

Experimental Inoculation of *Artibeus jamaicensis* Bats with Dengue Virus Serotypes 1 or 4 Showed No Evidence of Sustained Replication

Salomé Cabrera-Romo, Benito Recio-Tótoro, Ana C. Alcalá, Humberto Lanz, Rosa María del Ángel, Victor Sánchez-Cordero, Ángel Rodríguez-Moreno, and Juan E. Ludert*

Departament of Infectomics and Molecular Pathogenesis, Center for Research and Advanced Studies (CINVESTAV-IPN), Mexico City, Mexico; Center for Research on Infectious Diseases (CISEI), National Institute of Public Health (INSP), Cuernavaca, Mexico; Institute of Biology, Mexico City, Universidad Nacional Autónoma de México, Mexico

Abstract. Dengue is the most important mosquito-borne viral disease to humans. Bats are potential reservoirs for flaviviruses, including dengue virus (DENV). In this work, *Artibeus jamaicensis* bats were inoculated with two serotypes of DENV using different routes. For experimental inoculations (EI) 1 and 2, bats were inoculated subcutaneously or intraperitoneally with DENV-4; for EI-3 bats were inoculated intraperitoneally with DENV-1. Mock inoculated bats were kept as controls. In EI-4, bats were bitten by *Aedes aegypti* mosquitoes infected with DENV-1 or 4. Reverse transcription-polymerase chain reaction assays in plasma and spleen tissue collected from Day 1 to Days 9–17 after inoculation failed to reveal the presence of viral RNA in any of the samples. No evidence of circulating NS1 or specific anti-DENV IgG was detected in the plasma of the inoculated bats. These results indicate that *A. jamaicensis* bats are incapable of sustaining dengue virus replication and are unlikely to act as reservoirs for this virus.

INTRODUCTION

Dengue virus (DENV) is classified within the family *Flaviviridae*, in the genus *Flavivirus*; a genus that groups several other important human pathogens associated with encephalitis and hemorrhagic fevers. Four antigenically different but genetically related serotypes of DENV (DENV-1 to DENV-4) are known. The DENV mature virion is composed of a small icosahedral capsid, of about 45 nm in diameter, made of a single protein (C), surrounded by a lipid envelope containing the E and M proteins. The virus presents a single-stranded RNA genome, of positive polarity, of ~10 kb. In addition to the three structural proteins, the genome encodes for seven non-structural proteins named NS1, NS2A, NS2B, NS3, NS4A, NS4B, and NS5, which all participate in virus replication.^{1–3} In addition, NS1 is secreted to the extracellular milieu and circulates in patients' sera early during infection along with infectious virus.⁴ Thus, NS1 detection is the basis of several commercial diagnostic assays.⁵

Dengue currently is considered the most important mosquito-borne viral disease for humans. The disease is endemic in more than 100 countries where an estimated 100 million cases are reported annually, associated with more than 25,000 fatalities, mainly in children < 5 years of age.⁶ Mexico is a country endemic for dengue, where all 4 DENV serotypes are known to circulate. Infection with any of the 4 DENV serotypes usually results in subclinical or mild dengue; however, a small fraction of the patients can evolve to severe dengue, the life threatening form of the disease. Mild dengue is characterized by high fever, headache, muscle and joint pain, and rash, whereas patients with severe dengue (also known as dengue hemorrhagic fever) present hemorrhages, plasma leakage, and respiratory distress.^{2,3} Despite the great disease burden associated with dengue, currently there are neither vaccines nor spe-

cific treatment of this disease.⁷ In addition, no widely available animal model for dengue virus infection is available.⁸

The DENV is maintained in nature in two ecologically and evolutionary distinct transmission cycles: the human and the sylvatic cycle.⁹ The mosquito *Aedes aegypti* is the main vector for the human cycle, which involves only humans as reservoirs or amplification hosts. The sylvatic cycle involves non-human primates as reservoirs and amplification hosts and is maintained by mosquitoes in the genus *Aedes*. The DENV sylvatic cycles have been well documented only in West Africa and in Southeast Asia.¹⁰ However, there is scattered evidence suggesting that a sylvatic cycle for DENV among several species of small mammals, may also exist in the Americas. A serological survey conducted by de Thoisy and others,¹¹ comprising 27 wild forest mammal species collected in the French Guiana, found sporadic evidence of DENV infection in several of the species sampled. Serologic evidence for DENV infection in several species of chiroptera collected in Ecuador, Costa Rica, and Mexico has also been reported.^{12–14} In addition, reverse transcription-polymerase chain reaction (RT-PCR) testing of spleen or heart tissue samples from bats captured in dengue-endemic areas in Mexico, detected the presence of DENV-2 in frugivorous (*Artibeus jamaicensis*, *Artibeus lituratus*, *Glossophaga soricina*) and insectivorous (*Myotis nigricans*) bats.^{13,15} In an elegant study carried out in the French Guiana, RNA corresponding to the 4 DENV serotypes was detected in tissue samples collected from several species of wild mammals. Interestingly, evidence of DENV infection was found in mammals, including several species of Chiroptera, captured in remote sylvatic areas, where no evidence of dengue cycle exists.¹¹ All these results suggest that wild animals, including several species of bats, may carry and presumably sustain DENV infection in nature.

Bats belong to the order Chiroptera, one of the most abundant, species diverse and widely disseminated mammal orders.¹⁶ Over 350 bats species are found in the Americas and the Caribbean.^{17,18} In recent years, bats are increasingly being recognized as reservoirs for a variety of important human viruses^{19,20} including flaviviruses,¹¹ and have become an important study subject from a public health point of view. Because mounting evidence indicates that bats may be a reservoir for

*Address correspondence to Juan E. Ludert, Departamento de Infectómica y Patogénesis Molecular, Centro de Investigación y Estudios Avanzados (CINVESTAV-IPN), Av. IPN 2508, Colonia San Pedro Zacatenco, Mexico, D.F. 07360, Mexico. E-mail: jludert@cinvestav.mx

DENV, the main objective of this study was to directly test the susceptibility of the very wide spread species of bat *A. jamaicensis* to DENV infection. Experimental inoculations with DENV-1 and DENV-4, using several inoculation routes suggest that *A. jamaicensis* bats are incapable of sustaining DENV replication.

MATERIALS AND METHODS

Bats capture. *Artibeus jamaicensis* bats were captured at Morelos State (located in Central Mexico). Bats were trapped in mist nets opened for 4 hours after sunset. Bat species were identified according to the guide of Medellín and others.²¹ Lactating and pregnant females were released immediately without any further manipulation. Captured bats were placed in cloth bags for transportation to the enclosure at the animal facility of the National Public Health Institute (INSP, Cuernavaca, Mexico). No bat died or suffered injury as a result of capture or handling procedures. The collection of specimens was performed according to the guidelines of the American Society for the Use of Mammalogists of Wild Mammals in Research and under a collecting permit issued by the General Direction of Wildlife of Mexico (permission number SGPA/DGVS/00471/11).

Bats enclosure and handling. Enclosures were constructed using PVC pipes of 1.5 inch in diameter and PVC connectors to build a structure of 1 cubic foot, covered with shade netting tied to the base frame tubes. The enclosure had two apertures on the top side for introduce nourishment. After placement in the cages, bats were left in quarantine for 1 week and fed fresh fruits daily *ad libitum*.

Virus propagation and titration. The DENV serotype 1, strain Hawaii (a kind gift of Dr. Juan Salas, Instituto Politecnico Nacional, Mexico), and DENV serotype 4, strain H241 were both propagated in C6/36 HT as previously described.²² Briefly, monolayers grown in 75 mm² flasks were inoculated with DENV at multiplicity of infection of 1 for 1 h. After inoculation, cells were maintained in minimum essential medium (MEM) (Gibco, Grand Island, NY) supplemented with 7% fetal calf serum (FCS) and maintained at 34°C for 1 week. Cells were freeze-thawed twice, the media clarified by low speed centrifugation and stored in aliquots (0.6 mL) at -70°C until used. The DENV titers were determined by plaque assay inoculating 10-fold dilutions on confluent monolayers of BHK-21 cells grown in 24-well plates, following protocol described by Ludert and others.²³

Bat experimental inoculations. A total of 52 *A. jamaicensis* bats were used in four experimental inoculations as follows. For experimental inoculation (EI) 1, 10 bats were inoculated subcutaneously in the chest with 200 µL of DENV-4 inoculum. For EI-2, 8 bats were inoculated intraperitoneally, also with 200 µL of DENV-4. For EI-3, 10 bats were inoculated intraperitoneally with 200 µL of DENV-1. The inoculum titers were 2.4×10^4 plaque-forming units (PFU)/mL and 8.4×10^4 PFU/mL, for DENV-1 and DENV-4, respectively. The viability of the viral inoculum was tested in parallel by inoculation of suckling mice in the brain; inoculated mice were watched for the appearance of neurological signs 5–7 days after inoculation and their brains collected to test for the presence of viral RNA by RT-PCR. For each EI, five bats were mock inoculated under identical conditions with supernatant of uninfected C6/36 cells, and kept as controls. Mock

inoculated bats were maintained in separate enclosures. Finally, for EI-4, mosquitoes from group “a” and “b”, taken at Day 9 after the infection with DENV²⁴ were deprived of sucrose for 4 hours and placed in a small container with a mosquito net. Bats ($N = 5$) were settled inside the container but outside of the mosquito net and five mosquitoes allowed to feed on each bat until the mosquito engorged (~20 min).²⁵ Following exposure to the mosquito bites, bats were returned to the enclosures until sampled or euthanized. Calendars and experimental program of each EI are shown in Table 1.

Collection of spleen tissue and plasma. Before their enrollment in any of the experimental inoculations, all bats were bled by puncture of the cephalic vein using a needle of 21G Becton Dickinson (BD), 3 or 4 days after capture and just the day before the experimental inoculation. Blood (~80 µL) was collected with heparinized capillary tubes and stored at -70°C until RNA extraction to test for the presence of genomic DENV RNA by RT-PCR.

Following the experimental program (Table 1), bats were anesthetized with Pentobarbital (0.2 mL/mL total volume) and euthanized for exsanguination by intracardiac puncture using a sterile syringe with 100 µL of heparin per 1 mL of blood. Blood was centrifuged at $1,300 \times g$ for 5 minutes to obtain the plasma. After obtaining blood samples, all animals were necropsied and the spleen collected. Organs and plasma were frozen at -70°C and kept until RNA extraction to test for the presence of genomic DENV RNA by RT-PCR.

An aliquot of ~0.5 mL of whole blood from each of the euthanized animals from EI-3 was immediately put in mosquito feeders, and used to feed a pool of 10 mosquitoes. Mosquitoes were maintained under controlled conditions, and 7 days after the blood meal, each pool was macerated in phosphate buffered saline, the mixture clarified by low speed centrifugation and the clarified supernatant tested for the presence of soluble NS1 by a commercial enzyme-linked immunosorbent assay (ELISA).²⁶

Mosquito strains. *Aedes aegypti* mosquitoes (Rockefeller strain), were maintained under controlled conditions at 26 to 28°C and 70–80% relative humidity with 12-h/12-h light-dark cycle at the insectary of the National Public Health Institute (INSP), Cuernavaca, Mexico. Mosquito eggs were hatched and larvae were reared in water containers at a density of 200–300 larvae per liter and daily fed with high quality food as described by Moreno-Garcia and others.²⁷ Pupae were transferred to screened cages and emergent adults were maintained with an *ad libitum* diet of 10% sucrose solution.

Oral mosquito infections. Adults mosquitoes (4–6 days old) were deprived of sucrose for 4 hours, and afterward fed with an infectious blood meal for 40 minutes through a piece of Parafilm M (Chicago, IL) clamped to mosquito feeders. Infection meals consisted of 500 µL of fresh heparinized rabbit blood mixed with an equal volume of viral inoculum. During the meal, the feeding mixture was maintained at 37°C by circulating warm water through the feeders. Group “a” mosquitoes were fed with DENV-4 (4.2×10^4 PFU/mL) and group “b” mosquitoes with DENV-1 (1.2×10^4 PFU/mL). Doses in this range have proven to be infectious for *A. aegypti* females²⁸. Three days after the blood meal and up to day 12, five female mosquitoes from each group were dissected daily and the thorax and abdomen each pooled and macerated in 100 µL of TRIzol Reagent (Invitrogen, Carlsbad, CA) following the manufacturer’s protocol. Extracted RNA was

TABLE 1
Experimental inoculations schedule

Experimental inoculation	DENV serotype	Route of inoculation	Day 0		Day 1 to Day 17																		
			Mock infected	Bats infected	Day 1	Day 2	Day 3	Day 4	Day 5	Day 6	Day 7	Day 8	Day 9	Day 10	Day 11	Day 12	Day 13	Day 14	Day 15	Day 16	Day 17		
EI-1	4	Subcutaneous	N = 5	A	N = 10	E	BS	E	BS	E	BS	E	BS	E	BS	E	BS	E	BS	E	BS	E	
				1, 2																			
				3, 4																			
				5, 6																			
				7, 8																			
9, 10																							
EI-2	4	Intraperitoneal	N = 4	F	N = 8	E	BS	E	BS	E	BS	E	BS	E	BS	E	BS	E	BS	E	BS	E	
				11, 12																			
				13, 14																			
				15, 16																			
				17, 18																			
EI-3	1	Subcutaneous	N = 5	J	N = 10	E	BS	E	BS	E	BS	E	BS	E	BS	E	BS	E	BS	E	BS	E	
				19, 20																			
				21, 22																			
				23, 24																			
				25, 26																			
				27, 28																			
EI-4	1 and 4	Mosquito bite	N = 5	Group "a" DENV-4	N = 5	BS	E	BS	E	BS	E	BS	E	BS	E	BS	E	BS	E	BS	E	BS	
				1a																			1b
				2a																			2b
				3a																			3b
				4a																			4b
				5a																			5b

Procedure: **E**: Euthanized; **BS**: Blood sampled.
 Experiment 4: Group a: Bats bitten for mosquitoes infected with DENV-4.
 Group b: Bats bitten for mosquitoes infected with DENV-1.

immediately tested for the presence of DENV RNA by RT-PCR. In agreement with previous results Salazar and others,²⁴ the maximum infections rates of the mosquitoes were observed at Day 5 in the abdomen and at Day 8 in the thorax.

Detections of dengue virus genome (RT-PCR). Total RNA was extracted from all plasma or organs with TRIzol Reagen (Invitrogen) following the manufacturer's protocol. The RNA was reverse transcribed into cDNA for 50 min at 42°C using hexamer random primers (New England, Biolabs, Ipswich, MA) and Invitrogen SuperScript II Reverse Transcriptase (Invitrogen) according to the manufacturer's recommendations. Two microliters of cDNA were used in a PCR carried out using the Taq DNA polymerase (Kapa Biosystems, Wilmington, MA) and primers D1 (forward) and D2 (reverse) as reported in Lanciotti and others.²⁹ Water instead of nucleic acids and nucleic acids extracted from mice brain inoculated with DENV-1 or 4 were used in the RT-PCR reactions as negative and positive controls, respectively. As an additional positive control, the cDNA generated with the random primers was amplified in parallel by PCR using primers specific for the housekeeping gene GAPDH (forward 5'-GCAGGGGG GAGCCAAAAGGG-3'; reverse 5'-TGCCAGCCCCAGCG TCAAAG 3'). Both amplifications were carried out as follows: initial incubation at 94°C for 10 min, then 30 cycles of denaturation at 94°C for 30 s, annealing at 55°C for 30 s, and extension at 72°C for 30 s, followed by a final incubation at 72°C for 10 min. The PCR products were analyzed by 1.5% agarose gel electrophoresis and visualized by staining with ethidium bromide.

Enzyme immune assay (ELISA). All plasmas collected from the DENV inoculated bats ($N = 38$) were tested in duplicate for the presence of DENV NS1 protein, using the commercial assay Platelia Dengue NS1 Ag (Bio-Rad, Hercules, CA). This diagnostic test is a one-step sandwich format microplate enzyme immunoassay designed for the qualitative or semi-quantitative detection of DENV NS1 antigen in human sera or plasma.⁵ The assay was carried and the cut-off values were established following the manufacturer's protocol.

Plasma samples from the DENV ($N = 22$) and mock ($N = 5$) inoculated bats were evaluated for the presence of specific immunoglobulin G (IgG) using the reagents and materials provided in the Panbio Diagnostics Dengue IgG capture ELISA. This commercial assay is intended as an aid in the diagnosis of DENV infection and is based on the capture of human IgG by anti-human IgG antibodies coated onto the plate. Because bat IgG was not captured in the plate (data not shown), a competition ELISA was set-up. Briefly, bat plasma (1:25 final dilution) was reacted with the DENV 1-4 antigen provided with the kit for 1 h at 37°C before the Mab tracer (anti-E Mab labeled with horse radish peroxidase) provided with the assay was added to the mixture and further incubated for 1 h at 20–25°C. In parallel, the assay plate was reacted with the human serum positive for DENV IgG provided with the assay as a positive (reactive) control for 1 h at 37°C. The antigen mixture was added to the plate, previously washed 6 times, and the plate covered and incubated for 1 h at 37°C. Finally, the plate was washed again ($6\times$) and 100 μ L of the tetramethylbenzidine (TMB) chromogen reagent added to each well. After 10 min. incubation at 20–25°C, the colorimetric reaction was stopped by adding 100 μ L of acid stop solution. Absorbance of each well was read immediately at a wavelength of 450 nm. As competition controls, one human serum, tested negative for anti-dengue IgG by the Panbio

Dengue capture ELISA and by plaque reduction neutralization titer (PRNT) ($\text{PRNT}_{50} < 20$) and one human serum positive for DENV IgG by the Panbio Dengue ELISA and by PRNT ($\text{PRNT}_{50} = 320$) were used. As an additional control, a mouse monoclonal anti-E antibody (4G2) ($\text{PRNT}_{50} = 1,250$) was also used. The calibrator, reactive and negative controls were also run in parallel in a standard assay as quality controls. Any reading below the mean optical density (O.D.) plus three standard deviations, obtained with the negative human serum was taken as indicative of the presence of anti-DENV IgG in the bat's plasma.

RESULTS

No clinical signs were observed in any of the bats experimentally inoculated with DENV virus by any of the inoculation routes used. In addition, no evident changes in the behavior or in the feeding pattern of the bats were observed. Upon necropsies, no skin damages at the inoculation site, hemorrhages, or apparent pathological signs were observed.

The RT-PCR analysis of RNA extracted from plasma and spleen of any of the bats experimentally inoculated by any of the routes or directly exposed to mosquito bites, failed to show evidence of viremia or the presence of DENV nucleic acid in the tested organs (Table 2). The spleen sample from one bat from EI-1 killed on Day 7 post-inoculation showed a faint amplicon of ~500 base pairs after the PCR assay. Similar amplicons were also observed for the spleens of bats killed on Days 4, 7, and 9 from EI-2 and Days 10 and 12 from EI-3. However, these results could not be reproduced after using either the original cDNA or new tissue samples. Moreover, direct sequencing of the amplicons confirmed their spurious nature (data not shown). No positive RT-PCR results were obtained from bats of EI-4. In contrast, the amplification of the GAPDH mRNA carried out in parallel as a control of nucleic acid extraction for each sample clearly showed the 600 base pairs of the expected amplicon. These results indicate that no viral replication took place in the DENV inoculated animals. In view of these results and the limited amount of plasma available, assessing viremia by plaque assay was not attempted.

The presence of circulating NS1 was evaluated in the plasma collected from the DENV inoculated animals by ELISA (Table 2). All plasma were tested for the presence of NS1 regardless of the RT-PCR results, because they expect kinetics in bats of NS1 antigenemia with respect to viremia could not be anticipated. Plasma collected from mock infected bats ($N = 14$) were also analyzed as controls. Readings slightly above the positive cut-off value of the assay were observed for one plasma collected from EI-1 and one from EI-2 on Day 7 post infection (p.i.) and also for plasmas collected each on p.i. Days 16 and 17 of EI-4, but these results were not reproducible. In addition, strong O.D. readings were observed for plasma collected from the two animals killed on Days 1 (bats 19 and 20) and 4 (bats 21 and 22) p.i. of EI-3. However, levels of circulating NS1 were undetectable in plasma collected at later times (p.i. Days 7, 10, and 12) from the same EI-3. These results suggest that not NS1 sustained antigenemia appeared after the experimental inoculation of the bats.

One of the 10 pools of mosquitoes fed with the fresh blood collected from EI-3 inoculated bats, tested positive for the presence of NS1. This pool corresponded to a bat euthanized

TABLE 2
Results of experimental inoculation in *Artibeus jamaicensis* bats

Days post inoculum	EI-1			EI-2		
	Bats positive to the following tests			Bats positive to the following tests		
	Platelia NS1	RT-PCR Lanciotti	RT-PCT GAPDH	Platelia NS1	RT-PCR Lanciotti	RT-PCT GAPDH
Day 1	–	–	1, 2	–	–	11, 12
Day 2	–	–	No samples	–	–	No samples
Day 3	–	–	3, 4	–	–	No samples
Day 4	–	–	No samples	–	13 (suspected)	13, 14
Day 5	–	–	5, 6	–	–	No samples
Day 6	–	–	No samples	–	–	No samples
Day 7	8	8 (suspected)	7, 8	15	16 (suspected)	15, 16
Day 8	–	–	No samples	–	–	No samples
Day 9	–	–	9, 10	–	18 (suspected)	17, 18
Days post inoculum	EI-3			EI-4		
	Bats positive to the following tests			Bats positive to the following tests		
	Platelia NS1	RT-PCR Lanciotti	RT-PCT GAPDH	Platelia NS1	RT-PCR Lanciotti	RT-PCT GAPDH
Day 1	19, 20	–	19, 20	–	–	No samples collected
Day 2	–	–	No samples	–	–	–
Day 3	–	–	No samples	–	–	–
Day 4	21, 22	–	21, 22	–	–	–
Day 5	–	–	No samples	–	–	1a, 1b
Day 6	–	–	No samples	–	–	No samples
Day 7	–	–	23, 24	–	2a (suspected)	2a, 2b
Day 8	–	–	No samples	–	–	No samples
Day 9	–	–	No samples	–	–	No samples
Day 10	–	26 (suspected)	25, 26	–	–	No samples
Day 11	–	–	No samples	–	–	3a, 3b
Day 12	–	28 (suspected)	27, 28	–	–	No samples
Day 13	–	–	–	–	–	No samples
Day 14	–	–	–	–	–	No samples
Day 15	–	–	–	–	–	No samples
Day 16	–	–	–	4b	–	4a, 4b
Day 17	–	–	–	5a	5a (suspected)	5a, 5b

Results obtained with Mock inoculated bats are not present for simplicity. The reverse-transcription-polymerase chain reaction (RT-PCR) and NS1 assays carried out the samples collected from these animals all resulted as negatives.

on Day 1 after inoculation (bat 19). Mosquitoes fed with blood from mock infected bats or with blood collected from inoculated bats afterward were negative for the presence of NS1 (data not shown).

None of the bat plasma samples tested caused a reduction in the O.D. readings in the competition ELISA set to detect the presence of DENV specific IgG (mean O.D. reading 1.168 ± 0.066 versus 1.061 ± 0.049 for the negative human serum). In contrast, over 98% reduction in the O.D. reading was observed with the human serum (0.022 ± 0.010) and anti-E Mab (0.009 ± 0.004) included as positive controls. These results suggest that no serological immune response was induced in the DENV inoculated bats and no anti-DENV IgG was present in the plasma of the mock inoculated animals.

DISCUSSION

Bats are a natural host for a large number of viruses, some of which are important human pathogens. Recent metagenomic analyses have identified viruses belonging to the *Retroviridae*, *Herpesviridae*, *Poxviridae*, *Flaviviridae*, *Reoviridae*, *Picobirnaviridae* families, among others, as part of the bat viromes.^{30–34} Moreover, the existence of several yet unknown mammalian viruses in bats is indicated by these studies. Dengue is the most important viral disease transmitted by mosquitoes to humans. The implementation of control measures for this disease is of public health interest and a clear understanding of the life cycle of the DENV in nature,

including identification of possible natural hosts is a necessity. The aim of our study was to test the susceptibility of the species of bat *A. jamaicensis* to DENV infection. We decided to use this species as an experimental model because it is an abundant species of fruit-eating habits and very tolerant to habitat disturbance,¹⁸ which allows *A. jamaicensis* to cohabit with humans in rural and semi-rural areas where DENV is endemic. Moreover, evidence for DENV infection in wild *Artibeus* sp. bats has been reported.^{11,13,15}

For EI-1, DENV-4, and the subcutaneous route were used in an attempt to generate an initial localized infection of the skin resident cells of the immune system.² Skin resident dendritic cells and macrophages are supposed to be first cells encountered by the DENV after a natural infection by a mosquito bite.^{35,36} The subcutaneous route has been used successfully for the humanized mouse models of DENV infection³⁶ and also has been used in the experimental inoculation of bats with West Nile virus (WNV) and Japanese *encephalitis virus* (JEV).^{37,38} In view of the results obtained after EI-1 and in an attempt to augment tissue virus adsorption, for EI-2 the same DENV-4 strain was used as inoculum but inoculated using the intraperitoneal route. This route has been used in experimental protocols to inoculate mice with DENV³⁹ and bats with DENV and other flaviviruses.^{40,41} However, no evidence of viremia, antigenemia or serological immune response was found after EI-1 or EI-2. Analysis of spleen tissue, an organ found to be infected in humans⁴² and bats¹⁵ also failed to show evidence of DENV replication.

In a study conducted by Barr and others⁴³ cell lines of different species, including the TB Lu cell line I, which is a pulmonary epithelium cell line derived from the *Tadarida brasiliensis* bat, were infected *in vitro* with the 4 DENV serotypes. Evidence of infection in the bat cell line was found by qRT-PCR 72 h p.i., only with the DENV-1 inoculum. Furthermore, DENV-1 sequences have been isolated from wild bats.¹¹ Therefore, a DENV serotype 1 strain was used as inoculum for EI-3. In addition, because no data about the possible kinetics a DENV infection in bats are available, the follow-up time after inoculum was extended from 9 to 12 days in EI-3. Although no evidence of viremia or viral replication in the spleen was found in the inoculated bats, the plasma collected on Days 1 and 4 after inoculation showed positive values for NS1 (antigemia). Dengue NS1 protein levels are known to rapidly rise in sera during acute phase and to decrease about 7 days after fever onset.^{44,45} However, antigenemia levels usually parallel viremia levels in patients. Thus, in the absence of any evidence for viremia, these results are difficult to interpret. In addition, because the DENV-1 inoculum titer was three times higher than the DENV-4 titer and the Platelia Dengue NS1 Ag assay is much more sensitive detecting DENV-1 NS1 than DENV-4 NS1,⁴⁶ the possibility that the NS1 detected in the plasma of bat from EI-3 correspond to the inoculum is highly plausible. Of note, mosquitoes fed from the blood of the bats that tested positive for NS1 also tested positive for this antigen.

The importance of the mosquito saliva to enhance dengue infection of the vertebrate host is well established.^{25,47-49} Therefore for EI-4, infections of the bats by mosquito bite was attempted. Both DENV serotypes 1 and 4 were used and the bats followed up for 17 days. *Aedes aegypti* mosquitoes of the Rockefeller strain, known to be highly susceptible to DENV infection, were used. The kinetics of mosquito infection, followed by RT-PCR, was in agreement with previous reports²⁴ and no differences in kinetics were observed between DENV-1 and 4. The number of infected mosquitoes per bat ($N = 5$) was as reported by Cox and others²⁵ to infect humanized mice and mosquitoes were retired only after observed to have had a blood meal. Despite all these precautions and controls, no evidence of viremia or organ viral replication was observed in any of the EI-4 bats.

The results from all the EIs taken together strongly suggest that *A. jamaicensis* bats are incapable of sustaining DENV replication. This conclusion is supported by the fact that highly sensitive and specific methods such as RT-PCR for virus genome detection and ELISA detection of circulating NS1 were used to follow-up infection in the bats. In addition, mosquitoes highly susceptible to DENV infections were used to detect or to induce infection in EI-3 and EI-4, respectively. Deficiencies in the quality or the amount of the inoculum to explain these results are unlikely. The viability of the inoculum was corroborated by parallel titrations in neonate mice and the amount of viruses used in each inoculum (10^3 to 10^4 PFU/bat) is similar or comparable to the inoculum used in the mouse animal models.⁸ The parallel detection of mRNA from housekeeping genes (GAPDH) from spleen tissue indicates appropriate nucleic acid extraction and amplification protocols. In addition, no apparent changes in bats feeding habits or behavior were observed during the experiments, suggesting that neither the manipulation procedure nor the captivity affected significantly the animals. Our results are in agree-

ment with previous results obtained by Perea-Martinez and others⁴¹ after the intraperitoneal experimental inoculation of 27 *Artibeus intermedius* bats with DENV serotype 2. Our results are also in agreement with previous reports of experimental inoculation of 2 species of North American bats with WNV or Australian black flying foxes with JEV, two flavivirus closely related to DENV, and one species of Asian bats with yokose virus, a flavivirus distantly related to DENV.^{37,38,40} In contrast, low to relatively high levels of susceptibility were observed in three species of insectivorous bats experimentally inoculated with several strains of JEV or St. Louis encephalitis virus (SLEV).⁵⁰

One limitation of this work is that field-collected bats were tested only for active DENV infection and not for anti-DENV antibodies that may have prevented infection, before the EIs. However, if these animals presented past DENV infections, a robust memory IgG response was to be expected after the EIs and this was not observed. The same rationale will hold true if putative cross-reactive flaviviral antibodies caused by past infection with other arthropod-borne flavivirus were to be present.

The results obtained in this and other experimental inoculations of flaviviruses in bats and the several reports of DENV infection in wild bats are not easy to reconcile. However, the current evidence of DENV infection in bats is based on serology or the amplification of DENV-related sequences. Cross-reactivity among the more than 70 members of the *Flaviviridae* family is well established⁵¹ thus adding confusion to the serological data, and the significance of isolated DENV sequences is hard to assess. On the other hand, despite their abundance, many things about the biology and ecology of bats remain unknown, and the possibility that conditions in captivity fail to fully reproduce the conditions in which bats encounter viruses in nature cannot be excluded. Further experiments, including *bonafide* DENV isolates and robust flavivirus metagenomic analysis from bats, are still required to fully evaluate the role played by bats in DENV ecology.

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Authors' addresses: Salomé Cabrera-Romo, Ana C. Alcalá, and Rosa María del Ángel, CINVESTAV - Infectomics and Molecular Pathogenesis, Mexico City, Mexico, E-mails: gorilagorila@yahoo.com.mx, ana.acal1711@gmail.com, and rmangel@cinvestav.mx. Benito Recio-Tótoro and Humberto Lanz, Instituto Nacional de Salud Pública, Center for Research on Infectious Diseases, Cuernavaca, Mexico, E-mails: enitor@gmail.com and humberto@insp.mx. Victor Sánchez-Cordero and Angel Rodríguez-Moreno, Institute of Biology - UNAM, Mexico City, Mexico, E-mails: victor@ibunam2.ibiologia.unam.mx and tanicandil@hotmail.com.

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