

# Detection of Pathogenic *Leptospira* Bacteria in Pinniped Populations via PCR and Identification of a Source of Transmission for Zoonotic Leptospirosis in the Marine Environment<sup>∇</sup>

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**Leptospirosis, caused by the spirochete *Leptospira*, is a geographically widespread disease that affects a broad range of mammals, including marine mammals. Among pinniped populations, periodic epizootics of leptospirosis are responsible for significant die-offs. Along the west coast of North America, the most recent leptospirosis epizootic occurred in 2004, during which samples were collected from cases ranging from California to British Columbia. The primary objective of this study was to use this well-defined sample set to determine the feasibility of using PCR techniques to diagnose *Leptospira* infection among pinniped populations in comparison with diagnostic methodologies commonly used for marine mammals. Successful amplification was achieved from a variety of samples, including freshly collected urine, urine stored at  $-80^{\circ}\text{C}$  for less than 6 months, and kidney (freshly collected, frozen, and decomposed), as well as feces- and urine-contaminated sand collected in the vicinity of a live-stranded animal. Pathological examination of tissue collected from *Leptospira*-infected animals revealed the presence of leptospiral antigen in the kidneys. The use of species-specific primer pairs revealed a pattern of host specificity for *Leptospira interrogans* in sea lions and *Leptospira kirschneri* in elephant seals. These studies indicate PCR is a sensitive and specific diagnostic tool for the detection of *Leptospira* infection in pinnipeds and reveal a potential source for epizootic, enzootic, and zoonotic spread of leptospirosis in a marine environment.**

Leptospirosis is a ubiquitous disease with a global distribution that affects humans and a wide variety of domestic and wild animal species, including marine mammals such as California sea lions (*Zalophus californianus*) (7), northern fur seals (*Callorhinus ursinus*) (10), northern elephant seals (*Mirounga angustirostris*) (2), and harbor seals (*Phoca vitulina*) (11). Leptospirosis typically presents in these animals as interstitial nephritis with clinical signs of impaired renal function, including dehydration, polydipsia, vomiting, and depression (7). Disease outbreaks have occurred repeatedly in California sea lions off central and northern California, with hundreds of animals stranding and subsequently dying in each outbreak (1, 7). These large-scale epizootics are cyclic and have been recognized since the early 1970s, with a distinct 3- to 4-year periodicity that is separated by enzootic maintenance of the disease (3, 4, 7, 14). During the most recent outbreak in 2004, over 300 sea lions died along the central California coast and further mortalities were tracked off the coasts of Oregon, Washington, and British Columbia.

Current methods for diagnosing leptospirosis among live

marine mammals rely upon a combination of the microscopic agglutination test (MAT) performed on sera, clinical observations, and detection of serum biochemistry changes typical of renal failure (1). If animals are dead, additional diagnostic tools include histopathology and immunohistochemistry, both of which may indicate the presence of *Leptospira* and, in conjunction with clinical chemistry, clinical signs, or necropsy findings, determine the extent of disease. Culture of organisms is possible from urine and harvested kidney of live and dead animals; however, as these organisms are fastidious and require selective media, bacterial isolation is not used as a routine diagnostic tool. Furthermore, without paired serum samples, MAT cannot distinguish an acute active infection from a previous recent infection. The use of molecular analytical techniques such as PCR has been established in terrestrial animals (5, 15, 16) and is ideally suited for the detection of *Leptospira* infection, in that PCR technology is sensitive, specific, widely available, and can be reliably performed on a range of templates, including urine and renal tissue. The studies reported herein evaluate the use of PCR technology for assessing *Leptospira* infection among marine mammal populations compared to conventional diagnostic methodologies.

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## MATERIALS AND METHODS

**Animals and samples.** Live animals that stranded along the California coast were admitted to The Marine Mammal Center and examined clinically and, if

they died, at postmortem as described by Greig et al. (6). Fresh urine samples were collected from stranded, captive, and wild animals by established methods (free flow capture, cystocentesis, or catheterization), separated into 1- to 3-ml aliquots, and stored at 4°C prior to analysis. Serum samples were collected from the caudal gluteal vein and used for serum biochemistry and MAT testing as described by Colagrass-Schouten et al. (1). The cutoff for a positive agglutination reaction was defined as a titer of  $\geq 1:400$  in a single sample. If animals died or were euthanized due to poor prognosis, gross necropsies were performed within 12 h of death. Gross necropsy findings consistent with leptospirosis included swollen kidneys, loss of renicular differentiation, pale tan cortices, or subcapsular hemorrhages and serum biochemistry results indicative of renal failure (blood urea nitrogen of  $>100$  mg/dl, creatinine of  $>2$  mg/dl, sodium of  $>155$  meq/liter, and phosphorus  $>$  calcium) (1, 7). At necropsy, kidney tissue and urine samples were collected aseptically and frozen immediately at  $-80^{\circ}\text{C}$ ; samples 53 to 61 (Table 1) remained at  $-80^{\circ}\text{C}$  for greater than 6 months. Tissue samples from multiple organs were fixed in 10% neutral buffered formalin, processed routinely for paraffin embedding, sectioned at 5  $\mu\text{m}$ , and stained with hematoxylin and eosin. Representative slides were also prepared with Warthin-Starry and Gram stain (9). Immunohistochemical staining was performed on kidney sections using an established streptavidin-biotin staining protocol (12). Sections were incubated at room temperature with an anti-*Leptospira* polyclonal antibody (National Veterinary Services Laboratory, Ames, IA) at a 1:40,000 dilution. The antibody was directed against *L. interrogans* serovars Bratislava, Canicola, Hardjo, Icterohemorrhagiae, and Pomona and *L. kirschneri* serovar Grippotyphosa. Appropriate positive and negative controls were included in all cases.

Animals that live-stranded along the Washington coast were observed for clinical signs of leptospirosis (dehydration, polydipsia, emaciation, and depression), and, if possible, agonal or postmortem urine and serum samples were collected. In one live-stranded sea lion that returned to the ocean prior to sample collection, feces- and urine-contaminated sand in the vicinity of the stranded animal was collected and analyzed. For animals along the British Columbia and Washington coasts that were found dead or were euthanized due to poor prognosis, tissue, serum, and urine samples were collected. In three instances, dead animals found along the Washington coast had moderate levels of decomposition. Histopathology and serology analyses and interpretations were performed as described above.

**DNA isolation.** Total DNA was prepared from samples collected from the captive, wild, live-stranded, dead, and euthanized animals. Isolation of total DNA from renal tissue and feces- and urine-contaminated sand was accomplished using the Qiagen DNeasy kit (Qiagen, Valencia, CA). Isolation of total DNA from urine was performed by the method described by Zuerner et al. (15). Briefly, 1 ml of urine was concentrated by centrifugation ( $16,100 \times g$ , 20 min). The resulting pellet was resuspended in an equal volume of 1 mM EDTA, concentrated, washed with an equal volume of distilled water, and subjected to a final concentration step prior to resuspension in 50  $\mu\text{l}$  of distilled water.

**PCR analyses. (i) Species-specific *Leptospira* PCR amplification.** To specifically amplify the species *Leptospira interrogans*, primers unique to the IS1500 insertion sequence (16) were used (P1, 5'-TTCGATTCAAAGCATGGCTAAC G-3'; M16, 5'-AAAGAAGGACTCAGCGACTGCG-3') with a two-step amplification protocol: 7 cycles of 94°C for 30 s and 72°C for 2 min and 35 cycles of 94°C for 30 s and 67°C for 2 min. To specifically amplify the species *Leptospira kirschneri*, the flagellum-specific primers B64-I (5'-ACTAAGTGAAGAACTTC TAC-3') and B64-II (5'-TCCTTAAGTCGAACCTATGA-3') were used (5). The conditions for this amplification protocol were as follows: 1 cycle of 94°C for 2 min and 72°C for 3 min and 35 cycles of 94°C for 1.5 min, 55°C for 1 min, and 72°C for 2 min. Control amplification templates included water as a negative control and *L. interrogans* and *L. kirschneri* genomic DNA as positive controls. Amplified products were separated on 1% agarose gels, stained with ethidium bromide, and viewed using a UV light source.

## RESULTS

Table 1 summarizes the data collection and sample analyses performed on the animals in our study. Cases 1 to 35 represent test samples. Cases 36 to 52 include samples collected from healthy captive animals (cases 36 and 37) and healthy wild animals (cases 38 to 52) to determine the rate of false-negative PCR amplifications, and test cases 53 to 61 comprise urine samples stored at  $-80^{\circ}\text{C}$  for greater than 6 months to test for sample stability during long-term storage.

**Clinical observations, serology, pathology, and immunohistochemistry.** Clinical observations recorded for cases 1 to 5, 7, 9, 10, 12 to 30, 35, and 53 to 61 were consistent with leptospirosis, and serology performed on cases 2 to 4, 6 to 14, 16 to 22, 24 to 27, 29, and 35 were positive, with the exception of cases 6, 8, 11, and 35. The cause of death in sea lions 6, 8, 11, and 32 was determined to be secondary to trauma (cases 8, 11, and 32) and encephalomalacia (case 6). The cause of death in three animals was unknown (cases 33 to 35), with the latter case demonstrating clinical observations and pathological findings consistent with renal failure. Gross necropsy findings in cases 1 to 5, 7, 9, 10, 12 to 17, 27, 54, 55, and 59 to 61 included markedly swollen kidneys with pale tan cortices. There was loss of renicular and corticomedullary differentiation with occasional infarcts, consistent with nephritis. Histopathologic examination of samples collected from cases 1 to 5, 7, 9, 10, 12 to 17, 22, 24 to 27, 29, 31, and 35 suggested that leptospirosis was the cause of death as lesions were similar to those reported in past leptospirosis outbreaks in pinnipeds (1, 2, 11) (lymphoplasmacytic tubulointerstitial nephritis of various severities, with tubular degeneration, necrosis, and regeneration) (Fig. 1A). In sea lions 6, 8, and 11, renal lesions were limited to a few scattered aggregates of lymphocytes in the cortical interstitium and renal pelvis. Immunohistochemistry results for cases 1 to 17 and 22 were positive, with the exception of cases 6, 8, and 11, where no antigen staining was observed. Positive samples showed antigen within the lumen renal tubules, within the cytoplasm of renal tubular epithelia, and interspersed within associated peritubular inflammatory cells and directly correlated with sea lions that exhibited kidney tubulointerstitial nephritis. In some cases, whole spirochetes could be visualized within the lumina of renal tubules or the microvasculature (Fig. 1B).

**Molecular analyses.** For the molecular analyses, *Leptospira*-specific PCR analyses were performed on DNA isolated from either urine (collected from live-stranded, dead, captive, and wild animals) and renal tissue samples (collected from dead animals) or, in one case, urine- and feces-contaminated sand samples (collected from a transiently stranded pinniped). Leptospiral DNA was successfully amplified from all sample sources, and the majority of samples that demonstrated positive amplification were obtained from animals dying from leptospirosis (cases 1 to 5, 7, 9, 10, and 12 to 31). Amplification was also observed in samples collected from three animals that were negative by clinical observations, immunohistochemistry, and serology but displayed mild interstitial nephritis upon histopathologic analysis (cases 6, 8, and 11). However, since immunohistochemistry and serological analyses detected neither leptospiral antigen in the renal tubules nor an antibody response in these potential carrier animals, respectively, an alternative, yet unlikely, possibility is that the positive PCR amplicons could have resulted from cross-contamination of the necropsy facility. Three animal carcasses in a moderate state of decomposition (cases 32 to 34), including two animals which died from unknown causes (cases 33 and 34), were positive by PCR. An additional sample (case 35) from an animal exhibiting clinical symptoms and histopathology consistent with renal failure was negative by both PCR analysis and serology; this animal likely died from a condition unrelated to leptospirosis. It should be noted that samples from all healthy animals in-

TABLE 1. Comparison of diagnostic methodologies for identification of *Leptospira* infection

Case <sup>a</sup>	Accession no.	Species	Sample origin	PCR result/template	Clinical observation(s) consistent with leptospirosis	Serology	Histopathology	Immunohistochemistry	Status/primary cause of death or working diagnosis
Test samples									
1	ES2339	Elephant seal	CA	+ ( <i>L. kirschneri</i> )/urine	Yes	NA <sup>b</sup>	Tubulointerstitial nephritis	Positive	Dead/leptospirosis
2	CSL6119	CA sea lion	CA	+ ( <i>L. interrogans</i> )/urine	Yes	Positive	Tubulointerstitial nephritis	Positive	Dead/leptospirosis
3	CSL6146	CA sea lion	CA	+ ( <i>L. interrogans</i> )/urine	Yes	Positive	Tubulointerstitial nephritis	Positive	Dead/leptospirosis
4	CSL6144	CA sea lion	CA	+ ( <i>L. interrogans</i> )/urine/kidney	Yes	Positive	Tubulointerstitial nephritis	Positive	Dead/leptospirosis
5	CSL6139	CA sea lion	CA	+ ( <i>L. interrogans</i> )/urine	Yes	NA	Tubulointerstitial nephritis	Positive	Dead/leptospirosis
6	CSL6079	CA sea lion	CA	+ ( <i>L. interrogans</i> )/urine	No	Negative	Minimal nephritis	Negative	Dead/encephalomalacia
7	CSL6257	CA sea lion	CA	+ ( <i>L. interrogans</i> )/urine	Yes	Positive	Tubulointerstitial nephritis	Positive (marked)	Dead/leptospirosis
8	CSL6138	CA sea lion	CA	+ ( <i>L. interrogans</i> )/urine	No	Negative	Minimal pyelitis	Negative	Dead/trauma
9	CSL6175	CA sea lion	CA	+ ( <i>L. interrogans</i> )/urine	Yes	Positive	Tubulointerstitial nephritis	Positive	Dead/leptospirosis
10	CSL6214	CA sea lion	CA	+ ( <i>L. interrogans</i> )/urine	Yes	Positive	Tubulointerstitial nephritis	Positive	Dead/leptospirosis
11	CSL6263	CA sea lion	CA	+ ( <i>L. interrogans</i> )/urine	No	Negative	Minimal nephritis	Negative	Dead/trauma
12	CSL6218	CA sea lion	CA	+ ( <i>L. interrogans</i> )/urine	Yes	Positive	Tubulointerstitial nephritis	Positive	Dead/leptospirosis
13	CSL6250	CA sea lion	CA	+ ( <i>L. interrogans</i> )/urine	Yes	Positive	Tubulointerstitial nephritis	Positive	Dead/leptospirosis
14	CSL6210	CA sea lion	CA	+ ( <i>L. interrogans</i> )/urine	Yes	Positive	Tubulointerstitial nephritis	Positive	Dead/leptospirosis
15	CSL6187	CA sea lion	CA	+ ( <i>L. interrogans</i> )/urine	Yes	NA	Tubulointerstitial nephritis	Positive	Dead/leptospirosis
16	CSL6176	CA sea lion	CA	+ ( <i>L. interrogans</i> )/urine	Yes	Positive	Tubulointerstitial nephritis	Positive	Dead/leptospirosis
17	CSL6205	CA sea lion	CA	+ ( <i>L. interrogans</i> )/urine	Yes	Positive	Tubulointerstitial nephritis	Positive	Dead/leptospirosis
18	CSL6308	CA sea lion	CA	+ ( <i>L. interrogans</i> )/urine	Yes	Positive	NA	NA	Dead/leptospirosis
19	CSL6338	CA sea lion	CA	+ ( <i>L. interrogans</i> )/urine	Yes	Positive	NA	NA	Released/leptospirosis
20	CSL6344	CA sea lion	CA	+ ( <i>L. interrogans</i> )/urine	Yes	Positive	NA	NA	Released/leptospirosis
21	CSL6378	CA sea lion	CA	+ ( <i>L. interrogans</i> )/urine	Yes	Positive	NA	NA	Released/leptospirosis
22	NA	CA sea lion	British Columbia	+ ( <i>L. interrogans</i> )/urine	Yes	Positive	Tubulointerstitial nephritis	Positive	Dead/leptospirosis
23	ZL-NPP-04-01	CA sea lion	WA	+ ( <i>L. interrogans</i> )/urine	Yes	NA	NA	NA	Dead/leptospirosis
24	WDFW 0904-04	CA sea lion	WA	+ ( <i>L. interrogans</i> )/urine	Yes	Positive	Interstitial nephritis	NA	Dead/leptospirosis
25	WDFW 0804-10	Steller/CA sea lion cross	WA	+ ( <i>L. interrogans</i> )/urine	Yes	Positive	Interstitial nephritis	NA	Dead/leptospirosis
26	WDFW 0904-09	CA sea lion	WA	+ ( <i>L. interrogans</i> )/urine	Yes	Positive	Interstitial nephritis	NA	Dead/leptospirosis
27	WDFW 1004-01	CA sea lion	WA	+ ( <i>L. interrogans</i> )/urine	Yes	Positive	Interstitial nephritis	NA	Dead/leptospirosis

28	WDFW 1004-02	Steller sea lion	WA	+ ( <i>L. interrogans</i> )/ feces/urine/sand	Yes	NA	NA	NA	Unknown/leptospirosis
29	WDFW 1004-14	CA sea lion	WA	+ ( <i>L. interrogans</i> )/ urine	Yes	Positive	Interstitial nephritis	NA	Dead/leptospirosis
30	WDFW 1004-10	CA sea lion	WA	+ ( <i>L. interrogans</i> )/ urine	Yes	NA	NA	NA	Dead/leptospirosis
31	WDFW 1004-15	CA sea lion	WA	+ ( <i>L. interrogans</i> )/ kidney	NA	NA	Interstitial nephritis	NA	Dead/leptospirosis
32	CWB-04-ZC-02	CA sea lion	WA	+ ( <i>L. interrogans</i> )/ kidney	NA	NA	NA	NA	Dead/trauma (moderate decomposition of carcass)
33	WDFW 1004-16	CA sea lion	WA	+ ( <i>L. interrogans</i> )/ kidney	NA	NA	NA	NA	Dead/unknown decomposition of carcass
34	WDFW 1004-11	CA sea lion	WA	+ ( <i>L. interrogans</i> )/ kidney	NA	NA	NA	NA	Dead/unknown decomposition of carcass
35	WDFW 1004-06	CA sea lion	WA	-/urine	Yes	Negative	Tubulointerstitial nephritis	NA	Dead/unknown decomposition of carcass
Samples from healthy animals									
36	CSL LML male	CA sea lion	CA	-/urine	No	NA	NA	NA	Alive, captive born
37	CSL LML female	CA sea lion	CA	-/urine	No	NA	NA	NA	Alive, captive born
38	WCSL ZC939	CA sea lion	CA	-/urine	No	NA	NA	NA	Alive, wild sea lion
39	WCSL ZC1043	CA sea lion	CA	-/urine	No	NA	NA	NA	Alive, wild sea lion
40	WCSL ZC1072	CA sea lion	CA	-/urine	No	NA	NA	NA	Alive, wild sea lion
41	WCSL ZC1074	CA sea lion	CA	-/urine	No	NA	NA	NA	Alive, wild sea lion
42	WCSL ZC1077	CA sea lion	CA	-/urine	No	NA	NA	NA	Alive, wild sea lion
43	WCSL ZC1079	CA sea lion	CA	-/urine	No	NA	NA	NA	Alive, wild sea lion
44	WCSL ZC845	CA sea lion	CA	-/urine	No	NA	NA	NA	Alive, wild sea lion
45	WCSL ZC870	CA sea lion	CA	-/urine	No	NA	NA	NA	Alive, wild sea lion
46	WCSL ZC952	CA sea lion	CA	-/urine	No	NA	NA	NA	Alive, wild sea lion
47	WCSL ZC954	CA sea lion	CA	-/urine	No	NA	NA	NA	Alive, wild sea lion
48	WCSL ZC956	CA sea lion	CA	-/urine	No	NA	NA	NA	Alive, wild sea lion
49	WCSL ZC958	CA sea lion	CA	-/urine	No	NA	NA	NA	Alive, wild sea lion
50	WCSL ZC960	CA sea lion	CA	-/urine	No	NA	NA	NA	Alive, wild sea lion
51	WCSL ZC962	CA sea lion	CA	-/urine	No	NA	NA	NA	Alive, wild sea lion
52	WCSL ZC963	CA sea lion	CA	-/urine	No	NA	NA	NA	Alive, wild sea lion
Stored urine samples									
53	CSL6707	CA sea lion	CA	-/urine	Yes	Positive	NA	NA	Released/leptospirosis
54	CSL6797	CA sea lion	CA	+ ( <i>L. interrogans</i> )/ urine	Yes	Positive	NA	NA	Dead/leptospirosis
55	CSL6809	CA sea lion	CA	+ ( <i>L. interrogans</i> )/ urine	Yes	Positive	NA	NA	Dead/leptospirosis
56	CSL6807	CA sea lion	CA	-/urine	Yes	Positive	NA	NA	Released/leptospirosis
57	CSL6820	CA sea lion	CA	-/urine	Yes	Positive	NA	NA	Released/leptospirosis
58	CSL6645	CA sea lion	CA	-/urine	Yes	Positive	NA	NA	Released/leptospirosis
59	CSL6818	CA sea lion	CA	-/urine	Yes	Positive	NA	NA	Dead/leptospirosis
60	CSL6824	CA sea lion	CA	-/urine	Yes	Positive	NA	NA	Dead/leptospirosis
61	CSL6828	CA sea lion	CA	-/urine	Yes	Positive	NA	NA	Dead/leptospirosis

<sup>a</sup> Cases 6, 8, and 11 represent potential carrier animals. Cases 36 to 52 represent samples taken from healthy captive (cases 36 and 37) and wild (cases 38 to 52) animals. Cases 53 to 61 represent samples stored at -80°C for >6 months.  
<sup>b</sup> NA, not available.



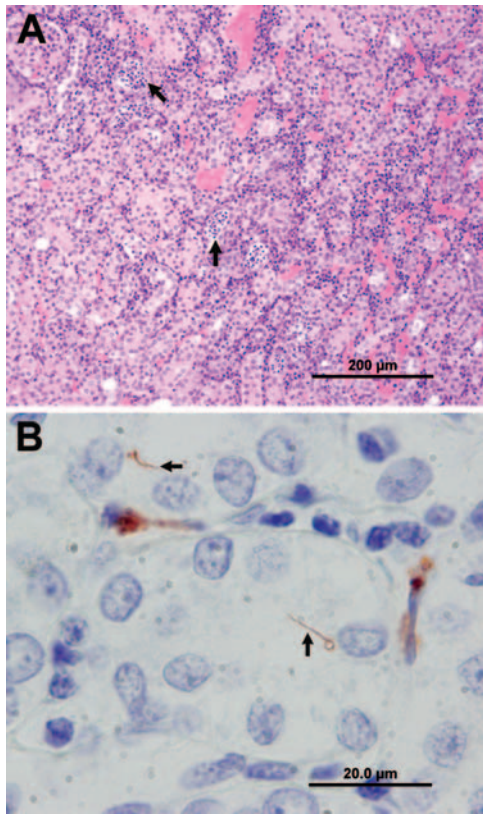


FIG. 1. Images taken from a California sea lion with leptospirosis. (A) Photomicrograph of a hematoxylin and eosin-stained section of kidney illustrating numerous lymphocytes and plasma cells infiltrating the renal cortical interstitium. Tubules are occasionally ecstatic, lined by flattened epithelial cells and contain eosinophilic fluid, necrotic epithelial cells, and neutrophils (arrows). (B) Photomicrograph of immunohistochemistry for *Leptospira* sp. demonstrating positive staining in renal interstitial inflammatory cells and of spirochetes within renal tubules (arrows). Images A and B were taken from case 16 (CSL6176).

investigated in this study (cases 36 to 52), including both captive and wild sea lions that were serologically negative by MAT, were also negative for *Leptospira* DNA by PCR.

Urine samples stored at  $-80^{\circ}\text{C}$  for extended periods (6 months or longer) demonstrated a reduced amplification potential, as exemplified by cases 53 to 61, which were positive by

clinical observations and serological investigations but demonstrated only a 22% amplification accuracy by PCR. No such reduction in amplification potential was observed in similarly stored renal tissue samples or in freshly isolated urine samples, as shown by the reliable amplification observed in cases 1 to 34.

Interestingly, samples collected from California sea lions and Steller sea lions exhibited positive amplification using the *L. interrogans*-specific primer pair, while the elephant seal sample (case 1) demonstrated positive amplification using the *L. kirschneri*-specific primer pair.

Table 2 shows a comparative summary of the methods used to detect the presence of *Leptospira* in our study. The use of PCR identified the highest number of *Leptospira* infections, with 34 samples showing positive PCR amplification. This number exceeds the number of cases deemed positive for leptospirosis via conventional methodology, which included clinical observations (27 cases) and serological (21 cases), histopathological (21 cases), and immunohistochemical (15 cases) investigations. An overall pattern of specificity was observed for the PCR analyses, with 0/17 healthy animals, 28/28 freshly isolated urine samples, and 7/7 kidney samples displaying amplification.

## DISCUSSION

The widespread 2004 leptospirosis outbreak, in which sea lion mortalities were observed from California to British Columbia, afforded an opportunity to evaluate and compare PCR-based methods to existing methods for diagnosis of *Leptospira* infection among marine mammal populations. These data clearly demonstrate that *Leptospira* DNA can be successfully and accurately amplified from samples collected from stranded marine mammals, using DNA templates prepared from a wide variety of sources, including urine, renal tissue, and feces- and urine-contaminated sand samples collected from the vicinity of the stranded animal. The positive amplifications observed in the latter two sources of sample material are particularly noteworthy in that this result allows biologists an opportunity to collect samples from live pinnipeds that have clinical signs consistent with leptospirosis but return to the ocean prior to urine collection. Furthermore, these results suggest leptospirosis could be transmitted via contamination of the coastal environment if the excreted bacteria remain viable

TABLE 2. Comparative summary of methods to detect the presence of *Leptospira* in pinnipeds using fresh samples

Animals ( <i>n</i> )	No. positive/no. of samples by:						
	PCR			Clinical observations	MAT	Histopathology	IHC <sup>a</sup>
	Urine	Kidney	Total				
Captive healthy (2)	0/2	0/0	0/2	0/2	0/2	0/0	0/0
Wild healthy (15)	0/15	0/0	0/15	0/15	0/15	0/0	0/0
Leptospirosis cases (28)	25/25 <sup>b</sup>	4/4 <sup>b</sup>	28/28	27/27	21/21	21/21	15/15
Other clinical cases (4)	3/3	1/1	4/4	0/3	0/3	0/3	0/3
Unknown (3)	0/1	2/2	2/3	1/1	0/1	1/1	0/0
Total positive/total cases examined	28/28 <sup>b</sup>	7/7 <sup>b</sup>	34/35	28/31	21/25	22/25	15/18

<sup>a</sup> IHC, immunohistochemistry.

<sup>b</sup> Includes case 4 (CSL6144), for which both urine and kidney were tested.

for a significant period of time. The suitability of PCR for detecting *Leptospira* infections in pinniped populations is further shown by the successful identification of infected animals that were unable to be analyzed by conventional diagnostic methodologies, including decomposed animal carcasses. Additionally, the sensitivity and specificity of PCR are useful in distinguishing renal failure due to leptospirosis from other causes of renal failure. A key finding revealed through these studies was the labile nature of isolated urine samples; templates prepared from fresh urine stocks provided reproducible amplifications, while templates prepared from urine samples stored at  $-80^{\circ}\text{C}$  for 6 months or longer resulted in sporadic amplification and unreliable results. Collectively, these investigations highlight the versatility of PCR over conventional diagnostic methodologies, in that analyses can be tailored to the available sample type collected from stranded animals, but attention must be given to proper sample storage and processing to ensure valid results are achieved using this technique.

The three animals that did not exhibit clinical symptoms of disease and displayed only mild nephritis upon necropsy, but displayed positive PCR amplicons, suggest that PCR may be useful in the detection of carrier animals. Carrier animals could shed *Leptospira* sp. in their urine and in this way function as reservoirs for *Leptospira* transmission. Detection of carrier animals is vital to the understanding of enzootic and epizootic leptospirosis in marine mammals, since pathogenic *Leptospira* serovars have been shown to survive for only short periods of time in seawater (8, 13) and therefore the mode of transmission of this organism in a marine species is not understood. Further investigation of the carrier status of California sea lions using PCR is warranted.

Use of two primer sets that discriminate between *L. interrogans* and *L. kirschneri* suggests these two species may have different host preferences. Samples from California and Steller sea lions were positive for *L. interrogans* but negative for *L. kirschneri*. In contrast, the one sample obtained from an elephant seal was positive for *L. kirschneri* but negative for *L. interrogans*. Expansion of these experiments to include additional marine and terrestrial mammal species, combined with the use of supplementary *Leptospira* species-specific primer pairs, will provide invaluable information on transmission routes, both within the marine environment and between terrestrial and marine environments, and may identify additional incidental and maintenance hosts for this disease. Overall, this study has shown that PCR represents a powerful diagnostic technique that has many advantages over classic methods of leptospirosis diagnosis.

The majority of animals included in this study stranded along the California coast, a region that is routinely frequented by recreational sporting enthusiasts and densely populated with humans, pinnipeds, rodents, and domestic animals, thus establishing an optimal milieu for zoonotic disease transmission and long-term disease maintenance. Many of the animals stranded near freshwater estuaries, increasing the potential for disease transmission to humans and domestic animals due to enhanced leptospiral survival in freshwater. Furthermore, detection of *Leptospira* in sand contaminated by fecal material and urine in this study suggests a potential environmental

source of pathogen exposure. Collectively, these observations reveal the significant zoonotic potential of leptospirosis within a marine environment. This investigation increases our understanding of potential routes of *Leptospira* transmission and will provide scientists and animal health experts with methodology to rapidly and accurately diagnose future outbreaks of leptospirosis.

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