

Detection of Antibodies to Lokern, Main Drain, St. Louis Encephalitis, and West Nile Viruses in Vertebrate Animals in Chihuahua, Guerrero, and Michoacán, Mexico

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Abstract

We conducted serologic surveillance for flaviviruses and orthobunyaviruses in vertebrate animals in Mexico in 2018–2019. Sera were collected from 856 vertebrate animals, including 323 dogs, 223 horses, and 121 cows, from 16 species. The animals were from 3 states: Chihuahua in northwest Mexico (704 animals) and Guerrero and Michoacán on the Pacific Coast (27 and 125 animals, respectively). Sera were assayed by plaque reduction neutralization test using four flaviviruses (dengue type 2, St. Louis encephalitis, West Nile, and Zika viruses) and six orthobunyaviruses from the Bunyamwera (BUN) serogroup (Cache Valley, Lokern, Main Drain, Northway, Potosi, and Tensaw viruses). Antibodies to West Nile virus (WNV) were detected in 154 animals of 9 species, including 89 (39.9%) horses, 3 (21.4%) Indian peafowl, and 41 (12.7%) dogs. Antibodies to St. Louis encephalitis virus (SLEV) were detected in seven animals, including three (0.9%) dogs. Antibodies to Lokern virus (LOKV) were detected in 22 animals: 19 (8.5%) horses, 2 (1.7%) cows, and a dog (0.3%). Antibodies to Main Drain virus (MDV) were detected in three (1.3%) horses. WNV and LOKV activity was detected in all three states, SLEV activity was detected in Chihuahua and Michoacán, and MDV activity was detected in Chihuahua. None of the animals was seropositive for Cache Valley virus, the most common and widely distributed BUN serogroup virus in North America. In conclusion, we provide serologic evidence that select flaviviruses and BUN serogroup viruses infect vertebrate animals in Chihuahua, Guerrero, and Michoacán. We also provide the first evidence of LOKV and MDV activity in Mexico.

Keywords: flavivirus, orthobunyavirus, Mexico, vertebrate animals, surveillance, serology

Introduction

MANY ARTHROPOD-BORNE VIRUSES (arboviruses) of veterinary significance occur in North America. Some of these viruses are classified in the genus *Flavivirus* and others

are classified in the Bunyamwera (BUN) serogroup of the genus *Orthobunyavirus*. The most prevalent and widely distributed flavivirus of veterinary importance in North America is West Nile virus (WNV), which occurs throughout Mexico, the contiguous United States and southern Canada (Elizondo-Quiroga

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and Elizondo-Quiroga 2013, Zheng et al. 2014, Petersen 2019). The most prevalent and widely distributed BUN serogroup virus of veterinary importance in North America is Cache Valley virus (CVV), with isolations made as far north as Canada and as far south as the state of Yucatan, Mexico (Thorsen et al. 1980, Calisher et al. 1986, Blitvich et al. 2012a).

The introduction of WNV into the Americas in 1999 was associated with extensive morbidity and mortality in horses in the United States and Canada (Castillo-Olivares and Wood 2004). Surprisingly, there have been few cases of WNV disease in horses in Mexico, despite serological evidence of widespread WNV activity (Blitvich 2008, Elizondo-Quiroga and Elizondo-Quiroga 2013). Several licensed WNV vaccines are now available for horses (Bosco-Lauth and Bowen 2019, Saiz 2020). These vaccines are routinely used in the United States and Canada, but their use in Mexico is sporadic. Domestic ruminants are also susceptible to WNV infection and there have been occasional reports of encephalitis in alpacas and sheep (Dunkel et al. 2004, Rimoldi et al. 2017). WNV infections in companion animals are usually subclinical but fatal cases have occurred (Read et al. 2005, Schwab et al. 2007). WNV also infects poultry, often progressing to clinical disease in geese (Komar et al. 2001, Swayne et al. 2001).

Other flaviviruses that occur in North America include St. Louis encephalitis virus (SLEV), dengue virus (DENV), and Zika virus (ZIKV). SLEV is maintained in a bird/mosquito transmission cycle, but other vertebrates can become infected (Reisen 2003, Diaz et al. 2018). Vertebrate animals susceptible to SLEV infection include chickens, cows, horses, pheasants, and turkeys (Morris et al. 1994, Ulloa et al. 2003, Marlenee et al. 2004). There have been no reports of fatal SLEV infections in vertebrate animals in North America, but the virus was isolated from brain tissue of a horse that died of encephalitis in Brazil (Rosa et al. 2013). DENV and ZIKV are not usually associated with infections of domestic animals, but antibodies to these viruses were recently identified in horses on islands in the South Pacific (Beck et al. 2019).

CVV infections in sheep are common and can result in embryonic and fetal death, stillbirths, and congenital defects (Noronha and Wilson 2017, Waddell et al. 2019). The first documented outbreak of CVV occurred in a sheep flock in Texas in 1987 (Chung et al. 1990). Nineteen percent of the 360 lambs born during the outbreak had musculoskeletal and central nervous system defects and the total neonatal loss was 26%. Other vertebrate animals susceptible to CVV infection include cows, goats, horses, poultry, and rabbits (Waddell et al. 2019). Additional BUN serogroup viruses that occur in North America are Lokern virus (LOKV), Main Drain virus (MDV), Potosi virus (POTV), Northway virus (NORV), and Tensaw virus (TENV) (Calisher et al. 1986, Heard et al. 1991). MDV has been isolated from brain tissue of a horse that died of encephalitis and caused musculoskeletal and nervous system malformations and death in ovine fetuses after experimental challenge (Emmons et al. 1983, Edwards et al. 1997, Wilson et al. 2015). LOKV, POTV, NORV, and TENV are not recognized pathogens of vertebrate animals but their ability to cause disease has not been widely investigated.

Many studies have been performed to estimate the seroprevalence of flaviviruses and BUN serogroup viruses in vertebrate animals in Canada and the United States, but relatively few studies have been performed in Mexico. BUN serogroup virus surveillance has been especially limited. In

the last 20 years, there have been two serological investigations performed in Mexico where vertebrate animals were tested for antibodies to BUN serogroup viruses and these were performed in the states of Chiapas, Quintana Roo, and Yucatan (Ulloa et al. 2003, Blitvich et al. 2012b). To increase our understanding of the host range and seroprevalence of BUN serogroup viruses and flaviviruses in Mexico, we serologically assayed vertebrate animals in Chihuahua, Guerrero, and Michoacán for select BUN serogroup viruses and flaviviruses known to occur in North America.

Methods

Study sites and sample population

The sample population consisted of vertebrate animals in three states of Mexico: Chihuahua in northwest Mexico and Guerrero and Michoacán on the Pacific Coast (Fig. 1). Study sites were located in 13 of the 67 municipalities of Chihuahua: Ahumada, Ascención, Batopilas de Manuel Gómez Morín, Buenaventura, Casas Grandes, Chihuahua, Dr. Belisario Domínguez, Galeana, Juárez, Namiquipa, Nuevo Casas Grandes, Ocampo, and Riva Palacio. In Michoacán and Guerrero, study sites were located in the municipalities of Lázaro Cárdenas and La Unión, respectively.

Sera were collected from 856 vertebrate animals of 16 species and 1 nothospecies and they are as follows: cat (*Felis silvestris catus*), cow (*Bos taurus*), dog (*Canis lupus familiaris*), domestic goose (*Anser anser domesticus*), goat (*Capra aegagrus hircus*), golden pheasant (*Chrysolophus pictus*), helmeted guinea fowl (*Numida meleagris*), horse (*Equus ferus caballus*), hybrid duck (interspecies breeding between *Anas platyrhynchos* and *Anser anser*), Indian peafowl (*Pavo cristatus*), mallard duck (*A. platyrhynchos*), pig (*Sus scrofa domesticus*), pigeon (*Columba livia*), sheep (*Ovis aries*), silver pheasant (*Lophura nycthemera*), white-sided jackrabbit (*Lepus callotis*), and wild turkey (*Meleagris gallopavo*) (Table 1). All of the cats, cows, dogs, goats, horses, pigs, rabbits, and sheep were from a convenience selection of ranches, farms and houses. All ducks, pigeons and geese were from the Parque Central Poniente Hermanos Escobar, Ciudad Juárez (Central Park, Juárez City). All peafowl, pheasants, guinea fowl and turkeys were from the Recreativo Zoológico San Jorge, Ciudad Juárez (St. Jorge Recreational Zoo, Juárez City). All animals were asymptomatic at the time of sampling, except for one horse in Ciudad Juárez (Juárez City) with ataxia and vision loss that eventually recovered. None of the animals had ever left the state or been vaccinated against WNV. Sera were collected from animals with the approval of the Animal Ethics Committees from each institution that participated in the study (Log No. A3236-01, 11-14-7897-G, 19-072, and 18-173).

Cell culture and viruses

African Green Monkey kidney (Vero) cells were cultured at 37°C with 5% CO₂ in Dulbecco's modified Eagle's medium (Thermo Fisher Scientific, Waltham, MA) supplemented with 10% fetal bovine serum, 2 mM L-glutamine, 100 U/mL penicillin, and 100 µg/mL streptomycin. Four flaviviruses and six BUN serogroup viruses were used in this study. The flaviviruses are as follows: dengue virus type 2 (DENV2; strain NGC), SLEV (strain TBH-28), WNV (strain NY99-35261-11), and ZIKV (strain PRVABC59). All

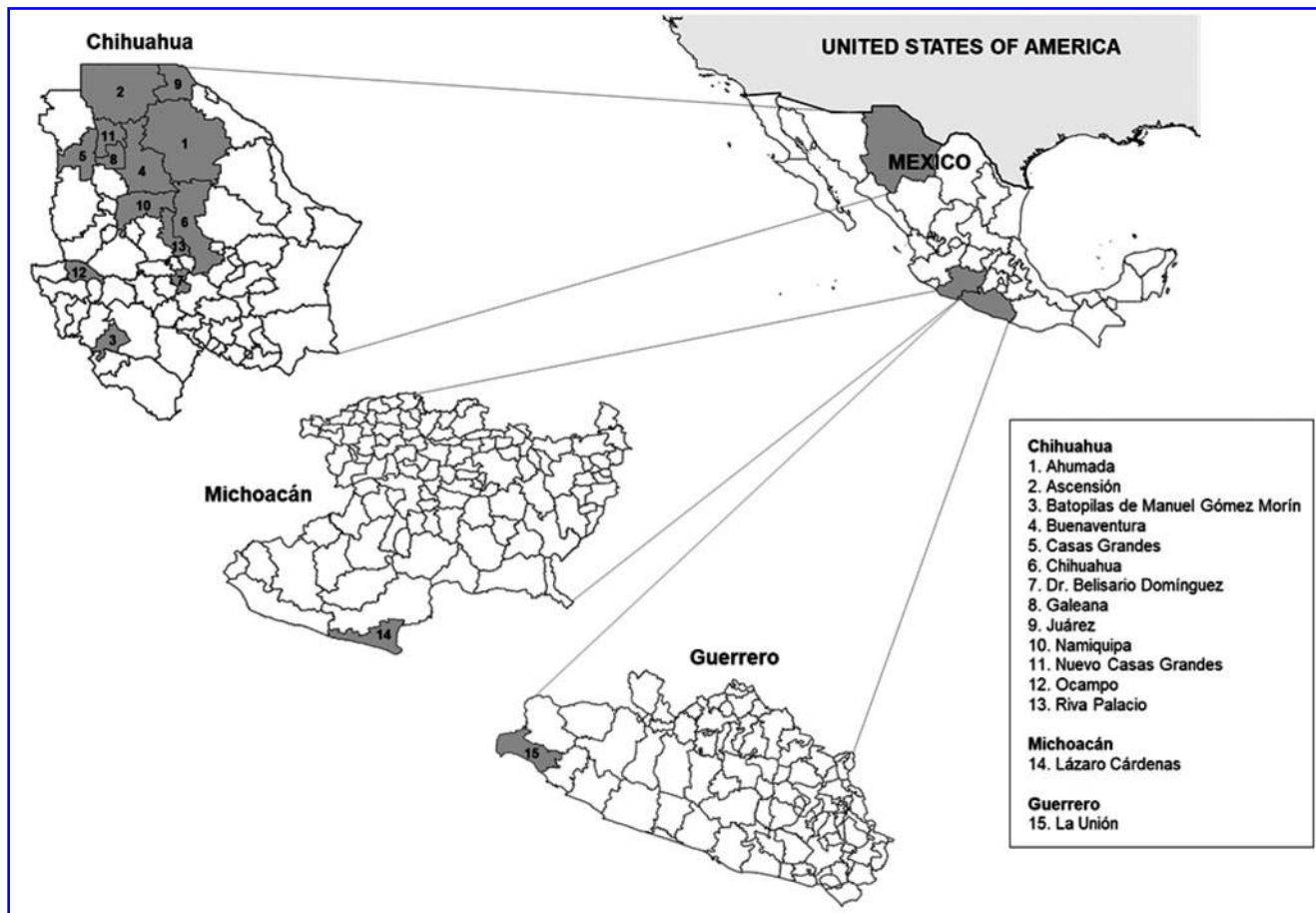


FIG. 1. Geographic location of the study sites. Municipalities in Chihuahua, Guerrero, and Michoacán from where animals were sampled are shaded.

TABLE 1. NUMBER OF VERTEBRATE ANIMALS IN THE STUDY POPULATION, MEXICO, 2018–2019

Common name (species name)	No. of animals sampled in each state			Total (%)
	Chihuahua	Guerrero	Michoacán	
Cat (<i>Felis silvestris catus</i>)	14	0	—	14 (1.6)
Cow (<i>Bos taurus</i>)	117	—	4	121 (14.1)
Dog (<i>Canis lupus familiaris</i>)	208	21	94	323 (37.7)
Domestic goose (<i>Anser anser domesticus</i>)	44	—	—	44 (5.1)
Goat (<i>Capra aegagrus hircus</i>)	2	—	—	2 (0.2)
Golden pheasant (<i>Chrysolophus pictus</i>)	1	—	—	1 (0.1)
Helmeted guinea fowl (<i>Numida meleagris</i>)	6	—	—	6 (0.7)
Horse (<i>Equus ferus caballus</i>)	197	6	20	223 (26.0)
Hybrid duck ^a ; interspecies breeding (<i>Anas platyrhynchos</i> , <i>Anser anser</i>)	31	—	—	31 (3.6)
Indian peafowl (<i>Pavo cristatus</i>)	14	—	—	14 (1.6)
Mallard duck (<i>A. platyrhynchos</i>)	12	—	—	12 (1.4)
Pig (<i>Sus scrofa domesticus</i>)	7	—	7	14 (1.6)
Pigeon (<i>Columba livia</i>)	9	—	—	9 (1.1)
Sheep (<i>Ovis aries</i>)	35	—	—	35 (4.1)
Silver pheasant (<i>Lophura nycthemera</i>)	1	—	—	1 (0.1)
White-sided jackrabbit (<i>Lepus callotis</i>)	4	—	—	4 (0.5)
Wild turkey (<i>Meleagris gallopavo</i>)	2	—	—	2 (0.2)
Total	704	27	125	856 (100)

^aHybrid ducks were not included in the species total.

flaviviruses were obtained from the World Health Organization Center for Arbovirus Reference and Research, which is maintained at the Division of Vector-Borne Infectious Diseases, Centers for Disease Control and Prevention in Fort Collins, Colorado. The BUN serogroup viruses are as follows: CVV (strain CVV-478), LOKV (strain FMS 4332), MDV (strain BFS 5015), NORV (strain 0234), POTV (strain BeAr7272), and TENV (strain A9-171b). CVV-478 was isolated from mosquitoes collected in Mexico (Blitvich et al. 2012a). All other BUN serogroup viruses were obtained from the World Arbovirus Reference Collection at the University of Texas Medical Branch in Galveston, Texas.

Plaque reduction neutralization test for flaviviruses

Plaque reduction neutralization tests (PRNTs) were performed in six-well plates containing confluent monolayers of Vero cells following standard methods (Beaty et al. 1995). Initially, all sera were screened at a single dilution of 1:20 using WNV. The initial screen was performed using WNV because it is the most widespread flavivirus in North America. These assays are not WNV specific; flaviviruses are antigenically similar and therefore, antibodies to other flaviviruses are also detected (Calisher et al. 1989). All sera with antibodies that neutralized WNV were further diluted and tested by comparative PRNT using DENV2, SLEV, WNV, and ZIKV to identify the flavivirus responsible for these infections. Titers were expressed as the reciprocal of highest serum dilutions yielding ≥90% reduction in the number of plaques (PRNT₉₀). For etiologic diagnosis, the PRNT₉₀ antibody titer to the flavivirus was required to be at least fourfold greater than that to the other flaviviruses tested. If neutralizing antibodies were detected, but there was not at least a fourfold difference in PRNT₉₀ antibody titers, the animal was considered to have antibodies to an undetermined flavivirus.

PRNTs for BUN serogroup viruses

PRNTs were performed in six-well plates containing confluent monolayers of Vero cells (Beaty et al. 1995). Initially, all sera were screened at a single dilution of 1:20 using CVV, the most prevalent and widespread BUN serogroup virus in North America. These assays are not CVV specific; BUN

serogroup viruses are antigenically similar and therefore, antibodies to other BUN serogroup viruses are also detected (Hunt and Calisher 1979). All sera with antibodies that neutralized CVV were further diluted and tested by PRNT using CVV, LOKV, MDV, NORV, POTV, and TENV. For etiologic diagnosis, the PRNT₉₀ antibody titer to the BUN serogroup virus was required to be at least fourfold greater than that to the other BUN serogroup viruses tested. If neutralizing antibodies were detected but there was not at least a fourfold difference in PRNT₉₀ antibody titers, the animal was considered to have antibodies to an undetermined BUN serogroup virus.

RNA extraction and RT-PCR

Total RNA was extracted from the serum of the symptomatic horse using the QIAamp Viral RNA Mini Kit (Qiagen) according to the manufacturer’s instructions and tested by RT-PCR using flavivirus-specific primers (Kuno et al. 1998). The IRB at each participating university waived the requirement that signatures be acquired from the owners and caregivers of animals.

Results

Sera were collected from 856 vertebrate animals of 16 species and 1 nothospecies in Mexico in 2018–2019 (Table 1). The majority (82.3%) of the animals were from Chihuahua, with the remainder from Michoacán (14.6%) and Guerrero (3.2%). The most commonly sampled animals were dogs (37.7%), horses (26.0%), and cows (14.1%). The municipality that comprised most (65.3%) of the sample population was Juárez in Chihuahua with 559 animals, followed by Lázaro Cárdenas in Michoacán (125 animals; 14.6%) and Buenaventura in Chihuahua (74 animals; 8.6%). All of the sheep, 112 of the 197 (56.9%) horses and 44 of the 117 (37.6%) cows were from the municipality of Juárez.

Sera were tested at a dilution of 1:20 by PRNT using WNV. Antibodies that neutralized WNV were detected in 296 animals: 19 (15.7%) cows, 80 (24.8%) dogs, 12 (27.3%) domestic geese, 1 (50.0%) goat, 1 (100%) golden pheasant, 1 (16.7%) helmeted guinea fowl, 153 (68.6%) horses, 5 (16.1%) hybrid ducks produced from crossbreeding, 12 (85.7%) Indian peafowl, 3 (25.0%) mallard ducks, 1 (11.1%)

TABLE 2. VERTEBRATE ANIMALS SEROPOSITIVE FOR WEST NILE VIRUS IN MEXICO, 2018–2019

Animal ^a	Geographic region			No. seropositive/total (%)
	Chihuahua	Guerrero	Michoacán	
Cow	8/117 (6.8)	0/0	1/4 (25.0)	9/121 (7.4)
Dog	34/208 (16.3)	3/21 (14.3)	4/94 (4.3)	41/323 (12.7)
Domestic goose	2/44 (4.5)	—	—	2/44 (4.5)
Goat	1/2 (50.0)	—	—	1/2 (50.0)
Golden pheasant	1/1 (100)	—	—	1/1 (100)
Horse	84/197 (42.6)	0/6 (0.0)	5/20 (25.0)	89/223 (39.9)
Hybrid duck	5/31 (16.1)	—	—	5/31 (16.1)
Indian peafowl	3/14 (21.4)	—	—	3/14 (21.4)
Mallard duck	1/12 (8.3)	—	—	1/12 (8.3)
Wild turkey	2/2 (100)	—	—	2/2 (100)
Total	141/704 (20.1)	3/27 (11.1)	10/125 (8.0)	154/856 (18.0)

^aAnimals are not listed if none was seropositive.

TABLE 3. VERTEBRATE ANIMALS SEROPOSITIVE FOR ST. LOUIS ENCEPHALITIS VIRUS IN MEXICO, 2018–2019

Animal ^a	Geographic region			No. of seropositive/total (%)
	Chihuahua	Guerrero	Michoacán	
	No. of seropositive/total (%)			
Dog	2/208 (1.0)	0/21	1/94 (1.1)	3/323 (0.9)
Domestic duck	1/44 (2.3)	0/0	—	1/44 (2.3)
Helmeted guinea fowl	1/6 (16.7)	—	—	1/6 (16.7)
Horse	1/197 (0.5)	0/6 (0.0)	0/20 (0.0)	1/223 (0.4)
Silver pheasant	1/1 (100)	—	—	1/1 (100)
Total	6/704 (0.9)	0/27	1/125 (0.8)	7/856 (0.8)

^aAnimals are not listed if none was seropositive.

pigeon, 4 (11.4%) sheep, 1 (100%) silver pheasant, 1 (25.0%) white-sided jackrabbit, and 2 (100%) wild turkeys. All sera that neutralized WNV in the initial screen were further diluted and tested by PRNT using DENV2, SLEV, WNV, and ZIKV. The comparative PRNTs revealed that 154 animals were seropositive for WNV, 7 animals were seropositive for SLEV, and 135 animals were seropositive for an undetermined flavivirus. Antibodies to WNV were detected in animals from nine species and the crossbreed (Table 2). Of the species/crossbreed where at least 10 individuals were sampled, seroprevalence for WNV was highest for horses (39.9%), Indian peafowl (21.4%), and hybrid ducks (16.1%). One horse seropositive for WNV developed symptoms several days before serum collection. Serum of the symptomatic horse was tested for flavivirus RNA by RT-PCR and shown to be negative. WNV activity was detected in all three states. Antibodies to SLEV were detected in five species (Table 3). Of the species where at least 10 individuals were sampled, seroprevalence for SLEV was highest for domestic ducks (2.6%). SLEV activity was detected in Chihuahua and Michoacán. Antibodies to DENV2 and ZIKV were not detected in any animals.

Sera were tested at a dilution of 1:20 by PRNT using CVV. Antibodies that neutralized CVV were detected in 42 animals: 2 (1.7%) cows, 5 (1.5%) dogs, 33 (14.2%) horses, 1 (2.9%) sheep, and 1 (25%) white-sided jackrabbit. All sera that neutralized CVV were further diluted and tested by PRNT using CVV, LOKV, MDV, NORV, POTV, and TENV. The comparative PRNTs revealed that 22 animals were seropositive for LOKV, 3 animals were seropositive for MDV, and 17 animals were seropositive for an undetermined BUN serogroup virus. Antibodies to LOKV were detected in animals from three species (Table 4). Seroprevalence for LOKV was highest for horses (8.5%) and cows (1.7%). LOKV activity was detected in all three states. All three animals seropositive for MDV were

horses in Chihuahua. The overall seroprevalence for MDV in horses was 1.3%. Antibodies to CVV, NORV, POTV, and TENV were not detected in any animals.

Twenty-five animals were seropositive for two viruses. Two horses from Chihuahua were seropositive for MDV and WNV, a horse from Guerrero was seropositive for LOKV and SLEV, 11 horses (4 from Chihuahua, 2 from Guerrero, and 5 from Michoacán) were seropositive for LOKV and WNV, and a cow from Michoacán was seropositive for LOKV and WNV. Five animals (four horses and one sheep) were seropositive for WNV and an undetermined BUN serogroup virus and five other animals (three dogs, one horse, and one rabbit) were seropositive for an undetermined BUN serogroup virus and an undetermined flavivirus.

Discussion

We provide serologic evidence of LOKV, MDV, SLEV, and WNV infections in vertebrate animals in Mexico. LOKV and MDV have never before been reported outside of the United States. LOKV has been isolated from arthropods and vertebrate animals in five states of the United States, including three (California, New Mexico, and Texas) that border Mexico (Crane et al. 1983, Calisher et al. 1986, Kramer et al. 1990). Isolations of MDV have been made from arthropods and vertebrate animals in six states, including three (Arizona, California, and Texas) than border Mexico (Calisher et al. 1986). Serological data suggest that LOKV and MDV also occur elsewhere in the United States (Sahu et al. 2002, Johnson et al. 2014, Meyers et al. 2015). In this study, we provide evidence that the geographic distributions of LOKV and MDV extend beyond the United States. The detection of LOKV activity as far north as Chihuahua and as far south as Guerrero indicates that this virus is widely distributed across Mexico.

TABLE 4. VERTEBRATE ANIMALS SEROPOSITIVE FOR LOKERN VIRUS IN MEXICO, 2018–2019

Animal ^a	Geographic region			No. of seropositive/total (%)
	Chihuahua	Guerrero	Michoacán	
	No. of seropositive/total (%)			
Cow	1/117 (0.9)	0/0	1/4 (25.0)	2/121 (1.7)
Dog	1/208 (0.5)	0/21 (0.0)	0/94 (0.0)	1/323 (0.3)
Horse	7/197 (3.6)	4/6 (66.7)	8/20 (40.0)	19/223 (8.5)
Total	9/704 (1.3)	4/27 (14.8)	9/125 (7.2)	22/856 (2.6)

^aAnimals are not listed if none was seropositive.

All viruses in the BUN serogroup have tripartite, single-stranded, negative-sense RNA genomes and the three genome segments are designated as small, (S), medium (M), and large (L) (Elliott et al. 1991, Horne and Vanlandingham 2014). The viral envelope glycoproteins are encoded by the M segment. Recent evidence suggests that LOKV is a reassortant that acquired its S and L segments from MDV and its M segment from an undiscovered, possibility extinct, virus (Tangudu et al. 2018). Because LOKV and its apparent unidentified donor share the same M segment, antibodies to these viruses cannot be differentiated by PRNT. Therefore, if the M segment donor still occurs in nature, some or all of the animals considered seropositive for LOKV could have instead been infected with the undiscovered donor virus.

CVV is the most common and widely distributed BUN serogroup virus in North America (Calisher et al. 1986, Waddell et al. 2019). CVV is also the most important BUN serogroup virus in North America in terms of its impact on human and veterinary animal health, having been associated with outbreaks of pregnancy loss and congenital malformations in sheep, in addition to six cases of severe disease in humans (Sexton et al. 1997, Campbell et al. 2006, Nguyen et al. 2013, Wilson et al. 2017, Yang et al. 2018b, Baker et al. 2021). None of the animals in our study was seropositive for CVV and the overall seroprevalence for BUN serogroup viruses was lower than other studies performed in North America. For example, antibodies to BUN serogroup viruses were detected in 169 (92.9%) horses and 29 (93.5%) sheep in Quintana Roo and Yucatán, Mexico in 2007–2008, with CVV identified as one of the most common causes of infection (Blitvich et al. 2012b). In another study, 1455 (28.3%) sheep across the United States in 2011 contained antibodies to BUN serogroup viruses (Meyers et al. 2015). CVV was the most common virus. Sera from 84 (64.6%) sheep, 40 (69.0%) horses, 8 (33.3%) goats, and 5 (20.0%) cows in Saskatchewan, Canada in 2013–2014 also contained antibodies to BUN serogroup viruses (Uehlinger et al. 2018).

One explanation for the relatively low seroprevalence for BUN serogroup viruses and apparent lack of CVV activity in our study is because competent reservoir hosts or vectors of CVV do not occur in the study area. White-tailed deer have been implicated as the principal reservoir hosts of CVV based on experimental infection data and seropositivity rates from field studies (Campbell et al. 1989, Neitzel and Grimstad 1991, Blackmore and Grimstad 1998, Dupuis et al. 2020). White-tailed deer are common in Mexico, although they are not found in parts of northern Chihuahua (Mandujano et al. 2010, Hewitt 2011). The majority (65.3%) of animals in our study population were from the municipality of Juárez in northern Chihuahua, including all sheep and many of the horses and cows. Experimental infections have revealed that mosquitoes of at least 10 species are competent vectors of CVV, including some that occur in the study area, and virus isolations have been made from several additional species (Yuill and Thompson 1970, Calisher et al. 1986, Blackmore et al. 1998, Andreadis et al. 2014, Armstrong et al. 2017, Anderson et al. 2018, Ayers et al. 2018, 2019, Yang et al. 2018a, Chan et al. 2020).

Many WNV and SLEV serological investigations were performed in Mexico shortly after the arrival of WNV into the Americas (Blitvich 2008, Elizondo-Quiroga and Elizondo-Quiroga 2013). However, relatively few have

been performed in recent years. Four of 78 (5.1%) humans in Ciudad Juárez (Juárez City), Chihuahua, in 2015 were seropositive for WNV (Palermo et al. 2019). SLEV testing was not performed. Antibodies to SLEV and WNV were not detected in any of the 639 humans in Guerrero in 2019 (Nunez-Avellaneda et al. 2021). There are no other recent studies where humans or vertebrate animals in Chihuahua, Guerrero, or Michoacán were tested for evidence of SLEV or WNV infection. In this study, we provide evidence of WNV activity in all three states and SLEV activity in Chihuahua and Michoacán. Of the 10 arboviruses included in our study, WNV had the broadest host range, with antibodies to the virus detected in animals of nine species. It is well documented that WNV infects vertebrate animals of numerous species (Root 2013, Root and Bosco-Lauth 2019, Habarugira et al. 2020).

Conclusion

We provide serologic evidence of LOKV, MDV, SLEV, and WNV infections in vertebrate animals in Mexico. LOKV and MDV have not been previously reported in Mexico and there are no recent reports of SLEV and WNV activity in Guerrero and Michoacán. This study increases our understanding of the geographic distribution, host range, and seroprevalence of BUN serogroup viruses and flaviviruses in North America.

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Author Disclosure Statement

No conflicting financial interests exist.

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References

- Anderson JF, Armstrong PM, Misencik MJ, Bransfield AB, et al. Seasonal distribution, blood-feeding habits, and viruses of mosquitoes in an open-faced quarry in Connecticut, 2010 and 2011. *J Am Mosq Control Assoc* 2018; 34:1–10.
- Andreadis TG, Armstrong PM, Anderson JF, Main AJ. Spatial-temporal analysis of Cache Valley virus (*Bunyaviridae: Orthobunyavirus*) infection in Anopheline and Culicine

- mosquitoes (Diptera: Culicidae) in the northeastern United States, 1997–2012. *Vector Borne Zoonotic Dis* 2014; 14:763–773.
- Armstrong PM, Andreadis TG, Shepard JJ, Thomas MC. Northern range expansion of the Asian tiger mosquito (*Aedes albopictus*): analysis of mosquito data from Connecticut, USA. *PLoS Negl Trop Dis* 2017; 11:e0005623.
- Ayers VB, Huang YS, Lyons AC, Park SL, et al. *Culex tarsalis* is a competent vector species for Cache Valley virus. *Parasit Vectors* 2018; 11:519.
- Ayers VB, Huang YS, Lyons AC, Park SL, et al. Infection and transmission of Cache Valley virus by *Aedes albopictus* and *Aedes aegypti* mosquitoes. *Parasit Vectors* 2019; 12:384.
- Baker M, Hughes HR, Naqvi SH, Yates K, et al. Reassortant Cache Valley virus associated with acute febrile, non-neurologic illness, Missouri. *Clin Infect Dis* 2021. [Epub ahead of print]; DOI: 10.1093/cid/ciab175.
- Beaty B, Calisher C, Shope R. Arboviruses. In: Lennette E, Lennette D, Lennette E, eds. *Diagnostic Procedures for Viral, Rickettsial, and Chlamydial Infections*. Washington, DC: American Public Health Association, 1995; 189–212.
- Beck C, Leparac-Goffart I, Desoutter D, Deberge E, et al. Serological evidence of infection with dengue and Zika viruses in horses on French Pacific Islands. *PLoS Negl Trop Dis* 2019; 13:e0007162.
- Blackmore CG, Blackmore MS, Grimstad PR. Role of *Anopheles quadrimaculatus* and *Coquillettidia perturbans* (Diptera: Culicidae) in the transmission cycle of Cache Valley virus (*Bunyaviridae: Bunyavirus*) in the midwest, USA. *J Med Entomol* 1998; 35:660–664.
- Blackmore CG, Grimstad PR. Cache Valley and Potosi viruses (*Bunyaviridae*) in white-tailed deer (*Odocoileus virginianus*): experimental infections and antibody prevalence in natural populations. *Am J Trop Med Hyg* 1998; 59:704–709.
- Blitvich BJ. Transmission dynamics and changing epidemiology of West Nile virus. *Anim Health Res Rev* 2008; 9:71–86.
- Blitvich BJ, Lirono-Pino MA, Garcia-Rejon JE, Farfan-Ale JA, et al. Nucleotide sequencing and serologic analysis of Cache Valley virus isolates from the Yucatan Peninsula of Mexico. *Virus Genes* 2012a; 45:176–180.
- Blitvich BJ, Saiyasombat R, Travassos da Rosa A, Tesh RB, et al. Orthobunyaviruses, a common cause of infection of livestock in the Yucatan peninsula of Mexico. *Am J Trop Med Hyg* 2012b; 87:1132–1139.
- Bosco-Lauth AM, Bowen RA. West Nile virus: veterinary health and vaccine development. *J Med Entomol* 2019; 56:1463–1466.
- Calisher CH, Francly DB, Smith GC, Muth DJ, et al. Distribution of Bunyamwera serogroup viruses in North America, 1956–1984. *Am J Trop Med Hyg* 1986; 35:429–443.
- Calisher CH, Karabatsos N, Dalrymple JM, Shope RE, et al. Antigenic relationships between flaviviruses as determined by cross-neutralization tests with polyclonal antisera. *J Gen Virol* 1989; 70 (Pt 1):37–43.
- Campbell GL, Eldridge BF, Hardy JL, Reeves WC, et al. Prevalence of neutralizing antibodies against California and Bunyamwera serogroup viruses in deer from mountainous areas of California. *Am J Trop Med Hyg* 1989; 40:428–437.
- Campbell GL, Mataczynski JD, Reisdorf ES, Powell JW, et al. Second human case of Cache Valley virus disease. *Emerg Infect Dis* 2006; 12:854–856.
- Castillo-Olivares J, Wood J. West Nile virus infection of horses. *Vet Res* 2004; 35:467–483.
- Chan KK, Auguste AJ, Brewster CC, Paulson SL. Vector competence of Virginia mosquitoes for Zika and Cache Valley viruses. *Parasit Vectors* 2020; 13:188.
- Chung SI, Livingston CW, Jr., Edwards JF, Crandell RW, et al. Evidence that Cache Valley virus induces congenital malformations in sheep. *Vet Microbiol* 1990; 21:297–307.
- Crane GT, Elbel RE, Francly DB, Calisher CH. Arboviruses from western Utah, USA, 1967–1976. *J Med Entomol* 1983; 20:294–300.
- Diaz A, Coffey LL, Burkett-Cadena N, Day JF. Reemergence of St. Louis encephalitis virus in the Americas. *Emerg Infect Dis* 2018; 24:2150–2157.
- Dunkel B, Del Piero F, Wotman KL, Johns IC, et al. Encephalomyelitis from West Nile flavivirus in 3 alpacas. *J Vet Intern Med* 2004; 18:365–367.
- Dupuis AP, Prusinski MA, Russell A, O'Connor C, et al. Serologic survey of mosquito-borne viruses in hunter-harvested white-tailed deer (*Odocoileus virginianus*), New York State. *Am J Trop Med Hyg* 2020; 104:593–603.
- Edwards JF, Karabatsos N, Collisson EW, de la Concha Bermejillo A. Ovine fetal malformations induced by in utero inoculation with Main Drain, San Angelo, and LaCrosse viruses. *Am J Trop Med Hyg* 1997; 56:171–176.
- Elizondo-Quiroga D, Elizondo-Quiroga A. West Nile virus and its theories, a big puzzle in Mexico and Latin America. *J Glob Infect Dis* 2013; 5:168–175.
- Elliott RM, Schmaljohn CS, Collett MS. Bunyaviridae genome structure and gene expression. *Curr Top Microbiol Immunol* 1991; 169:91–141.
- Emmons RW, Woodie JD, Laub RL, Oshiro LS. Main Drain virus as a cause of equine encephalomyelitis. *J Am Vet Med Assoc* 1983; 183:555–558.
- Habarugira G, Suen WW, Hobson-Peters J, Hall RA, et al. West Nile virus: an update on pathobiology, epidemiology, diagnostics, control and “One Health” implications. *Pathogens* 2020; 9:589.
- Heard PB, Niebylski ML, Francly DB, Craig GB, Jr. Transmission of a newly recognized virus (*Bunyaviridae, Bunyavirus*) isolated from *Aedes albopictus* (Diptera: Culicidae) in Potosi, Missouri. *J Med Entomol* 1991; 28:601–605.
- Hewitt DG. *Biology and Management of White-Tailed Deer*. Boca Raton, FL: CRC Press, 2011.
- Horne KM, Vanlandingham DL. Bunyavirus-vector interactions. *Viruses* 2014; 6:4373–4397.
- Hunt AR, Calisher CH. Relationships of Bunyamwera group viruses by neutralization. *Am J Trop Med Hyg* 1979; 28:740–749.
- Johnson GD, Bahnson CS, Ishii P, Cochrane ZN, et al. Monitoring sheep and *Culicoides* midges in Montana for evidence of Bunyamwera serogroup virus infection. *Vet Rec Open* 2014; 1:e000071.
- Komar N, Panella NA, Burns JE, Duszka SW, et al. Serologic evidence for West Nile virus infection in birds in the New York City vicinity during an outbreak in 1999. *Emerg Infect Dis* 2001; 7:621–625.
- Kramer WL, Jones RH, Holbrook FR, Walton TE, et al. Isolation of arboviruses from *Culicoides* midges (Diptera: Ceratopogonidae) in Colorado during an epizootic of vesicular stomatitis New Jersey. *J Med Entomol* 1990; 27:487–493.
- Kuno G, Chang GJ, Tsuchiya KR, Karabatsos N, et al. Phylogeny of the genus *Flavivirus*. *J Virol* 1998; 72:73–83.
- Mandujano S, Delfín-Alfonso CA, Gallina S. Comparison of geographic distribution models of white-tailed deer *Odocoileus virginianus* (Zimmermann, 1780) subspecies in Mexico: biological and management implications. *Therya* 2010; 1:41–68.
- Marlenee NL, Lirono-Pino MA, Beaty BJ, Blitvich BJ, et al. Detection of antibodies to West Nile and Saint Louis encephalitis viruses in horses. *Salud Publica Mex* 2004; 46:373–375.

- Meyers MT, Bahnson CS, Hanlon M, Kopral C, et al. Management factors associated with operation-level prevalence of antibodies to Cache Valley virus and other Bunyamwera serogroup viruses in sheep in the United States. *Vector Borne Zoonotic Dis* 2015; 15:683–693.
- Morris CD, Baker WG, Stark L, Burgess J, et al. Comparison of chickens and pheasants as sentinels for eastern equine encephalitis and St. Louis encephalitis viruses in Florida. *J Am Mosq Control Assoc* 1994; 10:545–548.
- Neitzel DF, Grimstad PR. Serological evidence of California group and Cache Valley virus infection in Minnesota white-tailed deer. *J Wildl Dis* 1991; 27:230–237.
- Nguyen NL, Zhao G, Hull R, Shelly MA, et al. Cache valley virus in a patient diagnosed with aseptic meningitis. *J Clin Microbiol* 2013; 51:1966–1969.
- Noronha LE, Wilson WC. Comparison of two zoonotic viruses from the order *Bunyavirales*. *Curr Opin Virol* 2017; 27:36–41.
- Nunez-Avellaneda D, Tangudu CS, Barrios-Palacios J, Machain-Williams C, et al. Co-circulation of all four dengue viruses and Zika virus in Guerrero, Mexico, 2019. *Vector Borne Zoonotic Dis* 2021; 21:458–465.
- Palermo PM, De la Mora-Covarrubias A, Jimenez-Vega F, Watts DM. Serological evidence of dengue and West Nile virus human infection in Juarez City, Mexico. *Vector Borne Zoonotic Dis* 2019; 19:134–141.
- Petersen LR. Epidemiology of West Nile virus in the United States: implications for arbovirology and public health. *J Med Entomol* 2019; 56:1456–1462.
- Read RW, Rodriguez DB, Summers BA. West Nile virus encephalitis in a dog. *Vet Pathol* 2005; 42:219–222.
- Reisen WK. Epidemiology of St. Louis encephalitis virus. *Adv Virus Res* 2003; 61:139–183.
- Rimoldi G, Mete A, Adaska JM, Anderson ML, et al. West Nile virus infection in sheep. *Vet Pathol* 2017; 54:155–158.
- Root JJ. West Nile virus associations in wild mammals: a synthesis. *Arch Virol* 2013; 158:735–752.
- Root JJ, Bosco-Lauth AM. West Nile virus associations in wild mammals: an update. *Viruses* 2019; 11:459.
- Rosa R, Costa EA, Marques RE, Oliveira TS, et al. Isolation of Saint Louis encephalitis virus from a horse with neurological disease in Brazil. *PLoS Negl Trop Dis* 2013; 7:e2537.
- Sahu SP, Pedersen DD, Ridpath HD, Ostlund EN, et al. Serologic survey of cattle in the northeastern and north central United States, Virginia, Alaska, and Hawaii for antibodies to Cache Valley and antigenically related viruses (Bunyamwera serogroup virus). *Am J Trop Med Hyg* 2002; 67:119–122.
- Saiz JC. Animal and human vaccines against West Nile virus. *Pathogens* 2020; 9:1073.
- Schwab S, Herden C, Seeliger F, Papaioannou N, et al. Non-suppurative meningoencephalitis of unknown origin in cats and dogs: an immunohistochemical study. *J Comp Pathol* 2007; 136:96–110.
- Sexton DJ, Rollin PE, Breitschwerdt EB, Corey GR, et al. Life-threatening Cache Valley virus infection. *N Engl J Med* 1997; 336:547–549.
- Swayne DE, Beck JR, Smith CS, Shieh WJ, et al. Fatal encephalitis and myocarditis in young domestic geese (*Anser anser domesticus*) caused by West Nile virus. *Emerg Infect Dis* 2001; 7:751–753.
- Tangudu CS, Charles J, Blitvich BJ. Evidence that Lokern virus (family *Peribunyaviridae*) is a reassortant that acquired its small and large genome segments from Main Drain virus and its medium genome segment from an undiscovered virus. *Virol J* 2018; 15:122.
- Thorsen J, Artsob H, Spence L, Surgeoner G, et al. Virus isolations from mosquitoes in southern Ontario, 1976 and 1977. *Can J Microbiol* 1980; 26:436–440.
- Uehlinger FD, Wilkins W, Godson DL, Drobot MA. Seroprevalence of Cache Valley virus and related viruses in sheep and other livestock from Saskatchewan, Canada. *Can Vet J* 2018; 59:413–418.
- Ulloa A, Langevin SA, Mendez-Sanchez JD, Arredondo-Jimenez JJ, et al. Serologic survey of domestic animals for zoonotic arbovirus infections in the Lacandon Forest region of Chiapas, Mexico. *Vector Borne Zoonotic Dis* 2003; 3:3–9.
- Waddell L, Pachal N, Mascarenhas M, Greig J, et al. Cache Valley virus: a scoping review of the global evidence. *Zoonoses Public Health* 2019; 66:739–758.
- Wilson MR, Suan D, Duggins A, Schubert RD, et al. A novel cause of chronic viral meningoencephalitis: Cache Valley virus. *Ann Neurol* 2017; 82:105–114.
- Wilson WC, Gaudreault NN, Hossain MM, McVey DS. Lesser-known bunyavirus infections. *Rev Sci Tech* 2015; 34:419–429.
- Yang F, Chan K, Marek PE, Armstrong PM, et al. Cache Valley virus in *Aedes japonicus japonicus* mosquitoes, Appalachian Region, United States. *Emerg Infect Dis* 2018a; 24:553–557.
- Yang Y, Qiu J, Snyder-Keller A, Wu Y, et al. Fatal Cache Valley virus meningoencephalitis associated with rituximab maintenance therapy. *Am J Hematol* 2018b; 93:590–594.
- Yuill TM, Thompson PH. Cache Valley virus in the Del Mar Va Peninsula. IV. Biological transmission of the virus by *Aedes sollicitans* and *Aedes taeniorhynchus*. *Am J Trop Med Hyg* 1970; 19:513–519.
- Zheng H, Drobot MA, Coulthart MB. West Nile virus in Canada: ever-changing, but here to stay. *Can Commun Dis Rep* 2014; 40:173–177.

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