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8	Evidence of Rapid Transfer and Bioaccumulation of Microcystin-LR Poses
9	Potential Risk to Freshwater Prawn Macrobrachium rosenbergii (de Man)
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26	Abstract
27	Microcystins accumulate in aquatic organisms and can be transferred to higher trophic
28	levels, eventually affecting vector animals and consumers. We examined three levels of
29	an aquatic food chain (Microcystis aeruginosa, Daphnia magna, and Macrobrachium
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rosenbergii) to identify the transfer efficiency and risk of microcystin on prawns. Samples were analyzed using UPLC-MS/MS and microcystin-LR (MC-LR) distributions in prawn tissues were studied. The results showed that prawns accumulate MC-LR both directly from M. aeruginosa and indirectly through D. magna which was pre-exposed to M. aeruginosa. MC-LR was detected in the gills, digestive tracts, and hepatopancreas of the prawns 2 h after exposure. MC-LR accumulated in prawns to 0.49±0.04 µg g⁻¹ dry weight in hepatopancreas within 24 h, while it was not detected in muscle samples, and rarely appeared in blood samples in such a short period. Although MC-LR was not detected in muscle, the head including hepatopancreas of the prawns accumulated troublesome amounts of MC-LR. These results demonstrate that microcystis blooms in prawn farming potentially pose a risk to human consumers, even though prawns may be exposed to the bloom for a very short time, hence regular monitoring of blue green algae population is recommended.

Introduction

Toxic cyanobacterial blooms are globally problematic as they may produce secondary metabolites such as microcystins (MC) (Vareli et al. 2012; Beaver et al. 2014). So far, more than 80 variants have been isolated and identified (Dietrich and Hoeger, 2005). Among these isoforms, Microcystin-LR (MC-LR) is the most common and most toxic to mammals. Studies have demonstrated that MCs can cause toxic effect on various aquatic organisms (Cazenave et al. 2005; Chen & Xie, 2005; Xie et al. 2005, 2007; Lance et al. 2006; Deblois et al. 2008; Ortiz-Rodríguez and Wiegand 2010; Deblois et al. 2011; Liu et al. 2011; Ziková et al. 2013; Liu et al. 2014). Aquaculture species are exposed to microcystins through the aquatic food web (Andersen et al. 1993; Ibelings et al. 2005; Smith and Haney 2006). Trophic transfer has also been demonstrated under laboratory conditions in which hepatotoxins were transferred from zooplankton to fish (Engström-Öst et al. 2002; Karjalainen et al. 2005; Smith and Haney 2006).

MC-LR also poses potential threats to human health (Codd et al. 2005a, b; Ibelings and Chorus 2007). The transfer of MC to higher trophic levels through the food web poses serious health implications (Ozawa et al. 2003). Recently, microcystins were

identified for the first time in the serum (average 0.228 ng mL⁻¹) of fishermen who were chronically exposed to cyanotoxins in Lake Chaohu, China, indicating hepatocellular damage (Chen et al. 2009). An important exposure route for humans is through the ingestion of contaminated aquatic food. The World Health Organization (WHO) published a guideline of 1 µg kg⁻¹ of MC in drinking water, and established a tolerable daily intake guideline (TDI) of 0.04 µg kg⁻¹ per day (Chorus and Bartram 1999).

In recent years, studies evaluating MC contamination in aquatic organisms from natural waters with cyanobacterial blooms have increased, but these studies mainly focused on fishes (Mohamed et al. 2003; Ibelings et al. 2005; Xie et al. 2005; Chen et al. 2006; Deblois et al. 2008; Wilson et al. 2008). Similar information is relatively rare for freshwater prawns. Freshwater prawns are commercially important because they are used for human consumption worldwide. In both intensive and extensive aquaculture, phytoplankton succession mimics natural systems, with cyanobacterial abundance reaching its maximum in summer (Smith et al. 2008).

Zooplankton can accumulate MC and therefore may act as vectors of the toxin up the aquatic food web; however, information on the transfer and bioaccumulation efficiency of MC through prawns is lacking. It is likely that consumption of prawns exposed to high MC levels could lead to risk of public health. Quantitative evaluation of the potential risk posed to food safety is needed. Therefore, we monitored transfer and bioaccumulation of MC-LR at three levels of an aquatic food chain, namely blue green alga *Microcystis aeruginosa*, daphnia *Daphnia magna* and freshwater prawns *Macrobrachium rosenbergii* in three experiments so as to assess its potential risk by human consumption of prawns.

Materials and methods

Chemicals and reagents

MC-LR (purity≥95.0%) was purchased from Alexis (Lausen, Switzerland). Standard stock solution was prepared in methanol at 0.50 μg mL⁻¹. HPLC-grade methanol and formic acid were supplied by Merck KGaA (Darmstadt, Germany). All other reagents and chemicals used were of analytical grade. Water was purified from a

Milli-Q deionization unit (Millipore, Molsheim, France).

M. aeruginosa was cultured in BG-11 medium in flasks at 25±1 °C, with irradiance at 56 μEm⁻² S⁻¹ and a 12 h light:12 h dark photoperiod. *D. magna* was provided by Key Laboratory of Marine Ecology and Environmental Science, Institute of Oceanology, Academy of Science (Qingdao City, China).

Experimental design

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Experiment A was an exposure experiment where prawns and D.magna were exposed to M. aeruginosa water. Prawns with an average weight of 12.92±2.79 g and body length of 8.36±0.57 cm were obtained from a local farm (Shanghai, China), acclimated for 1 week in PVC tanks (150 L) containing de-chlorinated water and fed with commercial prawn feed(Tongwei Group Co.Ltd, China) at a rate of 3% of body weight per day. The water temperature was 25±1 °C, and dissolved oxygen was 5.9 ± 0.6 mg L^{-1} . Fresh cells of M. aeruginosa were used for this experiment, after being concentrated by centrifugation (4 min, 7000 rpm, 4 °C) and resuspended in water at cell densities of 5×10⁷ cells mL⁻¹. D.magna at density of 2 individual mL⁻¹was maintained in 5 glass tanks each with 10 L in volume containing water with M. aeruginosa at density of 5×10^7 cells mL⁻¹ for 24 h. Each group of prawns (4 groups, n=10) was placed in a 30 L PVC tanks containing water with M. aeruginosa at the density of 5×10^7 cells mL⁻¹ for 24 h. During exposure the prawns were fed with commercial feed and aeration was provided. Prawns were sampled from tanks and dissected at 2, 8, 16 and 24 h of exposure. The gills, blood, digestive tract, muscle and hepatopancreas were frozen and freeze-dried separately. D. magna was collected and frozen for MC-LR content analysis and feeding experiment B.

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Experiment B was a feeding experiment where prawns were fed with D. magna which was exposed to M. aeruginosa in experiment A. The prawns (average weight of 13.87 ± 2.52 g and body length of 8.65 ± 0.47 cm) were acclimated to laboratory conditions for 1 week in PVC tanks with de-chlorinated water. During acclimation period commercial feed was supplied, while when the experiment began prawns were exclusively fed with frozen D. magna exposed to M. aeruginosa. Prawns were sampled and dissected at 8, 16 and 24 h post feeding and the gills, blood, digestive tract, muscle

and hepatopancreas were frozen and freeze-dried separately.

Experiment C was a food chain experiment where prawns (average weight of 13.25±2.85 g and body length of 8.43±0.86 cm) were acclimated to laboratory conditions for 1 week in 5 PVC tanks. Prawns along with *D. magna* (2 individuals mL⁻¹) were stocked in 30 L PVC tanks with *M. aeruginosa* at cell densities of 5×10⁷ cells mL⁻¹. Prawns from each tank were dissected at 2, 8, 16 and 24 h and the gills, blood, digestive tract, muscle and hepatopancreas were frozen and freeze-dried separately.

MC analyses

Extraction of the microcystins was carried out according to the method of Zhang et al. (2009) with some modification. The lyophilized samples (~0.1 g dry weight for each sample) were extracted two times with 25 mL of 0.01 M EDTA-Na₂-5 v/v % formic acid by homogenization with a high-speed blender (PT2000, Polytron, Switzerland) for 30 s at 20,000 rpm and followed by 5 min sonication. The samples were then centrifuged at 4500 rpm (Type 3K-18, Sigma, Germany) at 4 °C. The supernatant was applied to an Oasis HLB cartridge (500 mg, 6 mL, Waters, Milford, MA, USA) which had been preconditioned by washing with 5 mL 100% methanol and 10 mL distilled water. The column containing sample was washed with 5 mL water followed by 3 mL 10% methanol, and then eluted with 4 mL 100% methanol. The elutriant was dried under a nitrogen stream in a 45 °C water bath, and the residue was dissolved in 1.5 mL of 20% methanol with 0.1% formic acid. The resulting solution was filtered through a 0.45 μm nylon filter for analysis.

An Ultra Performance Liquid Chromatography (UPLC) and Mass Spectrometry system was used consisting of an Acquity UPLCTM system equipped with a XEVO triple quadrupole mass spectrometer (Waters, Milford, USA). The injection volume was full-loop (10 μ L) and the chromatographic separation was performed at 40 °C using a C₁₈ column (Waters, 100 mm×2.1 mm internal diameter, 1. μ m particle size), with the flow rate set at 0.35 mL min⁻¹. The mass spectrometer was operated in the positive ESI mode, with a capillary voltage of 3.5 kV. The source and desolvation temperatures were 145 and 450 °C, respectively. Gas desolvation and nebulization were carried out using

nitrogen at flow rates of 850 and 50 L h⁻¹, respectively. The signal acquisition was performed by multiple reactions monitoring mode (MRM). The divert valve was programmed to send the LC flow to waste for the first 2 min after injection and again after the analyte of interest had eluted. The gradient parameters are presented in Table 1. Mass spectrum tuning and optimization were achieved by infusing MC-LR and monitoring the [M+H]⁺ ion at m/z 995.6. The product ions at m/z 107.1 from the parent ion at m/z 995.6 were used for MC-LR qualitative analysis. For quantification purposes, mass chromatograms monitored the product ions at m/z 135.2 from the parent ion at m/z 995.6 (Table 2).

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Table 1

Table 2

The method was validated in terms of specificity, limit of detection (LOD), limit of quantification (LOQ), precision and recovery. Linearity of the method was assessed at MC-LR concentration ranging from 1 to 250 ng mL⁻¹. Five calibration curves with six concentration points were constructed. The linearity of the calibration curves was evaluated on the basis of linear regression analysis and the square correlation coefficients (r²) using SPSS 17.0. A correlation coefficient above 0.99 was desirable for all the calibration curves. LOD and LOQ were defined as concentrations in a sample resulting in signal-to-noise ratios of 3 and 10, respectively. Precision of the assay was expressed by percent relative standard deviation (%RSD).

The recoveries were obtained by analyzing MC-LR in prawn tissue at three spiked concentration (0.075, 1.5, 3.75 µg g⁻¹ for MC-LR). In all cases, samples were run in sextuplicate. The limit of detection (LOD) and limit of quantification (LOQ) were estimated sextuplicate. The matrix effect was assessed by comparing the peak areas of the neat analyte standards, standards spiked before and after extraction in prawn tissue samples. Mean and standard deviation of MC concentrations between treatments were performed using SPSS 17.0 for windows.

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Results

MC analyses

One possible way to remove the matrix interferences from prawn tissue is sample cleanup using an Oasis HLB extraction cartridge (Waters, Milford, USA). There are many interfering compositions unclear in complex prawn tissue, which ideally should be washed off before elution. We successfully used 3 mL of 10% methanol in water to wash off the interference of the matrix while leaving the targeted compounds on the cartridge. The washing curve is shown in Fig.1. Pure methanol was chosen for elution because of its high recovery and convenient dryness by nitrogen flush. The volume fraction of pure methanol in water was 4 mL and the elution curve is shown in Fig.2.

Figure 1

Figure 2

According to the analysis of 10 unexposed samples (including gills, blood, digestive tract, muscle and hepatopancreas), this UPLC-MS/MS method provided clean and background-free mass traces for the matrix studied, demonstrating that the method had good selectivity (Fig.3). Quantification matrix-fortified calibration curves were determined to compensate for the matrix effect and loss in sample preparation. Good linearity was obtained for MC-LR, with r^2 ranging from 0.9946 to 0.9990. The LOD and LOQ in this method were 0.0075 $\mu g g^{-1}$ and 0.01575 $\mu g g^{-1}$, respectively. The results are summarized in Table 3.

Figure 3

206 Table 3

Satisfactory MC-LR recoveries were obtained ranging between 84.8±7.6% and 128.4±11.3%. The precision was satisfactory since RSD of the mean recovery ranged from 4.2% to 10.9%. The results demonstrated that the accuracy and repeatability of the method were good for quantitative purposes.

Experimental results

MC-LR in the cells of *M. aeruginosa* was $119.67\pm14.01~\mu g~g^{-1}$ dry weight and in *D. magna* $2.72\pm0.09~\mu g~g^{-1}$ dry weight, respectively. The highest MC-LR peaks in the gills of prawns appeared at 8 h ($6.08\pm0.59~\mu g~g^{-1}$ dry weight) in experiment A. The digestive tracts of prawns showed remarkably high peaks at $16~h~(2.19\pm0.30~\mu g~g^{-1}$ dry weight), whereas the highest MC-LR content in the hepatopancreas appeared at $24~h~(0.49\pm0.04~\mu g~g^{-1}$ dry weight) in experiment C (Fig.4). MC-LR was not detected in the muscles, and rarely appeared in blood samples in this study (Table 4). Cyanobacteria (*M. aeruginosa*)

were observed in the gills of prawns in both experiments A and C.

Figure 4

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In all treated prawns, the average MC-LR content in the hepatopancreas rapidly increased during the experimental period (Figs. 4, 5). The highest MC-LR levels were recorded in the gills and digestive tracts of the prawns (Fig. 6). The average MC-LR content of experiment A in the gills and digestive tracts increased initially (around 8 h), then decreased until the end of the test (Fig. 4). However, the average MC-LR content in the gills and digestive tracts in experiment C increased initially then decreased after16 h. Prawns accumulated a maximum of 6.08±0.59 µg g⁻¹ dry weight in the gills in experiment A.

Figure 5

Figure 6

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Discussion

Aquatic organisms are generally considered more tolerant of cyanobacteria toxins than mammals as a result of their co-evolutionary history, which can reduce the likelihood for catastrophic losses of the cultured species but increase the potential for damaging human exposure to these toxins (Smith et al. 2008). Bioaccumulation of MCs in crustaceans has been reported in several publications. Vasconcelos et al (2001) detected a concentration of 2.9 µg g⁻¹ dry weight in crayfish *Procambarus clarkii* under laboratory conditions. The seasonal changes of mixed hepatotoxins (MC-LR, MC-LA, MC-RR, MC-YR, and NODLN) in the marine prawn Penaeus monodon was measured through the enzyme linked immunosorbent assays (ELISA) method and the total concentration of hepatotoxins in P. monodon hepatopancreas varied between 0.0064 and 0.0816 µg g⁻¹ DW (Kankaanpää et al. 2005). Zhang et al. (2009) studied seasonal variations of MC-LR contents in shrimp (Macrobrachium nipponensis) from Lake Taihu through liquid chromatography electrospray ionization mass spectrum (LC-ESI-MS), and found that the MC-LR concentrations ranged from 0.031±0.004 to 0.605±0.179 μg g^{-1} DW with averaging 0.244 \pm 0.220 μ g g^{-1} DW. In our study, the toxin concentration shown by freshwater prawns is consistent with the reported cases in the hepatopancreas

(maximum 0.49 μg g⁻¹ DW). Nevertheless, Zimba et al (2006) reported a case of cooccurrence of white shrimp, *Litopenaeus vannamei*, mortalities and microcystin toxin in ponds, where water samples from the affected pond contained high levels of microcystin-LR (45 μg g⁻¹), and free microcystin-LR concentrations in dead shrimp hepatopancreas determined by HPLC were 55 μg g⁻¹ total shrimp weight. However, we did not determine the total MC load in prawn. The common results from the above references demonstrated that crustacean muscle was not the primary organ for hepatotoxin bioaccumulation.

In this study, MC-LR was detected in the gills, digestive tracts, and hepatopancreas of the prawns after 2 h of exposure. The prawns accumulated MC-LR both directly from *M. aeruginosa* (Experiment A) and indirectly through *D. magna* (Experiment B) which was preexposed to *M. aeruginosa*. To our best knowledge, this might be the first evidence that freshwater prawns accumulate MC-LR in very short time, as well as the direct transfer of MC-LR from zooplankton to prawns and the subsequent accumulation of toxin in the hepatopancreas and digestive tract.

According to our results, MC appears to be absorbed through the digestive tract of prawns at a higher rate when the toxin is administered through a vector, *D. magna*, rather than through toxic cyanobacteria directly. In experiment C, as prawns and *D. magna* were exposed to *M. aeruginosa* together, *D. magna* ingested toxic cyanobacteria cells and prawns appeared to consume these toxic *D. magna*, based on the decrease of MC-LR content in prawn gills and the increase of MC-LR concentrations in hepatopancreas and digestive tracts. Thus, other aquaculture species may also be exposed to microcystins through the ingestion of cyanobacteria (Ortiz-Rodríguez and Wiegand 2010), consumption of contaminated food items (e.g., prey or detritus) (Deblois et al. 2011), and absorption of dissolved microcystins from the water column (e.g., after leakage from cells or cell lysis) (Vasconcelos et al. 2001).

Microcystins are usually taken up from intestine and transferred to liver tissue (Fischer and Dietrick 2000). In this study, we found that prawns ingested *M. aeruginosa* cells directly. We also observed that algal cells adhered to the gills of prawns. There was

no evidence that the gills absorbed microcystin through algae lysis, nevertheless, the possibility of this pathway should not be disregarded.

The UPLC-MS/MS analysis method used herein shortened analysis time significantly, with peak time for MC-LR appearing at 4.5-4.8 min. Comparatively, it took 18.51 minutes to analyze MC-LR content in *Litopenaeus vannamei* samples using HPLC/MS method (Zimba et al. 2006), while similar retention time (~21min) was recorded for two tilapia species *Oreochromis niloticus* and *Tilapia rendalli* using HPLC method (Deblois et al 2008).

The 24-h test was not sufficient to promote accumulation of MC-LR in the muscle, which is the edible part of the prawn for human consumption. Future studies should take this into consideration. The provisional tolerable daily intake (TDI) suggested by WHO is 0.04 µg kg⁻¹ of body weight or 2.4 µg for an adult human weighing 60 kg (Kuiper-Goodman et al. 1999). Assuming an adult human ingests 100 g of whole prawns per day from our experiment A and C, then daily uptake of MC-LR would range from 116 µg to 706 µg, much higher than TDI value proposed by WHO, which was unsafe for human consumption. Therefore, the risk of consuming prawns from aquaculture ponds and lakes during toxic cyanobacteria blooms cannot be overlooked, and regular monitoring of MC levels in prawns should be conducted to protect health of consumers. We recommend that future studies evaluate the total MC tissue load and the potentially harmful effects of MC on human health by multiple exposure routes through aquatic food.

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440	Table 1. UPLC Gradient parameters
441	Table 2. UPLC-MS/MS acquisition parameter for MC-LR analysis
442	Table 3. The calibration curve of MC-LR and recovery tests of HPLC-MS/MS method
443	(n=6)
444	Table 4. MC-LR content accumulated in different prawn tissues within 24h in the three
445	experiments (n=10) ^a
446	Figure 1. The loss rate of MC-LR in different volume fractions of methanol in water.

Figure 2. Curve of elution rate of MC-LR against volume of pure methanol.

Figure 3. Chromatograms of the MC-LR in the samples (a. chemical standard MC-LR at 100 ng/g; b. the blank hepatopancreas sample spiked at 100 ng/g MC-LR and

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real hepatopancreas samples; c. the blank digestive tract samples spiked at 100
ng/g MC-LR and real digestive tract samples (up to down); d. the blank muscle
sample spiked at 100 ng/g MC-LR and real muscle samples (up to down); e. the
blank gills sample spiked at 100 ng/g MC-LR and real gills samples (up to
down); f. the blank blood sample spiked at 100 ng/g MC-LR and real blood
samples (up to down)).
Figure 4. MC-LR contents in gills, hepatopancreas and digestive tract of prawns from
experiments A and C. Bars represent means \pm SD of four replicates.
Figure 5 MC I D concentrations in honotone and discative treat of many from
Figure 5. MC-LR concentrations in hepatopancreas and digestive tract of prawns from
experiment B. Bars represent means \pm SD of four replicates.
Figure 6. Distribution of MC-LR (%) in the prawn tissues at different times in
experiments A, B and C.

Table 1. UPLC Gradient parameters

Time (min)	A ^a	B^{b}	Curve
0.00	100	0	0
5.00	0	100	6
6.00	0	100	6
6.10	100	0	1
7.60	100	0	1

^a water containing 0.1% (v/v) formic acid, ^b methanol containing 0.1% (v/v) formic acid.

Table 2. UPLC-MS/MS acquisition parameter for MC-LR analysis

Parent ion/Product ion	Cone voltage(v)	Collision energy(v)	Dwell time(s)
995.6/107.1 ^a	42	70	0.2
995.6/135.2	42	76	0.2

^aquantification transition

Table 3. The calibration curve of MC-LR and recovery tests of HPLC-MS/MS method (n=6)

Tissue _	Standard curve	r^2	MC-LR	Recovery (%)	RSD
Tissue	Standard curve		$(\mu g/g)$	Recovery (70)	(%)
Blood	Y = 259.83X - 2.53	0.9990	0.075	91.7±5.2	5.7
	'		1.5	99.2±10.5	10.5
			3.75	103.7 ± 4.3	4.2
Muscle	Y = 262.45X - 2.71	0.9975	0.075	91.5±10.5	10.9
			1.5	84.8±7.6	9.0
			3.75	111.4±6.5	5.8
Gills	Y = 301.00X - 2.83	0.9952	0.075	125.2±12.2	9.7
			1.5	99.7±4.7	4.7
			3.75	100.7 ± 5.0	5.0
Hepatopancreas	Y = 451.70X - 2.84	0.9968	0.075	104.3±4.8	4.6
α			1.5	88.6±8.6	9.8
CO			3.75	99.2±5.8	5.8
Digestive tract	Y = 652.27X - 2.72	0.9946	0.075	128.4±11.3	8.8
			1.5	96.2±6.1	6.3

Table 4 MC-LR content accumulated in different prawn tissues within 24h in the three experiments (n=10) ^a

Exp	eriment/Time	MC-LR content (μg/g dry weight)								
(h)		Blood	Muscle	Gills	Hepatopancreas	Digestive				
		Diood	Widscie	Oms	Tiepatopanereas	tract				
A	2	ND	ND	2.16±0.19 ^a	0.02 ± 0.00^{a}	0.35 ± 0.02^{a}				
	8	0.15 ± 0.004	ND	6.08 ± 0.59^{b}	0.06 ± 0.01^{a}	0.77 ± 0.07^{b}				
	16	ND	ND	3.57 ± 0.44^{a}	0.19 ± 0.03^{b}	0.69 ± 0.08^{b}				
	24	ND	ND	0.89 ± 0.08^{c}	0.20 ± 0.04^{b}	0.56 ± 0.02^{ab}				
В	8	ND	ND	0.05±0.005	0.04 ± 0.02^{a}	0.05 ± 0.01^{a}				
	16	ND	ND	ND	0.05 ± 0.01^{a}	0.14 ± 0.01^{b}				
	24	0.02 ± 0.005	ND	ND	0.10 ± 0.02^{b}	0.40 ± 0.04^{c}				
C	2	ND	ND	0.37 ± 0.04^{a}	0.17 ± 0.02^{a}	0.62 ± 0.16^{a}				
	8	ND	ND	1.59 ± 0.09^{b}	0.18 ± 0.01^{a}	1.34 ± 0.08^{b}				
	16	ND	ND	1.66 ± 0.18^{b}	0.40 ± 0.04^{b}	2.19 ± 0.30^{c}				
	24	0.03±0.005	ND	1.11 ± 0.12^{b}	0.49 ± 0.04^{b}	1.29 ± 0.11^{b}				

^a Values represent means \pm SD of four replicates. Means of each experiment in the same column with different superscript letters are significantly different (P<0.05).

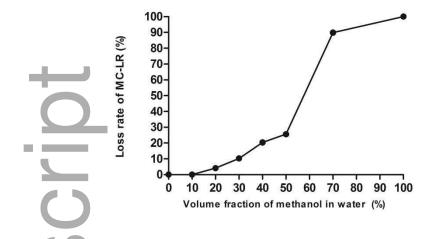


Figure 1. The loss rate of MC-LR in different volume fractions of methanol in water.

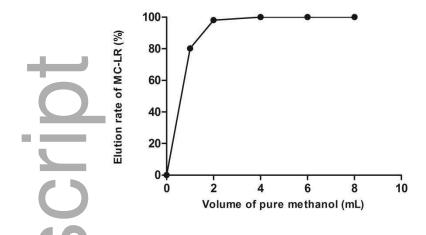


Figure 2. Curve of elution rate of MC-LR against volume of pure methanol.

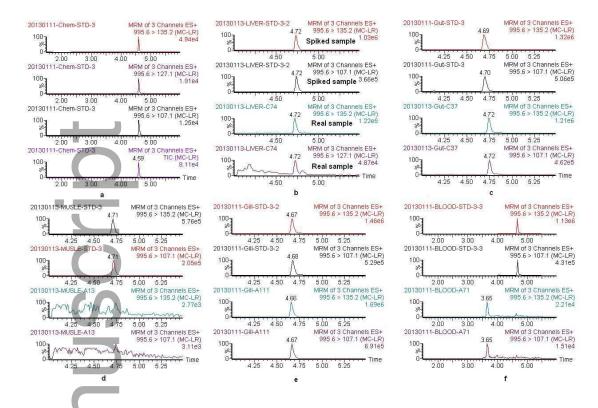


Figure 3. Chromatograms of the MC-LR in the samples (a. chemical standard MC-LR at 100 ng/g; b. the blank hepatopancreas sample spiked at 100 ng/g MC-LR and real hepatopancreas samples; c. the blank digestive tract samples spiked at 100 ng/g MC-LR and real digestive tract samples (up to down); d. the blank muscle sample spiked at 100 ng/g MC-LR and real muscle samples (up to down); e. the blank gills sample spiked at 100 ng/g MC-LR and real gills samples (up to down); f. the blank blood sample spiked at 100 ng/g MC-LR and real blood samples (up to down)).

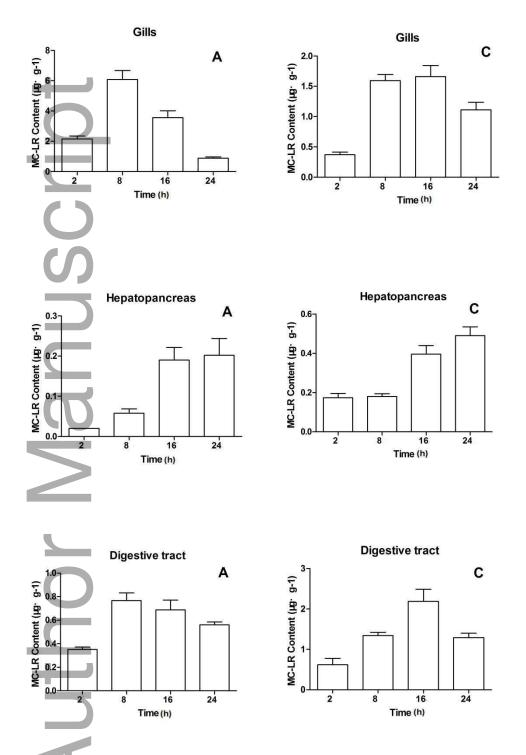
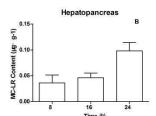


Figure 4. MC-LR contents in gills, hepatopancreas and digestive tract of prawns from experiments A and C. Bars represent means \pm SD of four replicates.



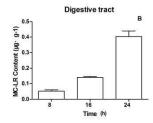


Figure 5. MC-LR concentrations in hepatopancreas and digestive tract of prawns from experiment B. Bars represent means \pm SD of four replicates.

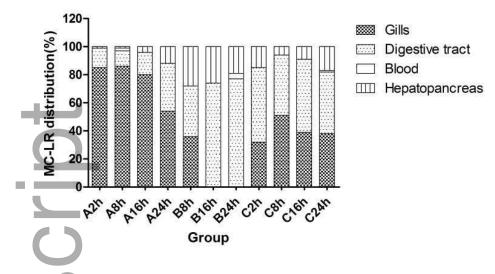


Figure 6. Distribution of MC-LR (%) in the prawn tissues at different times in experiments A, B and C.