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# **Determination of QLNC-3A6 in canine plasma by UHPLC-MS/MS and its application in pharmacokinetic studies**

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#### ABSTRACT

Multi-targeted tyrosine kinase inhibitor QLNC-3A6 Di-maleate, a structurally novel small molecule compound, has therapeutic efficacy for the treatment of canine cutaneous mast cell tumor (CMCT) caused by mutations in the c-Kit gene. Since pharmacokinetic (PK) information plays an important role in the development and application of new drugs, etc., a rapid, highly sensitive and selective UHPLC-MS/MS analytical method was developed and validated for the first time in this study for the quantitative detection of QLNC-3A6 in canine plasma. 100 µL of plasma was precipitated using 350µL of acetonitrile, and Chromatographic separation was performed on a Phenomenex Kinetex C18 column (50×2.1mm, 2.6µm) at a flow rate of 0.4mL/min, the mobile phases were set to 0.1% formic acid aqueous solution (A) and 0.1% formic acid acetonitrile (B). The calibration curve linear range was 0.5–100 ng/mL ( $R^2$ >0.99). The intraday and interday precision values (relative standard deviation, RSD) were 2.06–13.57% and 6.90–9.14%. Intraday and interday accuracies were −10.73 to 9.54% and −3.86 to 0.70% respectively. The dilution integrity RSD value and stability RSD value were less than 3.77 and 7.45%, respectively. Subsequently, the pharmacokinetics were investigated in canine after oral administration of QLNC-3A6 Di-maleate tablets at a dose of 3mg/kg BW using this method. The results showed that QLNC-3A6 showed fast absorption rate, rapid distribution and slow metabolic elimination in canine plasma. The results of the main PK parameters including *λ*z,  $T_{1/2\lambda z}$ ,  $C_{\text{max}}$ , *T<sub>max</sub>* and *AUC*<sub>last</sub> were 0.07±0.01/h, 11.00±2.57h, 50.88±31.94ng/mL, 9.08±11.57h and  $836.48 \pm 230.53$  ng h/mL, respectively.

# **Introduction**

<span id="page-0-14"></span>In recent years, with the rise in the number of pet canines and the rapid development of animal medical conditions, the lifespan of pet canines has been extended, however, the cancer rate of canines with extended lifespan has also risen (Vascellari et al. [2016\)](#page-10-0). Therefore, drugs for the treatment of canine tumors are urgently needed. Targeted therapy for different subtypes of tumors is the main direction for the future development of antitumor drugs, and targeted therapeutic agents are the main means of personalized treatment (Dobson and Scase [2007\)](#page-9-0). Currently, more and more pharmaceutical companies and research institutes are committed to the research and development of new antitumor targeted therapeutics (Yancey, Merritt, White, et al. [2010\)](#page-10-1) or different formulations (Liu et al. [2024](#page-9-1)), so as to achieve a more precise and rapid treatment. Meanwhile, the

<span id="page-0-13"></span><span id="page-0-12"></span><span id="page-0-11"></span><span id="page-0-7"></span>pathogenesis of some tumors is similar in humans and animals (Patruno et al. [2014](#page-10-2); Vascellari et al. [2016](#page-10-3); Simpson et al. [2017](#page-10-4)). For example, CMCT, the third most common tumor subtype and the most common malignant skin tumor in canines, accounts for 11% of canine skin cancer cases (de Nardi et al. [2022](#page-9-2)). Its malignancy rate can be as high as 19–39% (Ma et al. [2021](#page-10-5)). CMCT is a tumor driven by the cell surface protein tyrosine kinase c-Kit, and CMCT pathogenesis has been associated with mutations in the c-Kit gene (Yancey, Merritt, Lesman, et al. [2010](#page-10-6); Bonkobara [2015](#page-9-3); Ferguson et al. [2016\)](#page-9-4), which is similar to the mechanism of c-Kit mutations found in human gastrointestinal stromal tumors (GIST) (Patruno et al. [2014\)](#page-10-7). Therefore, the conversion of human drugs to veterinary drugs is now also becoming a new drug research direction.

<span id="page-0-15"></span><span id="page-0-9"></span><span id="page-0-6"></span>Qilu Animal Health Products Co., Ltd. has developed a small molecule compound of new structure,

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<span id="page-1-1"></span>QLNC-3A6 Di-maleate, a small molecule tyrosine kinase inhibitors (TKI) with a mechanism of action similar to that of the marketed drugs including imatinib mesylate (Gleevec, STI571), sunitinib malate (Sutent, SU11248), toceranib phosphate (Palladia, SU11654), and other drugs. The structure of QLNC-3A6 Di-maleate is similar to that drug structure of sunitinib malate, a molecularly modified version of sunitinib malate ([Figure 1\)](#page-1-0). However, QLNC-3A6 Di-maleate is a multi-targeted drug, compared with single-targeted drugs, multi-targeted drugs can avoid the disadvantages such as limited efficacy, drug resistance and having potential drug-drug interactions that occur when multiple single-targeted drugs are combined (Raghavendra et al. [2018](#page-10-8)). And QLNC-3A6 Di-maleate has a favorable safety profile, unlike anticancer drugs with cytotoxic classes (Liu et al. [2024\)](#page-9-5).

<span id="page-1-9"></span><span id="page-1-8"></span><span id="page-1-7"></span><span id="page-1-6"></span><span id="page-1-5"></span><span id="page-1-4"></span><span id="page-1-3"></span>The acquisition of PK information occupies an important role in the development and application of new drugs. For QLNC-3A6 analogues several assays are now available. Such methods as LC-MS/MS or UPLC-MS/MS (Tang et al. [2020](#page-10-9); Zhang et al. [2022\)](#page-10-10) can be used for the determination of imatinib (Caterino et al. [2013;](#page-9-6) Xu et al. [2019](#page-10-11)), sunitinib (AboulMagd and Abdelwahab [2021](#page-9-7)) and toceranib (TOC) (Yancey, Merritt, Lesman, et al. [2010\)](#page-10-6), among others. At the same time there are now a large number of studies on similar drugs for canine tyrosine kinase-driven different tumours such as insulinomas (Sheppard-Olivares et al. [2022\)](#page-10-12) and pancreatic adenocarcinoma (Musser and Johannes [2021\)](#page-10-13). To the best of our knowledge, there are now studies on pharmacokinetic properties, distribution, metabolism and excretion of the relevant components in CMCT affected dogs administered at a dose of 3.25mg/kg BW toceranib phosphate (Yancey, Merritt, Lesman, et al. [2010\)](#page-10-6). However, there is no method to detect QLNC-3A6 in canine plasma, so there is an urgent need to develop a rapid and accurate method to detect QLNC-3A6 in canine plasma. The present study aims to explore and establish a sensitive, accurate, and rapid method for the quantification of QLNC-3A6 in canine plasma, and to carry out a complete validation of the method, which was successfully applied to a single oral dose of QLNC-3A6 Di-maleate tablet 3mg/kg BW after a PK study of QLNC-3A6 in canine plasma. Among them, toceranib phosphate was selected as the internal standard (IS) drug for this study due to its better <span id="page-1-2"></span>stability and the fact that CMCT-related studies have been performed ([Figure 2\)](#page-2-0).

# **Materials and methods**

# *Chemicals and reagents*

QLNC-3A6 Di-maleate standard (Lot No.25271003GS, purity 99.0%) was supplied by Qilu Animal Health Products Co., Ltd. (China). Toceranib phosphate standard (CAS 874819-74-6, purity 95.07%, internal standard, IS) was purchased from Shandong Link Bio-technology Co., Ltd (China). HPLC grade acetonitrile (ACN) and formic acid (FA) were purchased from Fisher Scientific (USA). Water was purified with Milli-Q water purification system (USA).

# *Configuration of stock solutions, working solutions, calibration standards and QC samples*

<span id="page-1-10"></span>Precisely weigh a certain amount of QLNC-3A6 Di-maleate and toceranib phosphate control, respectively, and dissolve them with 50% aqueous acetonitrile to obtain 1.0mg/mL of stock standard solution (Stock-STD), stock internal standard solution (Stock-IS) and stock control stock solution (Stock-QC), respectively, and store them at −20°C temperature.

Stock-STD and Stock-QC were diluted with 50% aqueous acetonitrile to obtain QLNC-3A6 series standard sample working solution (W-STD) at concentrations of 5, 20, 50, 100, 200, 500 and 1000ng/mL, QLNC-3A6 Di-maleate series standard sample working solution (W-STD) at concentrations of 10, 400 and 800ng/mL, respectively. QLNC-3A6 series quality control sample working solution (W-QC) and internal standard working solution (W-IS) at a concentration of 100ng/mL.

A precision volume of 90µL of canine blank plasma was taken, and 10µL of W-STD and 10µL of W-QC were added serially to obtain corrected standard samples at concentrations of 0.5, 2, 5, 10, 20, 50, and 100ng/mL, and quality control samples of LLOQ, LQC, MQC, and HQC at concentrations of 1, 40, and 80ng/mL, respectively.

### *UHPLC-MS/MS analysis*

An ultra performance liquid chromatography tandem mass spectrometer (UHPLC-MS/MS, Agilent-6470-1260) was used for this analytical method. Chromatographic



<span id="page-1-0"></span>**[Figure 1.](#page-1-1)** Molecular structural formula of (a) QLNC-3A6 Di-maleate and (b) sunitinib malate.



Chemical Formula: C<sub>22</sub>H<sub>28</sub>FN<sub>4</sub>O<sub>6</sub>P Molecular Weight: 494.46

<span id="page-2-0"></span>

separation was performed on a Phenomenex Kinetex C18 column (50 $\times$ 2.1 mm, 2.6 µm) at a flow rate of 0.4mL/min and a temperature of 40°C. The samples were kept at 8°C in an autosampler. The mobile phase consisted of 0.1% formic acid aqueous solution (A) and 0.1% formic acid acetonitrile (B). The optimization gradient was as follows: 10% B, 0–0.5min; 10–50% B, 0.5–3.0min; 50–90% B, 3.0–3.3min; 90% B, 3.3–5.0min; 90–10% B, 5.0–5.3min; 10% B, 5.3–6.0min.

Drug quantification was performed by multiple reaction monitoring (MRM) in positive ion electrospray (ESI) mode. The optimized ESI source parameters were as follows: drying gas temperature: 320°C; sheath gas temperature: 320°C; nebulizing gas pressure: 45 psi; capillary voltage: 3500V; and sheath gas flow rate: 11L/min. The MRM precursor-to-product transitions for QLNC-3A6 and TOC were *m/z* 426.2>113.1\*, *m/z* 426.2>70.2 and *m/z* 397.3>283.3\*. (*Note*: \*is a quantitative ion)

# *Animals*

Six healthy male and female Beagles weighing 8.50– 11.50kg and aged 1.5–2years were purchased from the Laboratory Animal Center of China Agricultural University. All animals were acclimatized in an environmentally controlled rearing room (temperature 18–25°C, humidity 30–60%, light and dark cycle every 12h) for 7days with free access to food and water. Before the experiment, the canines were fasted for 12h but had free access to water. Animal experiments were conducted in accordance with the Guidelines for the Breeding and Use of Laboratory Animals and were approved by the Animal Breeding and Use Committee of China Agricultural University (NO.11359-18-E-001) (Beijing, China).

### *Sample preparation*

Frozen plasma samples were thawed at room temperature for about 20min, vortexed, 100μL of accurately aspirated plasma samples were transferred to Eppendorf centrifuge tubes, 10μL of W-IS was added accurately, vortexed, 350μL of ACN was added to precipitate the proteins, and 250μL of purified water was added and mixed for 5min after vortexing, and then centrifuged for 10min at 4°C with relative centrifugal force (RCF)  $11180 \times g$ , 10,000 rpm. The supernatant was passed through 0.22μm organic microporous membrane, vortexed and mixed well, and 3μL of the supernatant was used for UHPLC-MS/MS analysis.

### *Method validation*

Method validation was performed in accordance with the bioanalytical method validation guidelines of the U.S. Food and Drug Administration (FDA), the European Medicines Agency (EMA 2024), and the Ministry of Agriculture and Rural Affairs of the PRC (EMA 2024; FDA 2024; Veterinary Drug Evaluation Center MoAaRD 2024). At the same time, the International Council for Harmonization of Technical Requirements for Pharmaceuticals for Human Use (ICH) M10 Guideline "Validation of Bioanalytical Methods and Analysis of Research Samples" was used as reference.

#### *Selectivity*

To ensure that the established analytical method can be used to differentiate QLNC-3A6 and IS from endogenous components in the matrix or other components in the sample, selectivity should be demonstrated using a suitable blank matrix from at least 6 canines. The response of the interfering component should be less than 20% of the lower limit of quantitation response of the analyte and less than 5% of the internal standard response.

#### *Carry-over*

Residues are evaluated by determining the Upper Limit of Quantification (ULOQ) standard sample followed by a double blank (DBK). Residues of QLNC-3A6 should not exceed 20% of the Lower Limit of Quantification (LLOQ) and not exceed 5% of the IS.

#### *Standard curve and LLOQ*

After the calibration standard sample was processed, the best-fit curve was generated based on the measured peak area of QLNC-3A6, and linear regression was performed using the least squares method with the weights selected to be  $1/x^2$ , and at least 6 standard curves were selected for method validation. The calibration standards should be back-calculated to within ±15% of the labeled value, the LLOQ shall be within ±20%, and at least 75% of the calibration standards should contain a minimum of 6 valid concentrations, and the signal-to-noise ratio (S/N) of the LLOQ should be at least 10 times higher than that of the blank sample.

#### *Precision and accuracy*

Intra- and inter-batch precision and accuracy were analyzed within one day and on three consecutive days using LLOQ and QC samples (0.5, 1, 40 and 80ng/mL), respectively. Precision was expressed as the relative standard deviation (RSD) (coefficient of variation, CV) of the measured values and was calculated as the average of standard deviation/average estimated value  $\times$  100. Accuracy is expressed in terms of deviation and is calculated as: (measured value/true value) ×100%. For intra-batch precision/ accuracy, both require at least 4 concentrations from one analytical batch, with at least 5 samples per concentration; for inter-batch precision/accuracy, both require at least 4 concentrations from 3 analytical batches (at least 2days), with at least 5 samples per concentration. The CV shall not exceed 15% for QC samples within/between batches and 20% for LLOQ; intra/inter-batch accuracy shall be within ±15% for QC samples and ±20% for LLOQ.

# *Matrix effect*

HQC and LQC matrix effect samples of QLNC-3A6 were prepared using canine blank plasma from 6 batches of different sources. The peak areas of QLNC-3A6 and IS in the matrix were determined after treatment and compared with the corresponding peak areas of pure solution samples without matrix at the corresponding concentrations after treatment. The matrix factors (MF) of QLNC-3A6 and IS were calculated, and then the IS normalized matrix factor (ISnorm-MF) was calculated by dividing the MF of QLNC-3A6 by the MF of IS. The CV of the ISnorm-MF calculated from 6 batches of matrix should not be greater than 15%.

# *Recovery*

Three concentrations of QC samples, LQC (1ng/mL), MQC (40ng/mL) and HQC (80ng/mL), six parallels for each concentration, were processed and injected to determine the peak areas of QLNC-3A6 and IS, and the recoveries were calculated by comparing with the corresponding peak areas of the recovery samples injected for determination of the recoveries of QLNC-3A6 and IS, respectively.

# *Dilution reliability*

Canine plasma samples containing QLNC-3A6 were diluted 2-fold and 10-fold with blank canine plasma, respectively, and analytes need to be above the upper limit of quantification (LOQ) concentration, with a minimum of 5 determinations required for each dilution factor. The accuracy and precision of the dilution shall be within  $\pm 15\%$ .

# *Stability*

The stability of QLNC-3A6 was examined with QC samples that had been left at room temperature (25 °C) for 24 h after treatment, QC samples that had been left under an 8°C injector for 24h after treatment, QC samples that had been treated with

plasma assayed after three times freeze and thaw, QC samples that had been treated and frozen at −80 °C for 30 days after treatment, and QC samples that had been stored at −20 °C in a standard stock solution for 60 days. The stability of the QC samples were examined using a calibration curve that was based on the standard curve. Standard curves were obtained from freshly prepared calibration standards, and the QC samples were analyzed according to the standard curves, comparing the measured concentrations with the labelled concentrations, and the deviation of the mean value of each concentration from the labeled concentration shall be within ±15%.

# *Application to PK studies*

A single oral dose of QLNC-3A6 Di-maleate tablets at a dose of 3mg/kg BW was administered to canine after an overnight fast. A blood sample of 2mL was collected in sodium heparin anticoagulant tubes from the brachiocephalic vein of each canine before and 0.5, 1, 2, 4, 6, 8, 12, 24, 36, 48, and 72h after the administration of QLNC-3A6 Di-maleate tablets. Plasma samples were centrifuged with RCF 1788 $\times$ g at 4,000rpm for 10min and the supernatants were separated immediately and stored at −80°C until analysis and detection.

Data acquisition and automatic integration were performed using the Aglient MassHunter Workstation Data Acquisition System software configured for the mass spectrometer; the generated data were analyzed and calculated using the non-compartmental model in WinNonlin 6.4 PK software. The main plasma PK parameters included: elimination rate constant ( $\lambda$ z), elimination phase half-life ( $T_{1/2\lambda z}$ ), peak concentration ( $C_{\text{max}}$ ), time to peak ( $T_{\text{max}}$ ), area under the drug-time curve from 0 to the last detection time point  $(AUC<sub>last</sub>)$ , area under the drug-time curve from 0 to extrapolated infinity (AUC<sub>INF</sub>\_obs), area under the drug-time curve extrapolated as a percentage of the total area under the drug-time curve (AUC\_%Extrap\_obs), and mean retention time  $(MRT<sub>last</sub>)$ .

# **Results**

# *Method optimization*

# *UHPLC-MS/MS mass spectrometry method optimization*

<span id="page-3-0"></span>Firstly, MRM was selected as the detection method in this study to minimize the interference of endogenous molecules in biological matrices. (Tang et al. [2022](#page-10-14)) Secondly, the positive and negative ion detection modes were compared, and it was found that the signal response of the positive ion mode was significantly better than that of the negative ion mode. Subsequently, several columns were evaluated, including BEH-C18 (100 $\times$ 2.1 mm, 1.7 µm), BEH-HILIC (100×2.1mm, 1.7µm), HSS-T3 (150×2.1mm,

1.78µm), ZORBAX Eclipse Plus C18 (50×2.1mm, 1.8µm) and Phenomenex Kinetex C18 (50×2.1mm,  $2.6 \mu m$ ), the results showed that the Phenomenex Kinetex C18 (50×2.1mm, 2.6µm) column had highest response values ([Table 1](#page-4-0)).

<span id="page-4-3"></span>Finally, the pH and buffer of the mobile phase system were optimized. Since both drugs involved in this experiment are alkaline, the pH of the configured buffer can be determined directly from the pKa of the analyte, and then the total buffer correction factor can be calculated based on the correction factor of the alkaline analyte when the organic phase is acetonitrile, and different buffer types compatible with LC-MS systems can be selected according to the total correction factor. Among them, 0.1% formic acid produces a mobile phase with a pH of about 2.7, which has become the preferred buffer substance for LC-MS compatibility class under low pH conditions, and higher concentrations of formic acid can improve the response value, but not too high; ammonium acetate produces a mobile phase with a pH of 3.8–5.8; and although aqueous ammonia is a alkaline solution, for reversed-phase liquid chromatographic separations to retain alkaline compounds, a high pH mobile phase is usually selected to enhance reverse retention, but again, high ammonia concentrations should not be chosen because too high concentration of ammonia can inhibit the positive mode response. The results showed that when the mobile phase consisted of 0.1% formic acid aqueous solution  $(0.1\%FA\cdot H, O)$  (A) and 0.1% formic acid acetonitrile (0.1%FA•ACN) (B), the peak shapes of the substances to be measured were the best and the response was the highest ([Table 2](#page-4-1)).

<span id="page-4-4"></span><span id="page-4-0"></span>

	QLNC-3A6		
		$(426.2 > 113.1)$ TOC $(397.3 > 283.3)$	
	(area)	(area)	
BEH-C18 (100 $\times$ 2.1 mm, 1.7 µm)	8,432	10,542	
BEH-HILIC $(100 \times 2.1$ mm, $1.7$ µm)	2,346	7,500	
HSS-T3 $(150 \times 2.1$ mm, $1.78$ µm)	5,749	11,356	
ZORBAX Eclipse Plus C18	6.642	10,013	
$(50 \times 2.1 \text{ mm}, 1.8 \text{ }\mu\text{m})$			
Phenomenex Kinetex C18	9,912	13,836	
$(50 \times 2.1$ mm, $2.6$ µm)			

<span id="page-4-1"></span>**[Table 2.](#page-4-4)** Results of response value of different mobile phase system pH and buffers.



# *Optimization of sample preparation method*

<span id="page-4-8"></span>For sample preparation, biomarkers in blood are usually at low concentrations and can be masked by high levels of proteins, of which albumin is one of the most prominent, and ACN or MeOH solution (Tomascova et al. [2019\)](#page-10-15) is often utilized to remove albumin from plasma. In this study, different reagents used to precipitate proteins were compared and the results showed that the best results were obtained when 350 µL ACN was used to precipitate 100 µL of plasma with good recoveries ([Table 3](#page-4-2)).

<span id="page-4-5"></span>Therefore, 350µL ACN was used to precipitate 100µL plasma for sample preparation, and a Phenomenex Kinetex C18 (50×2.1mm, 2.6µm) column was selected for the detection, with  $0.1\%$  FA $\cdot$ H<sub>2</sub>O (A) and 0.1%FA•ACN (B) as the mobile phases, and the elution was run for 6 mins at a column temperature of 40°C according to the optimal gradient. The final chromatogram with good separation and low background noise could be obtained. [Figure 3](#page-5-0) shows the mass spectra of QLNC-3A6 and TOC from precursor ions to product ions.

#### <span id="page-4-6"></span>*Method validation*

#### *Selectivity*

The results showed that the peak area of the interfering peak at the retention time of QLNC-3A6 in blank plasma was less than 20% of the corresponding peak area of LLOQ, and the peak area of the interfering peak at the retention time of IS was less than 5% of the peak area of the IS of LLOQ, which was in accordance with the regulations, and the details are shown in [Figure 4](#page-6-0).

#### <span id="page-4-7"></span>*Carry-over*

After analytical determination, the peak areas of analyte QLNC-3A6 in the DBK samples after ULOQ were all less than 20% of the corresponding peak areas of LLOQ, and the peak areas of IS did not exceed 5% of the corresponding peak areas of LLOQ.

### *Calibration curves and LLOQ*

A calibration standard curve containing at least six points was constructed by plotting the peak area ratio (QLNC-3A6/TOC, Y) versus the nominal

<span id="page-4-2"></span>**[Table 3.](#page-4-5)** Results of response values of plasma proteins precipitated by different reagents.

	OLNC-3A6 (426.2 > 113.1) (area)	TOC (397.3 > 283.3) (area)	Recovery (%)
100µL Plasma + 300 µL ACN	5,474	11,729	59.48
100µL Plasma + 350 µL ACN	11,729	15,079	103.27
100µL Plasma + 400 µL ACN	14,751	13,546	133.38
100µL Plasma + 300 µL $0.1\%$ FA $\cdot$ ACN	17,035	15,643	141.02
100µL Plasma + 300 µL MeOH	4,932	9,058	70.64



<span id="page-5-0"></span>**[Figure 3.](#page-4-6)** Mass spectra of (a) QLNC-3A6 and (b) TOC.

concentration of QLNC-3A6 (X) over the range of 0.5– 100ng/mL. The representative regression calibration curve was *y*=0.580138x+0.013172 with a weighting factor of  $1/x^2$ . In each analytical run, good linearity was observed in the regression calibration curve  $(R<sup>2</sup>>0.99)$  with %CV < 15. The concentration of LLOQ was 0.5ng/mL, which showed sufficient precision and accuracy, and the detailed data are shown in [Table 4.](#page-6-1) Meanwhile, the S/N of LLOQ was greater than 10, which met the requirements, and the details are shown in [Figure 5](#page-7-0).

#### <span id="page-5-2"></span><span id="page-5-1"></span>*Precision and accuracy*

<span id="page-5-3"></span>[Table 5](#page-7-1) summarizes the intraday and interday precision and accuracy values. It is noteworthy that the intraday and interday %CV were 2.06–13.57% and 6.90–9.14%, respectively. Intraday and interday %DEV were −10.73 to 9.54% and −3.86 to 0.70% respectively. The results indicate that the analytical method is accurate, reproducible and meets the requirements.

### *Matrix effect*

In this experiment, the matrix had a certain inhibition or enhancement effect on the ionization of the analytes, but this effect was not significant (*p*>0.05), and had little effect on the measurement results. Moreover, the average value of normalized matrix factor of 6 LQC samples of QLNC-3A6 was 95.94%, and the %CV was 4.54%; the average value of normalized matrix factor of 6 HQC samples was 114.81%, and the %CV was 0.85%. The average %CV of the 12 matrix samples at low and high concentrations was 9.78%, which met the relevant requirements. Therefore, the presence of matrix had no effect on the determination of analyte QLNC-3A6.

#### *Recovery*

The mean recoveries of analyte QLNC-3A6 in low, medium and high concentration QC samples were 99.72, 102.72 and 102.48%, respectively; and the corresponding mean recoveries of IS were 99.81, 100.89 and 99.71%, respectively. The %CV of analyte and IS



<span id="page-6-0"></span>**[Figure 4.](#page-4-7)** (a) Mass spectra of QLNC-3A6 and TOC in blank canine plasma and (b) plasma spiked to the LLOQ level.

<span id="page-6-1"></span>**[Table 4.](#page-5-1)** Precision and accuracy of the calibration curve back-calculated concentrations and LLOQs (*n*=6).

Concentration(ng/mL)	%DEV	%CV
0.5	0.63	$-0.43$
2	2.57	0.37
5	3.88	2.34
10	4.59	0.02
20	3.84	2.63
50	2.49	$-2.32$
100	5.75	$-4.51$

for 18 QC samples at 3 concentrations ranged from 0.90 to 2.64% and 0.33 to 4.72%, respectively. The recoveries of analytes and IS at low, medium and high concentrations were not much different and basically the same, which satisfied the test requirements.

### *Dilution integrity*

<span id="page-6-2"></span>The results of dilution integrity are shown in [Table 6.](#page-7-2) The samples (150ng/mL) was analyzed with its 2-fold and 10-fold dilutions. The accuracy of analysis of diluted samples was −0.07 to 0.04% with %CV of 3.77 and 2.33%, respectively. The results showed that diluting the plasma samples up to 10-fold did not affect the accurate determination of QLNC-3A6.

### *Stability*

<span id="page-6-3"></span>[Table 7](#page-7-3) lists the stability results for QLNC-3A6 in canine plasma. Stability is defined as the accuracy and precision relative to freshly prepared calibrated plasma samples. It can be seen that the accuracy was -10.92 to 5.05% with %CV < 7.76% under five conditions. The results showed that QLNC-3A6 was stable in the samples that were left at room temperature (about 25°C) for 24h after treatment, in the samples that were left under the injector at 8°C for 24h after treatment, in the samples that were treated and determined by plasma after three freeze-thawing, and in the samples that were treated by freezing the samples for 30days at −80°C. In addition, appropriately diluted standard stock solutions were analyzed, and no significant degradation was observed after 60days of storage at −20°C.

### *PK study*

<span id="page-6-5"></span><span id="page-6-4"></span>The validated UHPLC-MS/MS analytical method was successfully applied to a preclinical PK study after a single oral dose of QLNC-3A6 Di-maleate tablets in canine at 3mg/kg BW. [Figures 6](#page-8-0) and [7](#page-8-1) show the



<span id="page-7-0"></span>**[Figure 5.](#page-5-2)** Signal-to-noise chromatogram of LLOQ.

<span id="page-7-1"></span>



*Note*: There were data deviations in the Day1 LLOQ, so the data was eliminated from processing.

<span id="page-7-2"></span>



<span id="page-7-3"></span>



<span id="page-8-3"></span>mean concentration-time curve and different animals' concentration-time curve of QLNC-3A6 in canine plasma under the Linear Trapezoidal Linear Interpolation (LTLI) calculation. [Table 8](#page-8-2) lists the main PK parameters calculated using the non-compartmental model. It can be seen from the results that QLNC-3A6 showed fast absorption rate, rapid distribution and slow metabolic elimination in canine plasma. The  $\lambda z$ ,  $T_{1/2\lambda z}$ ,  $C_{\text{max}}$ ,  $T_{\text{max}}$  and  $AUC_{\text{last}}$  were  $0.07 \pm 0.01/h$ , 11.00±2.57h, 50.88±31.94ng/mL, 9.08±11.57h and 836.48±230.53ng h/mL, respectively.

#### **Discussion**

From the concentration-time curves of QLNC-3A6 in different canine, it can be found that the curves in multiple animals showed the phenomenon of irregular double or multiple peaks.

According to the relevant literature, oral drugs are susceptible to bimodal absorption phenomenon during gastric emptying to small intestinal absorption



<span id="page-8-0"></span>**[Figure 6.](#page-6-4)** Plasma mean concentration-time curve of QLNC-3A6 di-maleate tablets after a single oral dose of 3 mg/kg BW in canine  $(n=6)$ .

<span id="page-8-15"></span><span id="page-8-13"></span><span id="page-8-10"></span><span id="page-8-8"></span>process (Mirfazaelian and Mahmoudian [2006](#page-10-16)) such as ranitidine (Yin et al. [2003\)](#page-10-17), phenazopyridine (Shang et al. [2005](#page-10-18)), talinolol, paracetamol (Weitschies et al. [2005\)](#page-10-19), and toceranib (Yancey, Merritt, White, et al. [2010\)](#page-10-1). There are many reasons for the bimodal or multimodal phenomenon for such drugs, such as delayed gastric emptying (Weitschies et al. [2005](#page-10-20); Metsugi et al. [2008;](#page-10-21) Ogungbenro et al. [2015](#page-10-22)), enterohepatic recirculation (EHC) (Marier et al. [2002](#page-10-23); Soulele and Karalis [2019](#page-10-24)), and uneven distribution of the drug efflux transporter protein, P-gp, in the small intestinal (Godfrey et al. [2011;](#page-9-8) Wada et al. [2013](#page-10-25)).

<span id="page-8-12"></span><span id="page-8-11"></span><span id="page-8-9"></span><span id="page-8-7"></span><span id="page-8-6"></span><span id="page-8-5"></span>For the phenomenon of sub-peak, non-linear PK or slow drug metabolism after the absorption time in the small intestine, it may be due to the individual liver metabolizing the drug to reach the saturation of the flip-flop phenomenon. In turn, the occurrence of flip-flop kinetics phenomenon is related to many factors such as the weak acid ionization equilibrium constant  $K_a$  of the drug, elimination rate constant  $K_{el}$  (Yáñez et al. [2011](#page-10-26)) and intestinal transporter substrate (Garrison et al. [2015](#page-9-9)). According to the literature, the presence of flip-flop

<span id="page-8-14"></span><span id="page-8-4"></span><span id="page-8-2"></span>**[Table 8.](#page-8-3)** PK parameters of QLNC-3A6 di-maleate tablets after a single oral dose of 3mg/kg BW in canine (*n*=6).

Parameters	Unit	$Mean \pm SD$
λz	$h^{-1}$	$0.07 \pm 0.01$
$T_{1/2\lambda z}$	h	$11.00 \pm 2.57$
$C_{\text{max}}$	ng $mL^{-1}$	$50.88 \pm 31.94$
$T_{\rm max}$		$9.08 \pm 11.57$
AUC <sub>last</sub>	ng h $mL^{-1}$	$836.48 \pm 230.53$
$\mathcal{AUC}_{\mathsf{INF\_obs}}$	ng h $mL^{-1}$	$851.74 \pm 230.17$
AUC_%Extrap_obs	%	$1.96 \pm 1.20$
$MRT_{last}$	h	$18.63 \pm 4.47$

 $λ_z$ : elimination rate constant;  $T_{1/2λz}$ : elimination phase half-life;  $C_{\text{max}}$ : peak concentration;  $T_{\text{max}}$ : peak time;  $AUC_{\text{last}}$ : area under the drug-time curve from 0 to the last detected time point;  $AUC_{\text{INF}}$  obs, area under the drug-time curve from 0 to extrapolation to infinity;  $AUC_{\text{MExp\_obs}}$ extrapolated drug-time curve area under the drug-time curve as a percentage of the total area under the drug-time curve percent; *MRT*<sub>last</sub>, mean retention time.



<span id="page-8-1"></span>**[Figure 7.](#page-6-5)** Plasma concentration-time curves in different animals following a single oral dose of QLNC-3A6 Di-maleate tablets at 3mg/kg BW in canine. A close-up view is shown for the first 36h post drug administration.

<span id="page-9-11"></span>phenomenon may also be accompanied by EHC (Gupta and Gupta [2016](#page-9-10)).

Based on the current data, it is initially hypothesized that QLNC-3A6 is more variable in canine plasma, and that the double or multiple peaks phenomenon in the drug-time graphs is most likely a result of the EHC, the gastrointestinal tract having multiple absorption points, and the flip-flop. There is also the phenomenon that the generated curve masks more peaks due to the lack of sampling time points.

Therefore, the sampling time points should be rationally adjusted in the following in-depth study of QLNC-3A6 in canine, and related studies such as drug excretion should be carried out.

### **Conclusion**

In this study, a rapid, highly sensitive and selective UHPLC-MS/MS analytical method was developed for the first time for the quantitative detection of QLNC-3A6 in canine plasma. The specificity and sensitivity, linearity and calibration curve, precision and accuracy, dilution integrity and stability of the method meet the requirements of the guidelines of the FDA, EMA and Ministry of Agriculture and Rural Affairs of the PRC. The developed method has significant advantages in terms of simple sample preparation, rapid chromatographic separation and high throughput efficiency. This new approach was also utilized in the present study to investigate the PK of QLNC-3A6 Di-maleate tablets after a single oral dose of 3mg/kg BW in canine. The results indicated that QLNC-3A6 showed fast absorption rate, rapid distribution and slow metabolic elimination in canine plasma, which is valuable for future exploration of the clinical PK of QLNC-3A6.

### **Ethical approval**

Animal experiments were conducted in accordance with the Guidelines for the Breeding and Use of Laboratory Animals and were approved by the Animal Breeding and Use Committee of China Agricultural University (NO.13303-24-E-001) (Beijing, China).

### **Author contributions**

Corresponding author S.C., Y.L., and Y.W. participated in study design, conceptualization and project management. Qilu Animal Health Products Co., Ltd. provided the QLNC-3A6 Di-maleate. S.C., Z.W. and J.M. assisted in sample collection and clinical observation. M.J., Y.Y., Y.Z. participated in data collection and organization. S.C., Y.L., M.J.,and X.C. responsible for outcome monitoring and evaluation. S.C., M.K., G.C. and X.C. involved in writing the original manuscript, surveys and study execution. All authors have read and agreed to the published version of the manuscript.

### **Disclosure statement**

No potential conflict of interest was reported by the author(s).

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### **Data availability statement**

The data supporting the findings of this study are available from Dr. Sumeng Chen (26769385[20@qq.com](mailto:20@qq.com)) upon request.

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