Cohabitation studies with common carp were conducted to determine whether the Cyprinid Herpes Virus 3 can infect and establish a productive infection in fish species that according to available data, are not susceptible to this virus. In order to examine if other fish species can contribute to further spreading of the virus, goldfish, silver carp, grass carp, prussian carp and tench were exposed to CyHV-3 through cohabitation with infected carp without clinical symptoms. After this period they cohabitated with naive carp for two weeks and were examined for CyHV-3 by PCR. Our results showed that CyHV-3 was present in the organs of these fish species and also in organs of naive carp after two weeks in cohabitation, suggesting that CyHV-3 may cause latent infection, and also that has a potential to infect a broader host range than it was believed before. Our study adds on better understanding of CyHV-3 transmission not just in its primary host, but also suggests the importance of common fish species in polyculture with carp in the epidemiology of CyHV-3.

Key words: carp polyculture, common carp, CyHV-3, koi herpes virus, KHV

INTRODUCTION

The disease of koi (Cyprinus carpio koi) and common carp (Cyprinus carpio carpio), caused by koi herpesvirus (KHV) has been observed since 1998 in many fish farms worldwide, causing mass mortality and significant financial losses. Herpes-like virus designated koi herpesvirus (KHV, CyHV-3) has been isolated from koi and common carp during outbreaks of the disease. The first occurrence of the disease with massive mortality has been reported in 1998 in Israel and USA, and since then, several countries have experienced outbreaks of the disease, including USA, South Africa, Japan, Israel and many European countries (Bretzinger et al., 1999; Hedrick et al., 2000; Neukirch and Kunz 2001; Perelberg et al., 2003; Chien et al., 2004; Sano et al., 2004; Tu et al., 2004). An important
The feature of herpesviruses is their ability to persist in their hosts, including those with natural or vaccine-induced immunity (Kucuktas and Brady, 1999). The virus remains dormant and inactive over a long period of time, but can be reactivated and become pathogenic and cause clinical symptoms and even death.

The host range and susceptibility of CyHV-3 is still not fully understood. Typical CyHV-3 clinical signs were described only in carp and koi carp. However, several lines of carp and their hybrids (e.g., Dor x Našice) have proved to be less susceptible to disease than others (Shapira et al., 2005). In addition, it was described that hybrids of golden carp and carp are partially resistant to CyHV-3 (Hedrick et al., 2006). Since the discovery of CyHV-3 infected fish in rivers and lakes in Japan (Haramoto et al., 2007; Ishioka et al., 2005), UK and USA (Grimmett et al., 2006), it has become very important to clarify the role of other fish species, cohabiting with common carp in the maintenance and transmission of CyHV-3.

**MATERIAL AND METHODS**

**Viruses and cells**

For all tests we used KHV-I isolated from a koi carp by Hedrick et al. (2000), generously provided by Dr SM Bergmann (Friedrich Loeffler Institute, Germany). The virus was propagated in CCB (Common carp brain) cell line, created by Neukirch et al. (1999), generously provided by Dr Olga Haenen (Central Veterinary Institute of Wageningen, The Netherlands). Cells were grown in minimal essential medium (MEM) with Earl salts and L-glutamine (Sigma), 10% fetal bovine serum (Gibco), 79.6 mg/L non-essential amino acids (PAA), 200 000 U/L penicillin and 200 mg/L streptomycin (PAA), incubated at 20°C.

**Animal experiments**

Fishes used in the experiment were obtained from a closed-system in the Center for Fishery and Applied Hydrobiology "Radmilovac" in Belgrade, with no history of KHV infection. The average weight of common carp (*Cyprinus carpio*), grass carp (*Ctenopharyngodon idella*), tench (*Tinca tinca*), goldfish (*Carassius auratus*), prussian carp (*Carassius auratus gibelio*) and silver carp (*Hypophthalmichthys molitrix*) was approximately 30 g.

In order to create an asymptomatic carrier state, the protocol described by Ronen et al. (2003) was used. In brief, 10 common carp were placed in the same tank with CyHV-3-infected koi carp showing symptoms of infection and kept at 23°C for 5 days, and thereon they were transferred to a new tank with a water temperature of 30°C and kept for 30 days. The surviving fish were kept at 23°C for 30 days, and used in the experiment three months after the initial cohabitation with sick fish. In order to examine if other fish species can contribute to further spreading of the virus, goldfish, silver carp, grass carp, prussian carp and tench were exposed to CyHV-3 through cohabitation with the asymptomatic carrier for a 14 and 21 day period. After this period they cohabitated with naive carp for two weeks and were examined for CyHV-3 by PCR.
Co-habitation procedure

Fifteen tench were in cohabitation with virus carriers for five days. After that period, they were transferred to another tank. On the 14th and 21th day after initial contact with the virus carrier, five tench fishes were euthanized and tested for CyHV-3. Fourteen and 21 days after initial contact with the carrier, five tenches were transferred to a new tank with five naive carp fishes, where they were in cohabitation for two weeks. After two weeks of cohabitation, tenches and carps were euthanized, the organs were sampled and tested for presence of CyHV-3. Before the experiment, and at the end, five tenches that were not exposed to the virus were euthanized and tested.

This procedure has been applied for testing of other fish species in experiment.

DNA extraction and PCR amplification

Portions (approximately 0.1 g) of the gill, kidney and spleen were sampled from individual fish and DNA was extracted using a DNeasy Qiagen kit, following the tissue extraction protocol (Qiagen).

The detection of KHV DNA from fish tissues was done using the oligonucleotide primer set TKf (5'-GGGTTACCTGTACGAG-3') and TKr 5'-CACCCAGTAGATTATGC-3') published in Bercovier et al. (2005), and the size of the amplified fragments was 409 bp. The Eppendorf Thermal Cycler was used for amplification using 1 cycle at 94°C for 5 min followed by 40 cycles of 1 min at 95°C, 1 min at 55°C, and 1 min at 72°C, followed by a final extension step of 10 minutes at 72°C. The inner pair of primers Intfw (5'-CGTCTGGAGGAATACGAG-3') and Intrev (5'-ACCGTACAGCTCGTACTGG-3') were used for nested PCR, and the size of the amplified fragments was 348 bp. The Eppendorf Thermal Cycler was used for amplification using 1 cycle of 94°C for 5 min followed by 40 cycles of 1 min at 95°C, 1 min at 55°C, and 1 min at 72°C, followed by a final extension step of 10 minutes at 72°C.

The PCR products were detected in 2% ethidium bromide stained agarose gel at 60V for 30 min and visualized under UV light.

RESULTS

Koi herpesvirus DNA was detected in three of 15 tench exposed to the virus. DNA was detected in one tench collected 14 days after the initial contact with carp virus carriers and in two tench fishes which cohabited from 14. to 28. day with five naive carp, of which in two carp the presence of CyHV-3 genome was detected (Figure 1). There were no clinical symptoms of the disease during the experiment, or any mortality.

In four of 15 goldfish in cohabitation with infected carp were detected parts of the CyHV-3 genome. DNA was detected in two goldfish sampled 14 days after the initial contact with carp virus carriers, in one goldfish which cohabited from 14. to 28. day with five carp, of which in one carp the presence of CyHV-3 DNA was detected, and in one goldfish which cohabited from 21st to 36th day with five carp, of which in one carp the presence of CyHV-3 DNA was detected. Other carp and
goldfish fishes did not show any symptoms of disease and they were CyHV-3 PCR negative.

Koi herpesvirus DNA was detected in two of 15 prussian carp that were exposed to the virus, in one prussian carp sampled 14 days after the initial contact with carp virus carrier, and in one prussian carp, which cohabited from 14. to 28. day with five carp, of which in one carp CyHV-3 DNA was detected. Other carp and prussian carp did not show any symptoms of disease and they were CyHV-3 PCR negative.

Part of the CyHV-3 genome was detected in one of 15 silver carp in cohabitation with the virus carrier. This silver carp cohabited from day 14 to day 28 with five naive carp, of which in one carp CyHV-3 DNA was detected. Other common carp and silver carp did not show any symptoms of disease and they were CyHV-3 PCR negative.

Koi herpesvirus DNA was detected in two of 15 grass carp that were exposed to the virus, in one grass carp sampled 14 days after the initial contact with the virus carrier, and in one which cohabited from the 14th to 28th day with five naive carp. In one of these carp CyHV-3 DNA was detected. Other common carp and grass carp did not show any symptoms of disease and they were CyHV-3 PCR negative.

**DISCUSSION**

The worldwide distribution of KHV within a few years of the first isolation (Bretzinger et al., 1999; Hedrick et al., 2000; Neukirch and Kunz 2001; Way et al., 2001; Perelberg et al., 2003; Sano et al., 2004; Tu et al., 2004) suggests that this virus may have an undetectable latent phase and, through the unregulated live fish trade, this may have contributed to its international spread.

It is known that many of herpesvirus infections of animals and fish can establish a latent infection (Fraser et al., 1981; Rock and Fraser 1983; van Nieuwstadt et al., 2001). The viral genome is integrated into the genome of specific host cells. These hidden infections can be activated by the action of stress or immunosuppression (Roizmann, 1996), which leads to the reactivation
of the virus. With improving diagnostic methods, evidence of the presence of KHV DNA in carp and koi carp without clinical symptoms were obtained (Gilad et al., 2004). Furthermore, Gilad et al., in 2004 showed that the detection of viral DNA is insufficient to establish whether a latent or persistent infection is present. Whether these fish can transmit virus upon contact with naïve fish or whether the virus can reactivate following some stressors proof is needed that these fish are true carriers. St-Hilaire et al., (2005) have provided the first evidence supporting this hypothesis when they detect reactivation of CyHV-3 in carp, several months after initial exposure.

In our study we confirmed that KHV may cause latent infections in KHV-infected fish and also that the koi herpesvirus transmission and epidemiology is more complex than it is thought before. We showed the potential of grass carp, prussian carp, goldfish and silver carp to interact with the virus, suggesting their importance in CyHV-3 transmission in nature. Furthermore, our in vivo study has demonstrated a strong potential of these species to serve as vector species for CyHV-3, without showing any apparent clinical signs for several weeks of the experiment. These findings contribute to better understanding of CyHV-3 infection in nature, indicating its interaction not just with its traditional hosts, but also with other carp species.

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Adress for correspondence:
Vladimir Radosavljević
Department of Fish Diseases
The Institute of Veterinary Medicine of Serbia
University of Belgrade
Vojvode Toze 14
11000 Belgrade
Serbia
E-mail: vladimiradosavljevic@gmail.com

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U eksperimentalnom istraživanju smo želeli da utvrdimo da li ciprinidni herpes virus 3 (CyHV-3) može izazvati infekciju kod vrsta riba koje, prema raspoloživim podacima, nisu prijemčive za ovaj virus. U cilju utvrđivanja da li ove vrste riba mogu doprineti širenju virusa, linjak, tolstolobik, amur, zlatni i srebrni karaš, su eksponirani CyHV-3 putem kohabitacije sa inficiranim šaranom bez kliničkih simptoma. Nakon toga, su kohabitirale sa zdravim šaranima tokom naredne dve nedelje, nakon čega je vršeno ispitivanje prisustva CyHV-3 pomoću PCR. Utvrđeno je prisustvo CyHV-3 u organima svih ispitivanih vrsta riba, kao i u organima šaranine tokom dvonedeljne kohabitacije sa njima. U našem istraživanju je potvrđeno da CyHV-3 može izazvati latentnu infekciju kod inficiranih riba, kao i da CyHV-3 ima više domaćina nego što se ranije mislilo. Naše istraživanje doprinosi boljem razumijevanju prenošenja CyHV-3 i ukazuje na značaj drugih vrsta riba koje se najčešće gaje u polikulturi sa šaranom u epizootiologiji CyHV-3.