



## Simultaneous determination by UPLC-MS/MS of seven bioactive compounds in rat plasma after oral administration of *Ginkgo biloba* tablets: application to a pharmacokinetic study\*

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**Abstract:** A rapid, reliable, and sensitive method was developed using ultra-performance liquid chromatography-tandem mass spectrometry (UPLC-MS/MS) with an electrospray ionization (ESI) source for determination of seven bioactive compounds in rat plasma after oral administration of *Ginkgo biloba* tablets (GBTs). The method simultaneously detects bilobalide (BB), ginkgolide A (GA), ginkgolide B (GB), ginkgolide C (GC), quercetin (QCT), kaempferol (KMF), and isorhamnetin (ISR) for pharmacokinetic study. The analytes and internal standard (IS) were extracted from rat plasma by acetic acid. An MS/MS detection was conducted using multiple reaction monitoring (MRM) and operating in the negative ionization mode. The calibration curve ranges were 5–500, 5–500, 2.5–250, 1–100, 1–100, 1–100, and 1–100 ng/ml for BB, GA, GB, GC, QCT, KMF, and ISR, respectively. The mean recovery of the analytes ranged from 68.11% to 84.42%. The intra- and inter-day precisions were in the range of 2.33%–9.86% and the accuracies were between 87.67% and 108.37%. The method was used successfully in a pharmacokinetic study of GBTs. The pharmacokinetic parameters of seven compounds were analyzed using a non-compartment model. Plasma concentrations of the seven compounds were determined up to 48 h after administration, and their pharmacokinetic parameters were in agreement with previous studies.

**Key words:** *Ginkgo biloba* tablet, Ultra-performance liquid chromatography-tandem mass spectrometry (UPLC-MS/MS), Pharmacokinetics

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### 1 Introduction

Products of *Ginkgo biloba* leaf extracts are used routinely in prescription drugs and dietary supple-

ments in many countries. They play an important role in treating cardiovascular diseases and neurological disorders (Diamond *et al.*, 2000). Many dosage forms of *G. biloba* leaf extracts have been documented in Chinese Pharmacopoeia (Chinese Pharmacopoeia Commission, 2010), such as tablets, capsules, and dripping pills. *Ginkgo biloba* tablets (GBTs) have been used for the clinical treatment of many conditions and diseases, such as stroke (Loh *et al.*, 2006; Zhang *et al.*, 2012), Alzheimer's disease (Stackman *et al.*, 2003), and cardiac angina (Xu and Xu, 2011).

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Flavonols and terpene lactones are considered to be the main bioactive compounds in GBTs (Deng and Zito, 2003). Bilobalide (BB), ginkgolide A (GA), ginkgolide B (GB), ginkgolide C (GC), quercetin (QCT), kaempferol (KMF), and isorhamnetin (ISR) are the main compounds for quality control (QC) of GBTs. Flavonols in *G. biloba* leaf extracts have been proven to have scavenging free radical effects (Ellnain-Wojtaszek *et al.*, 2003), and terpene lactones are platelet-activating factor (PAF) antagonists (Korth *et al.*, 1988). Du *et al.* (2013) studied the biological effects of bioactive compounds in GBTs and showed that the flavonols, mainly including QCT, KMF, and ISR, have remarkable antioxidant effects, and terpene lactones have anti-platelet aggregation activity.

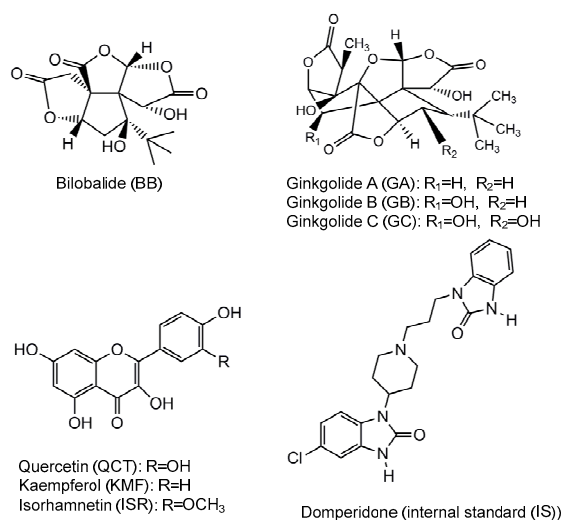
Recent studies have used various methods to determine the components of *G. biloba* extracts. Chen *et al.* (2010) and Hu *et al.* (2010) developed a high-performance liquid chromatography (HPLC)-ultra violet (UV) method to study the pharmacokinetics of QCT, KMF, and ISR in rat plasma. However, these studies could identify only the flavonol components. The HPLC-UV method is also not sensitive enough to apply to pharmacokinetic studies at clinical equivalent doses. Liquid chromatography-mass spectrometry (LC-MS) is an effective solution for complex matrix samples and is often used to study pharmacokinetics because of its high sensitivity and specificity. Xie *et al.* (2008) and Jin *et al.* (2011) successfully used an HPLC-tandem mass spectrometry (MS/MS) method to study the pharmacokinetics of terpene lactones. However, most methods usually focus only on flavonoid compounds or terpene lactones separately. Zhao *et al.* (2008) and Wu *et al.* (2013) reported that HPLC-MS/MS can be applied to the simultaneous determination of flavonoids and terpene lactones *in vivo*.

To depict the pharmacokinetic profiles of the main bioactive ingredients comprehensively, it is very important to develop a rapid and sensitive analytical method to quantify multiple components of GBTs in biological matrices. In this study, we aimed to develop an ultra-performance liquid chromatography (UPLC)-MS/MS method to quantify seven bioactive compounds simultaneously in rat plasma to evaluate their pharmacokinetic properties after a single oral administration of GBTs.

## 2 Materials and methods

### 2.1 Materials and reagents

BB (purity 100.0%, 110865-200605), GA (purity 95.4%, 110862-200305), GB (purity 99.9%, 110863-201209), GC (purity 97.1%, 110864-200906), QCT (purity 97.4%, 100081-200907), KMF (purity 99.0%, 110861-201209), ISR (purity 93.2%, 110860-201109), and internal standard (IS) domperidone (purity 100.0%, 100304-201103) were all purchased from the National Institute for the Control of Pharmaceutical and Biological Products, Beijing, China. The chemical structures of all analytes and the IS are shown in Fig. 1.



**Fig. 1** Chemical structures of bilobalide (BB), ginkgolide A (GA), ginkgolide B (GB), ginkgolide C (GC), quercetin (QCT), kaempferol (KMF), isorhamnetin (ISR), and internal standard (IS) domperidone

GBTs were supplied by Jiangsu Kanion Pharmaceutical Co., Ltd. (Lianyungang, China). The major components of each tablet were quantified by HPLC-evaporative light scattering detector (ELSD) for terpene lactones and HPLC-UV for flavonols (Table 1). Methanol and acetonitrile (MS grade) was supplied by Fisher Co., Inc. (USA) and formic acid (MS grade) by Sigma-Aldrich (USA). Deionized water was prepared using an ultrapure water system and further purified by the Millipore Simplicity<sup>®</sup> UV ultrapure water system (Germany). All other reagents were of analytical grade.

**Table 1** Contents of the major components in each *Ginkgo biloba* tablet (GBT) and dosage administered to each rat

Component	Content* (mg)	Dosage administered (mg/kg)
Bilobalide	2.22±0.003	3.33
Ginkgolide A	1.59±0.14	2.39
Ginkgolide B	0.41±0.04	0.62
Ginkgolide C	0.69±0.04	1.04
Quercetin	3.20±0.11	4.80
Kaempferol	3.68±0.08	5.52
Isorhamnetin	1.41±0.04	2.12

\* Data are presented as mean±SD ( $n=3$ ). The terpene lactones and flavonols in GBT were quantitated by HPLC-ELSD and HPLC-UV, respectively

## 2.2 UPLC-ESI-MS/MS conditions

A Waters<sup>®</sup> Xevo<sup>™</sup> TQ UPLC-MS/MS chromatographic system (USA) was used, interfaced with a triple quadrupole mass spectrometer and coupled with electrospray ionization (ESI). The analytes were separated on an Acquity BEH C18 column (2.1 mm×100 mm, 1.7 μm). The mobile phase was acetonitrile (A) and 0.1% formic acid in water (B). The gradient conditions of the mobile phase were: 0–0.5 min, A, held at 5%; 0.5–1.0 min, A, 5%–70%; 1.0–2.5 min, A, held at 70%; 2.5–3.5 min, A, 70%–30%; all at a flow rate of 0.4 ml/min and returned to the initial condition (acetonitrile-water 5:95, at a flow rate 0.4 ml/min) for a 4-min re-equilibration. The column temperature was maintained at 40 °C and the injection volume was 3 μl.

During the UPLC-ESI-MS/MS analysis, the negative ion mode was used with selected multiple reaction monitoring (MRM) for the quantitative analysis. The optimal MRM transitions for the

analytes to quantify were determined as follows: the mass-to-charge ratio ( $m/z$ ) for BB was 325.00→182.85, GA 407.21→350.91, GB 422.93→366.96, GC 438.99→382.92, QCT 300.85→150.85, KMF 285.12→142.97, ISR 314.77→299.88, and IS 423.63→166.74. Nitrogen was used as the collision gas. The nebulizer, cone, and de-solvation gas flows were set at 7.0 bar (1 bar=0.1 MPa), 150 L/h, and 800 L/h, respectively. The de-solvation temperature was set at 400 °C. Dwell time was set to 40 ms for each transition. The capillary voltage was set at 2.5 kV. Compound parameters are shown in Table 2.

## 2.3 Animals

Sprague-Dawley male rats (SPF (Beijing) Laboratory Animal Technology Co., Ltd., China) weighing (300±10) g were used. All experiments were carried out according to the guidelines of the Animal Care and Use Committee of Beijing University of Chinese Medicine, China. Before the experiment, all rats were acclimatized to the conditions of the laboratory over one week.

## 2.4 Plasma sample preparation

A liquid-liquid extraction method was used to extract all the plasma samples. To an aliquot of 100 μl of plasma, the same volume of 4 mol/L hydrochloric acid was added and vortexed for 3 min. The mixture was then hydrolyzed in a water bath at 80 °C for 30 min; afterwards, 1.0 ml acetic acid was added to the hydrolyzed and cooled sample, followed by vortexing for 3 min. After centrifugation at 10000 r/min for 10 min, the 800 μl supernatant was transferred to a clean centrifuge tube and evaporated to dryness under a gentle stream of nitrogen. The residue was

**Table 2** Optimum mass spectrometry conditions for the analytes in GBTs and the internal standard (IS)

Analyte	Scan type	Ionization mode	Precursor ion	Product ion	Cone (V)	Collision (V)
BB	Multiple reaction monitoring	Negative	325.00	182.85	30	20
GA	Multiple reaction monitoring	Negative	407.21	350.91	54	14
GB	Multiple reaction monitoring	Negative	422.93	366.96	22	16
GC	Multiple reaction monitoring	Negative	438.99	382.92	30	16
QCT	Multiple reaction monitoring	Negative	300.85	150.85	54	18
KMF	Multiple reaction monitoring	Negative	285.12	142.97	76	20
ISR	Multiple reaction monitoring	Negative	314.77	299.88	68	20
IS	Multiple reaction monitoring	Negative	423.63	166.74	2	36

BB: bilobalide; GA: ginkgolide A; GB: ginkgolide B; GC: ginkgolide C; QCT: quercetin; KMF: kaempferol; ISR: isorhamnetin

reconstituted in 100  $\mu$ l methanol, followed by centrifugation at 10 000 r/m for 10 min; finally, an aliquot of 3  $\mu$ l of the solution was injected into the UPLC-MS/MS system for analysis.

## 2.5 Preparation of reference solutions and samples for QC

Reference compounds were accurately weighed and dissolved in methanol to produce stock solutions. The concentrations of BB, GA, GB, GC, QCT, KMF, and ISR were 102, 96.0, 93.0, 69.0, 73.0, 26.0, and 27.0  $\mu$ g/ml, respectively. Seven different concentrations of a standard working solution of all analytes were obtained by diluting the solutions with methanol. All solutions were stored at 4 °C and then brought back to room temperature before use. A calibration standard at the seven different concentrations was obtained by spiking 100  $\mu$ l of the standard working solutions and 50  $\mu$ l of the IS solution (20 ng/ml) with 100  $\mu$ l blank rat plasma. For method validation, samples for QC were prepared in the same way.

## 2.6 Method validation

### 2.6.1 Selectivity and specificity

Chromatograms of the IS at the lowest limit of quantification (LLOQ), blank rat plasma from six sources, blank plasma spiked with the analytes, and plasma samples after oral administration were compared to investigate the specificity and selectivity of the method.

### 2.6.2 Calibration curve

For the seven different standard samples, the concentrations were: 5, 12.5, 25, 50, 100, 250, 500 ng/ml for BB; 5, 12.5, 25, 50, 100, 250, 500 ng/ml for GA; 2.5, 6.25, 12.5, 25, 50, 125, 250 ng/ml for GB; 1, 2.5, 5, 10, 20, 50, 100 ng/ml for GC; 1, 2.5, 5, 10, 20, 50, 100 ng/ml for QCT; 1, 2.5, 5, 10, 20, 50, 100 ng/ml for KMF; and, 1, 2.5, 5, 10, 20, 50, 100 ng/ml for ISR ( $n=3$ ). These sample preparation procedures were used to extract and analyze the samples. Plots of the peak area ratios of BB, GA, GB, GC, QCT, KMF, and ISR to IS ( $Y$ ) vs. the concentrations of BB, GA, GB, GC, QCT, KMF, and ISR ( $X$ , in ng/ml) using linear regression, were used to confirm the calculation curves.

### 2.6.3 Recovery and matrix effect

The extraction recoveries (ERs) of BB, GA, GB, GC, QCT, KMF, and ISR from plasma were determined at three different concentrations of BB (12.5, 50, 250 ng/ml), GA (12.5, 50, 250 ng/ml), GB (6.25, 25.0, 125 ng/ml), GC (2.5, 10, 50 ng/ml), QCT (2.5, 10, 50 ng/ml), KMF (2.5, 10, 50 ng/ml), and ISR (2.5, 10, 50 ng/ml), respectively. The samples were then extracted and analyzed by the same procedure. To determine the recovery of analytes, the mean peak areas ( $n=5$ ) of the extracted samples and the corresponding concentrations of the analyte neat standard solutions were compared. The matrix effect was calculated by comparing the mean peak areas of prepared samples with the mean peak areas of the neat standard solutions.

### 2.6.4 Precision and accuracy

Precision and accuracy were assessed on three consecutive days by using mixtures containing low, medium, or high concentrations of the QC samples. Comparisons of the calculated average measured concentrations using calibration curves with added concentrations were used to determine accuracy, expressed as the relative standard deviation (RSD).

### 2.6.5 Stability

For testing the stability of the analytes in rat plasma, the same concentrations of the following stored and freshly prepared samples were compared. Analysis of the extracted samples, stored for 4 h maintained at room temperature, was used to determine their short-term stability. To determine the stability of samples in the autosampler, the extracted samples were analyzed after being left in the autosampler at 10 °C for 8 h. Freeze and thaw stability was determined by subjecting the QC samples to three cycles of freezing at -20 °C for 12 h and thawing at room temperature for 2 h.

## 2.7 Pharmacokinetic studies

Five male rats were orally administered a single dose of GBTs. The rats were given samples equivalent to 3.33, 2.39, 0.62, 1.04, 4.80, 5.52, and 2.12 mg/kg of BB, GA, GB, GC, QCT, KMF, and ISR, respectively (Table 1). After dosing for 0, 0.166, 0.5, 0.75, 1, 1.5, 2, 4, 6, 8, 12, 24, 36, and 48 h, blood samples were collected in heparinized tubes to

quantify the drug concentrations. The blood samples were centrifuged at 10000 r/min for 10 min, and then the supernatants were collected in tightly sealed plastic tubes (sodium citrate anticoagulation) and kept frozen at  $-20^{\circ}\text{C}$  until analyzed. The plasma concentrations of BB, GA, GB, GC, QCT, KMF, and ISR were expressed as mean $\pm$ standard deviation (SD), and the mean concentration-time curve was plotted. Non-compartmental pharmacokinetic parameters were computed using the software Kinetica 4.4.

### 3 Results and discussion

#### 3.1 UPLC-MS/MS conditions and sample preparation optimization

To optimize the mass spectra conditions, abundant deprotonated molecules of the seven components that formed the base peak of each mass spectrum were observed from parent scans. According to the recent literature, the negative mode was suitable to ionize the seven components (Zhao *et al.*, 2008; Wu *et al.*, 2013). The instrument parameters were optimized to

detect parent and daughter ions. So the parent ions  $[\text{M}-\text{H}]^{-}$ ,  $m/z$  325.00 for BB, 407.21 for GA, 422.93 for GB, and 438.99 for GC, were subjected to collision. The most abundant fragments were then selected to be daughter ions (Fig. 2). The mass transition patterns,  $m/z$  325.00 $\rightarrow$ 182.85, 407.21 $\rightarrow$ 350.91, 422.93 $\rightarrow$ 366.96, 438.99 $\rightarrow$ 382.92, 300.85 $\rightarrow$ 150.85, 285.12 $\rightarrow$ 142.97, and 314.77 $\rightarrow$ 299.88, were selected to monitor BB, GA, GB, GC, QCT, KMF, and ISR, respectively. In the same way, an MS/MS channel of  $m/z$  423.63 $\rightarrow$ 166.74 was chosen to monitor the IS.

To optimize sample preparation, protein precipitation and liquid-liquid extraction methods were tested in our study. Using protein precipitation by methanol and acetonitrile, the recovery of four ginkgolides was not satisfactory. Therefore, we tested a liquid-liquid extraction method and acetidin was proved suitable for extracting all analytes from rat plasma. Chandra *et al.* (2011) reported that ginkgo flavonoids include flavonols and their glycosides. Some studies have shown that flavonoid glycosides can be absorbed in the intestine by bacteria to facilitate absorption (Bokkenheuser *et al.*, 1987; Kim *et al.*,

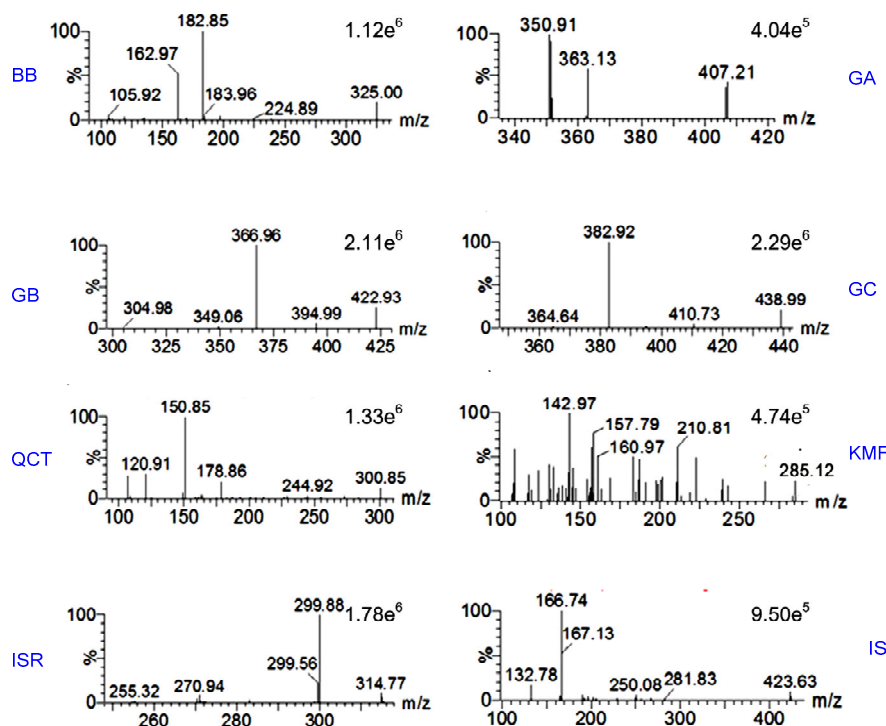


Fig. 2 Daughter ion mass spectra of the analytes in GBTs and the internal standard (IS)

BB: bilobalide; GA: ginkgolide A; GB: ginkgolide B; GC: ginkgolide C; QCT: quercetin; KMF: kaempferol; ISR: isorhamnetin

1998; Wu *et al.*, 2013). However, other studies have shown that flavonoids exist as aglycones, glucuronic acid conjugates, and sulfonic acid conjugates in plasma (Manach *et al.*, 1996; Yodogawa *et al.*, 2003; Silberberg *et al.*, 2005). So the plasma samples should be acid hydrolytic to transform conjugates to aglycones.

### 3.2 Validation of the assay

#### 3.2.1 Selectivity and specificity

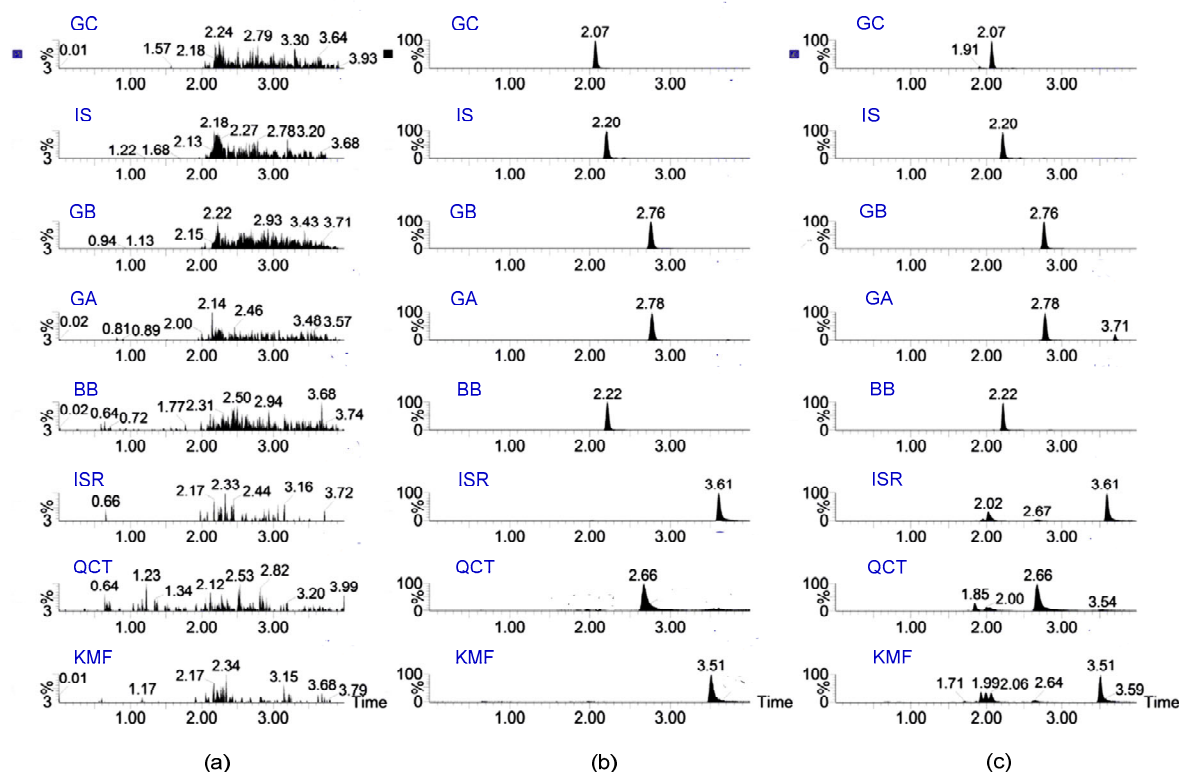
Under the UPLC-MS/MS conditions, a specific voltage at the optimum value was set to obtain the best sensitivity and specificity for each analyte (Fig. 3). No obvious endogenous interference was observed and a good separation of the analytes was achieved. The chromatographic peaks were at the retention time of 2.07, 2.20, 2.76, 2.78, 2.22, 3.61, 2.66, and 3.51 min for GC, IS, GB, GA, BB, ISR, QCT, and KMF, respectively.

#### 3.2.2 Linearities of calibration curves and the lowest limit of quantification (LLOQ)

The method was linear over the concentration range of 5–500 ng/ml for BB, 5–500 ng/ml for GA, 2.5–250 ng/ml for GB, 1–100 ng/ml for GC, 1.0–100 ng/ml for QCT, 1–100 ng/ml for KMF, and 1–100 ng/ml for ISR. The mean values of the linear regression equations of the analytes are listed in Table 3. The LLOQ was defined as the lowest concentration on the calibration curve with acceptable precision and accuracy.

#### 3.2.3 Matrix effect and recovery

The matrix effect in different mobile phases was investigated during method development. Using an electrolyte-free mobile phase, a significant matrix effect was observed. A mobile phase modified with 0.1% HCOOH minimized the matrix effects, but



**Fig. 3** Representative multiple reaction monitoring (MRM) chromatograms of the analytes and the internal standard (IS)

(a) Blank plasma sample; (b) Blank plasma sample spiked with the analytes at LLOQ and IS; (c) Plasma sample of orally administrated rat at 6 h post dose. GC: ginkgolide C; GB: ginkgolide B; GA: ginkgolide A; BB: bilobalide; ISR: isorhamnetin; QCT: quercetin; KMF: kaempferol

flavonols were still affected slightly. We then adjusted the eluent gradient, and a smaller matrix effect was observed (Table 4). This phenomenon may due to the co-eluting of residual matrix components interfering with the ionization of the target analyte (Dams *et al.*, 2003). Zhao *et al.* (2008) reported that inclusion of ammonium formate (0.2 mmol/L) in the mobile phase generated beneficial LC-electrolyte effects, including increased ESI efficiency and capacity. While 0.1% HCOOH in the mobile phase was found to improve the peak shape of the analytes, ammonium formate had no significant effect on the analytes in our study. Thus, 0.1% HCOOH was chosen as the mobile phase.

### 3.2.4 Precision and accuracy

The intra-day precision and accuracy were determined by the replicate analyses of QC samples ( $n=5$ ) at three concentrations during the three separate days. One replicate of the QC samples at each of three concentrations from three separate validation batches was used to evaluate the inter-day precision. The intra-day precision and accuracy of the method were determined by analyzing QC samples ( $n=5$ ) at three concentrations on the same day, and inter-day precision and accuracy by analyzing QC samples ( $n=5$ ) at three concentrations on each of three validation days.

**Table 3** Linearities, the lowest limits of quantification (LLOQs), and the lowest limits of detection (LLODs) for analytes of GBTs in rat plasma ( $n=3$ )

Analyte	Retention time (min)	Linear range (ng/ml)	Linear equation	$R^2$	LLOQ (ng/ml)	LLOD (ng/ml)
BB	2.22	5–500	$Y=3.1341X-39.786$	0.9960	5	0.018
GA	2.78	5–500	$Y=0.2459X-3.0339$	0.9969	5	0.032
GB	2.76	2.5–250	$Y=8.7944X-21.578$	0.9945	2.5	0.018
GC	2.07	1–100	$Y=8.0653X+12.650$	0.9997	1	0.023
QCT	2.66	1–100	$Y=1.9149X+3.2370$	0.9934	1	0.360
KMF	3.51	1–100	$Y=0.2868X+0.8961$	0.9984	1	0.270
ISR	3.61	1–100	$Y=9.8024X-5.6094$	0.9982	1	0.017

BB: bilobalide; GA: ginkgolide A; GB: ginkgolide B; GC: ginkgolide C; QCT: quercetin; KMF: kaempferol; ISR: isorhamnetin

**Table 4** Extraction recovery and matrix effect for the analytes of GBTs in rat plasma

Analyte	Concentration added (ng/ml)	Matrix effect		Extraction recovery	
		Value* (%)	RSD (%)	Value* (%)	RSD (%)
GC	2.5	103.94±9.63	8.25	79.87±6.59	9.26
	10	96.50±5.08	4.24	77.26±3.27	5.26
	50	95.27±4.61	4.84	68.29±8.99	8.08
BB	12.5	102.97±10.14	9.84	76.67±4.13	5.39
	50	97.33±6.98	7.17	75.68±4.03	5.32
	250	97.53±6.19	6.34	71.01±1.98	2.79
QCT	2.5	102.13±7.55	7.39	84.42±8.51	10.08
	10	99.01±8.61	8.70	80.17±2.12	2.64
	50	101.63±14.85	14.61	74.38±10.57	14.21
GB	6.25	101.51±12.26	12.08	77.32±4.82	6.23
	25	104.49±10.99	10.52	75.17±4.33	5.76
	125	102.35±13.92	13.60	79.84±2.27	2.85
GA	12.5	94.69±9.38	9.90	71.94±3.72	5.18
	50	94.91±10.81	11.39	72.36±5.16	7.14
	250	94.40±11.11	11.77	75.32±0.98	1.31
KMF	2.5	97.07±8.02	8.26	71.33±2.52	3.53
	10	100.87±6.48	6.42	68.11±2.92	4.29
	50	93.89±3.40	3.62	70.46±2.81	3.99
ISR	2.5	94.70±4.98	5.26	83.03±4.57	5.50
	10	96.04±5.14	5.35	83.03±4.57	5.50
	50	90.94±4.17	4.59	83.17±3.24	3.89

\* Data are expressed as mean±SD ( $n=5$ ). GC: ginkgolide C; BB: bilobalide; QCT: quercetin; GB: ginkgolide B; GA: ginkgolide A; KMF: kaempferol; ISR: isorhamnetin; RSD: relative standard deviation

The results are summarized in Table 5. The intra-day precision was within 10.13% and the inter-day precision value was within 9.82%. The assay mean accuracy was 93.20%–107.23% of the nominal values.

### 3.2.5 Stability

During the short-term and autosampler stability assays, all analytes were stable. None of the analytes in rat plasma was significantly affected after being subjected to three freezing and thawing cycles. The RSD values for long-term stability were less than 11.11% (Table 6). The results showed that all analytes were stable in the above conditions.

### 3.3 Pharmacokinetics studies

The UPLC-ESI-MS/MS method was successfully applied to the pharmacokinetic study of GA, BB, GB, GC, QCT, KMF, and ISR in rat plasma after oral administration of GBTs. The mean plasma concentration-time profiles of the terpene lactones and the flavonols in rats after a single oral dose of GBTs are shown in

Figs. 4 and 5, respectively. The pharmacokinetic parameters are shown in Tables 7 and 8. Note that there were marked differences between the pharmacokinetic behavior of terpene lactones and flavonols. After oral administration of GBTs, terpene lactones were eliminated faster than flavonols. This is consistent with Chen *et al.* (2013). Moreover, the flavonols presented double peaks in curves of mean plasma concentration. This phenomenon was also reported previously and indicates that these components might have enterohepatic recirculation (Tang *et al.*, 2009; Chen *et al.*, 2010).

### 4 Conclusions

In this study, a UPLC-ESI-MS/MS method was developed and validated for simultaneous quantification of plasma concentrations of BB, GA, GB, GC, QCT, KMF, and ISR. A simple and high recovery extraction method was carried out for sample

**Table 5 Precision and accuracy for the analytes in rat plasma**

Analyte	Conc. added (ng/ml)	Inter-day			Intra-day		
		Conc. found (ng/ml)	Accuracy (%)	RSD (%)	Conc. found (ng/ml)	Accuracy (%)	RSD (%)
GC	2.5	2.35±0.19	94.14	8.12	2.39±0.16	95.45	6.83
	10	10.74±0.42	107.41	3.95	10.15±0.82	101.52	8.11
	50	53.85±2.50	107.70	4.64	51.01±3.41	102.52	6.68
BB	12.5	11.86±1.16	94.84	9.82	11.65±0.88	93.20	7.58
	50	50.57±1.59	101.15	3.14	51.54±3.60	103.08	6.99
	250	259.44±14.54	103.77	5.60	256.17±15.69	102.47	6.13
QCT	2.5	2.44±0.21	97.78	8.72	2.57±0.26	102.77	10.13
	10	10.22±0.77	102.46	7.49	10.02±0.81	100.22	8.05
	50	52.24±5.24	104.13	9.66	50.84±4.96	101.68	9.75
GB	6.25	5.84±0.36	93.45	6.15	6.05±0.35	96.77	5.86
	25	24.14±0.56	96.54	2.33	25.51±1.87	102.04	7.32
	125	128.53±4.81	102.83	3.74	127.34±7.20	101.87	5.65
GA	12.5	12.55±0.71	100.41	5.67	11.88±1.02	95.06	8.69
	50	45.93±1.94	91.86	4.22	48.89±4.71	97.77	9.64
	250	238.99±11.23	95.59	4.70	243.44±14.08	97.38	5.78
KMF	2.5	2.64±0.16	105.49	5.89	2.68±0.13	107.23	5.02
	10	8.77±0.37	87.72	4.22	9.74±0.95	97.36	9.79
	50	48.66±2.85	97.32	5.85	48.91±3.95	97.83	8.07
ISR	2.5	2.43±0.20	97.67	8.35	2.49±0.25	99.57	9.86
	10	8.77±0.30	87.67	3.44	9.42±0.78	94.22	8.27
	50	54.18±1.71	108.37	3.16	50.15±4.59	100.30	9.15

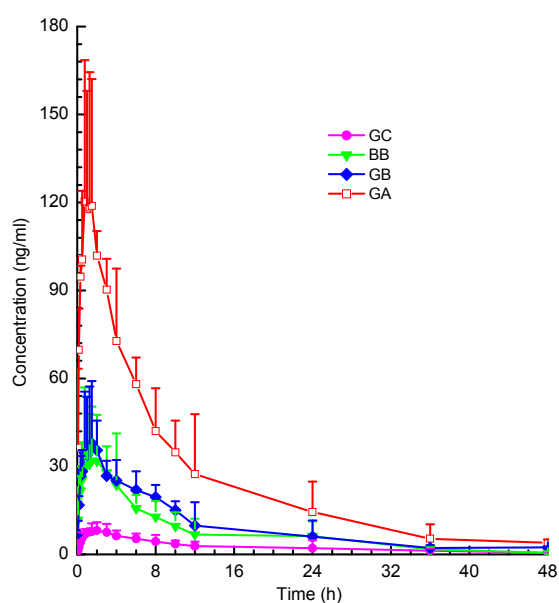
The values for the concentration found are expressed as mean±SD, five replicates per day for three days. GC: ginkgolide C; BB: bilobalide; QCT: quercetin; GB: ginkgolide B; GA: ginkgolide A; KMF: kaempferol; ISR: isorhamnetin; RSD: relative standard deviation



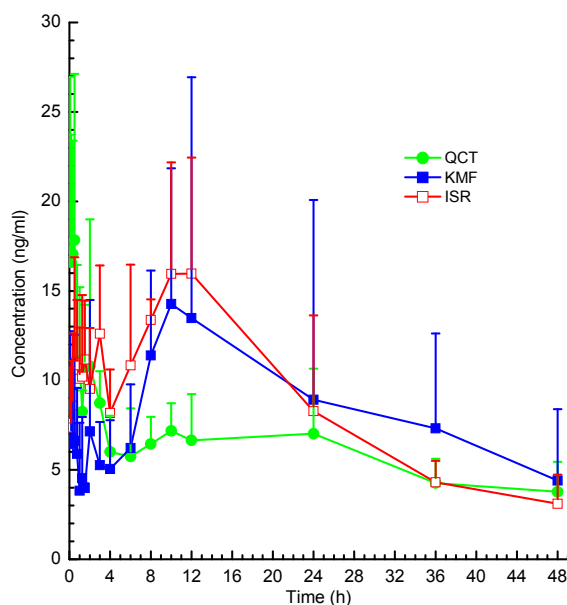
**Table 6 Stability of the analytes in rat plasma**

Analyte	Conc. added (ng/ml)	Short-term stability		Autosampler stability		Freeze-thaw stability (three cycles)		Four-week stability	
		Conc. found (%)	RSD (%)	Conc. found (%)	RSD (%)	Conc. found (%)	RSD (%)	Conc. found (%)	RSD (%)
GC	2.5	98.42±5.67	5.76	95.28±9.00	9.45	97.15±10.03	10.32	98.10±7.21	7.35
	10	101.02±4.35	4.31	104.95±2.84	2.71	102.94±3.48	3.38	103.76±8.29	7.99
	50	103.40±9.78	9.46	97.40±5.16	5.29	98.37±4.81	4.89	97.57±3.42	3.51
BB	12.5	87.60±4.52	5.16	84.59±10.79	12.76	90.10±5.27	5.85	105.67±10.12	9.58
	50	97.56±4.66	4.78	122.30±2.19	1.79	109.64±5.00	4.56	96.74±9.01	9.31
	250	105.12±10.24	9.74	100.84±5.90	5.85	102.29±5.30	5.18	97.11±10.08	10.38
QCT	2.5	101.65±3.41	3.35	104.43±7.73	7.40	98.72±6.81	6.90	101.31±11.09	10.95
	10	104.42±4.56	4.37	113.21±7.98	7.05	101.21±8.75	8.65	97.10±7.23	7.45
	50	101.74±7.88	7.75	98.97±9.05	9.14	105.31±7.64	7.25	102.35±8.22	8.03
GB	6.25	93.72±5.67	6.05	105.71±3.57	3.37	105.20±9.27	8.81	98.70±8.34	8.45
	25	98.67±6.78	6.87	114.43±1.55	1.35	104.52±5.65	5.41	100.78±5.48	5.44
	125	104.14±7.99	7.67	108.76±5.10	4.69	110.19±3.39	3.07	95.77±5.63	5.88
GA	12.5	106.24±10.34	9.73	79.73±5.88	7.37	89.05±4.91	5.51	97.70±6.80	6.96
	50	109.12±4.01	3.67	118.25±3.20	2.71	103.68±6.47	6.24	101.21±5.84	5.77
	250	90.31±6.74	7.46	89.14±3.13	3.52	90.97±2.98	3.28	98.81±4.45	4.50
KMF	2.5	107.34±5.43	5.06	96.83±9.89	10.21	96.55±7.32	7.58	97.99±7.22	7.37
	10	100.72±2.45	2.43	98.73±4.82	4.88	104.93±7.54	7.19	102.31±6.78	6.63
	50	98.97±3.21	3.24	102.3±7.15	6.99	99.74±8.71	8.73	108.98±12.11	11.11
ISR	2.5	104.54±4.54	4.34	87.98±5.75	6.54	89.74±5.25	5.85	101.16±9.12	9.02
	10	101.23±6.77	6.69	112.32±5.78	5.15	101.72±5.67	5.57	103.45±7.76	7.50
	50	98.78±7.68	7.77	100.44±6.91	6.88	105.50±9.78	9.27	98.79±6.34	6.42

The values for the concentration found are expressed as mean±SD,  $n=3$ . GC: ginkgolide C; BB: bilobalide; QCT: quercetin; GB: ginkgolide B; GA: ginkgolide A; KMF: kaempferol; ISR: isorhamnetin; RSD: relative standard deviation



**Fig. 4 Plasma concentration-time profiles of bilobalide (BB), ginkgolide A (GA), ginkgolide B (GB), and ginkgolide C (GC) in rats after a single oral dose of GBTs**  
Data are presented as mean±SD,  $n=5$



**Fig. 5 Plasma concentration-time profiles of quercetin (QCT), kaempferol (KMF), and isorhamnetin (ISR) in rats after a single oral dose of GBTs**  
Data are presented as mean±SD,  $n=5$

**Table 7 Pharmacokinetic parameters of BB, GA, GB, and GC following a single oral administration of GBTs to rats**

Analyte	$C_{\max}$ (ng/ml)	$T_{\max}$ (h)	$AUC_{0-t}$ (ng/(ml·h))	$AUC_{0-\infty}$ (ng/(ml·h))	$T_{1/2}$ (h)	MRT (h)
BB	40.62±20.53	2.08±1.81	303.02±66.64	346.69±104.26	3.58±1.86	5.65±2.60
GA	130.10±39.73	1.15±0.42	1125.74±283.31	1243.46±386.35	3.66±1.20	7.15±3.46
GB	38.88±19.77	0.92±0.49	428.37±134.55	457.32±148.72	4.60±2.80	6.72±2.32
GC	9.36±1.85	3.20±1.89	132.44±63.55	140.48±61.59	5.48±0.86	8.52±1.50

Data are presented as mean±SD,  $n=5$ . BB: bilobalide; GA: ginkgolide A; GB: ginkgolide B; GC: ginkgolide C;  $C_{\max}$ : peak concentration;  $T_{\max}$ : peak time;  $AUC_{0-t}$ : area under the curve from 0 h to  $t$  h;  $AUC_{0-\infty}$ : area under the curve from 0 h to infinite;  $T_{1/2}$ : half-life; MRT: mean residence time

**Table 8 Pharmacokinetic parameters of QCT, KMF, and ISR following a single oral administration of GBTs to rats**

Analyte	$C_{1\max}$ (ng/ml)	$C_{2\max}$ (ng/ml)	$T_{1\max}$ (h)	$T_{2\max}$ (h)	$AUC_{0-t}$ (ng/(ml·h))	$AUC_{0-\infty}$ (ng/(ml·h))	$T_{1/2}$ (h)	MRT (h)
QCT	26.40±5.74	13.45±6.60	0.22±0.19	11.50±11.54	285.04±78.20	444.80±174.77	26.74±9.38	45.48±11.10
KMF	12.26±2.37	14.81±7.54	0.18±0.18	9.00±8.49	297.22±209.13	552.27±283.01	14.23±3.38	29.70±6.19
ISR	11.55±4.93	21.04±8.40	0.25±0.23	12.00±11.02	704.86±328.49	978.37±465.73	22.46±2.00	37.29±5.18

Data are presented as mean±SD,  $n=5$ . QCT: quercetin; KMF: kaempferol; ISR: isorhamnetin;  $C_{\max}$ : peak concentration;  $T_{\max}$ : peak time;  $AUC_{0-t}$ : area under the curve from 0 h to  $t$  h;  $AUC_{0-\infty}$ : area under the curve from 0 h to infinite;  $T_{1/2}$ : half-life; MRT: mean residence time

pretreatment. Moreover, no interference was found between endogenous substances and the analytes. The developed method has proven to be very sensitive and reliable, and offers shorter analysis time over existing methods, with a single run taking only 4 min. In conclusion, the established UPLC-ESI-MS/MS method demonstrated good performance including specificity, linearity, detection and quantification limits, precision and accuracy, and was successfully used to quantify the seven bioactive compounds in rat plasma to support pharmacokinetic studies.

### Compliance with ethics guidelines

Wen-ping WANG, Na LIU, Qian KANG, Pei-pei DU, Yi LAN, Bo-chen ZHAO, Yan-yan CHEN, Qing ZHANG, Hui LI, Ye-wen ZHANG, and Qing WU declare that they have no conflict of interest.

All institutional and national guidelines for the care and use of laboratory animals were followed.

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## 中文概要:

**本文题目:** UPLC-MS/MS 法测定口服银杏叶片的大鼠血浆中七种活性成分及药代动力学研究

**Simultaneous determination by UPLC-MS/MS of seven bioactive compounds in rat plasma after oral administration of *Ginkgo biloba* tablets: application to a pharmacokinetic study**

**研究目的:** 建立一个快速灵敏且具有专属性的体内分析方法用于银杏叶片的药代动力学研究。

**创新要点:** 运用超高压液相色谱-二级质谱连用法 (UPLC-MS/MS) 进行大鼠血浆中七种活性成分的测定, 方法简单快速, 适合于银杏叶制剂的药代动力学研究。

**研究方法:** 采用液液萃取进行血浆样品处理, 该方法简单且快速。采用超高压液相色谱 (UPLC) 进行七种活性成分分离, 达到了多种成分快速分离的目的。采用二级质谱 (MS/MS) 进行检测, 优化质谱条件, 减小了内源性干扰并提高检测灵敏度。经过专属性、线性、回收率、基质效应、精密度、准确度和稳定性考察, 验证了该方法的适用性。经验证, 该方法适用于银杏叶片的药代动力学研究。

**重要结论:** 液液萃取处理血浆样品简单可靠, 回收率稳定, 无基质效应 (表 4)。UPLC-MS/MS 能够快速、灵敏、专属性地分析大鼠血浆七种活性成分的血药浓度 (表 3 和图 3)。该方法适用于银杏叶片的七种活性成分的药代动力学研究 (图 4 和 5)。

**关键词组:** 超高压液相色谱-二级质谱连用法 (UPLC-MS/MS); 银杏叶片; 药代动力学