salidroside	RG	179.0552 119.0488 101.0230
18	345.1556	C ₁₆ H ₂₆ O ₈	5.52	0.316	[M-H] ⁻	Rehmapicroside	RG	179.0552 165.0907 119.0488
19	487.1462	C ₂₁ H ₂₈ O ₁₃	5.56	0.998	[M-H] ⁻	Isomer of 3-O-(6-Deoxy-α-L-mannopyranosyl)-4-O-[(2E)-3-(3,4-dihydroxyphenyl)-2-propenoyl]-D-glucose	RG	179.0341 161.0234
20	474.2602	C ₂₅ H ₃₅ N ₃ O ₆	6.11	0.712	[M + H] ⁺	N1, N10-bis (dihydro-caffeoyl) spermidine	LC	457.2328 293.1862 236.1283 222.1126 165.0547
21	487.1462	C ₂₁ H ₂₈ O ₁₃	6.24	-0.998	[M-H] ⁻	3-O-(6-Deoxy-α-L-mannopyranosyl)-4-O-[(2E)-3-(3,4-dihydroxyphenyl)-2-propenoyl]-D-glucose	RG	179.0341 161.0234
22	472.2451	C ₂₅ H ₃₃ N ₃ O ₆	6.25	1.88	[M + H] ⁺	N5-caffeoyl-N10-caffeoylspermidine	LC	454.2336 310.2125 293.1858 220.0968 163.0389
23 [#]	785.2518	C ₃₅ H ₄₆ O ₂₀	6.45	0.947	[M-H] ⁻	Echinacoside	RG	623.2195 477.1616 459.1498 392.5579 315.1089 179.0342 161.0235
24	785.2513	C ₃₅ H ₄₆ O ₂₀	6.55	0.424	[M-H] ⁻	Purpureaside C	RG	623.2187 459.1502 161.0234
25	283.055	C ₁₅ H ₂₄ O ₅	6.63	-0.343	[M-H] ⁻	Hydroxy-acetic acid	RG	183.1018 153.0909 139.1116
26	451.2189	C ₁₉ H ₃₄ O ₉	6.67	0.902	[M + COOH] ⁻	Oxyrehmaionoside B	RG	243.1595 213.1493 179.0556 161.0447
27	403.1248	C ₁₇ H ₂₄ O ₁₀	6.74	0.534	[M-H] ⁻	Gardenoside	RG	371.0966 223.0605 165.0546
28 [#]	799.2673	C ₃₆ H ₄₈ O ₂₀	6.84	0.855	[M-H] ⁻	Jionoside A1	RG	623.2185 605.2119 477.1590 459.1524 422.5963
29	435.2236	C ₁₉ H ₃₄ O ₈	7.38	0.068	[M + COOH] ⁻	Rehmannioside B	RG	179.0557 161.0443
30 [#]	623.1985	C ₂₉ H ₃₆ O ₁₅	7.66	0.572	[M-H] ⁻	Acteoside	RG	461.1662 315.1082 161.0234
31	183.1019	C ₁₀ H ₁₆ O ₃	7.73	-4.192	[M-H] ⁻	(25R)-5β-Spirostane-1β,3β,25-triol	RG	139.1116 137.0959
32	813.283	C ₃₇ H ₅₀ O ₂₀	7.8	0.901	[M-H] ⁻	Jionoside B1	RG	637.2353 619.2266 491.1768 175.0391
33	435.2236	C ₁₉ H ₃₄ O ₈	7.95	0.068	[M + COOH] ⁻	Rehmaionoside A/B	RG	389.2188 179.0557 161.0442
34	193.0497	C ₁₀ H ₈ O ₄	7.96	0.853	[M + H] ⁺	Scopoletin	LC	178.0262 165.1270 149.1328 133.1013
35	193.0499	C ₁₀ H ₁₀ O ₄	7.98	-3.792	[M-H] ⁻	Ferulic acid	AC	178.0263 149.0596 134.0361
36	435.2236	C ₁₉ H ₃₄ O ₈	8.16	0.068	[M + COOH] ⁻	Rehmaionoside A/B	RG	389.2184 179.0552 161.0445
37 [#]	623.1984	C ₂₉ H ₃₆ O ₁₅	8.3	0.412	[M-H] ⁻	Isoacteoside	RG	461.1658 315.1091 161.0234
38	497.1667	C ₂₃ H ₃₀ O ₁₂	8.48	0.504	[M-H] ⁻	6-O-vanillate ajugol	RG	317.1029 167.0341 152.0105 123.0439
39	623.1984	C ₂₉ H ₃₆ O ₁₅	8.52	0.412	[M-H] ⁻	Forsythoside A	RG	461.1689 161.0234
40	637.2142	C ₃₀ H ₃₈ O ₁₅	9.24	0.638	[M-H] ⁻	Jionoside C	RG	526.9341 461.1667 315.1086 193.0498 175.0392
41	1007.5448	C ₄₈ H ₈₂ O ₁₉	9.42	1.556	[M + COOH] ⁻	20-Glc-Ginsenoside Rf	PG	961.5378 799.4826 637.4280 619.4224 475.3815 391.2862
42	874.3734	C ₄₂ H ₅₁ N ₉ O ₁₂	9.73	0.464	[M + H] ⁺	Lyciumin A	LC	856.3506 486.1976 468.1875 442.1821
43	977.5332	C ₄₇ H ₈₀ O ₁₈	9.75	0.544	[M + COOH] ⁻	Notoginsenoside R1/Fp1/Ginsenoside Re4	PG	931.5276 799.4855 637.4318 475.3791 391.2839

(continued on next page)

Table 1 (continued)

NO.	Observed (m/z)	Formula	t _R (min)	Mass error (ppm)	Adducts	Identification	Source	ESI-MS ²
44	1007.5446	C ₄₈ H ₈₂ O ₁₉	9.82	1.357	[M + COOH] ⁻	Notoginsenoside N	PG	961.5381 799.4863 637.4366 475.3795 391.2864
45	977.5338	C ₄₇ H ₈₀ O ₁₈	9.96	1.158	[M + COOH] ⁻	Notoginsenoside R1/Fp1/Ginsenoside Re4	PG	931.5278 799.4862 637.4304 475.3769 391.2843
46	1007.5447	C ₄₈ H ₈₂ O ₁₉	10.14	1.457	[M + COOH] ⁻	Ginsenoside Re3	PG	961.5371 799.4874 637.4311 475.3807 391.2851
47	977.5339	C ₄₇ H ₈₀ O ₁₈	10.18	1.26	[M + COOH] ⁻	Notoginsenoside R1/Fp1/Ginsenoside Re4	PG	931.5272 799.4841 637.4321 475.3808 391.2865
48 [#]	991.5497	C ₄₈ H ₈₂ O ₁₈	10.47	1.394	[M + COOH] ⁻	Ginsenoside Re	PG	945.5430 799.4824 783.4907 637.4328 475.3794
49	651.2297	C ₃₁ H ₄₀ O ₁₅	10.67	0.394	[M-H] ⁻	Martynoside/Isomartynoside	RG	475.1817 329.1227 175.0391
50 [#]	845.4914	C ₄₂ H ₇₂ O ₁₄	10.67	1.172	[M + COOH] ⁻	Ginsenoside Rg1	PG	799.4850 637.4317 475.3789
51	312.1242	C ₁₈ H ₁₉ NO ₄	10.96	0.22	[M-H] ⁻	N-trans-feruloyltyramine	OJ	190.0502 176.0345 148.0519
52	1031.5449	C ₅₁ H ₈₄ O ₂₁	11.18	1.617	[M-H] ⁻	Malonylfloralginsenoside Re1	PG	945.5433 799.4834 783.4877 637.4315 475.3795 391.2852
53	651.2301	C ₃₁ H ₄₀ O ₁₅	11.21	1.008	[M-H] ⁻	Martynoside/Isomartynoside	RG	505.1704 475.1822 339.8273 175.0392
54	897.3893	C ₄₄ H ₅₂ N ₁₀ O ₁₁	11.27	0.358	[M + H] ⁺	Lyciumin B	LC	879.3740 689.3036 468.1874 442.2082
55	344.1492	C ₁₉ H ₂₁ NO ₅	11.31	-0.143	[M + H] ⁺	N-feruloyl-3-methoxytyramine	LC	177.0546 145.0284 117.0337
56	887.5012	C ₄₄ H ₇₄ O ₁₅	11.36	0.255	[M + COOH] ⁻	Acetyl-Ginsenoside Rg1	PG	841.5014 781.4736 637.4306 475.3818 161.0446
57	1079.5288	C ₅₀ H ₈₂ O ₂₂	11.5	0.763	[M + COOH] ⁻	Parisaponin I	OJ	1033.5266 901.4800 887.4641 755.4255 688.1104
58	183.1019	C ₁₀ H ₁₆ O ₃	11.64	-4.192	[M-H] ⁻	1-hydroxy-2,6,6-trimethylcyclohex-2-enecarboxylic acid	RG	137.096
59	815.4808	C ₄₁ H ₇₀ O ₁₃	11.79	1.172	[M + COOH] ⁻	Pseudoginsenoside Rt3	PG	769.4763 475.3808
60	815.4808	C ₄₁ H ₇₀ O ₁₃	12.01	1.172	[M + COOH] ⁻	Notoginsenoside A3	PG	637.4323 475.3784 391.2861
61	964.4206	C ₄₉ H ₅₇ N ₉ O ₁₂	12.21	0.68	[M + H] ⁺	Lyciumin C	LC	558.2376 477.3723 389.1823
62	535.2554	C ₂₈ H ₄₀ O ₁₀	12.33	0.989	[M-H] ⁻	Neorehmannioside C	RG	163.0390 145.0283 117.0332
63	887.502	C ₄₄ H ₇₄ O ₁₅	13.17	1.156	[M + COOH] ⁻	Notoginsenoside Rt	PG	841.4960 799.4748 781.4729 637.4343 619.4168 475.3756
64 [#]	845.4911	C ₄₂ H ₇₂ O ₁₄	14	0.817	[M + COOH] ⁻	Ginsenoside Rf	PG	799.4850 637.4315 475.3798 391.2856
65	799.4843	C ₄₂ H ₇₂ O ₁₄	14.04	-0.788	[M-H] ⁻	Ginsenoside La	PG	637.4324 475.3801 457.3684 391.2860
66	1285.6432	C ₅₉ H ₁₀₀ O ₂₇	14.29	-0.2	[M + COOH] ⁻	Notoginsenoside R4	PG	1239.6381 1107.5948 945.5484 783.4958 621.4349 459.3830
67	1325.6381	C ₆₂ H ₁₀₂ O ₃₀	14.77	-0.162	[M-H] ⁻	Malonylnotoginsenoside R4	PG	1239.6378 1221.6266 1077.5881 763.5875 459.3802
68	887.5019	C ₄₄ H ₇₄ O ₁₅	14.91	1.044	[M + COOH] ⁻	Vinaginsenoside R1	PG	841.4951 799.4790 637.4281 475.3795 391.2851
69	815.4808	C ₄₁ H ₇₀ O ₁₃	14.98	1.172	[M + COOH] ⁻	Notoginsenoside R2	PG	769.4744 637.4310 475.3793 391.2857
70	815.481	C ₄₁ H ₇₀ O ₁₃	15.2	1.417	[M + COOH] ⁻	Ginsenoside F5	PG	769.4747 637.4305 475.3791 391.2854
71	267.1603	C ₁₅ H ₂₄ O ₄	15.2	0.44	[M-H] ⁻	Aeginetic acid	RG	205.1593 153.0911
72	815.4808	C ₄₁ H ₇₀ O ₁₃	15.38	1.172	[M + COOH] ⁻	Ginsenoside F3	PG	769.4753 637.4321 475.3788 391.2855
73 [#]	829.4965	C ₄₂ H ₇₂ O ₁₃	16.21	1.212	[M + COOH] ⁻	20(S)-Ginsenoside Rg2	PG	783.4904 637.4402 475.3792 457.3681 391.2857
74	1255.6327	C ₅₈ H ₉₈ O ₂₆	16.28	-0.108	[M + COOH] ⁻	Ginsenoside Ra2	PG	1209.6273 1077.5823 915.5300 783.4872 621.4367 459.3846 375.2906
75	683.438	C ₃₆ H ₆₂ O ₉	16.33	0.606	[M + COOH] ⁻	20(S)-Ginsenoside Rh1	PG	637.4297 475.3807 161.0449
76	1285.6429	C ₅₉ H ₁₀₀ O ₂₇	16.43	-0.389	[M + COOH] ⁻	Ginsenoside Ra3	PG	1239.6377 1107.5944 945.5367 783.4913 621.4368 459.3864
77 [#]	1153.6017	C ₅₄ H ₉₂ O ₂₃	16.5	0.484	[M + COOH] ⁻	Ginsenoside Rb1	PG	1107.5952 945.5409 783.4985 621.4354 459.3831
78	683.4879	C ₃₆ H ₆₂ O ₉	16.54	0.46	[M + COOH] ⁻	20(R)-Ginsenoside Rh1	PG	637.4308 475.3776 243.5933 161.0444
79	989.5336	C ₄₈ H ₈₀ O ₁₈	16.58	0.942	[M + COOH] ⁻	Quinquefoloside Le/Ld	PG	943.5275 781.4734 618.6047 355.3042
80	829.4962	C ₄₂ H ₇₂ O ₁₃	16.61	0.705	[M + COOH] ⁻	20(R)-Ginsenoside Rg2	PG	783.4894 637.4335 475.3798 391.2856
81	1325.6384	C ₆₂ H ₁₀₂ O ₃₀	17.15	0.065	[M-H] ⁻	Malonylginsenoside Ra3	PG	1239.6387 1221.6277 1107.5898 1077.5800 945.5466 789.2095 621.4324 375.2910
82	1153.6022	C ₅₄ H ₉₂ O ₂₃	17.16	0.918	[M + COOH] ⁻	Ginsenoside Rb4/Re8	PG	1107.5955 945.5380 459.3853
83	683.4376	C ₃₆ H ₆₂ O ₉	17.4	0.021	[M + COOH] ⁻	Ginsenoside F1	PG	637.4374 475.3787 391.2866 359.4645
84	1193.5973	C ₅₇ H ₉₄ O ₂₆	17.55	1.042	[M-H] ⁻	Malonylginsenoside Rb1	PG	1107.5951 1089.5854 945.5410 783.4872 621.4354 537.3389 459.3838

(continued on next page)

Table 1 (continued)

NO.	Observed (m/z)	Formula	t _R (min)	Mass error (ppm)	Adducts	Identification	Source	ESI-MS ²
85	955.491	C ₄₈ H ₇₆ O ₁₉	17.7	0.206	[M-H] ⁻	Ginsenoside Ro	PG	793.4349 613.3726 569.3881 523.3810 455.3516
86	1209.6276	C ₅₈ H ₉₈ O ₂₆	17.98	0.202	[M-H] ⁻	Ginsenoside Ra1	PG	1077.5856 945.5435 783.4880 621.4350 459.3854
87 [#]	1123.5918	C ₅₃ H ₉₀ O ₂₂	18.09	1.089	[M + COOH] ⁻	Ginsenoside Rc	PG	1077.5848 945.5372 783.4900 621.4371 459.3848 375.2893
88	1209.6277	C ₅₈ H ₉₈ O ₂₆	18.12	0.284	[M-H] ⁻	Notoginsenoside Fp2/Fc/FZ	PG	1077.5834 945.5252 915.5354 783.4905 621.4402 459.3853
89	1193.5962	C ₅₇ H ₉₄ O ₂₆	18.2	0.121	[M-H] ⁻	Malonylfloralginsenoside Rb1/Rb2	PG	1107.5944 1089.5839 987.5690 945.5416 783.4865 459.3839
90	1163.5859	C ₅₆ H ₉₂ O ₂₅	18.82	0.351	[M-H] ⁻	Malonylginsenoside Rc	PG	1077.5844 945.5419 915.5381 783.4935 621.4328 459.3820
91	1255.6333	C ₅₈ H ₉₈ O ₂₆	18.86	0.37	[M + COOH] ⁻	Notoginsenoside Fp2/Fc/FZ	PG	1209.6272 1077.5790 908.5573 621.4341 459.3869
92 [#]	1123.5919	C ₅₃ H ₉₀ O ₂₂	19.55	1.178	[M + COOH] ⁻	Ginsenoside Rb2	PG	1077.5853 945.5428 783.4907 621.4377 459.3840
93	1193.5956	C ₅₇ H ₉₄ O ₂₆	19.62	-0.382	[M-H] ⁻	Malonylfloralginsenoside Rb1/Rb2	PG	1107.5959 1048.1158 945.5400 783.4918 459.3834
94	925.4812	C ₄₇ H ₇₄ O ₁₈	19.91	1.039	[M-H] ⁻	Pseudoginsenoside Rt1	PG	763.4271 613.3738 455.3530
95	1123.5916	C ₅₃ H ₉₀ O ₂₂	20.09	0.911	[M + COOH] ⁻	Ginsenoside Rb3	PG	1077.5848 945.5406 915.5377 783.4989 621.4362 459.3835
96	1163.5867	C ₅₆ H ₉₂ O ₂₅	20.24	1.039	[M-H] ⁻	Malonylginsenoside Rb2	PG	1077.5848 945.5446 915.5311 783.4886 621.4409 459.3847
97	1255.6327	C ₅₈ H ₉₈ O ₂₆	20.35	-1.108	[M + COOH] ⁻	Notoginsenoside Fp2/Fc/FZ	PG	1209.6271 1077.5830 945.5359 783.4948 621.4354 459.3856
98	989.5336	C ₄₈ H ₈₀ O ₁₈	20.57	0.942	[M + COOH] ⁻	Quinquefoloside Le/Ld	PG	943.5278 781.4705 457.3697 373.2743
99	1327.6542	C ₆₁ H ₁₀₂ O ₂₈	20.71	0.177	[M + COOH] ⁻	Vesanchinoside J	PG	1281.6461 1239.6394 1221.6276 1107.5925 945.5411 745.9520 564.1544
100	1195.6125	C ₅₆ H ₉₄ O ₂₄	20.97	0.664	[M + COOH] ⁻	Quinquenoside R2/Yesanchinoside F	PG	1149.6060 1107.5952 1089.5842 945.5493 783.4852 621.4340 459.3822
101	821.3975	C ₄₂ H ₆₂ O ₁₆	21.08	1.207	[M-H] ⁻	Glycyrrhizic acid	LC	561.8256 351.0570 193.0345
102	1163.5867	C ₅₆ H ₉₂ O ₂₅	21.18	1.039	[M-H] ⁻	Malonylginsenoside Rb3	PG	1077.5851 945.5137 915.5367 783.4911 621.4338 459.3853
103 [#]	991.5497	C ₄₈ H ₈₂ O ₁₈	21.22	1.394	[M + COOH] ⁻	Ginsenoside Rd	PG	945.5433 783.4912 621.4348 459.3830 375.2910
104	1297.6437	C ₆₀ H ₁₀₀ O ₂₇	21.22	0.231	[M + COOH] ⁻	Ginsenoside Ra5	PG	1251.6377 1191.6173 1077.5901 945.5460 915.5298 783.4849 621.4379 459.3847
105	793.4388	C ₄₂ H ₆₅ O ₁₄	21.22	1.034	[M-H] ⁻	Chikusetsusaponin VIa	PG	631.3864 569.3839 455.3526
106	1165.6019	C ₅₅ H ₉₂ O ₂₃	21.39	0.651	[M + COOH] ⁻	Acetylginenoside Rc	PG	1119.5957 1077.5850 945.5319 783.4935 621.4355 459.3840
107	1031.5448	C ₅₁ H ₈₄ O ₂₁	21.51	1.52	[M-H] ⁻	Malonylfloralginsenoside Rd5	PG	945.5433 783.4907 621.4376 459.3843
108	991.5499	C ₄₈ H ₈₂ O ₁₈	21.61	1.596	[M + COOH] ⁻	Notoginsenoside K	PG	945.5428 783.4898 621.4308 459.3827
109	1165.6019	C ₅₅ H ₉₂ O ₂₃	21.69	0.651	[M + COOH] ⁻	Acetylginenoside Rb2	PG	1119.5962 1077.5852 945.5417 783.4894 621.4349 459.3828
110	1195.6127	C ₅₆ H ₉₄ O ₂₄	21.72	0.831	[M + COOH] ⁻	Quinquenoside R4/Yesanchinoside F	PG	1149.6067 1107.5955 945.5417 783.4963 621.4375 459.3866
111	1165.6021	C ₅₅ H ₉₂ O ₂₃	21.9	0.823	[M + COOH] ⁻	Acetylginenoside Rb3	PG	1119.5956 1077.5846 945.5437 783.4910 621.4347 459.3874
112	1221.6277	C ₅₈ H ₉₆ O ₂₄	21.98	0.282	[M + COOH] ⁻	Ginsenoside Ra6	PG	1175.6254 1107.5951 945.5496 740.3268 459.3838
113	1165.6023	C ₅₅ H ₉₂ O ₂₃	22.09	0.994	[M + COOH] ⁻	Ginsenoside Rs1/Rs2	PG	1119.5957 1077.5852 1059.5737 945.5395 915.5432 783.4889 621.4313 459.3859
114	1165.6021	C ₅₅ H ₉₂ O ₂₃	22.42	0.832	[M + COOH] ⁻	Ginsenoside Rs1/Rs2	PG	1119.5957 1077.5847 1059.5741 945.5426 915.5307 783.4898 765.4787 621.4352 459.3841
115	797.47	C ₄₁ H ₆₈ O ₁₂	22.89	0.903	[M + COOH] ⁻	Notoginsenoside LY	PG	751.4631 619.4218
116	811.486	C ₄₂ H ₇₀ O ₁₂	23.07	1.319	[M + COOH] ⁻	Ginsenoside Rg4/Rg6/Rz1	PG	765.4791 619.4196 537.1358
117	797.4699	C ₄₁ H ₆₈ O ₁₂	23.18	0.778	[M + COOH] ⁻	Notoginsenoside T5	PG	751.4630 619.4222 457.3689
118	811.486	C ₄₂ H ₇₀ O ₁₂	23.29	1.319	[M + COOH] ⁻	Ginsenoside Rg4/Rg6/Rz1	PG	765.4789 619.4229 471.2258 313.2404
119	811.4861	C ₄₂ H ₇₀ O ₁₂	23.5	1.442	[M + COOH] ⁻	Ginsenoside Rg4/Rg6/Rz1	PG	765.4792 619.4230
120	665.4278	C ₃₆ H ₆₀ O ₈	23.62	1.17	[M + COOH] ⁻	Ginsenoside Rk3	PG	619.4216 161.0446
121	665.4275	C ₃₆ H ₆₀ O ₈	24.04	0.719	[M + COOH] ⁻	Ginsenoside Rh4	PG	619.4204 161.0446
122	793.4384	C ₄₂ H ₆₅ O ₁₄	24.05	0.53	[M-H] ⁻	Zingibroside R1	PG	613.3741 569.3845 455.3533
123	829.4968	C ₄₂ H ₇₂ O ₁₃	24.69	1.574	[M + COOH] ⁻	Ginsenoside F2	PG	783.4902 621.4359 459.3830 375.2095

(continued on next page)

2.3.4. Method validation

According to the above optimal analysis conditions and extraction conditions, the selectivity, linearity, limit of detection (LOD), limit of quantification (LOQ), accuracy, precision, stability and recovery of 13 compounds were determined.

The selectivity of the method was evaluated by comparing the chromatograms of each analyte in the blank solutions, standard solutions and sample solutions. Linearity was evaluated by plotting the calibration curves, and the curves were plotted with the concentration X (ng/mL) of the standard as the abscissa and the corresponding peak area Y of each standard as the ordinate. For each target compound, LOD and LOQ were determined at single to noise ratios (S/N) of 3 and 10 by continuous dilution of standard solutions. The QC samples of low, medium and high concentrations were analyzed for three consecutive days, accuracy was obtained from the relative error expressed as percentage (RE%), and precision was calculated using the relative standard deviation (RSD%). The stability of QC samples of low, medium and high concentrations was studied after being placed at room temperature for 0, 2, 4, 8, 12, 24 h. The test solutions and the standard solutions were added at a ratio of 1:1 to calculate the recovery.

2.4. Pharmacokinetics analysis by UPLC/QQQ-MS/MS

2.4.1. KSLP preparation

KSLP powder was repeatedly extracted twice under the optimal extraction process conditions, and the extract was concentrated and freeze-dried to obtain crude components of KSLP. The equal crude components of KSLP were further eluted with water and different proportions of ethanol/ethanol/methanol by resin D101/ resin AB-8/ODS, the water-eluting fractions were discarded, other fractions were combined respectively, concentrated and freeze-dried to obtain refined components of KSLP.

0.1 g powders of KSLP, crude components of KSLP and refined components of KSLP were added to 10 mL of 50 % ethanol, and ultrasonically extracted for 90 min. Extracts of KSLP and its crude components were centrifuged and diluted to obtain 100 μ g/mL sample solutions. Due to the large difference of the compound content in the refined components, 1000 μ g/mL and 10 μ g/mL of sample solutions were obtained. The UPLC/QQQ-MS/MS quantitative analysis of sample solutions was the same as in part 2.3. The refined components of KSLP obtained by the optimal refining process were used for pharmacokinetic studies.

2.4.2. Animals and experimental procedure

A total of 6 male Sprague-Dawley (SD) rats (200–220 g) were purchased from Beijing Huafukang Bio-Technology Co., Ltd. All of the rats were kept in pathogen-free animal laboratory in a 12 h light/dark cycle, with a feeding temperature of 26 °C and relative humidity of 50 %. Animal experiments were consistent with the Animal Ethics Committee of Tianjin University of Traditional Chinese Medicine, and the animal ethics approval number was TCM-LAEC2022124.

Rats were acclimatized for 7 days to laboratory environments before the experiment was directed. Diet was prohibited for 12 h before the experiment, but the water was freely available. Appropriate refined components of KSLP were dispersed in distilled water as a suspension with a concentration of 0.4 g/mL, and the oral administration dose was set at 10 mL/kg of rat weight. The blood samples were collected through canthus into heparinized tubes before administration and at 0.083, 0.25, 0.5, 1, 2, 4, 6, 8, 10, 12, 24, 36, 48, 60 and 72 h after oral administration of refined components of KSLP (4 g/kg). Plasma samples were obtained by centrifuging the blood samples immediately at 1600g for 10 min and stored at -80°C until analysis.

2.4.3. Plasma sample preparation

100 μ L of each sample was processed by adding 300 μ L of acetonitrile, 100 μ L of internal standard solution (IS) and 100 μ L of 50 % acetonitrile. The supernatant was collected after vortex mixing 1 min and the following centrifugation (15000 g, 10 min, 4°C). Then, the supernatant was transferred and dried under a gentle stream of nitrogen gas at room temperature. The resulting residues were redissolved in 100 μ L of 50 % acetonitrile, and 5 μ L was injected into the UPLC/QQQ-MS/MS system.

2.4.4. Standard solutions preparation

The stock solutions of ginsenoside Rb1, ginsenoside Rb2, ginsenoside Rc, ginsenoside Rd, ginsenoside Re and ginsenoside Rg1 were prepared in 50 % acetonitrile, which were then diluted with 50 % acetonitrile to obtain mixed standard solutions of the 6 compounds at different concentrations. Digoxin was weighed accurately, dissolved in 50 % acetonitrile and diluted gradually to 100 ng/mL as the internal standard solution. 100 μ L of blank plasma was processed by adding 300 μ L of acetonitrile, 100 μ L of IS and 100 μ L of mixed standard solutions. Similar to post-administration plasma processing, a range of working standard solutions were obtained for analysis. The QC samples were prepared in a similar manner at three different concentrations (10, 100 and 800 ng/mL) of ginsenoside Rb1, ginsenoside Rb2, ginsenoside Rc, ginsenoside

Table 1 (continued)

NO.	Observed (m/z)	Formula	t_R (min)	Mass error (ppm)	Adducts	Identification	Source	ESI-MS ²
124 [#]	829.4969	C ₄₂ H ₇₂ O ₁₃	24.87	1.694	[M + COOH] ⁻	20(S)-Ginsenoside Rg3	PG	783.4899 621.4357 459.3828 375.2894
125	829.4965	C ₄₂ H ₇₂ O ₁₃	25.05	1.005	[M + COOH] ⁻	20(R)-Ginsenoside Rg3	PG	783.4897 621.4373 459.3828 375.2913
126	341.1028	C ₁₉ H ₁₈ O ₆	26.47	-0.262	[M-H] ⁻	Methylpogonane A	OJ	206.0576 178.0626 163.0388
127	293.2119	C ₁₈ H ₃₀ O ₃	26.51	-1.085	[M-H] ⁻	HOTrE	LC	275.2014 171.1015
128	811.4854	C ₄₂ H ₇₀ O ₁₂	27.15	0.579	[M + COOH] ⁻	Ginsenoside Rk1	PG	765.4797 603.4258 319.0105
129	811.486	C ₄₂ H ₇₀ O ₁₂	27.33	1.319	[M + COOH] ⁻	Ginsenoside Rg5	PG	765.4791 603.4272 453.0886
130	667.4434	C ₃₆ H ₆₂ O ₈	27.73	1.092	[M + COOH] ⁻	20(S)-Ginsenoside Rh2	PG	621.4393 459.3858
131	667.4432	C ₃₆ H ₆₂ O ₈	28.1	0.792	[M + COOH] ⁻	20(R)-Ginsenoside Rh2	PG	504.3103
132	485.3641	C ₃₁ H ₅₀ O ₄	28.72	0.692	[M-H] ⁻	Tumulosic acid	PC	423.3291 331.3891 180.2139
133	853.4957	C ₄₄ H ₇₂ O ₁₃	28.75	0.241	[M + COOH] ⁻	Ginsenoside Rs5	PG	807.4891 765.4779 603.4246
134	485.3274	C ₃₀ H ₄₆ O ₅	28.75	0.314	[M-H] ⁻	Poricoic acid G	PC	441.3332 423.3265 351.2659 254.0528
135	853.4957	C ₄₄ H ₇₂ O ₁₃	29.17	0.241	[M + COOH] ⁻	Ginsenoside Rs4	PG	807.4896 765.4786 603.4275
136	481.3328	C ₃₁ H ₄₆ O ₄	30.63	0.97	[M-H] ⁻	Polyphenolic acid C	PC	403.9587 253.2169 152.9946
137	455.3533	C ₃₀ H ₄₈ O ₃	32.24	0.508	[M-H] ⁻	Oleanic acid	PC	261.8651 246.0458 201.1751 146.1933 129.5593
138	527.3739	C ₃₃ H ₅₂ O ₅	32.5	-0.565	[M-H] ⁻	Pachymic acid	PC	444.3078 371.6741 228.7377

RG, Rehmanniae radix; PG, Ginseng radix et rhizoma rubra; AC, Asparagi radix; OJ, Ophiopogonis radix; LC, Lych cortex; PC, Poria.

Rd, ginsenoside Re and ginsenoside Rg1.

2.4.5. Chromatographic and mass spectrographic conditions

The conditions of chromatographic and mass spectrographic were the same as in part 2.3.3, the remaining specific parameters of 7 compounds were shown in Table S2.

2.4.6. Method validation

According to the above optimal analysis conditions, the selectivity, linearity, lower limit of quantification (LLOQ), accuracy, precision, extraction recovery, matrix effect and stability were validated.

The chromatograms of blank plasma, standard spiked plasma and sample plasma were used to assess the selectivity of the method. Linear regression equations were obtained by least squares linear regression on the ratio Y of the compound peak area to IS peak area as the ordinate, the compound concentration X (ng/mL) in plasma as the abscissa, and LLOQ was the lowest concentration of linearity. Intra-day and inter-day accuracy and precision were estimated by analyzing a calibration curve and QC samples of low, medium and high concentrations on three days. The extraction recovery was evaluated through the ratio of the mean concentration between regularly prepared QC samples of low, medium and high concentrations and spike-after-extraction plasma samples. Similarly, the matrix effect was assessed through the ratio of concentration between post-extraction samples spiked with analytes and ultrapure water spiked with analytes at the same concentration. The stability of each analyte at different conditions (autosampler for 24 h and three freeze-thaw cycles from -80°C to room temperature) was assessed by analyzing at QC levels.

2.5. Data processing and analysis

Data was expressed as mean \pm standard deviation, pharmacokinetic parameters were statistically calculated using the pharmacokinetic software (DAS version 2.0).

3. Results

3.1. Qualitative analysis of chemical constituents in KSLP by UPLC/Q-Orbitrap-MS/MS

The base peak ion chromatograms (BPI) of KSLP samples and standards were shown in Fig. 1, according to the experimental conditions described in part 2.2, combined with standard comparison and literature comparison, a total of 138 compounds were inferred (Table 1), including 79 ginsenosides, 14 phenylethanol glycosides, 10 iridoids, 10 alkaloids, 5 ionones and 20 others, 13 compounds were further identified by comparison with standards.

3.1.1. Identification of ginsenosides in KSLP

Ginsenosides were the main active components of Ginseng radix et rhizoma rubra, which were mainly divided into protopanaxadiol (PPD), protopanaxatriol (PPT) and others according to their structural differences (Yang et al., 2014). The sapogenin fragments at m/z 459.38 (PPD) and m/z 475.38 (PPT) could be used as the diagnostic product ions to rapidly characterize ginsenosides of these two subtypes (Qiu et al., 2015, Li et al., 2021a, 2021b). Taking ginsenoside Rd as an example to illustrate the cleavage pathway of PPD, the excimer ion peak of m/z 991.5497 $[\text{M} + \text{COOH}]^{-}$ was easily generated in the negative ion mode. Typical neutral losses (NL) of HCOOH (46 Da) and Glu (162 Da) produced fragments of m/z 945.5433 $[\text{M}-\text{H}]^{-}$, m/z 783.4912 $[\text{M}-\text{H}-\text{Glu}]^{-}$, m/z 621.4348 $[\text{M}-\text{H}-2\text{Glu}]^{-}$ and m/z 459.3830 $[\text{M}-\text{H}-3\text{Glu}]^{-}$. Taking the PPT type ginsenoside Re as an example, the quasimolecular ion peak of m/z 991.5497 $[\text{M} + \text{COOH}]^{-}$ was easily generated in the negative ion mode. Due to NL of HCOOH (46 Da), Glu (162 Da) and Rha (146 Da), fragments of m/z 945.5433 $[\text{M}-\text{H}]^{-}$, m/z 799.4824 $[\text{M}-\text{H}-\text{Rha}]^{-}$, m/z 783.4907 $[\text{M}-\text{H}-\text{Glu}]^{-}$ and m/z

Table 2

The linear regression equations, R^2 , linear ranges, LOD and LOQ of multi-components quantitative analysis method of KSLP.

Compound	Regression equations	R^2	Linear ranges (ng/mL)	LOD (ng/mL)	LOQ (ng/mL)
Acteoside	$y = 734.92x + 57.755$	0.9999	0.4 ~ 100	0.02	0.08
Isoacteoside	$y = 937.50x - 6.9475$	0.9999	0.4 ~ 100	0.02	0.08
Echinacoside	$y = 139.96x + 62.214$	0.9994	0.4 ~ 100	0.06	0.20
Jionoside A1	$y = 142.74x + 55.328$	0.9996	0.4 ~ 100	0.06	0.20
Ginsenoside Rb1	$y = 29.248x + 99.106$	0.9994	4 ~ 1000	0.06	0.20
Ginsenoside Rb2	$y = 39.981x + 7.6629$	0.9999	4 ~ 1000	0.06	0.20
Ginsenoside Rc	$y = 19.805x - 1.3026$	0.9998	4 ~ 1000	0.60	2.00
Ginsenoside Rd	$y = 26.910x - 119.27$	0.9997	4 ~ 1000	0.24	0.80
Ginsenoside Re	$y = 24.074x + 22.586$	0.9995	4 ~ 1000	0.60	2.00
Ginsenoside Rf	$y = 112.71x + 165.65$	>0.9999	4 ~ 1000	0.60	2.00
Ginsenoside Rg1	$y = 14.476x - 49.816$	>0.9999	4 ~ 1000	0.24	0.80
Ginsenoside Rg2	$y = 111.03x + 40.289$	0.9997	0.4 ~ 100	0.24	0.80
Ginsenoside Rg3	$y = 87.747x + 17.923$	0.9998	0.4 ~ 100	0.24	0.80

637.4328 $[\text{M}-\text{H}-\text{Glu}-\text{Rha}]^{-}$ and m/z 475.3794 $[\text{M}-\text{H}-2\text{Glu}-\text{Rha}]^{-}$ were detected.

3.1.2. Identification of phenylethanol glycosides in KSLP

Phenylethanol glycosides, one of the bioactive components from Rehmanniae radix, possessed various pharmacological activities for human health. The fragmentation pathway was mainly related to breaking of ester bond or C-O bond, resulting in a series of degradation products. Taking acteoside for a witness, the excimer ion peak of m/z 623.1985 $[\text{M}-\text{H}]^{-}$ was easily generated in the negative ion mode. Acteoside produced ions at m/z 461.1662 $[\text{M}-\text{H}-\text{caffeoyl}]^{-}$ and m/z 315.1082 $[\text{M}-\text{H}-\text{caffeoyl}-\text{Rha}]^{-}$ through the loss of caffeoyl and Rha moiety. The caffeoyl moiety was produced ion at m/z 161.0234 $[\text{Caffeoyl}-\text{H}-\text{H}_2\text{O}]^{-}$ by the loss of H_2O (Qi et al., 2013).

3.2. Quantitative analysis of multi-components in KSLP by UPLC/QQQ-MS/MS

Combined with the qualitative analysis of chemical constituents in KSLP, and comprehensively considering its active components and characteristic components, the UPLC/QQQ-MS/MS method for the simultaneous determination of acteoside, isoacteoside, echinacoside, jionoside A1, ginsenoside Rb1, ginsenoside Rb2, ginsenoside Rc, ginsenoside Rd, ginsenoside Re, ginsenoside Rf, ginsenoside Rg1, ginsenoside Rg2 and ginsenoside Rg3 in KSLP was established, in order to provide reference for the quality control of KSLP.

3.2.1. Validation of analytical method

The chromatograms of blank solutions, standard solutions and sample solutions were displayed in Fig. S1. It could be seen from the chromatograms that 13 compounds had good peak shapes and no endogenous interference, indicating that the method was suitable. The linear regression equations, correlation coefficients (R^2), linear ranges, LOD and LOQ of the 13 target compounds were listed in Table 2. The R^2 of analytes were greater than 0.999, indicating good linearity, the LOD and LOQ of the 13 target compounds were the range of 0.02 ~ 0.6 ng/mL and 0.08 ~ 2.0 ng/mL, indicating that the method had high sensitivity. The test results of accuracy, precision and stability were summarized in Table 3. Intra-day and inter-day accuracy (RE%) ranged from -4.78% to 4.74% , intra-day and inter-day precision (RSD%) were

Table 3The accuracy, precision and stability of multi-components quantitative analysis method of KSLP (Mean \pm SD, $n = 3$).

Compound	QC (ng/mL)	Intra-day			Inter-day			Stability	
		$\bar{X} \pm S$ (ng/mL)	RE (%)	RSD (%)	$\bar{X} \pm S$ (ng/mL)	RE (%)	RSD (%)	$\bar{X} \pm S$ (ng/mL)	RSD (%)
Acteoside	1	0.96 \pm 0.06	-3.70	7.16	0.96 \pm 0.04	-3.59	4.76	0.97 \pm 0.02	2.40
	10	9.52 \pm 0.11	-4.78	1.47	9.54 \pm 0.09	-4.65	1.00	10.08 \pm 0.39	4.75
	80	82.77 \pm 1.06	3.46	1.57	83.3 \pm 1.53	4.13	2.01	81.97 \pm 2.34	3.50
Isoacteoside	1	0.96 \pm 0.04	-4.42	4.94	0.96 \pm 0.03	-4.47	3.32	0.99 \pm 0.03	3.20
	10	9.56 \pm 0.05	-4.37	0.65	9.61 \pm 0.07	-3.93	0.84	9.52 \pm 0.05	0.63
	80	79.83 \pm 0.60	-0.22	0.93	78.92 \pm 1.12	-1.35	1.56	79.28 \pm 0.75	1.15
Echinacoside	1	0.97 \pm 0.02	-3.23	2.50	0.97 \pm 0.02	-3.33	2.02	0.95 \pm 0.01	1.12
	10	9.55 \pm 0.05	-4.47	0.64	9.70 \pm 0.32	-3.02	3.57	10.15 \pm 0.09	1.04
	80	80.12 \pm 1.42	0.14	2.17	81.53 \pm 2.91	1.91	3.91	80.76 \pm 1.55	2.36
Jionoside A1	1	0.96 \pm 0.02	-3.66	2.71	0.97 \pm 0.03	-3.18	2.96	0.99 \pm 0.03	3.92
	10	10.47 \pm 0.36	4.74	4.21	10.31 \pm 0.38	3.06	4.00	10.28 \pm 0.20	2.42
	80	81.67 \pm 2.02	2.09	3.03	82.12 \pm 2.33	2.66	3.11	80.24 \pm 0.86	1.32
Ginsenoside Rb1	10	9.57 \pm 0.05	-4.28	0.68	9.58 \pm 0.10	-4.17	1.11	9.67 \pm 0.13	1.69
	100	97.13 \pm 3.50	-2.87	4.41	96.33 \pm 2.66	-3.67	3.03	102.65 \pm 1.42	1.70
	800	814.85 \pm 12.49	1.86	1.88	811.38 \pm 10.06	1.42	1.36	806.27 \pm 3.25	0.49
Ginsenoside Rb2	10	9.80 \pm 0.18	-2.01	2.27	9.84 \pm 0.15	-1.58	1.72	9.70 \pm 0.21	2.70
	100	96.76 \pm 3.10	-3.24	3.92	96.82 \pm 3.49	-3.18	3.95	104.31 \pm 0.59	0.69
	800	810.68 \pm 10.44	1.34	1.58	807.40 \pm 9.54	0.93	1.29	801.42 \pm 1.43	0.22
Ginsenoside Rc	10	9.85 \pm 0.22	-1.49	2.74	9.73 \pm 0.24	-2.73	2.71	9.63 \pm 0.26	3.35
	100	102.35 \pm 1.82	2.35	2.17	102.44 \pm 2.39	2.44	2.56	104.61 \pm 1.23	1.44
	800	814.97 \pm 4.95	1.87	0.74	809.70 \pm 6.46	1.21	0.87	808.47 \pm 3.03	0.46
Ginsenoside Rd	10	10.46 \pm 0.15	4.59	1.76	10.32 \pm 0.23	3.21	2.41	10.35 \pm 0.20	2.36
	100	96.07 \pm 1.25	-3.93	1.59	98.97 \pm 3.65	-1.03	4.04	100.17 \pm 2.74	3.35
	800	804.42 \pm 6.53	0.55	0.99	806.89 \pm 6.41	0.86	0.87	805.46 \pm 8.82	1.34
Ginsenoside Re	10	9.59 \pm 0.24	-4.14	3.12	9.65 \pm 0.22	-3.53	2.47	9.57 \pm 0.11	1.47
	100	96.19 \pm 4.05	-3.81	5.15	96.6 \pm 3.29	-3.40	3.73	103.49 \pm 1.00	1.18
	800	815.46 \pm 6.46	1.93	0.97	812.48 \pm 8.61	1.56	1.16	800.94 \pm 15.22	2.33
Ginsenoside Rf	10	9.61 \pm 0.29	-3.85	3.74	9.78 \pm 0.34	-2.18	3.75	9.75 \pm 0.16	2.07
	100	97.70 \pm 3.09	-2.30	3.87	96.55 \pm 3.56	-3.45	4.04	102.97 \pm 1.29	1.53
	800	823.39 \pm 2.63	2.92	0.39	816.5 \pm 7.94	2.06	1.07	804.55 \pm 2.31	0.35
Ginsenoside Rg1	10	10.27 \pm 0.15	2.70	1.77	10.18 \pm 0.17	1.76	1.82	10.37 \pm 0.38	4.50
	100	101.86 \pm 3.81	1.86	4.58	102.36 \pm 3.78	2.36	4.04	102.11 \pm 1.65	1.98
	800	803.33 \pm 0.70	0.42	0.11	805.61 \pm 4.82	0.70	0.65	812.23 \pm 5.56	0.84
Ginsenoside Rg2	1	0.98 \pm 0.02	-2.15	2.19	0.98 \pm 0.02	-2.25	2.24	0.94 \pm 0.03	3.85
	10	9.83 \pm 0.30	-1.70	3.72	9.72 \pm 0.35	-2.78	3.99	10.63 \pm 0.24	2.74
	80	83.29 \pm 2.07	4.11	3.05	82.56 \pm 2.00	3.20	2.65	80.69 \pm 0.54	0.81
Ginsenoside Rg3	1	0.97 \pm 0.04	-3.03	4.67	0.97 \pm 0.03	-3.22	3.32	0.96 \pm 0.03	3.22
	10	9.63 \pm 0.36	-3.71	4.54	9.92 \pm 0.39	-0.83	4.26	9.66 \pm 0.27	3.41
	80	82.56 \pm 2.50	3.21	3.70	81.52 \pm 2.14	1.90	2.88	81.41 \pm 1.92	2.89

within 7.16 % and stability (RSD%) was less than 4.75 %, indicating that the method was reproducible and accurate for the determination of analytes. The sample recovery rate was showed in Table S6, and it was between 90.27 ~ 108.22 % (RSD% less than 5.54 %), indicating that no significant loss of analyte occurred during the analysis.

3.2.2. Quantitative analysis of multi-components in KSLP

The effects of various extraction conditions on the yield of target compounds were represented in Table S3 and Table S4. The yield of crude components extracted by ultrasonic and reflux was not much different, and the total content of target compounds extracted by ultrasonic was slightly higher than that of reflux. With the increase of the ethanol content, the crude components yield showed a trend of first increasing and then decreasing. When the ethanol concentration reached 50 %, the total content of target compounds was higher than that of other concentration group. The yield of crude components increased slowly with increasing extraction time, eventually reached stability at 90 min, indicating that extraction time has a significant positive effect on yield when it is below 90 min. There was no significant change in the yield when extraction time was longer than 90 min. There was a gradual increase in the yield of crude components with increasing solid-to-liquid ratio. The yield increased slightly when the solid-to-liquid ratio was greater than 1:100 g/mL, however, a large solid-to-liquid ratio caused solvent wastage, 1:100 g/mL was selected as the optimal solid-to-liquid ratio. Therefore, 50 % ethanol, ultrasonic, 90 min and 1:100 g/mL were the optimal process for extracting crude components of KSLP based on the criterion of the maximum total content of the

13 target compounds, combined with economic considerations.

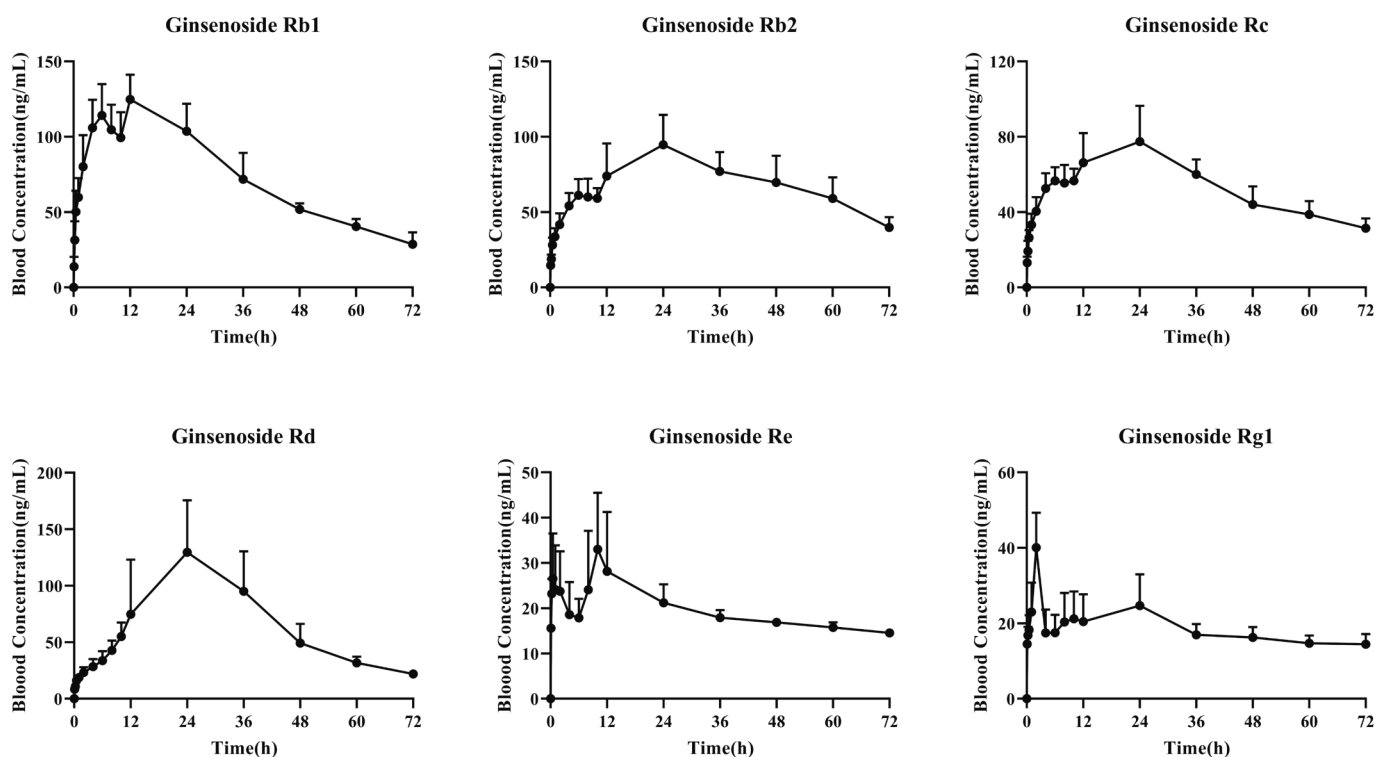
According to the established UPLC/QQQ-MS/MS quantitative analysis method and extraction process conditions, the content of 13 compounds in KSLP was determined. Among 13 analytes, the compounds with higher content (>1000 μ g/g) were ginsenoside Rb1, ginsenoside Rb2, ginsenoside Rc, ginsenoside Re and ginsenoside Rg1.

3.3. Pharmacokinetic analysis of multi-components in KSLP by UPLC/QQQ-MS/MS

The composition of TCM is complex, and the blood concentration in the body is low, which is not easy to detect. In the previous pilot experiment, the rats were gavaged with the largest dose of KSLP, and no compounds could be detected in the plasma. Therefore, KSLP was further refined to obtain the refined components in formal experiments. According to the UPLC/QQQ-MS/MS quantitative analysis method established in part 2.3, the effects of different fillers on the yield of refined components could be seen in Table S5. After being refined by resin D101, resin AB-8 and ODS, the total contents of the 13 target compounds were not much different in the refined components of KSLP, and the yields of refined components were 13.19 %, 10.10 % and 10.69 % respectively. Although resins D101 and AB-8 retained the ginsenosides well, most of the phenylethanol glycosides were lost due to its large polarity when eluted with water. ODS not only enriched the ginsenosides but also retained the phenylethanol glycosides with greater polarity. Therefore, ODS was selected to refine the crude components of KSLP. The total content of the 13 target compounds in the refined

Table 4The accuracy, precision, extraction recovery and matrix effect of multi-components pharmacokinetic analysis method of KSLP (Mean \pm SD, $n = 6$).

Compound	QC (ng/mL)	Intra-day			Inter-day			Extraction recovery and matrix effect	
		$\bar{X} \pm S$ (ng/mL)	RE (%)	RSD (%)	$\bar{X} \pm S$ (ng/mL)	RE (%)	RSD (%)	Extractionrecovery (%)	Matrixeffect (%)
Ginsenoside Rb1	10	11.19 \pm 0.48	11.89	4.32	11.09 \pm 0.40	10.92	3.56	98.68 \pm 3.47	100.46 \pm 5.83
	100	88.18 \pm 3.88	-11.82	4.40	89.03 \pm 3.67	-10.97	4.13	89.80 \pm 3.65	107.69 \pm 5.53
	800	766.31 \pm 26.96	-4.21	3.52	762.72 \pm 22.43	-4.66	2.94	89.68 \pm 3.09	103.81 \pm 5.55
Ginsenoside Rb2	10	10.44 \pm 0.59	4.45	5.66	10.20 \pm 0.54	1.97	5.32	95.60 \pm 3.30	105.43 \pm 1.89
	100	88.65 \pm 2.97	-11.35	3.35	87.78 \pm 3.11	-12.22	3.54	93.18 \pm 5.65	106.33 \pm 4.15
	800	766.08 \pm 28.71	-4.24	3.75	754.13 \pm 33.48	-5.73	4.44	89.93 \pm 3.93	99.72 \pm 3.97
Ginsenoside Rc	10	10.28 \pm 0.70	2.82	6.79	10.25 \pm 0.67	2.46	6.51	91.93 \pm 3.33	108.45 \pm 3.07
	100	91.32 \pm 0.86	-8.68	0.95	90.75 \pm 3.21	-9.25	3.54	97.20 \pm 5.10	103.49 \pm 8.33
	800	750.49 \pm 11.51	-6.38	1.99	749.00 \pm 14.89	-6.19	1.53	97.87 \pm 8.87	102.47 \pm 8.38
Ginsenoside Rd	10	10.21 \pm 0.31	2.08	3.03	10.58 \pm 0.16	5.79	1.53	92.09 \pm 4.69	107.14 \pm 6.85
	100	94.43 \pm 6.82	-5.57	7.22	94.26 \pm 5.57	-5.74	5.91	88.95 \pm 1.19	102.00 \pm 8.56
	800	749.05 \pm 10.39	-6.37	1.39	743.23 \pm 14.4	-7.10	1.94	91.94 \pm 6.50	109.78 \pm 5.44
Ginsenoside Re	10	10.60 \pm 0.42	6.02	3.95	10.48 \pm 0.51	4.81	4.89	99.07 \pm 5.81	89.97 \pm 1.18
	100	93.42 \pm 3.89	-6.58	4.16	96.24 \pm 2.44	-3.76	2.53	92.99 \pm 4.65	105.30 \pm 6.42
	800	778.67 \pm 21.17	-2.67	2.72	775.07 \pm 18.73	-3.12	2.42	96.45 \pm 11.02	95.03 \pm 5.54
Ginsenoside Rg1	10	10.25 \pm 0.61	2.47	5.95	10.73 \pm 0.19	7.26	1.78	102.09 \pm 3.93	88.02 \pm 3.57
	100	97.27 \pm 1.59	-2.73	1.63	98.67 \pm 3.49	-1.33	3.53	90.93 \pm 7.16	97.20 \pm 6.47
	800	778.68 \pm 14.15	-2.67	1.82	776.02 \pm 12.72	-3.00	1.64	90.34 \pm 4.77	95.87 \pm 1.17

**Fig. 2.** Plasma drug concentration and time curve of refined components of KSLP in rats after intragastric administration ($n = 6$).

components was 5 times higher than that of the crude components. The UPLC/QQQ-MS/MS was used to determine plasma after administration of refined components of KSLP, and a total of 6 compounds were detected, including ginsenoside Rb1, ginsenoside Rb2, ginsenoside Rc, ginsenoside Rd, ginsenoside Re and ginsenoside Rg1.

3.3.1. Validation of analytical method

The chromatograms of blank plasma solutions, standard plasma solutions and sample plasma solutions were plotted in Fig. S2, all the peaks of the analytes and IS were detected with excellent resolutions as well as shapes. The endogenous substances in the plasma did not interfere with the determination of each compound and IS, suggesting that the method developed in this study had good selectivity. The 6 target compounds had a good linear relationship within the corresponding concentration

range, and its LLOQ were 4.0 ng/mL. The statistical parameters of calibration were listed in Table S7. As exhibited in Table 4, intra-day and inter-day accuracy (RE%) was between $-12.22 \sim 11.89$ %, intra-day and inter-day precision (RSD%) were within 7.22 %, the extraction recovery and matrix effect of 6 compounds ranged from 88.02 % to 109.78 %, it showed that the instrument had good precision, the method had high repeatability. The stability results in Table S8 indicated that the 6 compounds in rat plasma were stable with RSD < 14.61 % for autosampler for 24 h, three freeze-thaw cycles.

3.3.2. Determination of the content of KSLP in plasma and pharmacokinetic parameter fitting

The plasma drug concentration and time curve of refined components of KSLP after intragastric administration for 72 h with a single

Table 5
Pharmacokinetic parameters of 6 analytes in rat plasma after intragastric administration of refined components of KSLP (Mean \pm SD, $n = 6$).

Compound	Ginsenoside Rb1	Ginsenoside Rb2	Ginsenoside Rc	Ginsenoside Rd	Ginsenoside Re	Ginsenoside Rg1
T _{1/2} (h)	31.672 \pm 17.692	47.631 \pm 10.293	37.464 \pm 5.061	22.997 \pm 10.598	148.83 \pm 64.087	67.019 \pm 38.726
T _{max} (h)	8.333 \pm 3.445	24 \pm 0	23 \pm 4.899	24 \pm 0	10.333 \pm 1.506	2 \pm 0
C _{max} (ng/mL)	131.342 \pm 13.055	94.667 \pm 19.846	79.147 \pm 15.999	129.622 \pm 46.008	41.023 \pm 12.471	40.079 \pm 9.236
AUC _{0-t} (ng/mL ^h)	5297.488 \pm 559.59	4935.666 \pm 876.227	3840.482 \pm 578.394	4700.077 \pm 1334.759	1402.143 \pm 178	1342.658 \pm 185.234
AUC _{0-∞} (ng/mL ^h)	6746.532 \pm 1047.45	8285.218 \pm 2023.524	5523.503 \pm 759.415	5445.909 \pm 1401.581	4500.245 \pm 1315.926	2539.865 \pm 574.874
V _z /F (L/kg)	26065.292 \pm 11112.873	33789.23 \pm 5310.405	40093.334 \pm 9621.342	25723.978 \pm 12221.349	184037.526 \pm 29842.507	144175.903 \pm 51476.184
CL _z /F (L/h/kg)	604.993 \pm 93.639	512.833 \pm 153.38	738.339 \pm 124.228	778.744 \pm 212.756	967.909 \pm 334.351	1643.17 \pm 367.034
MRT _{0-t} (h)	28.101 \pm 0.832	34.367 \pm 1.5	32.2 \pm 1.353	31.718 \pm 2.194	32.356 \pm 2.351	32.517 \pm 1.426
MRT _{0-∞} (h)	48.737 \pm 19.743	68.882 \pm 14.101	61.403 \pm 6.679	42.874 \pm 8.795	210.355 \pm 87.169	113.409 \pm 57.93

dose in rats was presented in Fig. 2, and the pharmacokinetic parameters were summarized in Table 5. The maximum plasma concentration (C_{max}) of ginsenoside Rb1, ginsenoside Rb2, ginsenoside Rc and ginsenoside Rd was about 100 ng/mL, while the C_{max} of ginsenoside Re and ginsenoside Rg1 was about 40 ng/mL. All compounds took long time to their elimination t_{1/2}, indicating that the elimination rate of the target compounds *in vivo* was slow. Also, the AUC_{0-∞} of ginsenoside Rb1, ginsenoside Rb2, ginsenoside Rc and ginsenoside Rd was greater than 4500 ng/ml·h, while the AUC_{0-∞} of ginsenoside Re and ginsenoside Rg1 was about 2500 ng/ml·h.

4. Discussion

With its rich resources and unique curative effects, TCM has attracted the attention of many countries around the world, and has been gradually accepted, researched, developed and utilized. The composition of TCM is complex and diverse, quality control is difficult, and traditional analysis methods can't longer meet the requirements of quality analysis and evaluation of TCM (Zhao et al., 2018). The chemical composition of KSLP was comprehensively characterized, and the results showed that it contained a large number of ginsenosides, phenylethanol glycosides and iridoids. Ginsenosides were the main chemical components of Ginseng radix et rhizoma rubra, and studies have shown that total ginsenosides of Ginseng radix et rhizoma rubra could produce an anti-aging effect by intervening in the lipid metabolism and correcting the amino acid metabolism disorders in aging rats (Sun et al., 2018). Phenylethanol glycosides and iridoids were the most abundant compounds in Rehmanniae radix. Studies have shown that echinacoside and catalpol could delay aging and prevent the development of age-related diseases (Zhang et al., 2008, Shen et al., 2017, Chen et al., 2018).

The complexity of TCM components determines that it is difficult to comprehensively characterize the quality of TCM in the determination of only a single compound (Luo et al., 2013, Xiong et al., 2020). The quantitative analysis of multi-components has become the development direction of quality evaluation of TCM (Wang et al., 2020, Li et al., 2021a, 2021b). The quality evaluation of TCM containing ginsenosides has always been difficult. Ginsenosides lack functional groups with strong ultraviolet absorption, resulting in inaccurate quantitative results of samples, so the detection has certain limitations. Therefore, the UPLC/QQQ-MS/MS technology equipped with electrospray ionization was used to establish a method for simultaneous detection of multi-components in KSLP. The process of extraction is the core link in the field of pharmacodynamic substances and quality control of TCM (Zhang et al., 2018). Optimizing the parameters of extraction of TCM is the premise of efficient and sufficient extraction of pharmacodynamic substances and ensuring clinical effectiveness (Tang et al., 2022, Chang et al., 2023). Under the optimal extraction conditions, 13 representative compounds of KSLP were simultaneously quantified by UPLC/QQQ-MS/MS.

By further refining the refined components in KSLP, it was used for the study of highly exposed *in vivo* components. The UPLC/Q-Orbitrap-MS/MS technology was used for the determination of plasma after administration, and the results showed that no compounds were detected. We speculate that ginsenosides have the characteristics of large molecular weight, poor membrane permeability, and unstable chemical structure that can be degraded by hydrolases in the gastrointestinal tract, these characteristics lead to its low bioavailability (Xiong et al., 2009, Li et al., 2011, Dai et al., 2016). MRM monitoring mode can detect and analyze specific compounds with strong specificity and high sensitivity (Ren et al., 2022). The MRM acquisition method of UPLC/QQQ-MS/MS technology was used to detect plasma after administration, and a total of 6 compounds were detected, including ginsenoside Rb1, ginsenoside Rb2, ginsenoside Rc, ginsenoside Rd, ginsenoside Re and ginsenoside Rg1. Studies have found that ginsenoside Rb1, ginsenoside Rb2, ginsenoside Rc, ginsenoside Rd, ginsenoside Re and ginsenoside Rg1 account for more than 90 % of the total

ginsenosides, and were called non-rare ginsenosides because of their higher content (Wang et al., 2006). The pharmacokinetic results showed that the absorption and elimination of ginsenosides in the refined components of KSLP were relatively slow. This may be related to the complexity of the components of TCM and their interaction transformation *in vivo* (Akao et al., 1998, Bae et al., 2002).

In the present study, a total of 138 compounds were identified, which preliminarily clarified the material basis of KSLP. Subsequently, 13 representative compounds of KSLP were simultaneously quantified, which could be beneficial to improve the quality control. Finally, a specific and sensitive method was established for the simultaneous quantification of 6 compounds in the rat plasma after gavage administration of refining components of KSLP, which provided a reference for its clinical application.

CRediT authorship contribution statement

Chengjuan Liu: Conceptualization, Methodology, Investigation, Writing – original draft. **Qibao Jiang:** Supervision, Data curation, Resources. **Zhirong Zhou:** Supervision, Data curation, Resources. **Peng Lei:** Supervision, Data curation, Resources. **Peng Zhang:** Data curation. **Xin Chai:** Data curation. **Guixiang Pan:** Data curation. **Yuefei Wang:** Data curation. **Miaomiao Jiang:** Project administration, Funding acquisition, Writing – review & editing.

Declaration of competing interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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Appendix A. Supplementary data

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