

Oral and cloacal microflora of wild crocodiles *Crocodylus acutus* and *C. moreletii* in the Mexican Caribbean

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ABSTRACT: Bacterial cultures and chemical analyses were performed from cloacal and oral swabs taken from 43 American crocodiles *Crocodylus acutus* and 28 Morelet's crocodiles *C. moreletii* captured in Quintana Roo State, Mexico. We recovered 47 bacterial species (28 genera and 14 families) from all samples with 51.1 % of these belonging to the family *Enterobacteriaceae*. Fourteen species (29.8 %) were detected in both crocodile species and 18 (38.3 %) and 15 (31.9 %) species were only detected in American and Morelet's crocodiles, respectively. We recovered 35 bacterial species from all oral samples, of which 9 (25.8 %) were detected in both crocodile species. From all cloacal samples, we recovered 21 bacterial species, of which 8 (38.1 %) were detected in both crocodile species. The most commonly isolated bacteria in cloacal samples were *Aeromonas hydrophila* and *Escherichia coli*, whereas in oral samples the most common bacteria were *A. hydrophila* and *Arcanobacterium pyogenes*. The bacteria isolated represent a potential threat to crocodile health during conditions of stress and a threat to human health through crocodile bites, crocodile meat consumption or carrying out activities in crocodile habitat. We especially warn about the presence of *Salmonella arizona* and *S. typhi*, which cause enteritis and septicemia in crocodiles and salmonellosis and typhoid fever in humans. The risk of bacterial contamination from crocodiles to humans could increase in the future because of the accelerated destruction of crocodile habitat, which could lead to an augmentation of human–crocodile interactions. Information on bacterial diversity reported here could help in the choice of antibacterial products in case of infections that are of crocodile origin.

KEY WORDS: Bacterial flora · *Crocodylus acutus* · *Crocodylus moreletii* · *Salmonella* · Cozumel · Río Hondo · Banco Chinchorro · Mexico

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INTRODUCTION

Crocodilians have the strongest bite of all living animals (Erickson et al. 2003) and can cause serious injuries. The gravity of wounds ranges from simple punctures or tearing skin wounds to fractures, amputations and death (Caldicott et al. 2005, Hertner 2006, Gruen 2009, Cupul-Magaña et al. 2010, Langley 2010). Conflicts between humans and crocodilians

have been reported in many countries and while in some regions attacks on humans are uncommon, in others they seem to occur frequently (Pooley et al. 1989, Lamarque et al. 2009, Wamisho et al. 2009). Those adverse encounters are expected to increase in the future because of the modification and destruction of crocodile habitat or the increase in crocodile numbers (Scott & Scott 1994, Caldicott et al. 2005, Vyas 2010).

Besides the physical damage, survivors of attacks from crocodilians can develop wound infections caused by bacteria or fungi (Flandry et al. 1989, Caldicott et al. 2005, Wamisho et al. 2009, Langley 2010). Several studies display the existence of many bacteria and fungi in the oral cavity of wild and captive crocodilians (Doering et al. 1971, Gorden et al. 1979, Flandry et al. 1989, Matushima & Ramos 1995, Anderson 1999, Cupul-Magaña et al. 2005, Silva et al. 2009). While several antibiotics provide protection from wound infections, in some cases patients can die from sepsis (Vanwersch 1998, Caldicott et al. 2005). Crocodilians show great resistance and healing capacity to injuries and illness, but several studies report diseases or symptoms in crocodilians related to bacterial infections, which can even lead to the death of individuals (Shotts et al. 1972, Novak & Seigel 1986, Hernández-Hurtado & Cupul-Magaña 1996, Mehrotra et al. 2000, Huchzermeyer 2003, Bishop et al. 2007, Garcia et al. 2008).

The Quintana Roo State is the major tourist region in Mexico, where increasing coastal development continues to destroy and fragment the habitats of the American crocodile *Crocodylus acutus* and Morelet's crocodile *C. moreletii* (Cedeño-Vázquez et al. 2006, Machkour-M'Rabet et al. 2009). This reduction of habitat, along with other factors, is causing the depletion of crocodile populations and could also increase human–crocodile encounters (Cupul-Magaña et al. 2010). Several cases of crocodile attacks have been reported in Quintana Roo and are often linked to fishing and spear-fishing activities (Lazcano-Barrero 1996, Cantera 2000).

Although the number of studies on crocodiles has increased in the last decade in Quintana Roo, they have yet to examine the bacterial flora of crocodiles in this region. The objective of the present study was to determine the bacterial flora in oral and cloacal cavities of wild American and Morelet's crocodiles in Quintana Roo. This information would then serve in the assessment of health risks to humans after encounters with crocodiles.

MATERIALS AND METHODS

Study site

Samples were obtained from American crocodiles captured on Cozumel Island and Banco Chinchorro Biosphere Reserve, and from Morelet's crocodiles captured in the Río Hondo. In Cozumel, crocodiles were captured at Colombia lagoon, Chunchaka'ab lagoon and Xtakún lagoon in the Ecological Reserve

of Punta Sur in the southern tip of the Island (Fig. 1). Banco Chinchorro is an atoll located 30 km east from the coast of Quintana Roo and crocodiles were captured at Cayo Centro, the largest Cay of the atoll (Fig. 1). These reserves have relatively well-conserved populations of American crocodiles with individuals of all size classes and with male-biased sex ratios (Charruau et al. 2005, 2010, González-Cortés 2007). The Río Hondo is a relatively deep river (mean depth of about 8 m) that forms the border between México and Belize in South Quintana Roo (Fig. 1). Cedeño-Vázquez et al. (2006) found a relatively high number of Morelet's crocodiles in the Río Hondo with individuals of all size-classes and equal sex ratio and encounter rates that ranged from 1.72 to 4.70 crocodiles km^{-1} .

Sample collection

Crocodiles were captured at night by hand or by using the break-away snare technique, depending on their size, during spotlight surveys in May 2008 and May 2009 in Cozumel, in August 2010 in Banco Chinchorro and from June 2009 to February 2010 in Río Hondo. Each captured individual was physically examined for signs of diseases, sex was determined by cloacal examination (Brazaitis 1968, Ziegler & Olbert 2007) and each was measured from the tip of the snout to the tip of the tail (± 0.5 cm). Crocodiles were classified by size class based on their total length (TL). We used the following classes for American crocodiles: hatchlings (TL < 30 cm), yearlings (TL = 30.1 to 60 cm), juveniles (TL = 60.1 to 120 cm), subadults (TL = 120.1 to 180 cm) and adults (TL > 180 cm). For Morelet's crocodiles, we used the following classes: hatchlings (TL < 30 cm), yearlings (TL = 30.1 to 50 cm), juveniles (TL = 50.1 to 100 cm), subadults (TL = 100.1 to 150 cm) and adults (TL > 150 cm). Samples were taken by passing a sterile culture swab (BBLTM CultureSwabTM Plus, Becton-Dickinson) in the oral cavity and the cloaca of each crocodile. Swabs were then preserved at 2 to 8°C in AMIES transport mediums (Becton, Dickinson), which are ideal for aerobic and anaerobic bacteria, and were sent for analyses to the Africam Safari Zoo laboratory in Puebla, Mexico.

Bacterial analyses

Bacterial cultures were grown on various bacterial-growth media (Cowan & Steel 1974, Quinn et al.

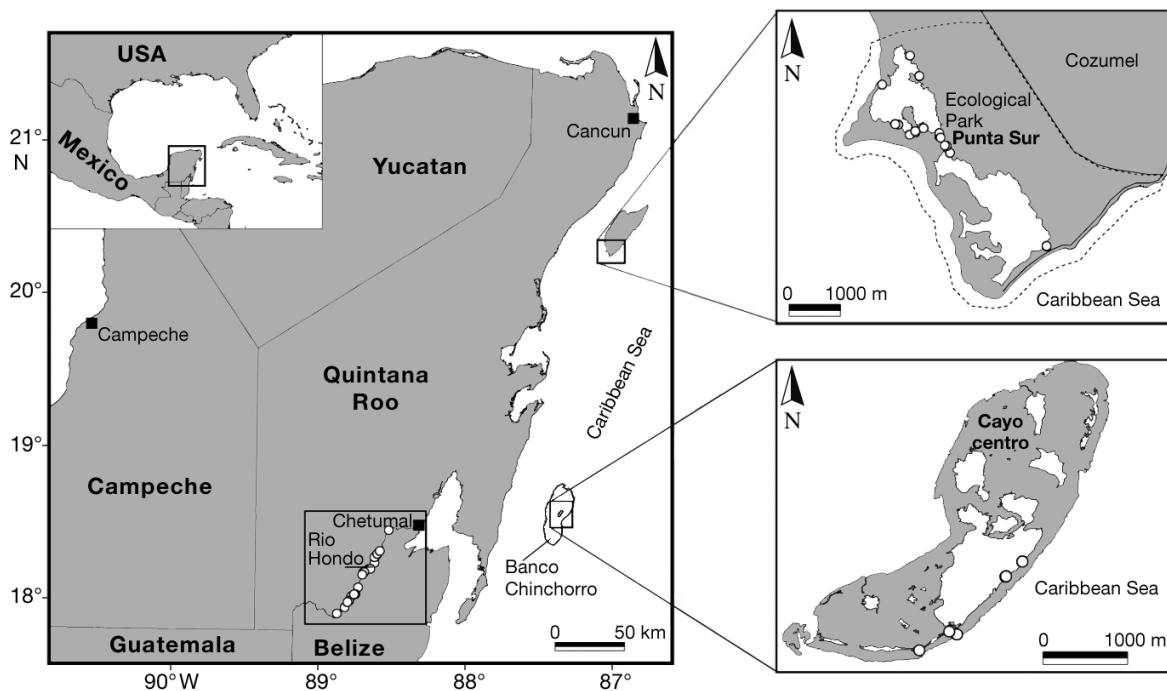


Fig. 1. *Crocodylus acutus* and *C. moreletii*. Location of study areas and capture sites (open circles) of American and Morelet's crocodiles in Quintana Roo State. Inset in left panel shows location of study areas in Mexico. Dotted line in upper right panel indicates boundary of ecological park

1994); blood agar 5%, McConkey agar, mannitol salt agar, brilliant green agar, and *Salmonella-Shigella* agar.

The quadrant streak method was used to inoculate whole plates. These plates were then incubated at 37°C for 24 h. Plates of blood agar were incubated in a carbon dioxide atmosphere of 5% and the other plates were incubated in aerobic conditions (Quinn et al. 1994). After 24 h of incubation, bacteria morphology was analyzed microscopically and by gross observation to record characteristics such as color, size, form, hemolysis, pigment production, smell and reaction to the indicators of the differential medium. Subsequently, each bacterial colony was submitted to catalase and oxidase tests and a smear from each colony was heat-fixed for Gram staining. If it was necessary to purify the strains, a second inoculation was performed and for the next 24 h these plates were subjected to the same procedures as were the first inoculations.

After gross observations, we proceeded to identify the Gram-negative bacteria by means of the following biochemical tests: mobility, presence of indole, decarboxylation of ornithine and lysine, fermentation of lactose, sucrose and dextrose, oxidation of iron, use of citrate as a source of carbon, hydrolysis of

urea, gas production, deamination of phenylalanine, nitrate reduction and differentiation of oxidative or reductive metabolism of bacteria (Kämpfer et al. 1991, MacFaddin 2000). All these reactions, along with the characteristics obtained during the primary identification, were compared with reference tables of biochemical reactions to obtain the final identification (Cowan & Steel 1974, Quinn et al. 1994, Murray et al. 1999).

For Gram-positive and Gram-negative bacteria requiring additional tests for identification we used a miniaturized method with the help of the BBL Crystal™ Enteric/Non-fermenter Identification System and the BBL Crystal™ Rapid Gram-positive Identification System (Becton-Dickinson), both of which contain 30 dried biochemical and enzymatic substrates. Fermentation reactions detect the ability of an isolate to metabolize carbohydrates in the absence of atmospheric oxygen, and oxidation reactions are based on the ability of an organism to metabolize the substrate with oxygen as the final electron acceptor. Both reactions are usually detected by means of a pH indicator in the test substrate (Killian & Bulow 1976, MacFaddin 2000). For *Salmonella* and *Shigella* species, samples were sent for serotyping to a laboratory (Asesores Especializados en Laboratorios S.A. de C.V.).

Crocodile attacks

We used the Google search engine to search for any publications (scientific articles, reports, press articles) on attacks by crocodiles on humans in Mexican states of the Yucatan Peninsula (Quintana Roo, Campeche and Yucatan) in order to evaluate the number of attacks and changes in their frequency over time in the region. We also searched in the publications found for any reference to bacterial infection of a victim's wounds.

Statistical analysis

We used linear regressions to test the relationship between the number of bacteria in cloacal and oral cavities and crocodile total length. Results were considered significant at $p < 0.05$.

RESULTS

Samples collected

We captured 43 American crocodiles (4 hatchlings, 8 yearlings, 18 juveniles, 4 subadults and 9 adults) and 28 Morelet's crocodiles (4 hatchlings, 8 yearlings, 5 juveniles, 7 subadults and 4 adults). Thirty-five American crocodiles were males (4 hatchlings, 8 yearlings, 16 juveniles, 3 subadults and 4 adults) and 8 were females (2 juveniles, 1 subadult and 5 adults). Of the Morelet's crocodiles captured, 22 were males (3 hatchlings, 7 yearlings, 2 juveniles, 6 subadults and 4 adults) and 6 were females (1 hatchling, 1 yearling, 3 juveniles and 1 subadult). We obtained oral samples from 38 American crocodiles (Table 1) and from all the 28 Morelet's crocodiles (Table 2). Five samples of juvenile American crocodiles were negative for bacteria after 72 h of incubation. We obtained cloacal samples from all American and Morelet's crocodiles captured (Tables 3 & 4).

American crocodile

From the American crocodiles we isolated 32 bacterial species, of which 5 (15.6%) were found in both oral and cloacal cavities and 17 (53.1%) and 10 (31.3%) were only found in the oral cavity and cloacal cavity, respectively. From oral samples of American crocodile we isolated 22 species of bacteria, of which 4 were determined to the genus level and 18 to the spe-

cies level (Table 1). Of these bacteria 12 (54.5%) were Gram-positive and 10 (45.5%) were Gram-negative (Table 1). *Arcanobacterium pyogenes* and *Aeromonas hydrophila* were the most frequent isolates from oral samples and were isolated from 18 and 11 crocodiles, respectively (Table 1). The other bacteria were isolated from 1 to 6 crocodiles (Table 1). The number of bacteria species per crocodile ranged from 0 to 3 and was not correlated with crocodile total length ($r^2 = 0.023$, $p = 0.36$, $n = 28$). From cloacal samples of American crocodiles we found 15 species of bacteria, of which 2 were determined to the genus level and 13 to the species level (Table 3). Two (13.3%) of these bacteria were Gram-positive and 13 (86.7%) were Gram-negative (Table 3). *Escherichia coli*, *A. hydrophila* and *Salmonella enterica* subsp. *arizona* were the most frequent bacteria in cloacal samples and were isolated from 29, 16 and 14 crocodiles, respectively (Table 3). The other bacteria were isolated from 1 to 8 crocodiles (Table 3). The number of bacteria species per crocodile ranged from 1 to 3 and was not correlated with crocodile total length ($r^2 = 0.058$, $p = 0.12$, $n = 43$).

Morelet's crocodile

From the Morelet's crocodiles we isolated 29 different bacteria, of which 7 (24.1%) were found in both oral and cloacal cavities and 15 (51.7%) and 7 (24.1%) were only found in the oral cavity and cloacal cavity, respectively. From oral samples we isolated 22 species of bacteria, of which 4 were determined to the genus level and 18 to the species level (Table 2). Eight (36.4%) of these bacteria were Gram-positive and 14 (63.6%) were Gram-negative (Table 2). The most frequent bacteria in the oral cavity of Morelet's crocodiles appeared to be *Aeromonas hydrophila* and *Klebsiella pneumoniae*, which were isolated from 13 and 7 crocodiles, respectively (Table 2). All the other bacteria species were isolated from 1 to 4 crocodiles only (Table 2). The number of bacteria species per crocodile ranged from 1 to 3 and was not correlated with crocodile total length ($r^2 = 0.163$, $p = 0.033$, $n = 28$). From the cloacal samples, we determined a total of 14 species of bacteria (Table 4). All of these bacteria were Gram-negative and the most frequent bacteria found in cloacal samples of Morelet's crocodiles was *Escherichia coli*, which was detected in 18 (64.3%) crocodiles (Table 4). The other bacteria species were isolated from 1 to 7 crocodiles (Table 4). The number of bacteria species per crocodile ranged from 0 to 3 and was not correlated with crocodile total length ($r^2 = 0.174$, $p = 0.027$, $n = 28$).

Table 1. *Crocodylus acutus*. Species of bacteria found in the oral cavity of wild American crocodiles and the number of swab samples from which each bacterial species was isolated, and the prevalence (% in parentheses) of that bacterial species within each crocodile size class. FA: facultative anaerobic; A: aerobic; AN: anaerobic; ND: not determined

Bacteria species	Gram/ type	Hatchlings (n = 4)	Yearlings (n = 7)	Juveniles (n = 19)	Sub-adults (n = 3)	Adults (n = 5)	Total (n = 38)
<i>Aeromonas hydrophila</i>	-/FA	1 (25.0)	0 (0.0)	7 (36.8)	1 (25.0)	2 (40.0)	11 (28.9)
<i>Arcanobacterium pyogenes</i>	+/AN	4 (100.0)	6 (85.7)	6 (31.6)	1 (25.0)	1 (20.0)	18 (47.4)
<i>Citrobacter freundii</i>	-/FA	0 (0.0)	0 (0.0)	1 (5.3)	0 (0.0)	0 (0.0)	1 (2.6)
<i>Corynebacterium</i> sp.	+/FA	0 (0.0)	0 (0.0)	1 (5.3)	0 (0.0)	0 (0.0)	1 (2.6)
<i>Enterococcus durans</i>	+/A	0 (0.0)	0 (0.0)	1 (5.3)	1 (25.0)	0 (0.0)	2 (5.3)
<i>Enterococcus faecium</i>	+/FA	0 (0.0)	1 (14.3)	2 (10.5)	0 (0.0)	0 (0.0)	3 (7.9)
<i>Escherichia coli</i>	-/FA	0 (0.0)	1 (14.3)	0 (0.0)	0 (0.0)	0 (0.0)	1 (2.6)
<i>Fusobacterium</i> sp.	-/AN	0 (0.0)	1 (14.3)	0 (0.0)	0 (0.0)	0 (0.0)	1 (2.6)
<i>Klebsiella pneumoniae</i>	-/FA	0 (0.0)	0 (0.0)	0 (0.0)	1 (25.0)	0 (0.0)	1 (2.6)
<i>Moraxella cuniculi</i>	-/A	0 (0.0)	1 (14.3)	3 (15.8)	2 (50.0)	0 (0.0)	6 (15.8)
<i>Moraxella catarrhalis</i>	-/A	0 (0.0)	1 (14.3)	1 (5.3)	0 (0.0)	0 (0.0)	2 (5.3)
<i>Pasteurella multocida</i>	-/FA	0 (0.0)	0 (0.0)	1 (5.3)	0 (0.0)	0 (0.0)	1 (2.6)
<i>Pseudomonas aeruginosa</i>	-/A	0 (0.0)	1 (14.3)	2 (10.5)	0 (0.0)	0 (0.0)	3 (7.9)
<i>Rhodococcus</i> sp.	+/A	0 (0.0)	1 (14.3)	1 (5.3)	0 (0.0)	0 (0.0)	2 (5.3)
<i>Serratia marcescens</i>	-/A	2 (50.0)	0 (0.0)	0 (0.0)	0 (0.0)	4 (80.0)	6 (15.8)
<i>Staphylococcus aureus</i>	+/FA	1 (25.0)	0 (0.0)	0 (0.0)	1 (25.0)	4 (80.0)	6 (15.8)
<i>Staphylococcus hyicus</i>	+/FA	0 (0.0)	0 (0.0)	1 (5.3)	0 (0.0)	0 (0.0)	1 (2.6)
<i>Streptococcus agalactiae</i>	+/FA	0 (0.0)	1 (14.3)	4 (21.1)	1 (25.0)	0 (0.0)	6 (15.8)
<i>Streptococcus intermedius</i>	+/AN	0 (0.0)	0 (0.0)	1 (5.3)	0 (0.0)	0 (0.0)	1 (2.6)
<i>Streptococcus pneumoniae</i>	+/FA	0 (0.0)	0 (0.0)	1 (5.3)	0 (0.0)	0 (0.0)	1 (2.6)
<i>Streptococcus pyogenes</i>	+/A	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	1 (20.0)	1 (2.6)
<i>Streptococcus</i> sp.	+/ND	0 (0.0)	1 (14.3)	0 (0.0)	0 (0.0)	0 (0.0)	1 (2.6)
Total species/size class		4	10	15	7	5	22
Mean (\pm SD) species per crocodile		2.0 \pm 0.0	2.1 \pm 1.1	1.7 \pm 1.1	2.0 \pm 1.4	2.4 \pm 0.6	2.0 \pm 1.0

Table 2. *Crocodylus moreletii*. Species of bacteria found in the oral cavity of wild Morelet's crocodiles and the number of swab samples from which each bacterial species was isolated, and the prevalence (% in parentheses) of that bacterial species within each crocodile size class. FA: facultative anaerobic; A: aerobic; AN: anaerobic

Bacteria species	Gram/ type	Hatchlings (n = 4)	Yearlings (n = 8)	Juveniles (n = 5)	Sub-adults (n = 7)	Adults (n = 4)	Total (n = 28)
<i>Aerococcus viridans</i>	+/A	0 (0.0)	0 (0.0)	1 (20.0)	2 (28.6)	1 (25.0)	4 (14.3)
<i>Aeromonas hydrophila</i>	-/FA	0 (0.0)	6 (75.0)	1 (20.0)	4 (57.1)	2 (50.0)	13 (46.4)
<i>Arcanobacterium pyogenes</i>	+/AN	0 (0.0)	1 (12.5)	0 (0.0)	2 (28.6)	0 (0.0)	3 (10.7)
<i>Citrobacter freundii</i>	-/FA	0 (0.0)	1 (12.5)	1 (20.0)	2 (28.6)	0 (0.0)	4 (14.3)
<i>Escherichia coli</i>	-/FA	1 (25.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	1 (3.6)
<i>Fusobacterium necrophorum</i>	-/AN	0 (0.0)	1 (12.5)	0 (0.0)	0 (0.0)	0 (0.0)	1 (3.6)
<i>Fusobacterium</i> sp.	-/AN	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	1 (25.0)	1 (3.6)
<i>Klebsiella pneumoniae</i>	-/FA	2 (50.0)	1 (12.5)	0 (0.0)	2 (28.6)	2 (50.0)	7 (25.0)
<i>Kluyvera</i> sp.	-/FA	0 (0.0)	0 (0.0)	1 (20.0)	0 (0.0)	0 (0.0)	1 (3.6)
<i>Listeria monocytogenes</i>	+/FA	0 (0.0)	1 (12.5)	0 (0.0)	1 (14.3)	0 (0.0)	2 (7.1)
<i>Obesumbacterium proteus</i>	-/FA	0 (0.0)	0 (0.0)	1 (20.0)	1 (14.3)	0 (0.0)	2 (7.1)
<i>Pantoea agglomerans</i>	-/FA	0 (0.0)	0 (0.0)	1 (20.0)	0 (0.0)	0 (0.0)	1 (3.6)
<i>Proteus vulgaris</i>	-/FA	0 (0.0)	1 (12.5)	0 (0.0)	0 (0.0)	0 (0.0)	1 (3.6)
<i>Pseudomonas fluorescens</i>	-/A	1 (25.0)	0 (0.0)	1 (20.0)	0 (0.0)	0 (0.0)	2 (7.1)
<i>Rhodococcus equi</i>	+/A	0 (0.0)	0 (0.0)	0 (0.0)	1 (14.3)	0 (0.0)	1 (3.6)
<i>Salmonella arizonae</i>	-/FA	0 (0.0)	0 (0.0)	0 (0.0)	1 (14.3)	1 (25.0)	2 (7.1)
<i>Serratia rubidaea</i>	-/FA	0 (0.0)	1 (12.5)	0 (0.0)	1 (14.3)	1 (25.0)	3 (10.7)
<i>Shigella</i> sp.	-/FA	0 (0.0)	0 (0.0)	0 (0.0)	1 (14.3)	0 (0.0)	1 (3.6)
<i>Staphylococcus aureus</i>	+/FA	0 (0.0)	0 (0.0)	1 (20.0)	0 (0.0)	0 (0.0)	1 (3.6)
<i>Staphylococcus</i> sp.	+/FA	0 (0.0)	3 (37.5)	0 (0.0)	0 (0.0)	0 (0.0)	3 (10.7)
<i>Streptococcus agalactiae</i>	+/FA	0 (0.0)	0 (0.0)	0 (0.0)	1 (14.3)	1 (25.0)	2 (7.1)
<i>Streptococcus pneumoniae</i>	+/FA	0 (0.0)	2 (25.0)	0 (0.0)	0 (0.0)	0 (0.0)	2 (7.1)
Total species/size class		3	10	8	12	7	22
Mean (\pm SD) species per crocodile		1.0 \pm 0.0	2.3 \pm 0.5	1.6 \pm 0.6	2.7 \pm 0.5	2.3 \pm 0.8	2.1 \pm 0.8

Table 3. *Crocodylus acutus*. Species of bacteria found in the cloacal cavity of wild American crocodiles and the number of swab samples from which each bacterial species was isolated, and the prevalence (% in parentheses) of that bacterial species within each crocodile size class. FA: facultative anaerobic; A: aerobic

Bacteria species	Gram/ type	Hatchlings (n = 4)	Yearlings (n = 8)	Juveniles (n = 18)	Sub-adults (n = 4)	Adults (n = 9)	Total (n = 43)
<i>Aeromonas hydrophila</i>	-/FA	2 (50.0)	3 (37.5)	9 (50.0)	0 (0.0)	2 (22.2)	16 (37.2)
<i>Alcaligenes faecalis</i>	-/A	0 (0.0)	0 (0.0)	2 (11.1)	0 (0.0)	0 (0.0)	2 (4.7)
<i>Citrobacter diversus</i>	-/FA	1 (25.0)	3 (37.5)	2 (11.1)	0 (0.0)	2 (22.2)	8 (18.6)
<i>Citrobacter freundii</i>	-/FA	0 (0.0)	1 (12.5)	3 (16.7)	0 (0.0)	0 (0.0)	4 (9.3)
<i>Escherichia coli</i>	-/FA	1 (25.0)	6 (75.0)	13 (72.2)	3 (75.0)	6 (66.7)	29 (67.4)
<i>Enterobacter cloacae</i>	-/FA	1 (25.0)	0 (0.0)	2 (11.1)	2 (50.0)	1 (11.1)	6 (14.0)
<i>Enterococcus faecalis</i>	+/FA	0 (0.0)	1 (12.5)	5 (27.8)	0 (0.0)	0 (0.0)	6 (14.0)
<i>Edwardsiella tarda</i>	-/FA	2 (50.0)	0 (0.0)	1 (5.6)	0 (0.0)	2 (22.2)	5 (11.6)
<i>Klebsiella pneumoniae</i>	-/FA	0 (0.0)	0 (0.0)	0 (0.0)	1 (25.0)	0 (0.0)	1 (2.3)
<i>Klebsiella sp.</i>	-/FA	0 (0.0)	1 (12.5)	1 (5.6)	0 (0.0)	0 (0.0)	2 (4.7)
<i>Proteus mirabilis</i>	-/FA	0 (0.0)	1 (12.5)	0 (0.0)	0 (0.0)	0 (0.0)	1 (2.3)
<i>Proteus vulgaris</i>	-/FA	1 (25.0)	0 (0.0)	5 (27.8)	0 (0.0)	2 (22.2)	8 (18.6)
<i>Rhodococcus</i> sp.	+/FA	0 (0.0)	2 (25.0)	2 (11.1)	1 (25.0)	0 (0.0)	5 (11.6)
<i>Salmonella arizona</i> e	-/FA	1 (25.0)	3 (37.5)	5 (27.8)	1 (25.0)	4 (44.4)	14 (32.6)
<i>Shigella sonnei</i>	-/FA	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	1 (11.1)	1 (2.3)
Total species/size class		7	9	12	5	8	15
Mean (\pm SD) species per crocodile		2.3 \pm 0.5	2.6 \pm 0.5	2.8 \pm 0.4	2.0 \pm 0.8	2.2 \pm 0.8	2.5 \pm 0.6

Table 4. *Crocodylus moreletii*. Species of bacteria found in the cloacal cavity of wild Morelet's crocodiles and the number of swab samples from which each bacterial species was isolated, and the prevalence (% in parentheses) of that bacterial species within each crocodile size class. FA: facultative anaerobic; AF: aerobic facultative

Bacteria species	Gram/ type	Hatchlings (n = 4)	Yearlings (n = 8)	Juveniles (n = 5)	Sub-adults (n = 7)	Adults (n = 4)	Total (n = 28)
<i>Aeromonas hydrophila</i>	-/FA	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	1 (25.0)	1 (3.6)
<i>Citrobacter diversus</i>	-/FA	0 (0.0)	1 (12.5)	0 (0.0)	1 (14.3)	0 (0.0)	2 (7.1)
<i>Edwardsiella tarda</i>	-/FA	0 (0.0)	1 (12.5)	1 (20.0)	0 (0.0)	0 (0.0)	2 (7.1)
<i>Enterobacter cloacae</i>	-/FA	0 (0.0)	2 (25.0)	2 (40.0)	0 (0.0)	1 (25.0)	5 (17.9)
<i>Enterobacter gergoviae</i>	-/FA	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	1 (25.0)	1 (3.6)
<i>Escherichia coli</i>	-/FA	0 (0.0)	8 (100)	2 (40.0)	6 (85.7)	2 (50.0)	18 (64.3)
<i>Hafnia alvei</i>	-/AF	0 (0.0)	0 (0.0)	1 (20.0)	0 (0.0)	0 (0.0)	1 (3.6)
<i>Klebsiella pneumoniae</i>	-/FA	0 (0.0)	2 (25.0)	0 (0.0)	2 (28.6)	2 (50.0)	6 (21.4)
<i>Pantoea agglomerans</i>	-/FA	0 (0.0)	0 (0.0)	1 (20.0)	0 (0.0)	1 (25.0)	2 (7.1)
<i>Proteus vulgaris</i>	-/FA	0 (0.0)	0 (0.0)	0 (0.0)	1 (14.3)	0 (0.0)	1 (3.6)
<i>Salmonella arizona</i> e	-/FA	0 (0.0)	2 (25.0)	1 (20.0)	3 (42.9)	1 (25.0)	7 (25.0)
<i>Salmonella typhi</i>	-/FA	0 (0.0)	1 (12.5)	0 (0.0)	0 (0.0)	0 (0.0)	1 (3.6)
<i>Serratia rubidaea</i>	-/FA	0 (0.0)	0 (0.0)	1 (20.0)	2 (28.6)	1 (25.0)	4 (14.3)
<i>Yersinia enterocolitica</i>	-/FA	0 (0.0)	1 (12.5)	0 (0.0)	2 (28.6)	0 (0.0)	3 (10.7)
Total species/size class		0	8	7	7	8	14
Mean (\pm SD) species per crocodile		0	2.3 \pm 0.7	1.8 \pm 0.8	2.4 \pm 0.5	2.5 \pm 1.0	2.3 \pm 0.7

Interspecific differences

In total, we isolated 47 species of bacteria from all samples, distributed within 28 genera and 14 families (Table 5). Species of the family *Enterobacteriaceae* represented 51.1% of the species. Fourteen species (29.8%) were detected in both croco-

dile species and 18 (38.3%) and 15 (31.9%) species were only detected in American crocodiles and Morelet's crocodiles, respectively. We found 35 species (23 genera, 13 families) of bacteria in all oral samples, of which 9 species (25.8%) were detected in both crocodile species. Thirteen species (37.1%) were only detected in Morelet's crocodiles

Table 5. *Crocodylus acutus* and *C. moreletii*. All bacteria species found in American and Morelet's crocodiles by capture location indicating the cavity in which each bacterium was isolated (O: oral; C: cloaca; -: not detected)

Family	Species	Río Hondo (<i>C. moreletii</i>)	Cozumel (<i>C. acutus</i>)	Banco Chinchorro (<i>C. acutus</i>)
<i>Actinomycetaceae</i>	<i>Arcanobacterium pyogenes</i>	O	O	O
<i>Aerococcaceae</i>	<i>Aerococcus viridans</i>	O	-	-
<i>Aeromonadaceae</i>	<i>Aeromonas hydrophila</i>	O, C	O, C	O, C
<i>Alcaligenaceae</i>	<i>Alcaligenes faecalis</i>	-	C	-
<i>Corynebacteriaceae</i>	<i>Corynebacterium</i> sp.	-	O	-
<i>Enterobacteriaceae</i>	<i>Citrobacter diversus</i>	C	C	C
	<i>Citrobacter freundii</i>	O	O, C	-
	<i>Edwarsiella tarda</i>	C	C	C
	<i>Enterobacter cloacae</i>	C	C	C
	<i>Enterobacter gergoviae</i>	C	-	-
	<i>Enterococcus durans</i>	-	O	-
	<i>Enterococcus faecalis</i>	-	C	-
	<i>Enterococcus faecium</i>	-	O	-
	<i>Escherichia coli</i>	O, C	O, C	C
	<i>Hafnia alvei</i>	C	-	-
	<i>Klebsiella pneumoniae</i>	O, C	O, C	-
	<i>Klebsiella</i> sp.	-	C	-
	<i>Kluyvera</i> sp.	O	-	-
	<i>Obesumbacterium proteus</i>	O	-	-
	<i>Pantoea agglomerans</i>	O, C	-	-
	<i>Proteus mirabilis</i>	-	C	-
	<i>Proteus vulgaris</i>	O, C	C	C
	<i>Salmonella enterica arizona</i> e	O, C	C	C
	<i>Salmonella enterica enterica</i> ser. <i>Typhi</i>	C	-	-
	<i>Serratia marcescens</i>	-	-	O
	<i>Serratia rubidaea</i>	O, C	-	-
	<i>Shigella sonnei</i>	-	-	C
	<i>Shigella</i> sp.	O	-	-
	<i>Yersinia enterocolitica</i>	C	-	-
<i>Fusobacteriaceae</i>	<i>Fusobacterium necrophorum</i>	O	-	-
	<i>Fusobacterium</i> sp.	O	O	-
<i>Listeriaceae</i>	<i>Listeria monocytogenes</i>	O	-	-
<i>Moraxellaceae</i>	<i>Moraxella catarrhalis</i>	-	O	-
	<i>Moraxella cuniculi</i>	-	O	-
<i>Nocardiaceae</i>	<i>Rhodococcus equi</i>	O	-	-
	<i>Rhodococcus</i> sp.	-	O, C	-
<i>Pasteurellaceae</i>	<i>Pasteurella multocida</i>	-	O	-
<i>Pseudomonadaceae</i>	<i>Pseudomonas aeruginosa</i>	-	O	-
	<i>Pseudomonas fluorescens</i>	O	-	-
<i>Staphylococcaceae</i>	<i>Staphylococcus aureus</i>	O	-	O
	<i>Staphylococcus hyicus</i>	-	O	-
	<i>Staphylococcus</i> sp.	O	-	-
<i>Streptococcaceae</i>	<i>Streptococcus agalactiae</i>	O	O	-
	<i>Streptococcus intermedius</i>	-	O	-
	<i>Streptococcus pneumoniae</i>	O	O	-
	<i>Streptococcus pyogenes</i>	-	-	O
	<i>Streptococcus</i> sp.	-	O	-

and another 13 species (37.1 %) were only detected in American crocodiles. From all cloacal samples, we found a total of 21 species (16 genera, 5 families) of bacteria, of which 8 species (38.1 %) were detected in both crocodile species, and 6 (28.6 %) and 7 (33.3 %) species were only detected in Morelet's crocodiles and American crocodiles, respectively.

Crocodile attacks

We found 12 publications that referred to attacks by crocodiles on humans in the Yucatan Peninsula. We also knew of 2 cases of attacks by American crocodiles on fishers in Banco Chinchorro during our field work (P. Charruau pers. obs.). From these publications and observations, we determined that a mini-

mum of 24 attacks occurred in the Yucatan Peninsula (21 in Quintana Roo, 2 in Campeche and 1 in Yucatan) from 1992 to 2011. Two were fatal (1 in Campeche and 1 in Quintana Roo) and evidence indicated that *Crocodylus moreletii* was responsible. Both species (*C. acutus* and *C. moreletii*) were involved in the attacks. Most of the attacks occurred when victims were in the water fishing, spear-fishing or swimming, and in 4 cases victims were under the influence of alcohol. Among the victims only 1 woman (4.2%) was involved. The number of attacks in Quintana Roo shows an increase of attacks during the last 6 yr, after almost a decade without incidents (Fig. 2). Information about injuries of victims is available in most of the press articles; injuries consisted of bites on various parts of the body (shoulders, hands, arms, foot, thorax, pelvis, back, legs, neck and head), lung perforation, partial loss of a foot and death (in 2 cases). However, the press articles do not present information about infection and bacteria involved.

DISCUSSION

Few studies on bacterial flora in cloacal and oral cavities of wild crocodilians have been done and, as far as we know, only 5 crocodilian species have been examined in wild conditions: *Alligator mississippiensis* (Flandry et al. 1989, Johnston et al. 2010), *Crocodylus acutus* (Cupul-Magaña et al. 2005), *C. niloticus* (Lovely & Leslie 2008), *C. porosus* (Anderson 1999) and *C. johnstoni* (Anderson 1999). To our knowledge, the present study is the first record of bacterial flora in oral and cloacal cavities of wild *C. moreletii*. The present study is the second to investigate oral flora and the first on cloacal flora for *C. acutus* in the wild. Before this study, Cupul-Magaña et al. (2005) examined the oral flora of wild *C. acutus*, and found 10

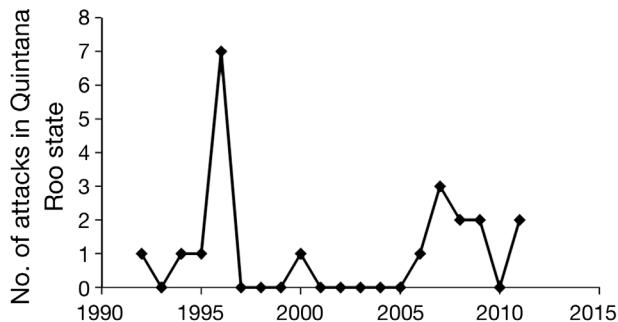


Fig. 2. *Crocodylus acutus* and *C. moreletii*. Number of attacks by American and Morelet's crocodiles on humans in Quintana Roo state, Mexico, from 1992 to 2011

bacteria species of which 5 were also found in this study either in the cloacal or oral cavity (*Aeromonas hydrophila*, *Citrobacter freundii*, *C. diversus*, *Escherichia coli* and *Klebsiella pneumoniae*).

The genera *Aeromonas*, *Citrobacter*, *Corynebacterium*, *Enterococcus*, *Escherichia*, *Fusobacterium*, *Klebsiella*, *Moraxella*, *Pasteurella*, *Proteus*, *Pseudomonas*, *Salmonella*, *Serratia*, *Shigella*, *Staphylococcus* and *Streptococcus* have been already identified in the oral cavity of crocodilians (Doering et al. 1971, Gorden et al. 1979, Flandry et al. 1989, Anderson 1999, Cupul-Magaña et al. 2005, Silva et al. 2009). To our knowledge, the present study documents for the first time the occurrence of the genera *Aerococcus*, *Arcanobacterium*, *Kluyvera*, *Listeria*, *Obesumbacterium*, *Pantoea* and *Rhodococcus* in the oral cavity of wild crocodilians. It is also the first record of occurrence of the following species in the oral cavity of crocodilians: *Aerococcus viridians*, *Arcanobacterium pyogenes*, *Enterococcus durans*, *E. faecium*, *Fusobacterium necrophorum*, *Listeria monocytogenes*, *Moraxella cuniculi*, *M. catarrhalis*, *Obesumbacterium proteus*, *Pantoea agglomerans*, *Pasteurella multocida*, *Rhodococcus equi*, *Salmonella enterica arizona*, *Serratia rubidaea*, *Staphylococcus aureus*, *S. hyicus*, *Streptococcus agalactiae*, *S. intermedius* and *S. pneumoniae*.

The genera *Aeromonas*, *Citrobacter*, *Edwardsiella*, *Enterobacter*, *Enterococcus*, *Escherichia*, *Hafnia*, *Klebsiella*, *Pantoea*, *Proteus*, *Salmonella*, *Serratia* and *Yersinia* have been already identified in cloaca of crocodilians (White et al. 1973, Madsen et al. 1998, Anderson 1999, Lovely & Leslie 2008, Johnston et al. 2010). To our knowledge, this is the first report of the genera *Alcaligenes* and *Rhodococcus* in the cloaca. The following species appear to be new reports for the cloacal flora of crocodilians: *Alcaligenes faecalis*, *Citrobacter diversus*, *Enterobacter gergoviae*, *Pantoea agglomerans*, *Salmonella enterica enterica* serotype Typhi, *Serratia rubidaea* and *Yersinia enterocolitica*.

Crocodilians are very resistant to disease with a great capacity for healing, principally owing to their high level of serum antimicrobial activity (Merchant & Britton 2006, Merchant et al. 2006). Huchzermeyer (2003) report that few bacteria cause specific diseases in crocodiles and even fewer are crocodile-specific, but many bacteria can cause non-specific septicemia. As all crocodiles captured during the present study were apparently healthy, we assume that all bacteria species found in the present study are part of the normal flora of *Crocodylus acutus* and *C. moreletii*. However, several of the bacteria iso-

lated could be pathogens of crocodilians when the host is subject to stress, particularly in captive conditions (Huchzermeyer 2003). Bacteria species carried by *C. acutus* and/or *C. moreletii* in the Yucatan Peninsula and isolated from cases of crocodilian septicemias are: *Aeromonas hydrophila*, *Arcanobacterium pyogenes*, *Citrobacter freundii*, *Corynebacterium* sp., *Edwardsiella tarda*, *Escherichia coli*, *Klebsiella* sp., *Pantoea agglomerans*, *Pasteurella multocida*, *Proteus* sp., *Pseudomonas aeruginosa*, *Serratia marcescens*, *Staphylococcus aureus*, *Staphylococcus* sp. and *Streptococcus* sp. (Huchzermeyer 2003). Moreover, *Streptococcus agalactiae* has been reported to cause necrotizing fasciitis in captive juvenile *Crocodylus porosus* (Bishop et al. 2007).

A large number of *Salmonella* serovars have been isolated from crocodiles (Huchzermeyer 2003) and are part of the normal intestinal tract flora of crocodilians. However, *Salmonella* can cause enteritis and septicemia in individuals subjected to severe stress conditions (Huchzermeyer 2003). In the present study, we identified 2 subspecies of *Salmonella enterica* (*S. enterica* subsp. *arizona* in the oral and cloacal cavities of *Crocodylus moreletii* and in the cloacae of *C. acutus*, and *S. enterica* subsp. *enterica* serotype Typhi from the cloacal cavity of *C. moreletii*), which in certain stressful conditions could cause enteritis and septicemia in *C. moreletii* and *C. acutus*.

Cloacal flora can also be responsible for infections of the egg shell, egg shell membrane and yolk during egg laying (Huchzermeyer 2003). Peucker et al. (2005) studied bacteria present in shell and yolk of farmed Australian freshwater crocodile *Crocodylus johnstoni* eggs and among the bacteria they found, 13 have been identified in *C. acutus* and/or *C. moreletii* in the present study: *Aeromonas hydrophila*, *Alcaligenes faecalis*, *Citrobacter freundii*, *Edwardsiella tarda*, *Enterobacter cloacae*, *Escherichia coli*, *Klebsiella* sp., *Pantoea agglomerans*, *Proteus vulgaris*, *Pseudomonas aeruginosa*, *P. fluorescens*, *Salmonella arizona* and *Serratia marcescens*. Schumacher & Cardeilhac (1990) also found *Enterobacter cloacae*, *Citrobacter* sp., *Proteus* sp. and *Pseudomonas aeruginosa* in lesions of infected eggs, and Thomas et al. (2001) isolated several *Salmonella* serotypes from the egg shell or yolk of *Crocodylus porosus*. Bacterial egg infection is a cause of egg mortality and can also lead to yolk-sac infections and omphalitis in hatchlings, which can result in their death (Huchzermeyer 2003). This information is important if we consider that *C. acutus* in Banco Chinchorro show a low nest number and a reduced

clutch size (Charruau et al. 2010). In this condition, bacterial infection of eggs could be a threat to this crocodile's conservation, and an assessment of bacteria present on *C. acutus* egg shell in Banco Chinchorro should therefore be made.

Bacteria species carried by crocodiles can also be transmitted to humans. The transmission of bacteria from crocodiles to humans can occur in 3 major ways: by consuming crocodile meat (Suárez et al. 2000, Magnino et al. 2009), from bites during crocodile attacks (Caldicott et al. 2005, Gruen 2009, Wamisho et al. 2009) or during activities in 'crocodile waters' (Johnston et al. 2010). Wounds from crocodile bites generally become infected by bacteria present in the oral cavity of crocodiles (Caldicott et al. 2005, Wamisho et al. 2009). Bacteria involved in wound infections after crocodilian attacks include *Aeromonas hydrophila*, *Burkholderia pseudomallei*, *Pseudomonas* spp., *Serratia* spp., *Citrobacter diversus*, *Enterococcus* spp., *Clostridium* spp. and *Pantoea agglomerans* (Raynor et al. 1983, Flandry et al. 1989, Mekisic & Wardill 1992). Most of these species and genera have been identified in the present study in oral and/or cloacal cavity of American or Morelet's crocodiles. Moreover, among bacteria identified in oral cavity of crocodiles in this study, *Aeromonas hydrophila*, *Corynebacterium* sp., *Citrobacter freundii*, *Enterococcus durans*, *E. faecium*, *Escherichia coli*, *Klebsiella pneumonia*, *Pantoea agglomerans*, *Proteus vulgaris*, *Salmonella enterica arizona*, *Serratia marcescens*, *S. rubidaea*, *Fusobacterium necrophorum*, *Fusibacterium* sp., *Moraxella catarrhalis*, *M. cuniculi*, *Rhodococcus* sp., *Pasteurella multocida*, *Pseudomonas aeruginosa*, *P. fluorescens*, *Staphylococcus aureus*, *S. hyicus*, *Staphylococcus* sp., *Streptococcus agalactiae*, *S. intermedius*, *S. pyogenes* and *Streptococcus* sp. have been reported in infected wounds caused by animal or human bites (Murphy et al. 1992, Blaylock 1999, Goldstein et al. 2002, Kunimoto et al. 2004, Jofré Morales et al. 2006, Brook 2009).

In the present study we found 24 reported cases of attacks by crocodiles on humans in the Yucatan Peninsula, most of them (87.5%) occurring in Quintana Roo and in the city of Cancun, which are the state and city in the peninsula, respectively, most frequented by tourists. Both crocodile species (*Crocodylus acutus* and *C. moreletii*) were involved in the attacks and *C. moreletii* was involved in 2 cases of fatal attacks. Attacks resulted in a variety of wounds on the victim from bites to different parts of the body to lung perforation, limb amputation and death. Bacteria found in oral cavities of crocodiles could cause

infections and septicemia of wounds reported in the attacks. However, no publications reporting the attacks present information about wound infection and bacteria involved. Furthermore, the number of attacks in Quintana Roo seems to have increased since 2005 (Fig. 2). In the country of Belize, adjacent to Quintana Roo, several studies have also reported an increase in the number of attacks on humans by crocodiles (Garel et al. 2005). These attacks involved *C. moreletii* and no case of attack by *C. acutus* was reported (Marlin et al. 1995, Finger et al. 2002, Garel et al. 2005). The principal factors of this increase in crocodile–human interactions in Quintana Roo and Belize are the increase in human and crocodile populations, the destruction and fragmentation of crocodile habitat, the deliberate and unintentional feeding of crocodiles in human-populated areas, the increased efforts in recording crocodile attacks, drought causing loss of water in lagoons that increase the density of crocodiles and the economic situation in Mexico that increases illegal and subsistence fishing, thereby increasing the probability of crocodile encounters (Lazcano-Barrero 1996, Finger et al. 2002, Garel et al. 2005).

Farmed crocodilian meat harvested for human consumption often shows bacterial contamination, which principally occurs during slaughter and dressing procedures (Revol 1995, Rickard et al. 1995, Thomas et al. 2001, Magnino et al. 2009). Microbiological analyses of meat of several farmed crocodilian species have shown the presence of the bacteria genera *Salmonella*, *Staphylococcus*, *Flavobacterium*, *Pseudomonas*, *Acinobacter*, *Enterobacter*, *Moraxella*, *Micrococcus*, *Streptococcus* and *Escherichia* (Madsen 1993, Rickard et al. 1995, Hoffmann & Romanelli 1998, Thomas et al. 2001). Furthermore, infection by *Bacillus cereus* after black caiman *Melanosuchus niger* meat consumption has been reported in the Amazonas (Suárez et al. 2000). A case of salmonellosis in South Africa has also been reported in a man who consumed crocodile meat infected by *Salmonella enterica diarizonae* (Narayana et al. 2008). Cases of salmonellosis in humans often occurred after direct or indirect contact with reptiles (Mermin et al. 2004). Although crocodiles are protected and their hunt has been banned since 1970 in Mexico, opportunistic killings occur occasionally in the vicinity of human settlements (Cedeño-Vázquez et al. 2006) and meat consumption of poached individuals could cause infection. Furthermore, during crocodile population surveys in the Yucatan Peninsula, we occasionally observed children who kept hatchling crocodiles as pets, and they could contract salmonel-

losis by contact with those crocodiles. We especially warn about the presence of *S. enterica enterica* ser. Typhi in the cloaca of Morelet's crocodile, which causes typhoid fever (Colomba et al. 2008). *Listeria monocytogenes*, *Yersinia enterocolitica*, *Shigella* spp. and *Escherichia coli* found in the present study are also important food-borne pathogens that can cause enteric infections (Levine & Vial 1988, Bottone 1997, Kotloff et al. 1999, Schlech 2000).

Furthermore, several of the bacteria found in the cloacal cavity of crocodiles in the present study are potential pathogens and could affect water quality and human health. Johnston et al. (2010) studied bacteria from the cloaca of the American alligator *Alligator mississippiensis* as well as bacteria from surface water samples from their habitat, and they found similar flora present in the cloaca of alligator and the water they inhabit. Johnston et al. (2010) concluded that alligators are a potential source of bacterial contamination of their aquatic habitat through the excretion of feces. Other studies have reported the presence of similar bacterial flora in crocodilians and the water they inhabit (Shotts et al. 1972, White et al. 1973, Flandry et al. 1989, Madsen 1994). In captivity, tank water was also contaminated with bacteria from crocodilian feces (Kennedy 1973, Flandry et al. 1989, Madsen 1994, Huchzermeyer 2003). Thus, bacteria found in the present study in American and Morelet's crocodiles can also be expected to be present in their respective environments, and several of these bacteria are opportunistic pathogens that may cause wound or enteric infections in humans during activities in water or mud (Pitlik et al. 1987, Keene et al. 1994, Vally et al. 2004, Noonburg 2005).

Therefore, the bacteria found in the oral and cloacal cavities in Morelet's and American crocodiles in this study are potential pathogenic agents for crocodiles, humans (and other victims of crocodile attacks such as domestic animals) and their environments. Presently, humans rely on a large array of antibiotics to fight bacterial infections, and several studies on crocodile attacks have presented a broad spectrum of prophylactic antibiotics that could be used for the treatment of crocodile bite wounds (Caldicott et al. 2005, Hertner 2006, Gruen 2009, Wamisho et al. 2009, Langley 2010). However, even though many bacteria are sensitive to antibiotics, some strains/species are unaffected by, or show resistance to, some antibiotics (Montgomery et al. 2002, Bergstrom & Feldgarden 2008). Moreover, bacteria are rapidly evolving organisms that can become resistant to antibiotics within several years after widespread use

of the antibiotic (Bergstrom & Feldgarden 2008). Thus, future studies should involve experiments on sensitivity and resistance of bacteria found in crocodiles in Quintana Roo in order to improve the treatment of wounds in victims of crocodile attacks. This would also provide important information to use in the treatment of wound infections in crocodiles and domestic animals that develop from crocodile bites.

CONCLUSION

This study provides important information about bacterial flora carried by American and Morelet's crocodiles in Quintana Roo State, Mexico. These bacteria can be a potential threat for crocodile health during conditions of stress but also represent a threat to human health through crocodile attacks and bites, consumption of crocodile meat or by simply carrying out activities in crocodile habitat. The risk of bacterial contamination from crocodiles to humans could increase in the future owing to several factors, principally the accelerated destruction and fragmentation of habitat, which could lead to an augmentation of human–crocodile interactions. Information on bacteria diversity reported in this study could help in the choice of antibacterial products for cases of infection from bacteria of crocodile origin.

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