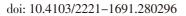


Short Communication

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Identification and investigation of Calodium hepaticum in rodents and insectivores from Wuhan section of the Yangtze River in China

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ABSTRACT

Objective: To investigate the prevalence of *Calodium hepaticum* (C. hepaticum) in rodents and insectivores from Wuhan section of the Yangtze River in China, and to provide evidence for the prevention and treatment of hepatic Calodium infection.

Methods: Rodents and insectivores were captured from three selected Yangtze River beaches using mousetraps. The three survey sites were divided into six areas according to natural conditions, with 60 mousetraps placed in each area. The liver lesions in the captured rodents were observed by the naked eye and the eggs in the liver tissue were observed by microscopic examination.

Results: A total of 1080 mousetraps were placed, and 1 075 mousetraps were retrieved, with the retrieve rate as 99.5%. A total of 101 Apodemus agrarius, 12 Rattus norvegicus, and 9 Crocidura attenuata were caught. The average density of rodents and insectivores was 10.5% and 0.8%, respectively. DNA of egg nodules from infected rodents showed 98% similarity with that of C. hepaticum 18S rRNA (LC425008.1). One Rattus norvegicus was infected with C. hepaticum, with an infection rate of 3.23% in the Ergi river beach; the other two beaches did not show the incidence of C. hepaticum.

Conclusions: The monitoring of C. hepaticum in the Yangtze River beaches should be strengthened to reduce the risk of human C. hepaticum infection.

KEYWORDS: Calodium hepaticum; Rodents; Insectivores; Yangtze River beach; PCR

1. Introduction

Calodium hepaticum (C. hepaticum) is a zoonotic nematode that is parasitic mainly on the liver of rodents. However, infections can spread to humans or other animals upon the ingestion of food or water contaminated with infective eggs, leading to hepatic Calodium infection[1]. The disease is easily transmitted and is a serious health

concern. In China, the infection rate was as high as 7.14%-76.83%[1,2]. In the current study, we investigated the infection status of C. hepaticum among rodents and insectivores in the Wuhan section of the Yangtze River in China and provide a basis for effective control.

2. Materials and methods

2.1. Basic overview of the survey point

Three Yangtze River beach areas were selected; the survey site was Wuhan wetland reserve, with a wide area of river beach and lush plants. The Jincheng and Laozihu river beaches were far from the residential area, and the distance was more than 1 km. The Erqi river beach was close to the residential area, and a small number of residents lived on the beach (Figure 1).

2.2. Survey method

The night trapping method was used to capture rodents and insectivores. The distance between mousetraps was about $10 \text{ m} \times 10$ m; a medium-sized mousetrap was used and peanuts were used as baits. The rat clips were placed according to the natural environment and rat habits and were placed at night and collected in the morning.

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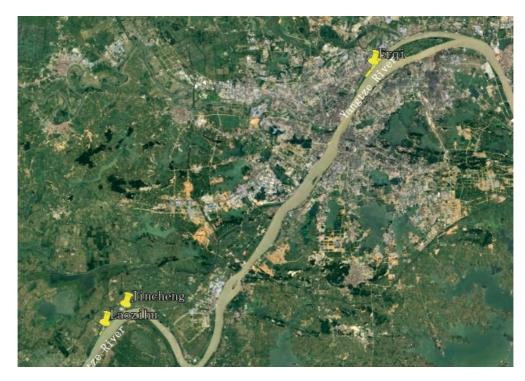


Figure 1. Distribution of the survey sites on the Yangtze River.

Each survey site was divided into six sections according to the natural conditions. In each survey area, 60 mousetraps were placed in the wild. The captured mammals were numbered and placed in white plastic bags, which were kept on ice packs and brought back to the laboratory for identification. The liver lesions were observed by the naked eye, and diseased liver tissues (nodule) were cut into slices. The infections in the liver tissue were observed by microscopic examination.

2.3. DNA extraction

Approximately 150 mg of diseased liver tissue was taken in a 2 mL centrifuge tube, ground for 2 min using RETSCH MM400 (Germany). Then, 1 mL double distilled water (ddH₂O) was added, and samples were filtered using a 100-mesh screen. The filtrate was incubated for 10 min and then the supernatant was removed. This process was repeated five times. Then, 1 mL ddH₂O was added and the samples were centrifuged at 1000 g for 5 min; this step was repeated twice. This was followed by the removal of supernatant and extraction of DNA using the OMEGA Tissue DNA Kit (OMEGA, USA). The extracted DNA was stored at -20 °C.

2.4. Molecular identification

Primer synthesis and sequencing were done by Sangon Biotech Co. Ltd. (Shanghai, China). The primer sequences of *C. hepaticum*[3] were as follows: F 5'-CCCTAGTTGCGACTTTAAACGA-3' and R 3'-TCCACCAACTAAGAACGGCC-5'. The PCR conditions were as follows: 94 °C for 3 min, followed by 40 cycles at 94 °C for 60 s, 50 °C for 60 s, 72 °C for 60 s, followed by extension step at 72 °C for 7 min. The sequencing results were analyzed using DNAstar 7.0 and

NCBI Blast.

2.5. Ethical statement

The study was approved by WHCDC Ethical Committee (WHCDCIRB-K-2016015) on 16 December, 2016 and followed the 3R's guideline.

3. Results

3.1. Distribution of rodents and insectivores

A total of 1080 mousetraps were placed at the three survey sites on the Yangtze River beach and 1075 mousetraps were retrieved (retrieve rate was 99.5%). As a result, 122 small mammals were caught, and these included 101 *Apodemus agrarius* (*A. agrarius*), 12 *Rattus norvegicus* (*R. norvegicus*), and 9 *Crocidura attenuata* (*C. attenuata*). Thus, we captured three species belonging to three genera, two families, and two orders (Table 1). The average density of rodents and insectivores was 10.5% and 0.8%, respectively.

3.2. Parasitic infections

Of the captured mammals, 77 *A. agrarius*, 9 *R. norvegicus*, 5 *C. attenuata* were captured in Jincheng and Laozihu; *C. hepaticum* was not found in these two sites. From Erqi, we captured 24 *A. agrarius*, 3 *R. norvegicus*, and 4 *C. attenuata*; the infection rate of *C. hepaticum* in these captured mammals was 3.23% (1/31) (Table 1).

Table 1. Distribution of rodents and insectivores in the Yangtze River beaches (n).

River beach position	Number of placed mousetraps	Number of recycled mousetraps	Apodemus agrarius	Rattus norvegicus	$Crocidura\ attenuata$	Total
Erqi	360	360	24	3	4	31
Jincheng	360	357	46	5	3	54
Laozihu	360	358	31	4	2	37
Total	1 080	1 075	101	12	9	122

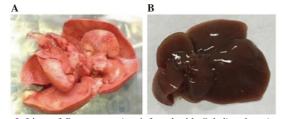


Figure 2. Liver of *Rattus norvegicus* infected with *Calodium hepaticum* (A) and normal liver of *Rattus norvegicus* (B).

3.3. Shape of the liver and egg of R. norvegicus infected with C. hepaticum

The liver of *R. norvegicus* infected with *C. hepaticum* was hard, and the liver was swollen with yellowish or white spots and filamentous or flaky lesions, which were prominent on the liver surface (Figure 2A). Microscopic examination showed that these eggs were colorless and shaped like a waist drum, and their average size was about 51 to 67 μ m in length and 30 to 35 μ m in width. The eggshell was thick and showed radial stripes. Transparent plugs were observed at both ends of the egg, which were not prominent outside the shell; a celllike structure could be seen inside (Figure 3A-C).

3.4. Molecular identification

Using PCR and *C. hepaticum*-specific primers, 288 bp target fragment was amplified for the molecular identification of *C. hepaticum* (Supplementary Figure 1). The target fragment, analyzed using DNAstar 7.0 and NCBI Blast, showed 98% similarity with the *C. hepaticum* 18S rRNA sequence (LC425008.1) (Figure 4).

4. Discussion

Hepatic *Calodium* infection is a parasitic zoonosis. More than 60 species of mammals are hosts of *C. hepaticum*, which causes severe

liver disease, leading to liver damage, liver function disorders, liver fibrosis, eosinophilia, and even death[1].

In this study, we investigated the prevalence of *C. hepaticum* at three Yangtze River beaches, from where 101 *A. agrarius*, 12 *R. norvegicus*, and 9 *C. attenuata* were caught; the average density was 11.3%. DNA extracted from the liver of these animals was subjected to PCR and the amplified target fragment sequence was analyzed using NCBI blast. As a result, we found 98% similarity with the *C. hepaticum* 18S rRNA sequence (LC425008.1), indicating that the infectious parasite could be *C. hepaticum*. Based on the shape of eggs and parasitic position in the infected animals, the infecting species was then confirmed to be *C. hepaticum*.

R. norvegicus seemed to be the main host species of *C. hepaticum*[4], as it was the only rodent infected in this study. This is consistent with other studies that reported a high infection rate in *R. norvegicus*, compared with that in other rodents.

The infection rate varies among the *R. norvegicus* living at different locations; the prevalence of *C. hepaticum* was higher in house *R. norvegicus* than in field *R. norvegicus*[1]. In this study, the *R. norvegicus* was caught from wild river beaches. The infection rate of *C. hepaticum* was 3.23%, which was lower than that reported in Henan Province[5] (13.5%), Dali City[2] (24.3%), and Wenzhou City[6] (18.3%), although the rodents in these cities were caught from indoor areas (such as slaughterhouses and factories).

Rodent is a major risk factor for the transmission of *C. hepaticum* to humans. In this study, we found no infected rodents in the two river beaches that were farther than 1 km away from where people lived; however, the *R. norvegicus* captured from the beach that was in close proximity to where humans were settled was infected with *C. hepaticum*. This is of concern, as it poses the risk of contamination of soil, environment and food, and a greater possibility of the infection transmission to humans.

Previously, 163 cases of humans infected with *C. hepaticum* were analyzed by Fuehrer[7], including 72 cases confirmed by liver biopsy, 13 cases confirmed by serology, 78 cases of spurious infection, and 72 cases of genuine infection with a high mortality rate of 61%. Since the

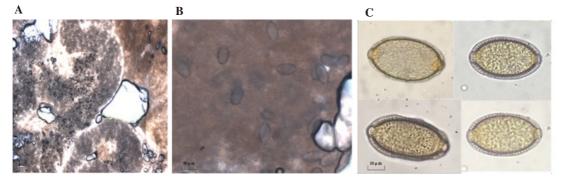


Figure 3. Result of microscopic examination. A: The egg nodule of *Calodium hepaticum*-infected liver $(100 \times)$; B: The eggs in *Calodium hepaticum*-infected liver tissue $(200 \times)$; C: The scattered eggs in *Calodium hepaticum*-infected liver tissue $(400 \times)$.

Score 494 bit	ts(267)	Expect) 1e-135	Identities 281/288(98%)	Gaps 0/288(0%)	Strand Plus/Plus
)uery	1	CCCTAGTTGCGACTTTA	AACGATGCCGACTAGCGATT	CGGCGACGTCTTTCTAA	TGACTC 60
Sbjct	1026	CCCTAGTTGCGACTTTA	AACGATGCCAACTAGCGATT	CGGCGATGTCTTTCTAA	TGACTC 1085
uery	61	GCTGGGCAGCTCCCGGG	AAACCAAAGTCTTTCGGTTC	CGGGGGGGAGTATGGTTG	CAAAGC 120
bjct	1086	GCTGAGCAGCTCCCGGG	AAACCAAAGTCTTTCGGTTC	CGGGGGGAAGTATGGTTG	CAAAGC 1145
uery	121	TGAAACTTAAAGGAATT	GACGGAAGGGCACCACCAGG	AGTGGAGCATGCGGCTT	AATTTG 180
bjct	1146	TGAAACTTAAAGGAATT	GACGGAAGGGCACCACCAGG	AGTGGAGCATGCGGCTT	AATTTG 1205
uery	181	ACTCAACACGGGAAACC	TCACCCGTCCCGAACACTGT	CAGGATTGACAGATTGA	TAGCTC 240
Sbjct	1206	ACTCAACACGGGAAAGC	TCACCCGTCCCGAACACTGT	CAGGATTGACAGATTGA	GAGCTC 1265
luery	241	TTTCTCGATTCAGTGGG	TAGTGGTGCATGGCCGTTCT	TAGTTGGTGGA 288	
Sbjct	1266	TTTCTTGATTCAGTGGG	TAGTGGTGCATGGCCGTTCT	TAGTTGGTGGA 1313	

Figure 4. Sequence alignment results: partial sequence of Calodium hepaticum NSMT.

eggs produced by adult worms are mainly deposited in the liver without being excreted, it is difficult to make a clinical diagnosis after human infection, and the final diagnosis must rely on a biopsy or autopsy. Moreover, the rate of misdiagnosis and missed diagnosis is high[8], with only a few human cases actually reported[9]. Hepatic *Calodium* infection should be considered when eosinophilia is found to increase clinically, the liver is enlarged and the patient complains of pain, and other parasitic infections are ruled out[10].

To summarize, we report the prevalence of *C. hepaticum* on the Yangtze River beaches. Our study indicates that the Yangtze River beach area could be an epidemic disease reservoir for hepatic *Calodium* infection. The beaches are wetland reserves, which are difficult to manage. Therefore, effective measures should be taken to exterminate the rodents, rodent carcasses, and their excrement should be disposed properly to avoid the environment pollution by infected eggs. Health education on the prevention and treatment of *C. hepaticum* infection should also be carried out to encourage residents to change their poor health and living habits so as to reduce the chances of infection.

Conflict of interest statement

We declare that there is no conflict of interest.

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Authors' contributions

ZSM and JXS conceived the study. ZSM, JXL and JXS undertook statistical analysis, and drafted the manuscript. WH and LHT undertook epidemiological investigations and sample collection. All authors contributed to the writing of the manuscript and approved the submitted version of the manuscript.

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