Antibody response of channel catfish after channel catfish virus infection and following dexamethasone treatment

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ABSTRACT: Channel catfish virus (CCV, *Ictalurid herpesvirus 1*) and CCV disease have been extensively studied. Yet, little is known about CCV–host interaction after resolution of the primary infection. In order to determine potential recrudescence of CCV from latency, we established latency by exposing channel catfish juveniles with CCV or a thymidine kinase-negative recombinant (CCVlacZ) at a dose that caused less than 20% mortality. Then, we evaluated antibody response by serially sampling the same fish at 0 (pre-infection), 30, 60 and 90 d post challenge (DPC). We then attempted to induce viral recrudescence by intramuscular administration of dexamethasone and sampled the fish at 2, 4, 7, or 10 d post treatment. Recrudescence was evaluated by leukocyte co-cultivation and cell culture of tissue homogenates but no virus was detected. Western blot data demonstrated the highest number of seropositive fish by 30 DPC and a secondary antibody induction after dexamethasone treatment. The antigen specificity of the secondary response corresponded to viral proteins with molecular masses similar to those recognized by the same fish by 30 DPC. The recognized proteins were predominantly large, ranging from ~90 to >200 kDa. Expression analysis of selected virus genes at 90 DPC and following dexamethasone treatment demonstrated occasional immediate-early virus gene expression in peripheral blood leukocytes. Early and late gene expression was rarely detected. The combined data suggest restricted re-activation of CCV in our experimental system. Primary and secondary responses and virus gene expression were demonstrated in CCVlacZ-exposed fish but were less frequent than in CCV-exposed fish.

KEY WORDS: *Ictalurus punctatus* · Teleost · Ictalurid herpesvirus 1 · Alloherpesviridae · Herpesvirus latency · Thymidine kinase-negative recombinant · Viral gene expression · Neutralizing antibody

INTRODUCTION

Channel catfish virus (CCV), also known as *Ictalurid herpesvirus 1*, is the causative agent of a severe hemorrhagic disease frequently associated with high mortality in channel catfish *Ictalurus punctatus* fry and fingerlings (Fijan et al. 1970). Survivors of CCV primary infection become latently infected and constitute potential carriers for vertical and horizontal transmission. Herpesvirus latency is characterized by the absence of lytic viral proteins and productive virions on infected host cells. Neither the latency site nor latency-associated transcripts have yet been identified for CCV. However, CCV DNA has been detected in several tissues including muscle, fins, anterior and posterior kidneys, spleen, gonads, brain, intestine and peripheral blood leukocytes (PBL) from asymptomatic carriers (Wise & Boyle 1985, Gray et al. 1999).

Although CCV is phylogenetically distant from herpesviruses of mammals and birds, many of its biological properties are similar to alphaherpesviruses. Recently, the taxonomy of herpesvirus has been updated with the incorporation of the genus *Ictalurivirus* into the new family *Alloherpesviridae* (Davison et al. 2009). The initial productive phase of infection of the most extensively studied alphaherpesvirus, herpes sim-
plex type 1 (HSV-1, *Human herpesvirus 1*), is characterized by expression of immediate-early (IE), early (E), and late (L) classes of viral genes in an ordered lytic cascade that culminates in the replication and packaging of viral DNA into infectious particles, and eventual lysis of the infected cells (Honess & Roizman 1974). With respect to the kinetics of gene expression, Dixon & Farber (1980) observed similar cascade regulation of the synthesis of 3 distinct classes of CCV proteins.

Basic components of the morphology, biology, and genomic structure of CCV (Wolf & Darlington 1971, Plumb & Gaines 1975, Chousterman et al. 1979, Davidson 1992) have been characterized since this fish herpesvirus was first isolated by Fijan (1968). CCV encodes thymidine kinase (TK), which phosphorylates thymidine and other deoxynucleotides and analogs (Hanson & Thune 1993). For HSV-1, TK is not required for growth in cell culture but is essential for virulence (Jamieson et al. 1974, Field & Wildy 1978, Elstathieu et al. 1989) and reactivation from latency (Tenser & Dunstan 1979). A potential live recombinant vaccine vector with a deletion in the TK gene of CCV was able to induce protective immunity and a humoral immune response to the expressed foreign gene product (Zhang & Hanson 1995, 1996). However, to be a viable candidate as a live virus vaccine, it is necessary to characterize the ability of this virus to spread, to establish, and to reactivate from latency.

Little is known about the molecular process that controls CCV latency and reactivation nor the role played by the immune system in controlling reactivated infection. We do know latently infected fry are common in catfish hatcheries and in fingerling ponds. In a survey of major hatcheries, latent CCV infection was prevalent and the infection was assumed to be due to vertical transmission since no virus could be cultured from the populations and no CCV disease (CCVd) occurred (Thompson et al. 2005). Also primary exposure to CCV appears to more frequently result in a latent carrier state than a lethal form of disease. In field studies, population prevalence of CCV latency increased over time with no overt disease (Thompson et al. 2005) and in experimental challenges, exposure to low levels of virus resulted in disseminated virus replication but low mortality (Kancharla & Hanson 1996). Since reactivation is a complex phenomenon that is infrequent unless an appropriate stimulus is applied, evaluation of the expression of RNA necessary for activation of the lytic cascade and production of infectious particles should be more sensitive during the early events of reactivation than culture. In reactivation studies with herpesviruses of mammals, detection of virus gene expression and increases in virus DNA are often more sensitive indicators than virus isolation in cell culture (Baxi et al. 1996, Bevan et al. 1996).

The objectives of this study were to establish an experimental model to study latency and recrudescence of CCV in its natural host and to evaluate the host–pathogen interaction in this process. We were not successful in inducing overt recrudescence, but we found an increase in CCV DNA derived from caudal fin samples of latently infected channel catfish 7 d post dexamethasone (DEX) treatment without reactivation of culturable virus. Therefore, we investigated additional indicators of reactivation, namely a rise of anti-CCV antibody responses and the expression of selected viral genes in fish that had been exposed to CCV or a less virulent TK-negative recombinant of the virus and injected with DEX.

### MATERIALS AND METHODS

**Viruses, fish, and cell lysates.** CCV (strain Auburn 1 Clone A, American Type Culture Collection no. VR-665) and the TK-negative recombinant CCV containing the *lacZ* gene replacing both TK loci in the genome (CCVlacZ) (Zhang & Hanson 1995, 1996) were propagated in channel catfish ovary cells (CCO) as previously described (Hanson et al. 1994). Virus stocks were quantified in a plaque assay on CCO cells using the method of Buck & Loh (1985). Research fish were hatched and reared indoors, measured ~15 cm in length, and had no previous history of CCVd. Cell lysates for western blots were produced by infecting confluent brown bullhead (BB) cells with 0.5 plaque forming units (PFU) CCV cell⁻¹. The infected cell cultures were maintained in serum-free Dulbecco’s modified Eagle’s medium (DMEM, Sigma-Aldrich) supplemented with 25 mM HEPES, 200 IU ml⁻¹ penicillin, and 200 µg ml⁻¹ streptomycin (GIBCO BRL). After ~12 h of incubation at 30°C, CCV and mock-infected BB cell monolayers were harvested and centrifuged at 700 × g for 10 min at 4°C. Pellets were resuspended in phosphate-buffered saline (PBS) (8 mM Na₂HPO₄, 2 mM KH₂PO₄, 138 mM NaCl, 3 mM KCl, pH 7.4), aliquoted, and stored at −80°C.

**Experimental protocol.** All work with channel catfish was registered and done under the supervision of Mississippi State University Institutional Animal Care and Use Committee using approved protocols to assure humane handling and use. Fish were anesthetized by immersion exposure to tricaine methane sulfonate (MS222, Argent Chemical Laboratories) prior to injection, bleeding or caudal fin sampling.

**Evaluation of dexamethasone and heat stress as inducers of recrudescence.** To evaluate a method to induce consistent viral recrudescence, 60 channel catfish fingerlings, ~7 cm in length, from a CCV-positive spawn of latently infected broodfish (identified by
positive results using CCV-specific PCR), were distributed into 4 tanks (N = 15 fish per treatment group). To monitor the presence of infectious virus, posterior kidney homogenates of 10 additional fish that originated from the same rearing tank were cultured for virus. The caudal fins of all experimental fish were biopsied for CCV DNA analyses by PCR. Then 2 control treatment groups were injected intramuscularly with either PBS (control) or DEX (Dexaject, Phoenix Scientific) at 0.55 mg kg\(^{-1}\) body weight, and the water temperature was kept constant at 30°C. The other 2 groups received a PBS or DEX injection (0.55 mg kg\(^{-1}\) body weight), but the water temperature was increased to 36°C for 3 h and returned to 30°C. We used this dose of DEX because Bowser et al. (1985) had shown it to be effective in inducing CCV recrudescence. At 7 d post treatment (DPT), all fish were euthanized, posterior kidneys were cultured for infectious virus and caudal fins were sampled for CCV DNA analyses by PCR.

**Evaluation of experimental latent infections and DEX induced recrudescence.** A total of 110 CCV-negative channel catfish (mean weight = 82 g; the fish used were obtained from an isolated population with no history of CCVD, produced and cultured in specific pathogen-free conditions and CCV status was confirmed by PCR on caudal fin clips and PBL) was distributed into 22 tanks (N = 5 fish per tank). To identify each fish, we used a body mark system that consisted of a combination of caudal fin and barbel clips. Fish were mock-infected (controls, N = 20) or immersion challenged with 3 × 10\(^5\) PFU of CCV (N = 40) or CCVlacZ (N = 50) per 10 l tank for 30 min at 30°C. This dose and challenge method was used because it had previously been shown to result in a low level of mortality and a high percentage of infection (Hanson et al. 2004). Fish from each challenge tank were then transferred to a respective 80 l aerated, flow-through tank held at 30°C with 12 h light:12 h dark cycle and fed 32% commercial catfish feed ad libitum once daily. Fish were bled by caudal vein puncture. Serum samples were harvested at 0, 30, 60, and 90 DPC, heat inactivated at 45°C for 30 min and stored at −20°C until screened by western blot assays. At 90 DPC, a subgroup of 15 mock-infected, 33 CCVlacZ- and 23 CCV-exposed fish were treated with a single intramuscular injection of DEX at 0.55 mg kg\(^{-1}\) body weight. Control groups included 10 CCVlacZ- and 7 CCV-exposed fish that received a single intramuscular injection of PBS. Subsamples of these groups were taken at 2, 4, 7, and 10 DPT, and evaluated for CCV production, viral gene expression and specific antibody response against CCV antigens.

**Polymerase chain reaction.** PCR was performed and products were evaluated as previously described (Thompson et al. 2005) with the following modifications. Each PCR used 50 µl of reaction mixture, 5 ng of DNA template and cycling conditions of 1 cycle of 94°C for 10 min, 45 serial cycles of 93°C for 30 s, 60°C for 30 s, and 72°C for 1 min followed by 1 cycle of 72°C for 5 min. Tissue culture-grade water (Sigma-Aldrich) was used for the negative control and 5 pg of purified CCV DNA served as positive control. DNA sample concentrations were determined by a Gene Spec I (Hitachi Genetic Systems) spectrophotometer.

**Western blot analysis.** Mock- and CCV-infected BB cell lysates were diluted (vol/vol) in a modified Laemmli sample buffer (Bio-Rad Laboratories), disrupted by boiling for 5 min, and centrifuged at 18 000 × g for 3 min (Laemmli 1970). Electrophoresis was performed using 12% sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) Tris-HCl gels in a Mini-gel unit (Bio-Rad) using 1.5 mm thick spacers and preparative gel combs. Electrophoresis was carried out at 100 V for the stacking gel and 200 V for the SDS-PAGE for ~40 min. Prestained SDS-PAGE broad range standards (Bio-Rad) and ECL protein standards (Amersham Pharmacia Biotech) were used as molecular weight markers. Separated proteins were visualized by incubating gels for 1 h with agitation with GelCode® blue stain reagent (Pierce) and destaining with distilled water. For the western blot analysis (Towbin et al. 1979, Burnette 1981), proteins were electroblotted onto nitrocellulose membranes (0.2 µm, 7 × 8.4 cm, Bio-Rad) using a Mini trans-blot apparatus (Bio-Rad) for 1 h at 100 V. Blotted proteins were visualized by adding 0.5% Ponceau S solution (Sigma-Aldrich) in acetic acid and subsequently destained with PBS. Blots were soaked overnight at 4°C in blocking buffer containing 5% non-fat dry milk (Johnson et al. 1984) in Tris-buffered saline (TBS) (10 mM Tris-HCl, 150 mM NaCl, pH 7.4). Blots were transferred to a Mini-protean II multiscreen apparatus (Bio-Rad) and were incubated overnight at 4°C with 25 µl of fish serum samples diluted 1:50 in half-strength blocking buffer under agitation. Negative control serum was obtained from a broodfish that had previously tested negative for CCV DNA by PCR and had a negative titer in a serum neutralization assay. Positive control serum was obtained by bleeding a juvenile catfish 21 d post immunization with a single intraperitoneal injection of 1 × 10\(^5\) PFU of CCV. After washing 4 times with 0.2% Tween 20 (Bio-Rad) in TBS, blots were incubated at room temperature for 2 h with 8 ml of the hybridoma supernatant from mouse anti-channel catfish IgM monoclonal 9E1 (Lobb & Clem 1982, Lobb et al. 1984) diluted 1:4 in half-strength blocking buffer. Following another series of washing steps with 0.2% Tween 20 in TBS, blots were incubated at room temperature for 1 h with goat anti-mouse immunoglobulins conjugated to horseradish peroxidase (Southern Biotechnology Associates) diluted 1:1500 in half-strength blocking buffer. After 4
washing steps as described above and 2 rinses with TBS, labeling was detected using the ECL chemiluminescence kit (Amersham) and hyperfilm ECL (Amersham) as described by the manufacturer.

**Serum neutralization index test (SNI).** The SNI was performed similarly to the protocol described by Crawford et al. (1999) with few modifications. Heat-inactivated fish serum samples collected at 60 DPC and after DEX treatment were diluted 1:25 in serum-free DMEM supplemented with 25 mM HEPES, 200 IU ml⁻¹ penicillin, and 200 µg ml⁻¹ streptomycin (diluent medium). The stock of CCV was serially diluted 10-fold with diluent medium, and 200 µl of each virus dilution was distributed into 7 replicates of 96-well microtiter plates (Corning, Fisher Scientific). Approximately 200 µl of diluted fish serum was added to each of 7 replicate wells containing respective virus dilution. Control wells received virus dilutions and diluent medium.

After incubation for 1 h at 30°C, 50 µl of CCO cell suspension containing ~3 × 10⁶ cells were added to each microtiter plate well. The plates were incubated for 5 d at 30°C and observed daily for plaque formation. The virus titers were determined on the basis of the 50% tissue culture infectious dose reduction point (TCID₅₀) and calculated by the method of Reed & Muench (1938). The neutralization index produced by each fish serum sample was expressed as the difference in log₁₀(TCID₅₀ ml⁻¹) between the virus dilutions incubated with diluent medium (control wells) and virus dilutions in the presence of fish serum samples. SNI values by treatment were examined using an unbalanced 1-way analysis of variance (α = 0.05) (SAS 1990). In addition, to test the null hypothesis of no differences between the sample means (i.e. treatments) an unpaired t-test (Steel & Torrie 1980) with a test of approximation for unequal variances was used. The Type I error probability was set at α = 0.05. Comparisons of SNI values to western blot results were made using Spearman’s correlation analyses.

**Peripheral blood leukocyte isolation and co-cultivation with CCO cells.** PBL were isolated from 150 µl of whole blood containing EDTA, overlaid onto 500 µl of Histopaque 1077 (Sigma-Aldrich), and centrifuged at 500 × g for 10 min at room temperature. The buffy coat layer containing PBL was collected, suspended in Hanks’ balanced salt solution, calcium-magnesium free (HBSS-CMF) (GIBCO BRL), at osmolarity 272 (pH 7.3), and centrifuged at 500 × g for 5 min. The pellet was resuspended in 500 µl of HBSS-CMF. Total DNA was extracted from 100 µl of PBL suspension and analyzed by PCR. The remaining 400 µl of PBL suspension were cultured in separate wells of a 6-well tissue culture plate containing 70% confluent CCO cell monolayers. Tissue cultures were maintained at 30°C in DMEM (without sodium bicarbonate) supplemented with 10% fetal calf serum and 25 mM HEPES, and observed daily for 10 d for cytopathic effect (CPE) as previously described by Kancharla & Hanson (1996). They were then blind-passaged by transferring supernatant fluids to a new plate of CCO cells at 70% confluency. After observation for 10 d without CPE, each tissue culture was subsequently blind-passaged 2 consecutive times by trypsinizing and transferring 50% of the CCO cell suspension to a new 6-well tissue culture plate.

**Isolation of nucleic acids from PBL and caudal fin samples.** Caudal fin clips and PBL were chosen as a non-invasive sampling method for PCR and gene expression analyses in live fish. Approximately 50 to 100 µg of caudal fin tissue and 100 µl of PBL suspension were collected and transferred to RNAlater tissue storage reagent with RNA stabilization solution (Ambion) at –20°C. Total RNA was extracted using TRIzol isolation reagent (GIBCO BRL) following the manufacturer’s protocol. The air-dried RNA pellet was dissolved in 50 µl of DEPC (diethyl pyrocarbonate)-treated RNAse- and DNase-free water (Ambion) and analyzed by RT-PCR. During the evaluation of a method to induce consistent viral recrudescence, total DNA was isolated from caudal fin samples by the Puregene DNA isolation kit (Gentra Systems) according to the manufacturer’s protocol and analyzed by PCR.

**RT-PCR.** First-strand cDNA synthesis was performed with ~0.6 µg of total RNA sample in 10 µl reactions as described by Chen et al. (2002). Samples were diluted in 40 µl of DEPC RNAse- and DNase-free water and stored at –20°C. The PCR reaction was performed in a 50 µl volume containing 20 pmol of each primer, 2.5 mM of each deoxynucleoside triphosphate, 25 mM of MgCl₂ and 1.5 units of AmpliTaq Gold DNA polymerase (PE Biosystems) with 5 µl of cDNA template. Specific PCR primer pairs were designed based on available sequences (GenBank accession no. NC_001493; Davison 1992) for open reading frame (ORF) 3, ORF 12, ORF 57, and ORF 46 regions of CCV, and are listed in Table 1. Cycling reactions were performed using an Omn-E thermal cycler (HyBaid, Pegasus Scientific). After a single cycle of 10 min at 94°C, 35 amplification cycles were performed, each consisting of 30 s at 93°C, 30 s at 56°C, and 1 min at 72°C, followed by a final cycle of 5 min at 72°C. Negative controls included tissue culture water (Sigma-Aldrich) and samples processed without reverse transcriptase. Purified CCV DNA (5 pg) served as positive control. To monitor the expression of the cellular housekeeping gene β-actin in fish samples, PCR was performed using actin primers as described (Chen et al. 2002).

**Kinetics of CCV gene expression in vitro.** To evaluate RT-PCR for the expression of IE, E, and L CCV genes, infected CCO cells were analyzed at various times post CCV inoculation with or without 100 µg ml⁻¹
of cycloheximide (Sigma-Aldrich). Tissue culture flasks containing 70% confluent monolayers of CCO cells were inoculated with CCV at a multiplicity of infection of 10 PFU cell\(^{-1}\). After incubation for 40 min at 30°C, cell monolayers were harvested at 0, 1, and 3 h post infection, and total RNA was extracted from cell lysates. RT-PCR was used to monitor expression of IE, E and L CCV genes.

**Slot-blot hybridization.** An oligonucleotide probe (5’-CCG GAC CGA TGT ACA CGT GTG-3’) was designed on the basis of an available sequence (Davison 1992) from the ORF 3 region of the CCV genome. The BrightStar psorallen-biotin nonisotopic labeling kit (Ambion) was used to biotinylate 500 ng of the probe according to the manufacturer’s instructions. For the slot-blot hybridization, 10 µl of PCR products amplified by ORF 3-specific primers were diluted with 140 µl of Tris-ethylenediaminetetraacetic acid buffer (pH 8.0) and mixed with 150 µl of denaturing buffer (0.4 M NaOH, 10 mM EDTA). Positive and negative controls included PCR products amplified from purified CCV DNA and β-actin, respectively. The mixture was placed in a water bath for 10 min and centrifuged at 7000 × g for 5 s. The cDNA samples were chilled on ice, applied to Zeta-probe GT genomic membranes (Bio-Rad) and immobilized on the membrane by adding 500 µl of 0.4 M NaOH to each well of the slot-blot apparatus. Upon removal from the manifold system, membranes were washed in 2× SSC and 0.1% SDS at 59°C for 15 min. The hybridization buffer component, 20× SSC, 10% SDS, and ECL liquid block, the membrane was hybridized for ~17 h with 20 µl of biotinylated probe at room temperature for 5 min and subsequently with 1× SSC and 0.1% SDS at 59°C for 15 min. The hybridization signal was detected using the BrightStar BioDetect nonisotopic detection system (Ambion) following the recommendations of the manufacturer. The membrane was wrapped in a single layer of plastic wrap, stored in the dark for 24 h, and exposed to hyperfilm ECL (Amersham) for ~10 min.

**RESULTS**

**Evaluation of dexamethasone and heat stress as inducers of recrudescence**

To evaluate conditions that induce CCV recrudescence, we treated carrier fish with DEX, elevated temperature or with both, then evaluated the fish for the presence of infectious virus and compared CCV DNA content by PCR. By Day 7 post treatment, we were not able to detect infectious virus in any of the posterior kidney samples (N = 70), even after performing 3 blind passages. However, the percentage of fish testing positive for CCV DNA by PCR increased from 40 to 100% in the group treated with DEX but not heat stress (Fig. 1). Such an increase was not observed in any other treatments. This suggested that some recrudescence was occurring in the DEX-treated fish. Therefore, we used this method in a more in-depth analysis of recrudescence which included viral gene expression and the antibody response of the host.

**Evaluation of experimental latent infections and DEX induced recrudescence**

In the second study, individually identified fish that were mock-infected or challenged with CCV or CCVlacZ were evaluated for changes in anti-CCV antibodies by serially sampling the same fish over 90 DPC. Then at 90 DPC, we attempted to induce viral recrudescence using DEX, and subsamples were evaluated at 2, 4, 7, and 10 DPT for changes in anti-CCV antibodies, infectious virus production and virus gene expression. At 4 DPC, 1 fish each died in the CCV and CCVlacZ treatment groups. Both dead fish demonstrated clinical signs of CCVD, and CCO cultures inoculated with kidney homogenates from these fish showed characteristic CPE. At 90 DPC, no productive virus was recovered from any of the PBL

![Table 1. Primer pairs used in RT-PCR for channel catfish virus (CCV) immediate-early (IE), early (E), and late (L) gene expression, and the housekeeping gene β-actin](image_url)
cultures. After terminal sampling of DEX and PBS control treated fish, no virus was recovered from PBL co-culture or from CCO cultures of kidney tissue homogenates.

Western blot analysis of seroconversion

Western blot analysis of CCV-infected BB cells with serum from CCV-infected fish demonstrated antibody recognition of multiple CCV-specific proteins. Fig. 2A demonstrates the protein profile of mock- and CCV-infected BB cell lysates and cell supernatants using a positive control serum from a juvenile catfish immunized with a single intraperitoneal injection of $1 \times 10^5$ PFU of CCV. We used this assay to evaluate the change in response of CCV- and CCVlacZ-exposed fish over time and after exposure to DEX. Western blot analyses revealed the immune response profile of each fish (Fig. 2B). Seroconversion occurred by 30 DPC in 17 of 22 (77.3%) CCV-exposed fish. Out of 22 CCVlacZ-exposed fish, only 2 (9.1%) produced a specific antibody response by the same time post challenge. These levels dwindled to 4.5% and 0% for CCV- and CCVlacZ-exposed fish by 90 DPC, respectively. Secondary induction of the antibody response was demonstrated in 10 of 22 (45.5%) CCV-exposed fish compared to 3 of 22 (13.6%) CCVlacZ-exposed fish after injection with either DEX or PBS, with the strongest induction occurring in the CCV-exposed fish that were treated with DEX (Fig. 3). The most prominent antibody reactivity was present in serum samples from CCV exposed fish and was directed against viral proteins with molecular masses ranging from ~90 to >200 kDa (Fig. 2B, Table 2). In 9 out of 14 CCV-exposed fish, the secondary induction of an antibody response coincided with the reappearance of at least 1 protein with molecular mass similar to that present at 30 DPC (Table 2). No secondary immune response was observed when fish were treated with PBS alone. Out of 48 fish screened, 2 (3C and 18E) had antibodies to CCV antigens prior to the experimental 194
challenge. Their western blots showed reactivity against viral proteins ranging in size from ~35 to 68 kDa. Immunogenicity against a 46 kDa viral protein was observed at 30 DPC in 1 mock-infected fish (16D), but no induction of a secondary response was observed 2 d post DEX treatment. In contrast, the reactivity to a CCV antigen (~139 kDa) was evident after administration of DEX to another mock-infected fish (16A) without prior seroreactivity to CCV (Table 2). Western blots from 9 CCVlacZ- (24.3%) and 15 CCV-exposed (57.6%) fish had antibodies that recognized cellular proteins present in mock-infected cell lysates. These cell-reactive antibodies recognized low molecular mass proteins with apparent masses ranging from 20 to 68 kDa. Fig. 4 illustrates the presence of cell-reactive antibodies in a western blot from a CCV-infected fish.

**Evaluation of serum neutralizing activity**

A serum neutralization test was performed with a subgroup of 29 fish representing 5 treatment groups: mock/DEX (N = 4), CCV/DEX (N = 7), CCVlacZ/DEX (N = 6), CCV/PBS (N = 6), and CCVlacZ/PBS (N = 6). Fish with SNI values >1 were considered positive for neutralizing activity. The SNI measured at 60 DPC identified 8 (62%) CCV- and 3 (25%) CCVlacZ-exposed fish as positive (Table 3). In these positive fish, the SNI values ranged from 1.12 to 3.95 and 1.28 to 2.37 in the CCV- and CCVlacZ-exposed groups, respectively. Comparing SNI to western blot data by Spearman’s correlation demonstrated a significant relationship (p < 0.01) with a rho value of 0.582. The mean ± SD SNI was 0.53 ± 0.65 (N = 38) and 1.75 ± 1.00 (N = 20) for western blot negative and western blot positive samples respectively and they were significantly different (p < 0.001). Of the 38 western blot negative samples (29 at 60 d and the same samples after treatment), 8 had SNI above 1.0. Of these, 3 samples had SNI above the 95% CI for the mean for that group. Of the 20 western blot positive samples, 4 had SNI below 1.0. The greatest number of fish identified as positive, following a stimulus to reactivate virus, were among those sampled at 2 DPT. The SNI values measured in serum samples from 4 CCV/DEX treated fish ranged from 1.54 to 3.11 whereas 4 CCV/PBS-treated fish had SNI values ranging from 1.12 to 2.93. In contrast, only 1 CCVlacZ/DEX-treated fish was considered positive with an SNI value no greater than 1.2. None of the serum samples from the mock-infected group had detectable levels of neutralizing activity. Overall, SNI values significantly differed between treatments (F = 7.46, df = 4, 28, p = 0.025). Regardless of the treatment, fish exposed to CCV had the highest SNI values (Fig. 5). The SNI values from serum samples obtained post DEX treatment were not significantly different from PBS-injected controls (F = 2.58, df = 5, 11, p = 0.317).

**Evaluation of the pattern of CCV gene expression**

To evaluate CCV gene expression in latently infected fish, we used RT-PCR to determine if the regions of representative immediate early (ORF 3 and ORF 12), early (ORF 57) and late (ORF 46) genes were transcribed. In cell culture, transcripts representing all regions were detected at 1 and 3 h post infection, and all except ORF 12 were blocked by cycloheximide exposure when sampled at 2 h post infection (Fig. 6). This experiment was repeated twice with the same outcome. No product was seen when reverse transcriptase was omitted from the reaction mix.

**CCV gene expression during latency and following dexamethasone treatment**

To elucidate the events occurring during the establishment of latency by CCV and CCVlacZ, as well as immediately after administration of DEX, we examined viral gene expression in PBL and caudal fin
Table 2. Molecular mass (kDa) of viral protein subunits recognized by channel catfish virus (CCV)-specific antibodies in channel catfish. Fish were analyzed by western blotting on different days post challenge with CCV or the thymidine kinase-negative recombinant CCVlacZ and following attempted reactivation treatment with dexamethasone (DEX, 0.55 mg kg⁻¹ body weight) or phosphate-buffered saline (PBS; control) at 90 d post challenge (DPC). DPT = 2, 4, 7, and 10 d post treatment. N = negative, ND = not done, Mock = mock-treated

Table 3. Comparison of western blot and neutralizing antibody data in channel catfish serum samples. Samples were taken at 60 d post challenge (DPC) with no virus (mock), channel catfish virus (CCV), or thymidine kinase-negative recombinant CCVlacZ, and after attempted reactivation treatment with dexamethasone (DEX, 0.55 mg kg⁻¹ body weight) or phosphate-buffered saline (PBS; control) at 90 DPC. Post = 2, 4, 7, and 10 d post treatment

Table 4. Summary of the mRNA expression of the ORF 3, ORF 12, ORF 57 and ORF 46 regions. Out of 10 CCV- and 10 CCVlacZ-exposed fish, we detected only 2 fish that were expressing ORF 3-containing transcripts (18A and 22D) by 90 DPC. Both were CCV exposed. Following treatment to induce reactivation, ORF 3-containing transcripts were detected in 1 CCV/Dex- (19E), 2 CCV/PBS- (18A and 22E), and 1 CCVlacZ/Dex-treated fish (6C).

The ORF 3 gene RT-PCR products were detected by slot-blot hybridization due to the presence of 2 bands in polyacrylamide gels, one of which may represent a non-specific amplification product (Fig. 6). Table 4 summarizes the mRNA expression of the ORF 3, ORF 12, ORF 57 and ORF 46 regions.
DISCUSSION

Our goals were to evaluate the immune response of channel catfish to CCV and CCVlacZ after the establishment of latency and to characterize viral recrudescence. In a previous study, channel catfish were infected with similar doses of CCV or a TK-negative mutant and demonstrated that 100% of 16 sampled fish developed disseminated viremia by 4 days post infection and this progressed to latent infections in the survivors (as defined by presence of detectable CCV genome but no infectious virus) by 12 days post exposure (Kancharla & Hanson 1996). However, the authors did not attempt to induce recrudescence or evaluate the immune response over time. We were unable to induce productive recrudescence with either strain, but gene expression data and the long-term immunological profile of channel catfish after exposure to CCV or CCVlacZ and following DEX treatment suggest partial recrudescence. Data from western blot analyses showed that CCV-specific antibodies were predominantly directed against high molecular mass proteins ranging from ~90 to >200 kDa. The antibody reactivity against CCV antigens varied over time and declined markedly around 90 days post challenge (DPC) in most seropositive fish. More fish exposed to CCV produced detectable CCV-specific antibodies and these antibodies recognized a broader spectrum of viral proteins compared to those produced by fish exposed to CCVlacZ. Likewise, fish exposed to CCV had higher neutralizing antibody activity than those exposed to CCVlacZ. Previous studies found that the ability of the TK-deleted recombinants to infect and replicate early during the infection...
was similar to the parent CCV, but then virus numbers dwindled more quickly (Zhang & Hanson 1995, Kanchcharla & Hanson 1996). This more rapid clearance may account for a lesser stimulation of specific antibody responses that we observed in the western blots.

Even though there was a good correlation between neutralizing antibody assays and western blot data, there were several cases where western blot negative fish had positive SNI and vice versa. This result is not unexpected since serum neutralization is often based on antibodies binding secondary or tertiary structural epitopes on the virus anti-receptor, whereas the SDS-PAGE treatment in western blot assays necessitates the evaluation of antibodies that mainly recognize primary structures. Furthermore, western blot assays evaluate responses to many more proteins in addition to those involved in the infection process. Similar results were documented by Crawford et al. (1999).

After experimental CCV exposure, they found that many fish had a negative SNI but were seropositive for CCV antigens when evaluated by ELISA. Also, they found that a similar percentage of fish (65%) seroconverted by 5 wk post exposure to a similar concentration of CCV as was used in our study.

The secondary induction of an antibody response was only demonstrated in fish that received DEX treatment. Serum samples from these fish demonstrated recognition of at least 1 viral protein with molecular mass similar to that recognized after the primary infection. No secondary immune response was observed in any fish injected with PBS alone for either CCV or CCVlacZ. Although these results imply that a recrudescent expression of CCV proteins was occurring, the attempted reactivation using DEX injection did not result in production of infectious virus in co-cultivation cultures from any PBL sample or tissue homogenates of the fish.

The injection of DEX resulted in elevated virus-specific antibodies. DEX is a synthetic glucocorticoid that regulates cellular and viral gene expression, induces apoptosis in nonspecific cytotoxic cells, and has immunosuppressive properties (Rousseau & Baxter 1979, Yamamoto 1985, Bishop et al. 2000). Glucocorticoids are the gold standard agents for anti-inflammatory and immunomodulatory therapy, whose main mechanism of action relies on inhibiting the activity of transcription factors, such as nuclear factor-kappa B (NF-kB) and activator protein-1 (AP-1) (De Bosscher et al. 2000). Therefore, a secondary antibody response is unexpected. However, glucocorticoids often induce reactivation of herpesviruses from latency. DEX treatment is a reliable method for reactivation of suid herpesvirus 1, equid herpesvirus 1, and caprine herpesvirus 1 when administrated alone (Gibson et al. 1992, Menzeling et al. 1992, Buonavoglia et al. 1996). The single DEX application likely induced sufficient CCV recrudescence to stimulate a secondary immune response and this response quickly suppressed virus production and prevented us from being able to culture the virus. Others have demonstrated an immune response to reactivated virus after DEX treatment. Tanaka & Mannen (2002) showed the activation of latent suid herpesvirus 1 in mice after DEX or acetylcholine treatment resulted in viral excretion and increased antibody titers. An alternative explanation is that the DEX treatment influenced antibody levels non-specifically. There are a few unusual cases where DEX has been shown to cause higher serum antibodies. In one case, a low dose of corticosteroid given orally to an immunodeficient patient with splenomegaly reduced the hypercatabolism of immunoglobulin by macrophages and this allowed administered immunoglobulin to build up in the blood (Spickett et al. 1996). Another possibility could be that the dosage of DEX administered in our study disrupted the activity of regulatory T lymphocytes (T_{reg}) or dendritic cells, allowing suppressed B-lymphocytes to generate antibodies. However, this possibility is unlikely. T_{reg} and dendritic cells are poorly characterized in fish, but in mammals DEX stimulates suppressive pathways in dendritic cells (Grohmann et al. 2007) and enhances the activity and numbers of T_{reg} cells (Karagiannidis et al. 2004).

Contrary to results reported by Bowser et al. (1985), we did not isolate infectious virus in co-cultivation assays in the initial trial with carrier fish and in the second trial with experimentally infected fish. Reactivation of herpesviruses is a complex phenomenon. Immunity plays an important role in modifying the outcome of primary and recurrent herpes simplex virus infections (Nash & Wildly 1983). In humans,
reactivation of HSV-1 is often influenced by immunity characterized by no detectable rise in circulating neutralizing antibodies (Douglas & Couch 1970). Simmons & Nash (1985) reported that the zosteriform spread of HSV-1 was completely inhibited by neutralizing antibodies. In a study on recrudescence of gammaherpesviruses in immunocompromised hosts, those that had antibodies to a lytic cycle viral protein were able to control virus re-expression (Gangappa et al. 2002). Furthermore, reactivation of herpesviruses in mammals sometimes requires administration of DEX in conjunction with cyclophosphamide (Minagawa et al. 1994) or anti-serum to lymphocytes to suppress antibody production (Bevan et al. 1996). We evaluated 2 potential methods of inducing recrudescence, heat stress and DEX injection, and their effect in combination. The heat stress response is characterized by the induction of heat shock proteins (HSPs) when cells are exposed to elevated temperatures. Stress-inducible HSPs, such as Hsp72, have anti-inflammatory properties. Tang et al. (2007) demonstrated an inhibition of NF-kB activity in cells expressing abundant Hsp72. This inhibition in turn decreased cellular levels of high-mobility-group box 1 induced cytokines (tumor necrosis factor [TNF]-α and interleukin [IL]-1β) and proinflammatory signaling in macrophages. Heat was evaluated here because heat exposure has been shown to induce HSV-1 in mammals (Sawtell & Thompson 1992, 2004). Moreover, Halford et al. (1996) postulated that pre-treatment of cultures with DEX facilitated the induction of HSV-1 lytic-phase mRNA transcription following heat stress. Furthermore, natural outbreaks of CCVD occur primarily during elevated summertime temperatures. We chose the dose of DEX based on the findings of Bowser et al. (1985). Contrary to what was expected, administration of DEX in our study did not result in productive infection in latently infected catfish. However, the PCR detection rate of CCV DNA increased from 40 to 100%, and this increase was not observed in any other treatments. There are 2 other studies where the researchers failed to isolate CCV from tissues of adult channel catfish suspected of carrying the virus after chemical or corticosteroid induction (Huston 1979, Plumb et al. 1981). The major difference between the findings of Bowser et al. (1985) and ours may relate to the physiology of the fish. In the study by Bowser et al. (1985), moribund, naturally infected CCV carriers were injected with DEX. These fish were adult broodfish, housed in outdoor ponds and exposed to winter conditions. Low temperature suppresses cellular and humoral immunity in channel catfish (Plumb et al. 1986, Bly & Clem 1991). A natural immunosuppressed state combined with the effect of the DEX may be needed to provide the level of recrudescence necessary to detect the virus in cell cultures. In future studies on CCV recrudescence, we plan to expose fish to low water temperatures and to decrease intervals of sample collection by adding time points at 6 and 12 h post DEX treatment.

The sporadic nature of expression of the selected gene regions precludes broad conclusions about gene expression profiles during latency according to expression class. However, in these analyses we looked at virus-exposed fish at 2 time points (90 DPC and post treatment) and we did detect virus gene transcription at one or both of these time points in 7 out of 10 CCV-exposed fish and in 6 out of 10 CCVlacZ-exposed fish. This suggests that sporadic CCV gene expression is relatively common in latently infected fish. Furthermore, serial expression was documented in 5 of these 13 fish. These findings indirectly support the suggestion that protein expression may induce increased antibody levels after DEX exposure.

Although not the focus of this research, we found a high incidence of antibodies reacting to proteins in mock-infected cell lysates in serum samples from channel catfish exposed to either CCV or CCVlacZ. The etiology of many chronic autoimmune diseases is still uncertain, but infectious agents have often been implicated in their pathogenesis. Cell-reactive antibodies have been reported in patients suffering from autoimmune diseases that in some cases also have specific antibodies to chronic herpesvirus infections including Epstein-Barr virus, human herpesvirus type 6, human cytomegalovirus, and herpes simplex virus (Vaughan et al. 1996, Galanakis et al. 2001, Nawata et al. 2001, Simmons 2001, Totani et al. 2001, Zhang et al. 2001, Alvarez-Lafuente et al. 2002).

The combined results obtained from western blot, SNI, and RT-PCR data suggest some viral recrudescence in channel catfish that received DEX treatment. This study illustrates the recalcitrant nature of CCV latency and suggests that recrudescence is rare and tightly regulated. Evaluating the extent and the physiological impact of virus reactivation on latently infected carriers will require a better understanding of the factors that restrain latent CCV from undergoing productive replication. Further investigation is necessary to examine the expression of CCV genes and to characterize the cell types involved during the latent and recrudescent phases of infection.

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LITERATURE CITED

- Houston MM (1979) Channel catfish virus disease: detection of latent virus and correlation with serology. PhD dissertation, Texas A&M University, College Station
- Kancharia SR, Hanson LA (1996) Production and shedding of channel catfish virus (CCV) and thymidine kinase negative CCV in immersion exposed channel catfish fingerlings. Dis Aquat Org 27:25–34
- Lobb CJ, Olson MOJ, Clem LW (1984) Immunoglobulin light...


Zhang HG, Hanson LA (1996) Recombinant channel catfish virus (Ictalurid herpesvirus 1) can express foreign genes and induce antibody production against the gene product. J Fish Dis 19:121–128