Elephant Herpesvirus References (alphabetical)
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Elephants, particularly Asian (Elephas maximus), are threatened by lethal elephant hemorrhagic disease (EHD) due to elephant endotheliotropic herpesviruses (EEHV). At least five of seven known EEHV types have been associated to EHD, with types 1, 4, and 5 predominantly affecting Asian elephants. In Switzerland, at least three Asian elephants have been lost due to EHD but nothing is known about the present EEHV1 circulation. Moreover, the prevalence of other EEHV types has never been assessed. Intermittent shedding of EEHV can be monitored through collecting trunk secretions and analyzing them by PCR methods that discriminate the different EEHV types. To identify EEHV shedders, seven of eight Asian elephants in a Swiss zoo were trained to provide trunk wash samples. These were collected at intervals over a period of four months and tested by PCR for presence of EEHV1 through 6. Moreover, the quality of each sample was assessed by testing for the elephant TNF-alpha gene. Overall, 57% of the samples were valid with five of seven participating elephants identified as EEHV shedders. Two of those shed virus only once, whereas the other three, all closely related among each other, shed virus on multiple occasions. One of the frequent shedders had been in very close contact to all of the three EHD victims. Therefore, we speculate that this particular animal may represent the virus source in all three cases. However, when subtyping was conducted, the presently circulating virus was identified as EEHV1B, while the virus subtype causing EHD had been 1A in all three cases. In addition to four animals excreting EEHV1, a recently introduced animal was observed to shed EEHV3/4. We suggest that the policy of trunk washing to identify and characterize EEHV-shedders is to be endorsed in zoos with ongoing or planned elephant breeding programs.


Elephant endotheliotropic herpesviruses (EEHVs) can cause acute hemorrhagic disease with high mortality rates in Asian elephants (Elephas maximus). Recently, a new EEHV type known as EEHV5 has been described, but its prevalence and clinical significance remain unknown. In this report, an outbreak of EEHV5 infection in a herd of captive Asian elephants in a zoo was characterized. In February 2011, a 42-yr-old wild-born female Asian elephant presented with bilaterally swollen temporal glands, oral mucosal hyperemia, vesicles on the tongue, and generalized lethargy. The elephant had a leukopenia and thrombocytopenia. She was treated with flunixin meglumine, famciclovir, and fluids. Clinical signs of illness resolved gradually over 2 wk, and the white blood cell count and platelets rebounded to higher-than-normal values. EEHV5 viremia was detectable starting 1 wk before presentation and peaked at the onset of clinical illness. EEHV5 shedding in trunk secretions peaked after viremia resolved and continued for more than 2 mo. EEHV5 trunk shedding from a female herd mate without any detectable viremia was detected prior to onset of clinical disease in the 42-yr-old elephant, indicating reactivation rather than primary infection in this elephant. Subsequent EEHV5 viremia and trunk shedding was documented in the other five elephants in the herd, who remained asymptomatic, except for 1 day of temporal gland swelling in an otherwise-healthy 1-yr-old calf. Unexpectedly, the two elephants most recently introduced into the herd 40 mo previously shed a distinctive EEHV5 strain from that seen in the other five elephants. This is the first report to document the kinetics of EEHV5 infection in captive Asian elephants and to provide evidence that this virus can cause illness in some animals.

Elephant endotheliotropic herpesviruses (EEHVs) are a continuous threat for young Asian elephants. We report a laboratory-confirmed infection of a 5-year-old female Asian elephant (AZ_2016) in the Berlin Zoologischer Garten. Initially, high EEHV-1 loads were detected in trunk swabs obtained from the young elephant during routine screening. The animal showed no clinical signs except for slight irritability. EEHV-1 was continuously shed for almost one year, with fluctuations in viral load from time to time. Our investigations highlight the continuous threat of EEHV-1 to young captive Asian elephants and stress the importance of routine monitoring of captive elephants to allow early detection of infection.


Elephant endotheliotropic herpesviruses (EEHVs) are the cause of acute hemorrhagic disease in endangered Asian and African elephants. In the present study, we report the incidence of EEHV infection and associated mortality in the captive elephant of Assam, India. Our result showed the gross morphology and histopathological changes of EEHV infection in the elephant. Moreover, the phylogenetic analysis of the polymerase, helicase, and GPCR genes from the infected tissue samples suggested the presence of EEHV1A virus.


Twenty-four species of South African wild animals were tested for the presence of antibodies to the viruses of 16 common diseases of domestic animals around 1993-5. Positive results were obtained for African horse sickness, equine encephalomyelitis virus, equine herpesvirus-1, bovine herpesvirus-1, Allerton disease (Herpes mammillitis; bovine herpesvirus 2), lumpy skin disease, parainfluenza, encephalomyocarditis, bluetongue, Wesselsbron disease, bovine ephemeral fever, and Akabane disease complex. No antibodies could be demonstrated against the viruses of equine influenza, equine infectious anaemia, equine viral arteritis or Rift Valley fever. The negative results support observations that the latter diseases, with the exception of equine viral arteritis, are absent in South Africa. The number of animal species found positive for a specific virus, ranged from 0-16. No antibodies were found in crocodiles or warthogs, whereas antibodies against Wesselsbron and bovine herpesvirus-1 were present in 16 species. Antibodies against viruses of horses were found almost exclusively in zebras and, although elephants reacted to African horse sickness, no neutralizing antibodies against it could be demonstrated in their sera. Zebras were also found to be positive for Wesselsbron and Akabane, which are usually regarded as viruses of ruminants. Antibodies against most viruses were encountered in all vegetation zones in South Africa, but most viruses were more prevalent in the high-rainfall zone in KwaZulu-Natal.


INTRODUCTION: EEHV-1 is a viral infection of elephants that has been associated with a fatal haemorrhagic syndrome in Asian elephants. Previous studies have suggested that pregnant animals may shed more virus than non-pregnant animals. METHODS: This study examined whether pregnancy affected the frequency or magnitude of shedding of elephant endotheliotropic herpesvirus 1 (EEHV1) using Taq man real-time PCR on trunk washes from four female elephants from a UK collection over three time periods between 2011 and 2014. These periods included pregnancies in two animals (period 1 and period 3). Behavioural observations made by keepers were also assessed. RESULTS: During period 1 there was a high degree of social hierarchical instability which led to a hierarchy change, and was associated with aggressive behaviour. Also during period 1 EEHV-1 shedding was of a higher magnitude and frequency than in the latter two time periods. CONCLUSIONS: These results suggest that
there is no clear relationship between shedding and pregnancy, and that behavioural stressors may be related to an increase in EEHV-1 shedding.


Antibodies were detected against bovine herpesviruses 1 (BHV 1) and 2 (BHV 2) in Asian elephants (Elephas maximus) using the passive haemagglutination (PHA) test. The study was conducted during May to December 1994 using sera collected from zoos and national parks in India. Four (4%) of 109 elephant sera had PHA titres ranging from 1:8 to 1:32 against BHV 1. 25 (23%) of the 109 elephant sera had PHA titres ranging from 1:8 to 1:64 against BHV 2. It is concluded that Asian elephants appear to be better reservoirs for herpesviruses which are serologically related to BHV 2.


Elephant endotheliotropic herpesviruses (EEHVs) can cause fatal hemorrhagic disease in elephants, especially young captive Asian elephants (Elephas maximus). Currently, seven EEHV types have been reported. In this study, EEHVs were examined in whole-blood samples derived from 56 captive Asian elephants from eight provinces in Thailand by nested PCR using primers specific to the viral DNA polymerase gene in an attempt to monitor EEHV elephant cases. After EEHV testing, one sample (1.78%) was positive and found to be closely related to EEHV4 with 99% amino acid identity. This sample was from a three-year-old female Asian elephant with no clinical signs. These data suggest that asymptomatic EEHV4 infection can occur in Asian elephants.


The elephant endotheliotropic herpesvirus (EEHV) is now recognized as one of the main causes of death of young Asian elephants (Elephas maximus) in North American zoos. Its impact in wild and domestic elephant populations in Asia is not clearly understood. This article describes the first case of EEHV infection in Lao People's Democratic Republic of a 2.5-yr-old domestic male Asian elephant. Clinical signs and pathological findings reported here are consistent with previous infections in Asian elephant calves. Phylogenetic analyses showed 100% homology with other EEHV-1A strains identified in Asia, Europe, and North America. Contamination of the molecular assays was ruled out, because the DNA polymerase sequence identified in this study differed from the positive control by two base pairs. © 2014 American Association of Zoo Veterinarians.


myocardium. Furthermore there have been ulcerations at various localizations of the whole digestive tract. Slightly basophilic intranuclear inclusion bodies have been found histologically in endothelial cells of different organ samples. An examination of altered organ-material by electronmicroscopy made some herpesvirus-like particles visible. A virological investigation first revealed evidence of giant cell formations with solitary basophilic intranuclear inclusion bodies in different cell cultures, however, without any distinct cytopathogenic effect. Supported by molecular biological procedures the infection of 'KIBA' could be verified as the elephants herpesvirus. By means of PCR and subsequent sequence analysis a DNA-sequence typical for the elephants herpesvirus could be obtained which showed an identity of 97% with the terminase sequence of the elephant herpesvirus described by American authors. The deduced amino acid-sequences were 100% identical. To the terminase of the human cytomegalovirus, the elephant sequence had an identity of 53% (similarity: 74%). Based on the cooperation of ILAT, Institute of Veterinary-Pathology/Free University Berlin, Robert-Koch-Institut Berlin, and Zoological Garden Berlin, the cause of 'KIBA's' death could be discovered immediately. Possible implications of this case especially on breeding and keeping elephants are discussed briefly.


The maintenance of wild animals in captivity in North America is regulated by a number of different laws and government agencies in each country. Member institutions of zoo and aquarium associations in Canada, the United States of America and Mexico experience an extra tier of regulation in the form of industry standards, which are sometimes stricter than those imposed by government. Climate, natural disasters and harmful pest species all contribute to the challenge of keeping animals in certain locales. Vigilance against zoonotic disease transmission is maintained through industry and government-mandated sanitation standards, which are fortified by reporting regulations of local, regional and Federal health agencies. Current controversies in the keeping of particular taxa in North America include the threat to non-human primate breeding programmes precipitated by strict new import regulations, the fear of herpesvirus B infection, and commercial airline transport bans. Successive human fatalities among elephant handlers have prompted the industry and governments to re-examine the manner in which these potentially dangerous creatures are maintained in captivity.


BACKGROUND: Elephant Endotheliotropic Herpesviruses (EEHVs) can cause acute haemorrhagic disease in young Asian elephants (Elephas maximus) and clinical EEHV infections account for the majority of their fatalities. The anti-herpesviral drug famciclovir (FCV) has been used routinely to treat viraemic at-risk elephants, but thus far without proven efficacy. This paper presents clinical and virological investigations of two EEHV-1A infected elephants treated with FCV, and discusses anti-herpesvirus therapies of viraemic elephants. CASES PRESENTATIONS: Two 1.5 year old male Asian elephants at a zoological collection in the UK developed clinical EEHV-1A infections. Case 1 showed signs of myalgia for the duration of 24 hours before returning back to normal. EEHV-1A DNAemia was confirmed on the day of clinical signs and continued to be present for 18 days in total. Trunk shedding of the virus commenced 10 days after detection of initial DNAemia. Case 2 tested positive for EEHV-1A DNAemia in a routine blood screening sample in the absence of clinical signs. The blood viral load increased exponentially leading up to fatal clinical disease seven days after initial detection of DNAemia. Both calves were treated with 15 mg/kg FCV per rectum on detection of DNAemia and penciclovir, the FCV metabolite, could be detected in the blood at assumed therapeutic levels. The early indicators for clinical disease were a marked absolute and relative drop in white blood cells, particularly monocytes prior to the detection of viraemia. The most prognostic haematological parameter at later stages of the disease was the platelet count showing a continuous sharp decline throughout, followed by a dramatic drop at the time of death. CONCLUSIONS: The EEHV-1A viraemic animals investigated here further highlight the ongoing threat posed by these viruses to juvenile Asian elephants. The findings call into question the efficacy of rectal FCV in
clinical cases and direct towards the use of alternative anti-herpesvirus drugs and complementary treatments such as plasma infusions if no improvement in either viral load or the above-mentioned blood parameters are observed in the initial days of viraemia despite anti-herpesvirus therapy.


A male Asian elephant (Elephas maximus) died at the Berlin zoological gardens in August 1998 of systemic infection with the novel endotheliotropic elephant herpesvirus (ElHV-1). This virus causes a fatal haemorrhagic disease in Asian elephants, the so-called endothelial inclusion body disease, as reported from North American zoological gardens. In the present work, ElHV-1 was visualized ultrastructurally in affected organ material. Furthermore, a gene block comprising the complete glycoprotein B (gB) and DNA polymerase (DPOL) genes as well as two partial genes was amplified by PCR-based genome walking and sequenced. The gene content and arrangement were similar to those of members of the Betaherpesvirinae. However, phylogenetic analysis with gB and DPOL consistently revealed a very distant relationship to the betaherpesviruses. Therefore, ElHV-1 may be a member of a new genus or even a new herpesvirus subfamily. The sequence information generated was used to set up a nested-PCR assay for diagnosis of suspected cases of endothelial inclusion body disease. Furthermore, it will aid in the development of antibody-based detection methods and of vaccination strategies against this fatal herpesvirus infection in the endangered Asian elephant.


Endotheliotropic elephant herpesvirus (elephantid herpesvirus 1; ElHV-1) is apathogenic for African elephants (Loxodonta africana), but causes fatal haemorrhagic disease in Asian elephants (Elephas maximus). This is thought to occur through transmission from African elephants in places where both species are housed, such as zoological gardens. The virus has caused considerable losses in North American and European zoological gardens and thus severely impedes breeding of the endangered Asian elephant. Previously, the ultrastructural and genetic characterization of ElHV-1 from a male Asian elephant that died from the disease at the Berlin zoological gardens in 1998 have been reported. Here, a partial characterization of the ElHV-1 genome is presented. A 60 kbp locus, spanning 34 open reading frames, was analysed. Most of the detected genes were found to be conserved among the herpesviruses and showed an overall arrangement most similar to that of betaherpesviruses, in particular Human herpesvirus 6 and Human herpesvirus 7. Most importantly, in addition to a protein kinase gene that is homologous to the human cytomegalovirus UL97 gene, a thymidine kinase (TK) gene was found, which is generally missing in betaherpesvirus genomes. Thus, ElHV-1 is the only known betaherpesvirus to encode a TK gene. This peculiarity might contribute to the fulminant pathogenicity of ElHV-1, but also provide a crucial enzymic activity for developing an efficient antiviral therapy with currently available nucleoside analogues.


Newly discovered, lethal elephant endotheliotropic herpesviruses (EEHV) have been identified in both Asian (Elephas maximus) and African (Loxodonta africana) elephants. Carried by otherwise healthy African elephants, they can be fatal, mainly for young Asian elephants. Since zoos often harbour both elephant species, we conducted a survey on the presence of EEHV in elephants (Asian elephants, n=57; African elephants, n=17) from 12 zoos and 3 circuses in Europe (Germany, Switzerland and the Netherlands), and 1 zoo in Israel [date not given]. Six of the 57 Asian elephants were positive for EEHV. Five elephants died of the infection, while one survived. EEHV was not detected in any of the 17 African elephants. All EEHV that affected the Asian elephants belonged to the EEHV1 group. We described the detection and the partial sequencing of an endotheliotropic
The recently described elephant endotheliotropic herpesviruses (EEHV) have been associated with the deaths of numerous captive elephants. A proposed tool for the detection of EEHV infection in elephants is the PCR-based screening for EEHV-DNA in whole blood samples. Unfortunately, this detection method has only been successful in post-mortem analyses or in animals already displaying clinical signs of EEHV disease, thus rendering this method unsuitable for identification of carrier elephants. Here, we focus on glycoprotein B (gB) for serologic assay development, since gB is an envelope protein known to induce a neutralising antibody response in other herpesvirus infections. We sequenced the entire gB gene from five Asian elephants with EEHV, representing four different gB variants. Computer-aided methods were used to predict functionally important regions within EEHVgB. An extra-cytoplasmic region of 153 amino acids was predicted to be under positive selection and may potentially contain antigenic determinants that will be useful for future serologic assay development.


Elephant endotheliotropic herpesvirus (EEHV) can cause lethal hemorrhagic disease in juvenile Asian elephants. A number of EEHV types and subtypes exist, where most deaths have been caused by EEHV1A and EEHV1B. EEHV4 has been attributed to two deaths, but as both diagnoses were made postmortem, EEHV4 disease has not yet been observed and recorded clinically. In this brief communication, two cases of EEHV4 infection in juvenile elephants at the Houston Zoo are described, where both cases were resolved following intensive treatment and administration of famciclovir. A quantitative real-time polymerase chain reaction detected EEHV4 viremia that correlated with clinical signs. High levels of EEHV4 shedding from trunk wash secretions of the first viremic elephant correlated with subsequent infection of the second elephant with EEHV4. It is hoped that the observations made in these cases— and the successful treatment regimen used—will help other institutions identify and treat EEHV4 infection in the future.


Elephant Endotheliotropic Herpesvirus (EEHV) can cause lethal hemorrhagic disease in juvenile Asian elephants, an endangered species. One hypothesis to explain this vulnerability of some juvenile elephants is that they fail to mount an effective T cell response to the virus. To our knowledge, there have been no studies of Asian elephant T cell responses to EEHV. To address this deficiency, we validated the IFN-gamma ELISpot assay for tracking antigen-directed T cell activity by monitoring rabies-specific responses in vaccinated elephants. Additionally, we generated monoclonal antibodies to Asian elephant CD4 and CD8 to facilitate phenotypic T cell profiling. Using these tools, we screened healthy elephants with a prior history of EEHV infection for reactivity against 9 EEHV proteins whose counterparts in other herpesviruses are known to induce T cell responses in their natural hosts. We identified glycoprotein B (gB) and the putative regulatory protein E40 as the most immunogenic T cell targets (IFN-gamma responses in 5 of 7 elephants) followed by the major capsid protein (MCP) (IFN-gamma responses in 3 of 7 elephants). We also observed that IFN-gamma responses were largely from CD4+ T cells. We detected no activity against the predicted major immediate early (E44) and large tegument (E34) proteins— both immunodominant T cell targets in humans latently infected with cytomegalovirus. These studies have identified EEHV-specific T cells in Asian elephants for the first time, lending insight into the T cell priming that might be required to protect against EEHV disease and will guide the design of effective vaccine strategies.IMPORTANCE Endangered Asian elephants are facing many threats, including lethal hemorrhagic disease from elephant endotheliotropic herpesvirus (EEHV). EEHV usually establishes chronic, benign infections in mature Asian
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Elephants but can be lethal to juvenile elephants in captivity and the wild. It is the leading cause of death in captive Asian elephants in North America and Europe. Despite availability of sensitive tests and protocols for treating EEHV-associated illness, these measures are not always effective. The best line of defense would be a preventative vaccine. We interrogated normal healthy elephants previously infected with EEHV for T cell responses to 9 EEHV proteins predicted to induce cellular immune responses. Three proteins elicited IFN-gamma responses, suggesting their potential usefulness as vaccine candidates. Our work is the first to describe T cell responses to a member of the proposed fourth subfamily of mammalian herpesviruses, the Deltaherpesvirinae, within a host species in the clade Afrotheria. An EEHV vaccine would greatly contribute to the healthcare of Asian and African elephants that are also susceptible to this disease.


Elephant endotheliotropic herpesvirus 1 (EEHV1), a member of the Betaherpesvirinae subfamily, has recently emerged as an important viral pathogen of Asian elephants that can cause a severe, often fatal, hemorrhagic disease. EEHV1 does not replicate in culture and little is currently known about the molecular biology of this emerging pathogen, with the notable exception of its genomic DNA sequence. Here, we have used small RNA deep sequencing to determine whether EEHV1, like other human and murine betaherpesviruses, expresses viral microRNAs in infected tissues in vivo. Our data provide evidence supporting the existence of at least three novel viral microRNAs encoded by EEHV1 and one of these, miR-E3-5p, is shown to repress target mRNA expression. Moreover, miR-E3-5p expression was readily detectable in tissue samples derived from two infected elephants, including in whole blood. These data shed new light on the biology of EEHV1 and identify small RNAs that have the potential to be useful in the diagnosis of sub-clinical infections in captive Asian and African elephants.


This month's Genome Watch highlights how deep sequencing was used to generate the first full genomes of herpesviruses associated with a fatal disease in elephants.


The first herpesviruses described in association with serious elephant disease were referred to as endotheliotropic herpesviruses (EEHV) because of their ability to infect capillary endothelial cells and cause potentially fatal disease. Two related viruses, EEHV1 and EEHV2, have been described based on genetic composition. This report describes the similarities and differences in clinico-pathologic features of 2 cases of fatal endotheliotropic herpesvirus infections in Asian elephants caused by a previously unrecognized virus within the betaherpesvirus subfamily. EEHV3 is markedly divergent from the 2 previously studied fatal probosciviruses, based on polymerase chain reaction sequence analysis of 2 segments of the viral genome. In addition to ascites, widespread visceral edema, petechiae, and capillary damage previously reported, important findings with EEHV3 infection were the presence of grossly visible renal medullary hemorrhage, a tropism for larger veins and arteries in various tissues, relatively high density of renal herpetic inclusions, and involvement of the retinal vessels. These findings indicate a less selective organ tropism, and this may confer a higher degree of virulence for EEHV3.


This study assessed the feasibility of identifying asymptomatic viral shedders using a novel TaqMan real-time PCR on trunk washes and swabs from the conjunctiva, palate and vulva of elephants. Six elephants from a UK collection were sampled weekly over a period of 11 weeks for this study. The herd prevalence of elephant endotheliotropic herpesvirus-1 (EEHV-1) was 100 per cent by PCR. The virus DNA was detected in all the
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sampling sites; however, the prevalence of virus DNA in the conjunctiva swabs was higher. In addition, Asian elephants from two continental European collections were sampled once and one animal tested positive on a trunk wash. The virus from this animal was phylogenetically typed as EEHV-1A based on 231 nucleotides of the terminase gene.


So far there are no valid diagnostic tools available for identifying latent carriers of endotheliotropic elephant herpes virus (EEHV). For this reason, the lateral retropharyngeal lymph node complex (LARELYNOC) of elephants, identified during postmortem studies as target organ for EEHV and suitable for transcutaneous biopsy, was grossly described. Transcutaneous ultrasound (3.5 MHz) was applied behind the ear region to identify the LARELYNOC containing up to four single lymph nodes on each side. The lymph node tissue is situated 20-50 mm below the skin surface. An ultrasonographic assessment of the LARELYNOC and two biopsies were performed on 39 healthy Asian elephants (Elephas maximus). Samples were tested for EEHV via PCR. Whole blood samples were also collected and tested for active EEHV infection. Lymph nodes were ultrasonographically classified as active (calculated mean volume=17.4 ± 6.9 cm3, P>0.001), inactive (calculated mean volume=3.1 ± 0.6 cm3, P<0.001), or chronic active (calculated mean volume=10.6 ± 1.0 cm3, P<0.05). Histology confirmed not only the presence of lymph tissue but also the ultrasonographically diagnosed reactivity status of the lymph node biopsies. Although all samples including whole blood were found to be negative for the EEHV DNA particles, the successful development of this procedure in elephants could prove beneficial for the screening of not only latent EEHV infections but might also be a less dangerous alternative method for the diagnosis of zoonotic infections such as tuberculosis.


Endotheliotropic herpesvirus causes a fatal disease in young Asian elephants, but there are no methods for identifying latent carriers of the virus. During the postmortem study of one female African elephant and three male and two female Asian elephants, a lymph node located bilaterally caudoventral to the parotid gland, approximately 1.5 to 5 cm below the skin, was identified as suitable for transcutaneous ultrasound-guided biopsy. An ultrasonographic assessment and two biopsies were performed on 39 Asian elephants, and these lymph nodes were classified ultrasonographically as active, inactive or chronically active. The calculated mean (se) volume of 10 active lymph nodes was 17.4 (6.9) cm3, and that of three chronically active lymph nodes was 10.6 (1.0) cm3, whereas the mean volume of 17 inactive lymph nodes was 3.1 (0.6) cm3. The presence of lymph node tissue in samples obtained by ultrasound-guided biopsy from three animals that were maintained under conditions that allowed for additional sampling was confirmed histologically. The dna extracted from the lymphoid tissue and the whole blood of all the elephants was negative for endotheliotropic herpesvirus by PCR.


Asian elephant (Elephas maximus) immunity is poorly characterized and understood. This gap in knowledge is particularly concerning as Asian elephants are an endangered species threatened by a newly discovered herpesvirus known as elephant endotheliotropic herpesvirus (EEHV), which is the leading cause of death for captive Asian elephants born after 1980 in North America. While reliable diagnostic assays have been developed to detect EEHV DNA, serological assays to evaluate elephant anti-EEHV antibody responses are lacking and will be needed for surveillance and epidemiological studies and also for evaluating potential treatments or vaccines against lethal EEHV infection. Previous studies have shown that Asian elephants produce IgG in serum, but they
failed to detect IgM and IgA, further hampering development of informative serological assays for this species. To begin to address this issue, we determined the constant region genomic sequence of Asian elephant IgM and obtained some limited protein sequence information for putative serum IgA. The information was used to generate or identify specific commercial antisera reactive against IgM and IgA isotypes. In addition, we generated a monoclonal antibody against Asian elephant IgG. These three reagents were used to demonstrate that all three immunoglobulin isotypes are found in Asian elephant serum and milk and to detect antibody responses following tetanus toxoid booster vaccination or antibodies against a putative EEHV structural protein. The results indicate that these new reagents will be useful for developing sensitive and specific assays to detect and characterize elephant antibody responses for any pathogen or vaccine, including EEHV.


Asian elephants (Elephas maximus) are susceptible to a unique infection caused by elephant endotheliotropic herpesvirus (EEHV).3,4 Worldwide, between the years 1983 and 2000, there have been 26 confirmed deaths from this virus in Asian elephants.2 Although most cases have been fatal, treatment with famciclovir (Famvir, SmithKline Beecham Pharmaceuticals, Philadelphia, PA 19101 USA) has been associated with survival in three cases of six cases of EEHV infection proven by PCR.2,5,6 Dose selections for surviving elephants (5.5 - 8.0 mg/kg, p.o. every 8 hr) were made without the benefit of elephant pharmacokinetics and were a direct extrapolation from recommended human dosages (7 mg/kg, p.o. every 8 hr).5,6 In this study, famciclovir was administered both orally and rectally in healthy young Asian elephants. The doses tested in this study were 5 mg/kg orally, 5 mg/kg rectally, and 15 mg/kg rectally. Blood samples were analyzed for famciclovir and penciclovir using a validated LC/MS assay. Famciclovir was absorbed well by both routes and underwent rapid biotransformation to the active compound penciclovir. None of the plasma samples had detectable famciclovir. Pharmacokinetic parameters for penciclovir were determined using non-compartmental analysis. After a single oral dose of 5 mg/kg the Cmax was 1.3 ìg/mL with a Tmax at 1.1 h. After a rectal dose of 5 mg/kg the Cmax was 1.2 ìg/mL with a Tmax at 0.34 hr. After a rectal dose of 15 mg the t½ was 2.6 h, with a Cmax of 3.6 ìg/mL at Tmax 0.66 h. These results were similar to those reported in humans where an oral dose of 500 mg (7 mg/kg) had a t½ of about 2 h with a Cmax of 3.3 ìg/mL. A dose range of 8 -15 mg/kg given orally or rectally every 8 hours should produce penciclovir concentrations in Asian elephants that are considered therapeutic in humans.

LITERATURE CITED

Proliferative cutaneous lesions developed in a herd of captive African elephants (33 from an animal importer in Texas, and 63 young elephants collected in Zimbabwe). Group-1 elephants were purchased 8 months before the arrival of the group-2 elephants. On arrival, 7 group-1 elephants had raised nodular fibrous growths, located predominantly on their trunks. Lesions were not observed in the group-2 elephants until approximately 3 months after they were acquired. Lesions on group-2 elephants began as small focal proliferative growths that regressed or that progressed into large nodular fibrous growths that were similar in appearance to those seen in the group-1 elephants. Lesions at various stages of development were biopsied and examined. Histologically, early lesions were inverted papillomas, with hyperplastic and hypertrophic epithelial cells containing amphoteric intranuclear inclusions in the lesion center. Older, large, nodular fibrous growths were ulcerated and were composed predominantly of a thickened dermis containing fibroblasts, collagen, and a mixed inflammatory cell infiltrate; inclusions were not observed in adjacent epidermal cells. Using a peroxidase-antiperoxidase technique, we did not detect group-specific papillomavirus antigens. Southern blot hybridization analysis of DNA from lesion specimens did not indicate papillomavirus-specific genomes. Electron-microscopically, inclusions consisted of aggregates of virus particles. The particles had electron-dense and electron-lucent cores and were 95 to 103 nm in diameter. Virions developed envelopes from nuclear membranes. Mature particles were seen within the cytoplasm and filled the intercellular spaces. On the basis of size, location, conformation, and envelopment, the particles most closely resembled those of herpesviruses.


Elephant endotheliotropic herpesvirus (EEHV) is one of the most devastating infections and causes of mortality in captive Asian elephant (Elephas maximus) populations. Eight confirmed fatal EEHV cases have occurred since 1995 within the captive Asian elephant population of the United Kingdom and Ireland. This report aims to review the impact of EEHV on the captive Asian elephant population in the United Kingdom and Ireland, document and compare fatal cases, and recommend a framework of monitoring within the United Kingdom and Ireland to increase the success of treatment of EEHV hemorrhagic disease (EEHV HD) in the future. Six zoologic institutions (which include zoos, safari parks, and wildlife parks) that currently house or have previously housed a captive Asian elephant group were included in this report. Medical records and postmortem results were collected from four of these institutions for each confirmed fatal case. EEHV HD was found to be responsible for 29.6% of fatalities in Asian elephants born in captivity in the United Kingdom and Ireland between 1995 and 2013. Following a review of all the cases, it is shown that although clinical signs may be associated with specific EEHV species, the swiftness of disease progression means that most body tissues are impacted 1-6 days following the presentation of visible clinical signs and treatment is less likely to succeed. Therefore, EEHV monitoring should consist of conducting regular polymerase chain reaction analysis of whole blood samples from at-risk, young Asian elephants aged 1-8 yr in order for subclinical viremia to be identified early and treatment to be started before the appearance of visible clinical signs. © Copyright 2016 by American Association of Zoo Veterinarians.


Systemic infections with elephant endotheliotropic herpesviruses (EEHV) cause a rapid onset acute hemorrhagic disease with an 85% mortality rate. More than 60 cases have been confirmed worldwide occurring predominantly in juvenile Asian elephants. Originally, three virus types EEHV1A, EEHV1B and EEHV2 were identified, all members of the Proboscivirus genus within the Betaherpesvirinae. However, four elephant gammaherpesviruses (EGHV) have also been found by DNA PCR approaches in eye and genital secretions of asymptomatic animals, and two more versions of the probosciviruses, EEHV3 and EEHV4, were recently detected in acute hemorrhagic disease cases. To ask whether even more species of elephant herpesviruses may
exist, we have developed several new diagnostic DNA PCR assays using multiple round primers in the DNA POL region. These have been used routinely for nearly three years to screen samples submitted to the Elephant Herpesvirus Laboratory for diagnosis of possible cases of EEHV disease in blood and necropsy tissue, as well as in biopsies of other suspicious lesions or growths. Several more cases of EEHV1-associated hemorrhagic disease were confirmed, but in addition, we describe here eleven examples of other known and novel herpesviruses detected and evaluated with these reagents. They include the prototypes of four new elephant herpesviruses, two more within the proboscivirus group EEHV5 and EEHV6, plus two more gammaherpesviruses EGHV3B and EGHV5. We also report initial semi-quantitative PCR assays demonstrating very high viral loads in the blood of the EEHV3 and EEHV4-associated hemorrhagic disease cases.


The study was aimed at characterizing elephant endotheliotropic herpesvirus (EEHV) that was detected in captive Asian elephants in Thailand from 2007 to 2013. Six tissue samples of dead elephants and two EDTA blood samples of surviving elephants in Thailand showed clinical signs or had lesions of the viral infection. Samples were extracted for DNA amplification using a PCR technique with strain specific primers based on terminase and DNA polymerase genes. Six samples gave positive amplicons for EEHV1 specific primers and two samples gave positive amplicons for EEHV3/4 specific primers. Nucleotide sequencing analysis was assured for strain identification. Five out of the six samples from EEHV1 PCR were positive for the EEHV1A strain and one sample was positive for the EEHV1B strain. The two samples of EEHV3/4 PCR positive products were revealed to be of the EEHV4 strain based on the sequencing of the partial terminase gene. Three strains of the EEHV including EEHV1A, EEHV1B and EEHV4 have been detected in Asian elephants in Thailand from 2007 to 2013. This study revealed the first EEHV1B isolate that has been detected in a captive Asian elephant in Thailand.


Elephant endotheliotropic herpesvirus 1A is a member of the Proboscivirus genus and is a major cause of fatal hemorrhagic disease in endangered juvenile Asian elephants worldwide. Here, we report the first complete genome sequence from this genus, obtained directly from necropsy DNA, in which 60 of the 115 predicted genes are not found in any known herpesvirus.


Lancing a finger elicits minimal pain in humans and is applied routinely to obtain small volumes of blood for clinical diagnostics. A modified lancet bleeding method and several blood sampling matrices were evaluated in this study for the purpose of routine elephant endotheliotropic herpesvirus (EEHV) surveillance in Asian elephants (Elephas maximus). The procedure enabled weekly sampling from elephants as young as 9 mo of age. The blood sampling matrices were evaluated for their sensitivity measuring beta-actin, tumor necrosis factor alpha, and/or EEHV-1 by quantitative polymerase chain reaction assays. Foam and flocked swabs produced significantly (P < 0.05) lower quantitation cycles, ie, increased analytical sensitivity, than filter papers, Whatman(R) FTA cards, or conventional cotton-tipped swabs. The two swab types also demonstrated comparable analytical sensitivity to that of a similar volume of EDTA whole blood for the detection of EEHV-1 DNA. This lancet-and-swab technique proved satisfactory for the detection of EEHV-1 viremia in two Asian elephant calves, and in one instance viremia could be detected 5 days prior to the development of clinical signs. Low blood yield from the lancet application may reduce sensitivity and compromise early detection of viremia. Therefore, standard venipuncture remains the recommended blood sampling method, and training for consistent and regular vein access should continue to be the priority for collections holding elephants. However, if appropriate measures are taken to collect an optimum blood volume, this lancet-and-swab technique offers a
suitable alternative for EEHV surveillance in situations where venipuncture may not be practical.


Full text: The lungs of the majority of 50 elephants examined recently in the Kruger National Park had several lymphoid nodules varying in size from 3 to 30 mm (Fig. 1 arrow). The alveolar lining cells within these nodules were metaplastic and contained Type A intranuclear inclusions (Fig. 2). Electronmicroscopy revealed the presence of a virus in the epithelial cells. It was subsequently successfully isolated in tissue culture and characterized as a herpes virus.


Lymphoid nodules associated with Cowdry Type A intranuclear inclusions in epithelial and syncytial cells were found in the lungs of 74% of 50 African elephants in the Kruger National Park. Subsequent studies proved these were caused by a herpes virus (Erasmus, McCully, Pienaar, Young, Pieterse & Els, 1971). The disease appears to be subclinical or latent. This virus, in common with other herpes viruses, might be more pathogenic in some other host. The pathogenesis of the lymphoid nodules and the various stages of their formation are given and the detailed characteristics are illustrated.


In mid 1988 a 3-yr-old Asian elephant (Elephas maximus) from a circus in Switzerland died following generalized manifestation of a herpesvirus infection. In an effort to determine prevalence of infection with the herpesvirus, and due to lack of a corresponding virus isolate, it was decided to evaluate contact animals and elephants from a second herd for antibody to bovine herpesvirus 1 (BHV1) and bovine herpesvirus 2 (BHV2). Of 15 sera tested four displayed low neutralizing antibody titers to BHV2. None of the sera neutralized BHV1. However, as evidenced by protein A-mediated immunoprecipitation of metabolically radio-labeled virus-infected and mock-infected cell antigens, followed by separation of precipitation products in SDS-polyacrylamide gels, the 15 sera precipitated multiple antigens from both viruses. Similar results were obtained when using BHV4 antigens. The extent of reaction was most distinct with respect to BHV2 antigens, less prominent with BHV1 antigens, and least with BHV4 antigens. The respective protein patterns, although less marked, matched well with those obtained with bovine reference sera. Additional evaluation of sera from six elephants from two zoos in the Federal Republic of Germany gave essentially identical results. It was concluded that at least one herpesvirus, immunologically related to BHV2, may be widely distributed among captive Asian elephants, and that this virus apparently does not cause overt disease in the majority of animals.


Elephant endotheliotropic herpesvirus (EEHV) infections and tuberculosis have emerged as causes of illness and mortality in captive elephants. Twenty-six confirmed EEHV cases are documented. Since 1995, 7 have occurred in North America, 10 in Europe and 2 in Asia. A PCR test was used to detect the virus in symptomatic animals; a serological test to identify carrier elephants is under development. The African elephant is a potential source of the EEHV that is lethal for Asian elephants. Fatal infections have also occurred in Asian elephants without African elephant contacts. Three of 6 elephants recovered after treatment with antiviral famciclovir; however, more research is needed to improve the usefulness of this drug. Asian elephants that are less than 10-years old and have been moved to another facility and/or have had contact with African elephants are at increased risk for contracting EEHV. Animals traveling between facilities with a history of EEHV cases may be at greater risk. All young elephants should be monitored daily for anorexia, lethargy, body swellings and blue discoloration.
(bruising) of the tongue, and be trained for blood sampling and potential oral and rectal treatment with famciclovir.

Since 1996, *Mycobacterium tuberculosis* has affected about 3% of Asian elephants in North America. Most were from 5 U.S. States with some contacts between private herds. Mandatory annual testing for tuberculosis by trunk wash cultures was established in 1998, and 22 culture-positive *M. tuberculosis* elephants were identified between 1996-2001. Fifteen were treated with anti-tuberculosis drugs and 7 that died or were euthanized were proven to have tuberculosis at necropsy. Antemortem sera was available from 4/7 4 (75%) were strongly ELISA positive. Tuberculosis is uncommon in African elephants but was recently associated with *M. bovis* in the U.S. and *M. tuberculosis* in Germany. Conversely, *M. bovis* tuberculosis, apparently unrecognized in Asian elephants, recently occurred in Germany. Management issues of elephant tuberculosis will be discussed relative to its complex epidemiology and clinical-pathological correlations.


A 21-year-old male African elephant (*Loxodonta africana*) died suddenly with no previous medical history. Grossly, there were severe multifocal epicardial and endocardial hemorrhages of the atria and ventricles, hydropericardium, multifocal pleural hemorrhages, and severe pulmonary congestion and edema. Histologically, there was fibrinoid vasculitis and thrombosis in the heart and lung and myocardial necrosis. *Citrobacter freundii* was isolated in abundance in pure culture from liver and heart samples. Low levels of multiples types of elephant endotheliotropic herpesvirus (EEHV-6, EEHV-2B, and EEHV-3A) were detected in spleen samples, but not in heart samples. The levels of EEHV DNA found were much lower than those usually associated with acute EEV hemorrhagic disease, and many other genomic loci that would normally be found in such cases were evidently below the level of detection. Therefore, these findings are unlikely to indicate lethal EEHV disease. Polymerase chain reaction for encephalomyocarditis virus (EMCV) and toxicology for oleander (*Nerium oleander*) were negative. Stress, resulting from recent transport, and antimicrobial therapy may have contributed to the death of this animal.


Infections with herpesvirus may cause papillomatous lesions in the Asian and African elephant. In both species, the virus has been reported to localize only in the skin. Disseminated nodules of epithelial cells were found in the lungs of a high percentage of wild African elephants. In these cases, the proliferated cells contained intranuclear inclusion bodies in which herpesvirus particles were observed by electron microscopy. The virus in those cases caused no illness. This report documents the necropsy findings of a juvenile Asian elephant dying peracutely from massive generalized hemorrhage due to lesions in the endothelial cells of the capillaries. The cell nuclei frequently contained inclusion bodies in which herpesvirus particles were demonstrated. This has not been described in elephants before.


Elephant populations are under intense pressure internationally from habitat destruction and poaching for ivory and meat. They also face pressure from infectious agents, including elephant endotheliotropic herpesvirus 1 (EEHV1), which kills ~20% of Asian elephants (*Elephas maximus*) born in zoos and causes disease in the wild. EEHV1 is one of at least six distinct EEHV in a phylogenetic lineage that appears to represent an ancient but newly recognized subfamily (the Deltaherpesvirinae) in the family Herpesviridae.


Elephant endotheliotropic herpesviruses (EEHVs) can cause fatal hemorrhagic disease in Asian and African elephants. There are quantitative real time PCR (qPCR) tests that can detect seven known EEHVs (1A, 1B, 2-6) in
Herpesvirus References

Elephant Care International Database

mucosal secretions, tissue isolates, and blood samples. However, current qPCR tests are unable to distinguish between EEHV 1A and 1B or 3 and 4. To address these inadequacies, new qPCR assays were generated and validated to specifically detect EEHV 1A, 1B, and 4. Each assay demonstrated robust efficiency, a broad linear range, and low intra- and inter-assay variability. Each also proved to be specific for its EEHV target when tested against known banked samples from past EEHV cases. The EEHV1A and 1B assays were then used to characterize an eight-week, low level EEHV1 viremic event in a young Asian elephant. These new tests will allow veterinarians and researchers to pinpoint the specific species causing infection more rapidly. They will also allow veterinarians and elephant keepers to better characterize the EEHV status of each animal within their herd leading to more informed management strategies.


There are many case reports of elephant pathogenic bacteria and viruses that require quick and sensitive diagnostic techniques due to the impact they generate.

Out of these the occurrence of TB in elephants, especially in captivity, leading to zoonotic risk for humans who live at the animal-human interface and the different strains of elephant endotheliotropic herpes virus (EEHV) that pose a threat to Asian elephants are of extreme importance. Hence, this study aims to evaluate the PCR based molecular techniques for the rapid and direct detection of TB in captive elephants by primers targeting gene hsp65 and EEHV 1 strain by primers targeting the terminase gene. Serologically positive captive Asian elephants at Elephant orphanage, Pinnawala were screened for TB by specific primer PCR assay for hsp65 gene of M.tuberculosis using direct DNA isolates from trunk wash samples. Among 21 trunk washes, only a single amplification was observed, with a size closer to 441bp. Sequencing of this resulted a 415bp fragment which was not responsible for TB. Although, there have been no recorded cases of EEHV in Sri Lanka, many healthy Asian elephants are asymptotically infected by EEHV1 in the neighboring Indian region. Therefore, asymptomatic Asian elephants in captivity at ETH, Udawalawe were screened for 336bp partial EEHV1 terminase gene using direct DNA isolates from blood, eye swabs and buccal cavity swabs. All tested samples were negative for EEHV1. Since these elephants were closely monitored even after the study and none of them developed classical symptoms of either EEHV or TB, it is difficult to prove the fact that they were originally infected. The nonspecific amplification proves that it is possible to extract microbial DNA from elephant trunk washes.


Since 1995, 4 suspected cases of Endotheliotropic Elephant Herpes Virus (EEHV) infection, i.e. based on clinical presentation, have occurred in Asia without resulting in epidemic outbreaks as expected. In order to confirm the presence of EEHV on the continent of Asia, viral DNA particles from liver samples of a wild-caught 3-year-old elephant found dead at a Cambodian elephant sanctuary and clinically diagnosed with EEHV, were PCR processed using known EEHV strain primers. The presence of EEHV viral nucleic acids was confirmed and the nucleic acids had a 99% sequence similarity to the U.S.A strain (gene bank locus: AF117265) and 97% sequence similarity to the European strain (gene bank locus: AF354746) assigning this case to the EEHV-1 cluster. More than the confirmation of EEHV on the continent of Asia, is the phylogenic relationship to the USA and European strains with no corresponding contact or transport of USA or European elephants to Asia. Thus, this brings many of the traditional theories into question. Although almost forgotten, this disease is still ramped in captive elephant populations worldwide and continues to devastate particularly the neonatal and weaning-age population. Special attention and continued research are needed specifically in the area of basic virology and epidemiology.


There are two newly recognized herpesviruses that cause a fatal disease syndrome in elephants. They are known as the elephant endotheliotropic herpesviruses, of which one is fatal for Asian elephants (Elephas maximus) and the other for African elephants (Loxodonta africana) The disease syndrome affects predominantly young elephants and has been described in North America, Europe, and Israel. The predominant clinical signs for both species include lethargy, oedematous swellings of the head, neck, and thoracic limbs, oral ulceration, cyanosis of the tongue, and death of most elephants in 1-7 days. Three affected young Asian elephants recovered after a 3-4-week course of therapy with the anti-herpesvirus drug famciclovir. Additional reported herpesvirus-associated lesions in otherwise healthy elephants include localized skin papillomas in African elephants, proliferative vulval lymphoid patches in African elephants, and pulmonary nodules in African elephants. Recent findings suggest that these localized herpesvirus-associated lesions in healthy African elephants may be one source of the herpesvirus that causes disseminated disease and death in the Asian species and the African species. These findings have implications for management practices in facilities keeping both African and Asian elephants and in protecting natural elephant habitats from virulent forms of the virus.


The unique clinical and pathological findings in nine Asian elephants (Elephas maximus) and two African elephants (Loxodonta africana) from North American Zoos with a highly fatal disease caused by novel endotheliotropic herpesviruses are described. Consensus primer polymerase chain reaction combined with sequencing yielded molecular evidence that confirmed the presence of 2 novel but related herpesviruses associated with the disease, one in Asian elephants and the other in African elephants. Disease onset was acute, with lethargy, edema of the head and thoracic limbs, oral ulceration and cyanosis of the tongue followed by death of most animals in 1 to 7 days. Pertinent laboratory findings in 2 of 3 clinically evaluated animals included lymphocytopenia and thrombocytopenia. Two affected young Asian elephants recovered after a 3- to 4-week course of therapy with famciclovir. PM examination in the fatal cases revealed pericardial effusion and extensive petechial hemorrhages in the heart and throughout the peritoneal cavity, hepatomegaly, cyanosis of the tongue, intestinal hemorrhage and ulceration. Histologically, there were extensive microhemorrhages and edema throughout the myocardium and mild, subacute myocarditis. Similar hemorrhagic lesions with inflammation were evident in the tongue, liver and large intestine. Lesions in these target organs were accompanied by amphophilic to basophilic intranuclear viral inclusion bodies in capillary endothelial cells. Transmission electron microscopy of the endothelial inclusion bodies revealed 80 to 92 nm diameter viral capsids consistent with herpesvirus morphology. The short course of the herpesvirus infections, with sudden deaths in all but the 2 surviving elephants, was ascribed to acute cardiac failure attributed to herpesvirus-induced capillary injury with extensive myocardial hemorrhage and edema.


A highly fatal haemorrhagic disease was identified in 10 young Asian (Elephas maximus) and African (Loxodonta africana) elephants at zoos in the USA between 1983 and 1997. In the affected animals there was ultrastructural evidence for herpesvirus-like particles in endothelial cells of the heart, liver, and tongue. Consensus primer polymerase chain reaction combined with sequencing yielded molecular evidence that confirmed the presence of 2 novel but related herpesviruses associated with the disease, one in Asian elephants and another in African elephants. Otherwise healthy African elephants with external herpetic lesions yielded herpesvirus sequences identical to that found in Asian elephants with endothelial disease. It is suggested that the Asian elephant deaths were caused by cross-species infection with a herpesvirus that is naturally latent in, but normally not lethal to, African elephants. A reciprocal relationship may exist for the African elephant disease.

A family of novel endotheliotropic herpesviruses (EEHVs) assigned to the genus Proboscivirus have been identified as the cause of fatal hemorrhagic disease in 70 young Asian elephants worldwide. Although EEHV cannot be grown in cell culture, we have determined a total of 378 kb of viral genomic DNA sequence directly from clinical tissue samples from six lethal cases and two survivors. Overall, the data obtained encompass 57 genes, including orthologues of 32 core genes common to all herpesviruses, 14 genes found in some other herpesviruses, plus 10 novel genes, including a single large putative transcriptional regulatory protein (ORF-L). On the basis of differences in gene content and organization plus phylogenetic analyses of conserved core proteins that have just 20% to 50% or less identity to orthologues in other herpesviruses, we propose that EEHV1A, EEHV1B, and EEHV2 could be considered a new Deltaherpesvirinae subfamily of mammalian herpesviruses that evolved as an intermediate branch between the Betaherpesvirinae and Gammaherpesvirinae. Unlike cytomegaloviruses, EEHV genomes encode ribonucleotide kinase B subunit (RRB), thymidine kinase (TK), and UL9-like origin binding protein (OBP) proteins and have an alphaherpesvirus-like dyad symmetry Ori-Lyt domain. They also differ from all known betaherpesviruses by having a 40-kb large-scale inversion of core gene blocks I, II, and III. EEV1 and EEHV2 DNA differ uniformly by more than 25%, but EEHV1 clusters into two major subgroups designated EEHV1A and EEHV1B with ancient partially chimeric features. Whereas large segments are nearly identical, three nonadjacent loci totaling 15 kb diverge by between 21 and 37%. One strain of EEHV1B analyzed is interpreted to be a modern partial recombinant with EEHV1A.

IMPORTANCE: Asian elephants are an endangered species whose survival is under extreme pressure in wild range countries and whose captive breeding populations in zoos are not self-sustaining. In 1999, a novel class of herpesviruses called EEHVs was discovered. These viruses have caused a rapidly lethal hemorrhagic disease in 20% of all captive Asian elephant calves born in zoos in the United States and Europe since 1980. The disease is increasingly being recognized in Asian range countries as well. These viruses cannot be grown in cell culture, but by direct PCR DNA sequence analysis from segments totaling 15 to 30% of the genomes from blood or necropsy tissue from eight different cases, we have determined that they fall into multiple types and chimeric subtypes of a novel Proboscivirus genus, and we propose that they should also be classified as the first examples of a new mammalian herpesvirus subfamily named the Deltaherpesvirinae.


Managers of cooperative breeding programs and re-introduction projects are increasingly concerned with the risk of disease transmission when specimens are transferred among facilities or between facilities and the natural environment. We used data maintained in North American studbooks to estimate the potential risks of disease transmission by direct and indirect contact of specimens in the American Zoo and Aquarium Association's Elephant Species Survival Plan. Histological evidence for a novel herpesvirus disease transmitted between and within elephant species housed in North American facilities prompted an examination of the scope of possible transmission routes within the captive population. We found that, compared with other species managed through Species Survival Plans, elephants experience relatively few transfers between zoos. Nevertheless, the number of direct contacts with other elephants born during the study period of 1983-1996 (excluding stillbirths) was much higher than we had anticipated (\(\bar{m}_{gr} = 25 +/ - 27; N = 59\)) and the number of potential indirect contacts was surprisingly large (\(\bar{m}_{gr} = 143 +/ - 92; N = 59\)). Although these high rates of potential contacts complicate exact identification of infection pathways for herpesvirus, we were able to propose potential routes of transmission for the histologically identified cases. Furthermore, the extraction of data from studbooks allowed us to readily identify other specimens that did not succumb to the disease despite similar exposure. Moreover, we were able to identify other possible cases to recommend for histological examination. Herein we reveal the possibilities of multiple disease transmission pathways and demonstrate how complex the patterns of transmission can be, confounded by the unknown latency of this novel herpesvirus. This emphasizes the need for zoo veterinarians and cooperative breeding programs to consider the full potential for
disease transmission associated with each and every inter-zoo transfer of specimens.


Elephant endotheliotropic herpesvirus 1 (EEHV1) can cause fatal hemorrhagic disease in Asian elephants (Elephas maximus). Several studies have described this virus as a major threat to young Asian elephants. A SYBR Green I-based real-time polymerase chain reaction (PCR) was developed to identify EEHV1 on trunk swabs and necropsied tissues. Two of 29 (6.9%) trunk swab samples from healthy Asian elephants were positive for EEHV1. The viruses were analyzed and classified as EEHV1A based on 231 nucleotides of the terminase gene. Necropsied spleen and heart tissue showed the highest level and second highest levels of DNA virus copy accumulation, respectively. The detection limit of the test was 276. copies/μl of DNA. There was no cross-reaction with other mammalian herpesviruses, such as herpes simplex virus 1 and equine herpesvirus 2. Inter- and intra-assay showed low coefficients of variation values indicating the reproducibility of the test. The results indicated that the test can be practically used for epidemiological study, clinical diagnosis, and management and control of EEHV1. © 2012 Elsevier B.V..


Several different strains of elephant endotheliotropic herpes virus-1 (EEHV-1) have been identified via polymerase chain reaction (PCR) techniques in both African and Asian elephants. EEHV-1 has been identified in both cutaneous lesions in healthy African elephants and fatal cases of hemorrhagic syndrome in Asian elephants. However, until now, no EEHV-1 strain has been identified or associated with otherwise healthy Asian elephants. This article describes recurrent nonendothelial lesions associated with EEHV-1 infection in a herd of Asian elephants not exhibiting fatal hemorrhagic syndrome. Genotypes of EEHV-1 strains, based on viral DNA polymerase and glycoprotein B, associated with fatal hemorrhagic syndrome, were compared to those identified in nonendothelial lesions. The same EEHV-1 genotypes were identified in fatal cases and mucosal lesions in otherwise healthy Asian elephants in this herd. Further studies of the Asian elephant immune system and virologic studies to determine the triggers of tissue tropism are needed before any conclusion can be reached. Copyright 2010 by American Association of Zoo Veterinarians.


Two juvenile Asian elephants (E. maximus) presented with an acute onset of facial oedema and lethargy. Examination of the oral cavity of each animal revealed cyanosis of the tip and distal margins of the tongue suggestive of endothelial inclusion body disease (EIBD) of elephants. Whole-blood samples were obtained, and polymerase chain reaction tests confirmed the presence of elephant herpesvirus. The animals were administered famciclovir (Flamvir; 500 mg/70 kg body weight, with a loading dose of 1000 mg/70 kg body weight) a potent human anti-herpesvirus drug, in the course of their disease, and recovery followed a treatment regime of 3-4 wk. These are the first known cases of elephants surviving EIBD.


Elephant Endotheliotropic Herpesviruses (EEHVs) are the cause of a highly fatal haemorrhagic disease in elephants primarily affecting young Asian elephants (Elephas maximus) in both captivity and in the wild. The viruses have emerged as a significant threat to Asian elephant conservation, critically affecting overall sustainability of their population. So far insight into the pathogenesis of EEHV infections has been restricted to examination of EEHV-infected tissues. However, little is known about distribution and burden of the viruses
within the organs of fatal cases, crucial elements in the understanding of the virus pathogenesis. This study was therefore undertaken to assess the extent of organ and cell involvement in fatal cases of EEHV-1A, 1B and 5 using a quantitative real-time PCR. EEHV-1 and 5 DNA were detectable in all the tissues examined, albeit with substantial differences in the viral DNA load. The highest EEHV-1A DNA load was observed in the liver, followed by the heart, thymus and tongue. EEHV-1B and 5 showed the highest DNA load in the heart, followed by tongue and liver. This study provides new insights into EEHV pathogenicity and has implications in choice of sample type for disease investigation and virus isolation.


Elephant endotheliotropic herpesvirus (EEHV) is a type of herpesvirus that causes acute hemorrhagic disease in Asian elephants (Elephas maximus) and is often fatal, especially in calves. This study describes the postmortem evaluation of two captive-born Asian elephants (2 and 3 yr of age, respectively) diagnosed with EEHV in Thailand. Both elephants presented only mild depression, lethargy, and anorexia before death within 24 hr of symptom onset. Necropsies were performed, and tissue samples were tested for EEHV viral presence using polymerase chain reaction. Molecular and phylogenetic evidence illustrated two types of EEHV, which were closely related to EEHV1A in Case 1 and EEHV4 in Case 2. Pathologic findings differed between the cases. More specific organ tropism was found in Case 1, where mainly the cardiovascular system was affected. In contrast, in Case 2, hemorrhages were noted in most organs, including in the gastrointestinal, respiratory, and cardiovascular systems. This report is the first to document EEHV4 in Asia and the second case of this strain to be identified in an elephant worldwide.


Elephant Endotheliotropic Herpesvirus (EEHV) is emerging as a new threat for elephant conservation, since being identified as the cause of severe, often fatal, haemorrhagic disease in young Asian elephants. To describe positive cases and the molecular relatedness of virus detected in elephants in Thailand, we re-examined all available of EEHV samples occurring in young elephants in Thailand between 2006 and 2014 (n=24). Results indicated 75% (18/24) of suspected cases were positive for EEHV by semi-nested PCR. Further gene analysis identified these positive cases as EEHV1A (72%, 13/18 cases), EEHV1B (11%, 2/18) and EEHV4 (17%, 3/18). This study is the first to phylogenetically analyse and provide an overview of most of the known EEHV cases that have occurred in Thailand. Positive individuals ranged in age from one to nine years, with no sex association detected, and occurred across geographical locations throughout the country. All individuals, except one, were captive-born. No history of direct contact among the cases was recorded, and this together with the fact that various subtype clusters of virus were found, implied that none of the positive cases were epidemiologically related. These results concur with the hypothesis that EEHV1 is likely to be an ancient endogenous pathogen in Asian elephants. It is recommended that active surveillance and routine monitoring for EEHV should be undertaken in all elephant range countries, to gain a better understanding of the epidemiology, transmission and prevention of this disease.


This article describes the treatment of clinical elephant endotheliotropic herpesvirus (EEHV) infection in a male Asian elephant (Elephas maximus; approximately 3 yr old), the dynamics of viral load during the active infection, and genetic analysis of the virus. Treatment included injectable acyclovir (12 mg/kg iv, bid), antibiotic, vitamin, and fluids. Quantitative polymerase chain reaction was used to measure the viral levels in blood, which decreased continuously after initiation of intravenous acyclovir. Low levels of virus were detected in the blood for 2 wk, and the virus was undetectable after 1 mo. No complication was observed during the treatment
period. This case report suggests that acyclovir, given parenterally, could potentially enhance survival of clinical EEHV-infected individuals.


Objective-To investigate the pathogenesis and transmission of elephant endotheliotropic herpesvirus (EEHV1) by analyzing various elephant fluid samples with a novel EEHV1-specific real-time PCR assay.

Animals-5 apparently healthy captive Asian elephants (Elephas maximus) from the same herd.

Procedures-A real-time PCR assay was developed that specifically detects EEHV1. The assay was used to evaluate paired whole blood and trunk-wash samples obtained from the 5 elephants during a 15-week period.

Deoxyribonucleic acid sequencing and viral gene subtyping analysis were performed on trunk-wash DNA preparations that had positive results for EEHV1. Viral gene subtypes were compared with those associated with past fatal cases of herpesvirus-associated disease within the herd.

Results-The PCR assay detected viral DNA to a level of 1,200 copies/mL of whole blood. It was used to detect EEHV1 in trunk secretions of 3 of the 5 elephants surveyed during the 15-week period. Viral gene subtyping analysis identified 2 distinct elephant herpesviruses, 1 of which was identical to the virus associated with a previous fatal case of herpesvirus-associated disease within the herd.

Conclusions and Clinical Relevance-EEHV1 was shed in the trunk secretions of healthy Asian elephants. Trunk secretions may provide a mode of transmission for this virus. Results of this study may be useful for the diagnosis, treatment, and management of EEHV1-associated disease and the overall management of captive elephant populations.


Elephant endotheliotropic herpesviruses (EEHVs) can cause lethal hemorrhagic disease in both African and Asian elephants. At least seven EEHV types have been described, and sensitive real-time PCR tests have been developed for EEHV1A and 1B, which are associated with the majority of characterized Asian elephant deaths. Despite growing knowledge of the different EEHV types, the prevalence of each type within African and Asian elephants remains to be determined and there is considerable need for diagnostic tests to detect and discriminate between each EEHV species for clinical management of African and Asian elephants that develop illness from one or more of these viruses. To begin to address these issues, we developed real-time PCR assays for EEHV2, 3, 4, 5, and 6. Overall, each assay had robust PCR efficiency, a dynamic linear range over 5log(10) concentrations, a limit of detection of 10 copies/test reaction with 100% sensitivity, and low intra- and inter-assay variability. Each assay proved to be specific for the EEHV targets for which it was designed, with the exception of EEHV3 and EEHV4, which was expected because of greater DNA sequence similarity between these two EEHV species than the others. These new tools will be useful for conducting surveys of EEHV prevalence within captive and range country elephants, for diagnostic testing of elephants with suspected EEHV-associated disease, and for managing the treatment of elephants with EEHV-induced illness.


OBJECTIVE: To investigate the pathogenesis and transmission of elephant endotheliotropic herpesvirus (EEHV1) by analyzing various elephant fluid samples with a novel EEHV1-specific real-time PCR assay. ANIMALS: 5 apparently healthy captive Asian elephants (Elephas maximus) from the same herd. PROCEDURES: A real-time PCR assay was developed that specifically detects EEHV1. The assay was used to evaluate paired whole blood and trunk-wash samples obtained from the 5 elephants during a 15-week period. Deoxyribonucleic acid
sequencing and viral gene subotyping analysis were performed on trunk-wash DNA preparations that had positive results for EEHV1. Viral gene subtypes were compared with those associated with past fatal cases of herpesvirus-associated disease within the herd. RESULTS: The PCR assay detected viral DNA to a level of 1,200 copies/mL of whole blood. It was used to detect EEHV1 in trunk secretions of 3 of the 5 elephants surveyed during the 15-week period. Viral gene subtyping analysis identified 2 distinct elephant herpesviruses, 1 of which was identical to the virus associated with a previous fatal case of herpesvirus-associated disease within the herd. CONCLUSIONS AND CLINICAL RELEVANCE: EEHV1 was shed in the trunk secretions of healthy Asian elephants. Trunk secretions may provide a mode of transmission for this virus. Results of this study may be useful for the diagnosis, treatment, and management of EEHV1-associated disease and the overall management of captive elephant populations.


Elephant endotheliotropic herpesviruses (EEHVs) can cause fatal hemorrhagic disease in juvenile Asian elephants (Elephas maximus); however, sporadic shedding of virus in trunk washes collected from healthy elephants also has been detected. Data regarding the relationship of viral loads in blood compared with trunk washes are lacking, and questions about whether elephants can undergo multiple infections with EEHVs have not been addressed previously. Real-time quantitative polymerase chain reaction was used to determine the kinetics of EEHV1 loads, and genotypic analysis was performed on EEHV1 DNA detected in various fluid samples obtained from five Asian elephants that survived detectable EEHV1 DNAemia on at least two separate occasions. In three elephants displaying clinical signs of illness, preclinical EEHV1 DNAemia was detectable, and peak whole-blood viral loads occurred 3-8 days after the onset of clinical signs. In two elephants with EEHV1 DNAemia that persisted for 7-21 days, no clinical signs of illness were observed. Detection of EEHV1 DNA in trunk washes peaked approximately 21 days after DNAemia, and viral genotypes detected during DNAemia matched those detected in subsequent trunk washes from the same elephant. In each of the five elephants, two distinct EEHV1 genotypes were identified in whole blood and trunk washes at different time points. In each case, these genotypes represented both an EEHV1A and an EEHV1B subtype. These data suggest that knowledge of viral loads could be useful for the management of elephants before or during clinical illness. Furthermore, sequential infection with both EEHV1 subtypes occurs in Asian elephants, suggesting that they do not elicit cross-protective sterilizing immunity. These data will be useful to individuals involved in the husbandry and clinical care of Asian elephants.


Elephant endotheliotropic herpesviruses (EEHVs) can cause fatal hemorrhagic disease in Asian (Elephas maximus) and African (Loxodonta africana) elephants. Of the seven known EEHV species, EEHV1 is recognized as the most common cause of hemorrhagic disease among Asian elephants in human care worldwide. Recent data collected from ex situ Asian elephants located in multiple North American and European institutions suggest that subclinical EEHV1 infection is common in this population of elephants. Although fatal EEHV1-associated hemorrhagic disease has been reported in range countries, data are lacking regarding the prevalence of subclinical EEHV infections among in situ Asian elephants. We used previously validated EEHV-specific quantitative real-time PCR assays to detect subclinical EEHV infection in three regionally distinct Asian elephant cohorts, totaling 46 in situ elephants in South India, during October and November 2011. Using DNA prepared from trunk washes, we detected EEHV1, EEHV3/4, and EEHV5 at frequencies of 7, 9, and 20% respectively. None of the trunk washes was positive for EEHV2 or 6. At least one EEHV species was detectable in 35% (16/46) of the samples that were screened. These data suggest that subclinical EEHV infection among in situ Asian elephants occurs and that Asian elephants may be natural hosts for EEHV1, EEHV3 or 4, and EEHV5, but not EEHV2 and...
EEHV6. The methodology described in this study provides a foundation for further studies to determine prevalences of EEHV infection in Asian elephants throughout the world. © Wildlife Disease Association 2014.


Infection of Asian elephants (Elephas maximus) with elephant endotheliotropic herpesvirus (EEHV) can be associated with rapid, lethal hemorrhagic disease and has been documented in elephant herds in human care and in the wild. Recent reports describe real-time quantitative polymerase chain reaction (qPCR) assays used to monitor clinically ill elephants and also to detect subclinical EEHV1 infection in apparently healthy Asian elephants. Acute phase proteins have been demonstrated to increase with a variety of infectious etiologies in domesticated mammals but have not yet been described in elephants. In addition, the immune response of Asian elephants to EEHV1 infection has not been described. In this study, whole blood and trunk wash samples representing repeated measures from eight elephants were examined for the presence of EEHV1 using a qPCR assay. Elephants were classified into groups, as follows: whole blood negative and positive and trunk wash negative and positive. Serum amyloid A (SAA) and haptoglobin (HP) levels were compared between these groups. A significant difference in SAA was observed with nearly a threefold higher mean value during periods of viremia (P = 0.011). Higher values of SAA were associated with >10,000 virus genome copies/ml EEHV1 in whole blood. There were no significant differences in HP levels, although some individual animals did exhibit increased levels with infection. These data indicate that an inflammatory process is stimulated during EEHV1 viremia. Acute phase protein quantitation may aid in monitoring the health status of Asian elephants. © American Association of Zoo Veterinarians.


The captive population of Asian elephants (Elephas maximus) is not self-sustaining. Poor reproduction and high juvenile mortality are key factors in the decreasing population. Infection with endotheliotropic elephant herpes virus (EEHV) is one of the major causes of death in the captive population, and has resulted in the loss of at least 40 captive animals. EEHV has been responsible for the peracute death of two juvenile males at Zurich Zoo, Switzerland. Mortality due to peracute infection with EEHV mainly is seen in juveniles. Early detection of characteristic clinical signs of EEHV and immediate initiation of therapy are of crucial importance due to its rapid progression. Based on past fatal EEHV experiences, Zurich Zoo modified its daily clinical health monitoring program to increase staff awareness of EEHV infection. Examinations have been incorporated into the daily routine and include daily evaluation of behaviour, appetite, colour of mucosal membranes and the measurement of body temperature; these examinations are performed by keepers. In our experiences, characteristic signs of acute EEHV infection are lethargy, anorexia, mild colic, and cyanosis of the mucosal membranes. Results of temperature measurements have shown that best estimations of body temperature are done by measurement of the temperature in the centre of a fecal ball 5-9 min after defecation. Mean values of 36.5°C (± 0.2°C SD) are within published reference values, although adult elephants have shown significantly lower body temperature than juveniles. Establishment of individual reference values for each elephant is essential to detect unusual temperature peaks that may indicate possible EEHV viremia. The present study has shown that daily health examinations increase the awareness of keepers for early signs of EEHV infection (e.g., peaks in body temperature and cyanotic mucosal membranes).

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LITERATURE CITED

Wellehan, J. F. X., A. J. Johnson and R. Isaza (2006). Identification of two novel herpesviruses associated with ocular inflammation in Asian elephants [Elephas maximus]. 2006 Proceedings American Association of Zoo Veterinarians. Disease caused by a herpesvirus (EEHV) is a serious concern in Asian elephant (Elephas maximus) calves. Herpesviruses are known for latency and life-long infections, with periodic shedding from mild inflammatory lesions in adapted adult hosts, and ocular disease has been seen with other herpesviruses in other species. Ocular inflammation is not uncommonly seen in Asian elephants. Degenerate PCR primers targeting a conserved region of herpesvirus DNA-dependent DNA polymerase were used to amplify products from eye swabs of eight Asian elephants with epiphora, blepharitis, and conjunctivitis. Nucleotide sequencing of the PCR products showed two novel herpesviruses distinct from EEHV. Comparative sequence analysis shows that these viruses are probable members of the subfamily Gammaherpesvirinae. The sequence phylogeny of these viruses has implications for both viral and host evolution. Further understanding and characterization of these viruses is needed to understand their role in elephant health.

Wilkie, G. S., A. J. Davison, K. Kerr, M. F. Stidworthy, S. Redrobe, F. Steinbach, A. Dastjerdi and D. Denk (2014). "First fatality associated with elephant endotheliotropic herpesvirus 5 in an asian elephant: Pathological findings and complete viral genome sequence." Scientific Reports 4. Infections of Asian elephants (Elephas maximus) with elephant endotheliotropic herpesvirus (EEHV) can cause a rapid, highly lethal, hemorrhagic disease, which primarily affects juvenile animals up to the age of four years. So far, the majority of deaths have been attributed to infections with genotype EEHV1 or, more rarely, EEHV3 and EEHV4. Here, we report the pathological characteristics of the first fatality linked to EEHV5 infection, and describe the complete viral DNA sequence. Gross post-mortem and histological findings were indistinguishable from lethal cases previously attributed to other EEHV genotypes, and the presence of characteristic herpesviral inclusions in capillary endothelial cells at several sites was consistent with the diagnosis of acute EEHV infection. Molecular analysis confirmed the presence of EEHV5 DNA and was followed by sequencing of the viral genome directly from post-mortem material. The genome is 180,800 bp in size and contains 120 predicted protein-coding genes, five of which are fragmented and presumably nonfunctional. The seven families of paralogous genes recognized in EEHV1 are also represented in EEHV5. The overall degree of divergence (37%) between the EEHV5 and EEHV1 genomes, and phylogenetic analysis of eight conserved genes, support the proposed classification of EEHV5 into a new species (Elephantid herpesvirus 5).

Wilkie, G. S., A. J. Davison, M. Watson, K. Kerr, S. Sanderson, T. Bouts, F. Steinbach and A. Dastjerdi (2013). "Complete genome sequences of elephant endotheliotropic herpesviruses 1a and 1b determined directly from fatal cases." Journal of Virology 87(12): 6700-6712. A highly lethal hemorrhagic disease associated with infection by elephant endotheliotropic herpesvirus (EEHV) poses a severe threat to Asian elephant husbandry. We have used high-throughput methods to sequence the genomes of the two genotypes that are involved in most fatalities, namely, EEHV1A and EEHV1B (species Elephantid herpesvirus 1, genus Proboscivirus, subfamily Betaherpesvirinae, family Herpesviridae). The sequences were determined from postmortem tissue samples, despite the data containing tiny proportions of viral reads among reads from a host for which the genome sequence was not available. The EEHV1A genome is 180,421 bp in size and consists of a unique sequence (174,601 bp) flanked by a terminal direct repeat (2,910 bp). The genome contains 116 predicted protein-coding genes, of which six are fragmented, and seven paralogous gene families are present. The EEHV1B genome is very similar to that of EEHV1A in structure, size, and gene layout. Half of the EEHV1A genes lack orthologs in other members of subfamily Betaherpesvirinae, such as human cytomegalovirus (genus Cytomegalovirus) and human herpesvirus 6A (genus Roseolovirus). Notable among these are 23 genes encoding type 3 membrane proteins containing seven transmembrane domains (the 7TM family) and seven genes encoding related type 2 membrane proteins (the EE50 family). The EE50 family appears to be under intense evolutionary selection, as it is highly diverged between the two genotypes, exhibits evidence of sequence duplications or deletions, and contains several fragmented genes. The availability of the genome sequences will facilitate future research on the epidemiology, pathogenesis, diagnosis, and treatment
Up to 65% of deaths of young Asian elephants (Elephas maximus) between 3 mo and 15 yr of age in Europe and North America over the past 20 yr have been attributed to hemorrhagic disease associated with a novel DNA virus called elephant endotheliotropic herpesvirus (EEHV). To evaluate the potential role of EEHV in suspected cases of a similar lethal acute hemorrhagic disease occurring in southern India, we studied pathologic tissue samples collected from field necropsies. Nine cases among both orphaned camp and wild Asian elephants were identified by diagnostic PCR. These were subjected to detailed gene subtype DNA sequencing at multiple PCR loci, which revealed seven distinct strains of EEHV1A and one of EEHV1B. Two orphan calves that died within 3 days of one another at the same training camp had identical EEHV1A DNA sequences, indicating a common epidemiologic source. However, the high level of EEHV1 subtype genetic diversity found among the other Indian strains matches that among over 30 EEHV1 strains that have been evaluated from Europe and North America. These results argue against the previous suggestions that this is just a disease of captive elephants and that the EEHV1 virus has crossed recently from African elephant (Loxodonta africana) hosts to Asian elephants. Instead, both the virus and the disease are evidently widespread in Asia and, despite the disease severity, Asian elephants appear to be the ancient endogenous hosts of both EEHV1A and EEHV1B.


The genomes of three types of novel endotheliotropic herpesviruses (elephant endotheliotropic herpesvirus 1A [EEHV1A], EEHV1B, and EEHV2) associated with lethal hemorrhagic disease in Asian elephants have been previously well characterized and assigned to a new Proboscivirus genus. Here we have generated 112 kb of DNA sequence data from segments of four more types of EEHV by direct targeted PCR from blood samples or necropsy tissue samples from six viremic elephants. Comparative phylogenetic analysis of nearly 30 protein-encoding genes of EEHV5 and EEHV6 show that they diverge uniformly by nearly 20% from their closest relatives, EEHV2 and EEHV1A, respectively, and are likely to have similar overall gene content and genome organization. In contrast, seven EEHV3 and EEHV4 genes analyzed differ from those of all other EEHVs by 37% and have a G+C content of 63% compared to just 42% for the others. Three strains of EEHV5 analyzed clustered into two partially chimeric subgroups EEHV5A and EEHV5B that diverge by 19% within three small noncontiguous segments totaling 6.2 kb. We conclude that all six EEHV types should be designated as independent species within a proposed new fourth Deltaherpesvirinae subfamily of mammalian herpesviruses. These virus types likely initially diverged close to 100 million years ago when the ancestors of modern elephants split from all other placental mammals and then evolved into two major branches with high- or low-G+C content about 35 million years ago. Later additional branching events subsequently generated three paired sister taxon lineages of which EEHV1 plus EEHV6, EEHV5 plus EEHV2, and EEHV4 plus EEHV3 may represent Asian and African elephant versions, respectively. IMPORTANCE: One of the factors threatening the long-term survival of endangered Asian elephants in both wild range countries and in captive breeding populations in zoos is a highly lethal hemorrhagic herpesvirus disease that has killed at least 70 young Asian elephants worldwide. The genomes of the first three types of EEHVs (or probosciviruses) identified have been partially characterized in the preceding accompanying paper (L. K. Richman, J.-C. Zong, E. M. Latimer, J. Lock, R. C. Fleischer, S. Y. Heaggans, and G. S. Hayward, J. Virol. 88:13523-13546, 2014, http://dx.doi.org/10.1128/JVI.01673-14). Here we have used PCR DNA sequence analysis from multiple segments of DNA amplified directly from blood or necropsy tissue samples of six more selected cases of hemorrhagic disease to partially characterize four other types of EEHVs from either Asian or African elephants. We propose that all six types and two chimeric subtypes of EEHV belong to multiple lineages of both AT-rich and GC-rich branches within a new subfamily to be named the Deltaherpesvirinae, which evolved separately from all other mammalian herpesviruses about 100 million years ago.

More than 80 cases of lethal hemorrhagic disease associated with elephant endotheliotropic herpesviruses (EEHVs) have been identified in young Asian elephants worldwide. Diagnostic PCR tests detected six types of EEHV in blood of elephants with acute disease, although EEHV1A is the predominant pathogenic type. Previously, the presence of herpesvirus virions within benign lung and skin nodules from healthy African elephants led to suggestions that African elephants may be the source of EEHV disease in Asian elephants. Here, we used direct PCR-based DNA sequencing to detect EEHV genomes in necropsy tissue from five healthy adult African elephants. Two large lung nodules collected from culled wild South African elephants contained high levels of either EEHV3 alone or both EEHV2 and EEHV3. Similarly, a euthanized U.S. elephant proved to harbor multiple EEHV types distributed nonuniformly across four small lung nodules, including high levels of EEHV6, lower levels of EEHV3 and EEHV2, and a new GC-rich branch type, EEHV7. Several of the same EEHV types were also detected in random lung and spleen samples from two other elephants. Sanger PCR DNA sequence data comprising 100 kb were obtained from a total of 15 different strains identified, with (except for a few hypervariable genes) the EEHV2, EEHV3, and EEHV6 strains all being closely related to known genotypes from cases of acute disease, whereas the seven loci (4.0 kb) obtained from EEHV7 averaged 18% divergence from their nearest relative, EEHV3. Overall, we conclude that these four EEHV species, but probably not EEHV1, occur commonly as quiescent infections in African elephants. © 2016, American Society for Microbiology.


Foot problems potentially represent the single most important clinical disease of captive elephants. Predisposing factors include obesity, lack of exercise, nail or sole overgrowth, improper foot care, poor hygiene, inappropriate enclosure surfaces, poor conformation, malnutrition and secondary skeletal disorders such as degenerative joint disease. Furthermore, factors such as elephant management philosophy, disposition of elephants, facilities and competency of staff in caring for elephant feet will contribute significantly to the foot health of captive animals. It is important to note that these conditions are rarely reported in free-ranging elephants. The elephant toenail abscess is characterized grossly by proliferative outgrowth of "crab meat-like" tissue that may acutely rupture through the surface of the nail wall and/or adjacent cuticle or sole. True abscess formation with localized collections of suppurative material is not a consistent clinical feature. In most cases, the inciting cause of these lesions are typically not found and are likely due to one or more of the predisposing factors listed above. Once established, these frustrating lesions require extensive, intensive and prolonged medical attention. If not cared for properly, these wounds may progress to phalangeal osteomyelitis and the need for surgical intervention. Sole abscesses are equally frustrating and difficult to manage with proposed etiologies similar to toenail lesions. There are no reports in the literature describing the pathology of the classic proliferative abscess tissue of the elephant nail abscess. Although variously interpreted as fibrous or granulation tissue, the authors are unaware of previous histologic descriptions of this tissue. Biopsy samples of toenail abscess tissue from two Asian elephants (Elephas maximus) at the San Diego Wild Animal Park (SDWAP) consisted of stratified squamous epithelium arranged in columns resembling horn tubules. The predominant histologic finding was marked, near diffuse, hydroptic degeneration of keratinocytes. There were multifocal areas of suppurative inflammation with admixed bacterial colonies. Inflammatory foci comprised only a small portion of the lesion and were interpreted as the external surfaces of the biopsy with likely secondary bacterial colonization. Because descriptions of the normal histology of the elephant toenail could not be located, a grossly normal toenail from a different Asian elephant was obtained to compare histologic features with those of the toenail abscesses. Sections demonstrated formation of the toenail in a manner similar to that of the hoof of the horse and cattle with tubular, intertubular and laminar horn. Primary and secondary epidermal laminae
were identified. Proliferative lesions of horn-producing epithelium associated with ballooning degeneration and inadequate keratinization of keratinocytes, have been described in horses as equine "canker" and coronary band dystrophy. Equine canker is most commonly observed in the hind feet of draft horses and begins in the frog sometimes with extension to the sole and hoof wall. Grossly, lesions are characterized by soft white papillary to "cauliflower-like" tissue associated with a foul odor. Similar to what is noted in elephant foot problems, predisposing factors for the development of equine canker include poor hygiene or wet environmental conditions. There is a lack of gross and histologic description of the normal nail and sole tissue of the elephant and further investigations are warranted. A review of the anatomy and histology of the normal equine hoof may provide a basic understanding of the elephant nail until more specific and detailed elephant information is available. From our investigation, the authors offer that a more accurate description of the elephant toenail abscess would be proliferative pododermatitis, the term synonymous with equine canker. A more colloquial term such as "elephant canker" may be appropriate, as well. Canker in the horse is an uncommon but difficult to treat disease of the hoof. Historically, treatment options for elephant toenail abscesses include corrective trimming, superficial debridement and application of topical disinfectants or antibiotics. Others have constructed innovative sandals to treat and protect the affected sole or nail with success. The use of regional intravenous perfusion of the affected limb with antibiotics has also been successful. Since the elephant nail abscess now appears to be histologically and clinically comparable to equine canker, this novel characterization of an old disease may offer unique insight for treatment. In the least, it has provided our practice with a new list of treatment options and experienced equine clinicians for consultation who have been managing patients with a similar disease for many years. One of the Asian elephants at the SDWAP has had chronic toenail abscesses for over 2 yr. Radiographs of the affected digits, as reported by others to assess degree of involvement, have fortunately been negative for evidence of osteomyelitis. Several bacterial and fungal cultures of deep tissue biopsies and swabs of affected lesions have resulted in a mixture of organisms with no consistent single etiologic agent. Biopsies were found negative for presence of viral DNA (elephant papillomavirus and herpesvirus) by PCR. Typical elephant foot care at the SDWAP includes trimming and debriding with hoof knives, foot soaks and topical antibiotics. Although difficult, attempts are made in keeping the affected foot clean and dry. Following recommendations for the treatment of equine canker, we recently implemented the routine use of cryotherapy in all elephants with proliferative pododermatitis with improved success in the control and recession of exuberant nail lesions. The proliferative tissue of the nail is first cleaned then disinfected, debrided, trimmed with hoof knives and allowed to dry. Modified brass branding tools with contact surfaces of variable size (2-5 cm diameter) and shape (round or ovoid) are placed into liquid nitrogen (-196 C) for several minutes and then placed directly on the cankerous tissue for 30-60 sec. This process is then repeated 4-5 min later, following a complete thaw of tissue. Within 24 hr, the cryoburned tissue becomes macerated and necrotic and is readily removed with gentle scrubbing. Cryotherapy offers the advantage of destroying tissue to a deeper level than trimming alone and provides hemostasis, as well. Because of decreased sensation at the cryotherapy treatment site, a memorable painful event is avoided and the elephant patient is more routinely accepting of this technique. With the use of hoof knives, we typically remove 2-3 mm of proliferative tissue before the patient refuses further treatment, presumably due to discomfort. With cryotherapy, we are able to remove an additional 3-5 mm of tissue by cell freezing and necrosis. The result is quicker resolution of cankerous lesions without the need for aggressive, and potentially painful, interventions. In conclusion, it appears that elephant nail abscesses can best be described as proliferative pododermatitis, or canker, as is seen in other species. Further gross and microscopic descriptions of normal and pathologic nail or sole lesions are necessary. Routine cryotherapy has shown promise in the treatment of these chronic, frustrating and potentially devastating lesions of our captive elephants.