

RESEARCH ARTICLE

Isolation of a Seawater Tolerant *Leptospira* spp. from a Southern Right Whale (*Eubalaena australis*)

Sylvia Grune Loffler^{1,2☯*}, Virginia Rago^{3,4‡}, Mara Martínez^{1☯}, Marcela Uhart^{4,5‡}, Monica Florin-Christensen^{1,2☯}, Graciela Romero^{1☯}, Bibiana Brihuega^{1☯□}

1 Institute of Pathobiology, National Institute of Agricultural Technology, Hurlingham, Buenos Aires, Argentina, **2** National Research Council of Argentina (CONICET), Buenos Aires, Argentina, **3** Institute of Ecology, Genetics and Evolution, National Research Council of Argentina (CONICET), University of Buenos Aires, Buenos Aires, Argentina, **4** Southern Right Whale Health Monitoring Program, Puerto Madryn, Chubut, Argentina, **5** One Health Institute, School of Veterinary Medicine, University of California Davis, Davis, California, United States of America

☯ These authors contributed equally to this work.

□ Current address: Laboratory of Leptospirosis, Institute of Pathobiology, National Institute of Agricultural Technology, Hurlingham, Buenos Aires, Argentina

‡ These authors also contributed equally to this work.

* grune.sylvia@inta.gob.ar



OPEN ACCESS

Citation: Grune Loffler S, Rago V, Martínez M, Uhart M, Florin-Christensen M, Romero G, et al. (2015) Isolation of a Seawater Tolerant *Leptospira* spp. from a Southern Right Whale (*Eubalaena australis*). PLoS ONE 10(12): e0144974. doi:10.1371/journal.pone.0144974

Editor: Brian Stevenson, University of Kentucky College of Medicine, UNITED STATES

Received: September 9, 2015

Accepted: November 26, 2015

Published: December 29, 2015

Copyright: © 2015 Grune Loffler et al. This is an open access article distributed under the terms of the [Creative Commons Attribution License](https://creativecommons.org/licenses/by/4.0/), which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

Data Availability Statement: All relevant data are within the paper.

Funding: These authors have no support or funding to report.

Competing Interests: The authors have declared that no competing interests exist.

Abstract

Leptospirosis is the most widespread zoonotic disease in the world. It is caused by pathogenic spirochetes of the genus *Leptospira* spp. and is maintained in nature through chronic renal infection of carrier animals. Rodents and other small mammals are the main reservoirs. Information on leptospirosis in marine mammals is scarce; however, cases of leptospirosis have been documented in pinniped populations from the Pacific coast of North America from southern California to British Columbia. We report the isolation of a *Leptospira* spp. strain, here named Manara, from a kidney sample obtained from a Southern Right Whale (*Eubalaena australis*) calf, which stranded dead in Playa Manara, Península Valdés, Argentina. This strain showed motility and morphology typical of the genus *Leptospira* spp. under dark-field microscopy; and grew in Ellinghausen-McCullough-Johnson-Harris (EMJH) medium and Fletcher medium after 90 days of incubation at 28°C. Considering the source of this bacterium, we tested its ability to grow in Fletcher medium diluted with seawater at different percentages (1%, 3%, 5%, 7% and 10% v/v). Bacterial growth was detected 48 h after inoculation of Fletcher medium supplemented with 5% sea water, demonstrating the halophilic nature of the strain Manara. Phylogenetic analysis of 16S rRNA gene sequences placed this novel strain within the radiation of the pathogenic species of the genus *Leptospira* spp., with sequence similarities within the range 97–100%, and closely related to *L. interrogans*. Two different PCR protocols targeting genus-specific pathogenic genes (G1-G2, B64I-B64II and LigB) gave positive results, which indicates that the strain Manara is likely pathogenic. Further studies are needed to confirm this possibility as well as determine its serogroup. These results could modify our understanding of the epidemiology of this zoonosis. Until now, the resistance and ability to grow in seawater for long periods of

time had been proven for the strain Muggia of *L. biflexa*, a saprophytic species. To the best of our knowledge, this is the first isolation of a *Leptospira* sp. from cetaceans. Our phenotypic data indicate that strain Manara represents a novel species of the genus *Leptospira*, for which the name *Leptospira brihuegai* sp. nov. is proposed.

Introduction

Leptospirosis is a neglected zoonotic disease, endemic in most tropical and subtropical regions of the world. The causative agents of this zoonosis are pathogenic strains belonging to the order Spirochaetales, family Leptospiraceae and genus *Leptospira* spp. Leptospirosis is maintained in nature through chronic renal infection of carrier animals, with rodents and other small mammals as the most important reservoirs [1–3].

In marine animals, seropositivity has been reported in North America, in California sea lions (*Zalophus californianus*) [4], Northern fur seals (*Callorhinus ursinus*) [5], Northern elephant seals (*Mirounga angustirostris*) [6], and harbor seals (*Phoca vitulina*) [7,8]. Furthermore, *Leptospira* spp. can cause disease in pinnipeds [9–12], periodic large scale stranding and mortality events, every three to four years, of California sea lions along the Pacific coast of North America from southern California to British Columbia have been attributed to *L. interrogans* serovar Pomona infections [4, 10, 11, 12, 13,14]. Seroprevalence in marine mammals has been reported also in the Pacific coast of South America, in Peruvian Amazon manatees (*Trichechus inunguis*) [15] and in Chilean South American sea lions (*Otaria byronia*) [16].

Cameron et al. [10], describe clinical signs of Leptospirosis in sea lions as renal failure, dehydration, polydipsia, vomiting, and depression. Using species-specific primer pairs Cameron et al. [10] revealed host specificity of *L. interrogans* for sea lions and of *L. kirschneri* for elephant seals.

The genus *Leptospira* spp. displays a great genomic plasticity and several schemes for the genotyping of pathogenic species have been developed during the last years around the world [17–19]. Bacterial isolation followed by molecular characterization has recently been successfully applied to genotype pathogenic *Leptospira* spp. strains in Argentina. A considerable number of pathogenic *Leptospira* spp. from wildlife [20–22], domestic animals [23–25] and water samples [26] were thus characterized.

Environmental factors are key determinants in leptospiral distribution, as water is the vehicle by which leptospirae travel and disseminate into ecosystems. To date, only one strain of *Leptospira* spp., *L. biflexa* strain Muggia, has been isolated from seawater in a region near Trieste, Italy [27, 28]. Growth of other *Leptospira* spp., including *L. interrogans*, in the presence of diluted seawater has always yielded negative results [2, 3, 29, 30]. However, Saito et al. [31] isolated two leptospiral strains after a typhoon in Philippines in soil samples, and commented that both isolated strains could live in seawater for only three and four days respectively.

The present work describes the characterization of a *Leptospira* sp. strain, isolated from the kidney of a stranded Southern Right Whale in Patagonia, Argentina, that grows optimally in the presence of seawater. This is the second halophilic *Leptospira* spp. reported to date, and to the best of our knowledge, the first one ever isolated from a cetacean.

Materials and Methods

Sample collection

A total of 27 kidney samples from 27 dead stranded Southern Right Whales (*Eubalaena australis*) Península Valdés, Argentina, were collected by the Southern Right Whale Health

Monitoring Program (SRWHMP) between 2009 and 2010. Field permits for this work were issued annually by the Department of Wildlife of the Subsecretary of Tourism and Protected Areas, Chubut Province, Argentina. Necropsies were performed using a Right Whale necropsy protocol developed by the SRWHMP (Chirife et al., 2014 unpublished) based on the methods of McCellan et al. [32], F. Gulland (pers. comm.), A. Carribero (unpublished) and Geraci & Lounsbury [33]. Carcass decomposition was graded subjectively on a scale from condition code 2 to 5 (2 = fresh, 3 = decomposed but tissues largely intact, 4 = advanced decomposition, 5 = mummified or skeletonized) [33]. Samples were placed in Whirlpack® bags, stored in liquid nitrogen at the site, and later transferred on dry ice to the laboratory for diagnosis. Samples were stored at -70°C until processing.

Isolation and maintenance of cultures

All kidney samples were cultured in leptospire-specific Ellinghausen-McCullough-Johnson-Harris (EMJH) and Fletcher media (Difco Laboratories). Cultures were incubated at 28°C for 90 days and observed every 15 days under dark-field microscopy to evaluate possible bacterial growth. Once the culture was positive, subcultures were conducted to maintain the isolated strain alive.

Growth assay in the presence of seawater. After weak growth in both EMJH and Fletcher media and considering the source of isolation, we tested the ability of these bacteria to grow in each of these media after the addition of seawater at different percentages (1%, 3%, 5%, 7% and 10%, v/v) [30,31]. Seawater was collected from Puerto Madryn, Chubut, Argentina, and sterilized by filtration through 0.22 µm Millipore filters. Cultures were incubated at 28°C for up to 12 days. A control experiment was carried out, where filter-sterilized seawater was cultured with both EMJH and Fletcher Media. Immunofluorescence staining of the strain was performed with a multivalent FA, LEP-FAC (Seasinglab, Argentina) conjugate specific for *Leptospira* spp.

Molecular characterization

Two different PCR protocols were tested to indicate if the isolated strain was pathogenic, amplifying genus-specific genes that are considered pathogenicity determinants. DNA templates were obtained after nucleic acid purification using the Chelex-100 resin (Bio Rad) protocol [22].

Multiplex PCR. The combined primers G1-G2 and B64I-B64II previously described in Gravekamp et al., [34] were used in this study. The PCR reaction was carried out in a final volume of 50 µl, which contained 2 µl purified DNA template, and the PCR mixture and the cycling program used was the same as indicated in Gravekamp et al. [34]. PCR was carried out in a My Cycler™ thermocycler (Bio Rad). Amplification products were analyzed by electrophoresis in ethidium bromide stained 2% agarose gels, followed by exposure to UV light (Uvi Tec transilluminator BTS-20.M). Amplicon sizes were estimated using a 100 bp ladder (Emitect).

PCR for LigB. A 1 kb sequence of the adhesin Lig B gene, which is present in all pathogenic *Leptospira* spp. strains, was amplified by PCR using primers LigBpetF and LigBpetR [35]. The PCR mixture and cycling was carried out as indicated in Martínez et al. [35]. PCR was carried out in a My Cycler™ thermocycler (BIO RAD) and amplified samples were analyzed by electrophoresis in 1% agarose gels stained with Sybr® Safe (Invitrogen) and visualized under UV light.

16S rRNA sequencing. A PCR targeting the 16S rRNA gene was carried out for bacterial identification after sequencing. The following primers were used: 5'-

GGCGGCGCGTCTTAAACATG-3' and 5'-GTCCGCCTACGCACCCCTTTACG-3' [36]. These primers have the ability to amplify all pathogenic and nonpathogenic species of *Leptospira* spp. PCR was performed as indicated in Djadid et al. [36]. After verification of the amplicon by electrophoresis in an ethidium bromide-containing 2% agarose gel and visualization upon UV light exposure, PCR products were purified using a commercial kit (EMBIOTECH). The samples were sequenced at the Institute of Biotechnology, National Institute of Agricultural Technology using a 3130xl Genetic Analyzer (Applied Biosystems). For alignment and construction of the phylogeny, the program MEGA version 6.06 [37] was used. The dendrogram was constructed using Neighbor-joining with a bootstrap of 100, partial sequences of the 16S rRNA gene were used.

Results and Discussion

Isolation of *Leptospira* sp. and maintenance of the cultures

One positive culture of *Leptospira* sp. was obtained from one Southern Right Whale, a newborn female calf (ID 092410PV-Ea23), 5.05 meters length. The calf stranded dead on September 24th 2010 at Playa Manara in Golfo Nuevo, Península Valdés (42° 40' 18.9"S 64° 59' 22.9"W), and was necropsied the following day. Carcass condition at necropsy was 3.

The isolated strain was designated strain Manara. This strain showed typical motility and morphology of the genus *Leptospira* spp. under dark-field microscopy and grew in EMJH and Fletcher media after 90 days of incubation at 28°C. When seawater (1%, 3%, 5%, 7%, 9% and 10%, v/v) was added, no growth was observed in EMJH medium during the first 6 days. At day 7, positive growth based on the turbidity of the culture was detected at 1% seawater (Fig 1). A control experiment, where only seawater was incubated in the same media did not show any leptospiral growth. On the other hand, growth of strain Manara was observed at 48 h in Fletcher medium, with a typical dinger ring formation in the culture. After 72 h, growth was achieved at percentages of 1% and 3%, and after 96 h, all cultures were positive (Table 1). At day 8 of this assay, new subcultures were started in Fletcher medium supplemented with the same seawater dilutions as before (Table 2). Importantly, the same growth behavior was observed. These experiments confirmed the ability of strain Manara to grow *in vitro* in the presence of diluted seawater demonstrating its halophilic nature.

Immunofluorescence detection was positive and leptospire could be microscopically observed. (Fig 2).

To the best of our knowledge, there are no recognized halophilic pathogenic leptospiral strains. Previous studies have demonstrated that *Leptospira* spp. can grow in seawater [27, 28] but the identification of serogroup and serovar was not provided and the authors indicated that this unique strain Muggia could belong to the species *L. biflexa*. In the study of Cinco et al. [28], the isolated halophilic strain grew in Korthof-Babudieri seawater medium for about 16 to 18 h, and a maximal concentration of 2×10^8 cells/ml was reached after 10 days of incubation at 30°C. In their study, the strain was considered as halodependent to specific NaCl requirements for growth. On the other hand, Saito et al. [31], could isolate two leptospiral strains from soil samples and tested the ability to grow in seawater, one strain called MS422 could grow in seawater for four days and the second strain MS432 only for three days. Both strains grew also in media with PBS for six days. These studies did not extend for a longer time period. In the same study [31], both strains were sequenced using the 16S rRNA gene and both showed similarity to *L. kmetyi*.

In our study, the strain Manara was capable of growing in the presence of different concentrations of seawater (Table 1). In media that was not supplemented with seawater, growth of Manara strain was considerably slower and could not be detected until 30 days (results not

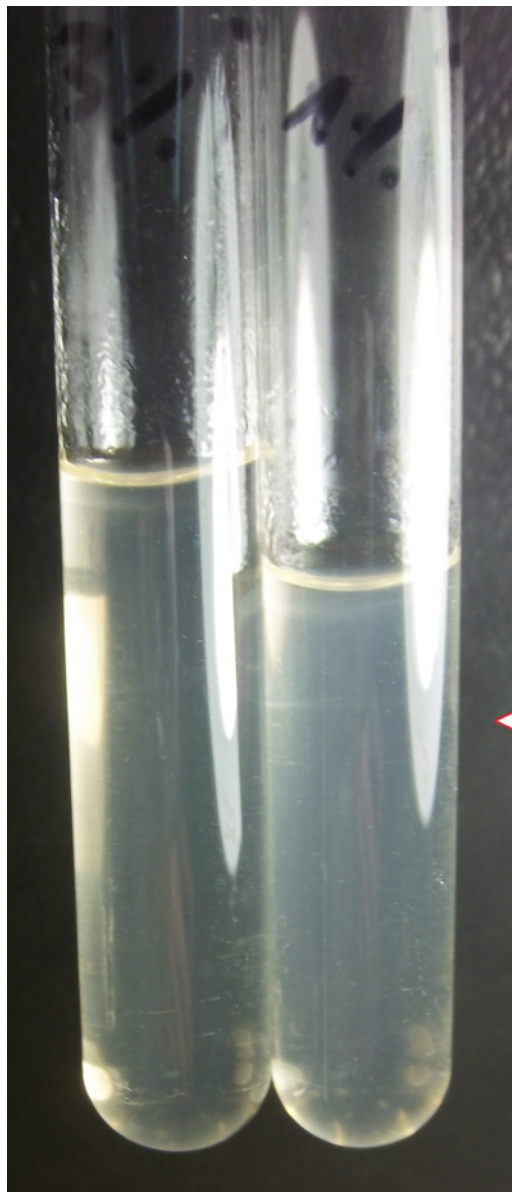


Fig 1. Picture of two cultures of *Leptospira* sp. strain Manara. Two cultures of *Leptospira* sp. strain Manara in Fletcher medium supplemented with 3% (left tube) and 1% (right tube) seawater after 48 h of incubation at 28°C. The dinger rings corresponding to *Leptospira* sp. growth are indicated with red arrows.

doi:10.1371/journal.pone.0144974.g001

shown), whereas growth could be detected as soon as 48 h after inoculation when cultured in media with 5% seawater in Fletcher media. Further studies are necessary to characterize the biological features of this novel strain.

Molecular characterization

Two different PCR protocols were used to explore the possible pathogenicity of the Manara strain, amplifying the genus-specific and virulence-associated genes G1-G2, B64I-B64II and LigB, both with positive results. Phylogenetic analysis of 16S rRNA gene sequences placed this novel strain within the radiation of the pathogenic species of the genus *Leptospira* spp., with

Table 1. Growth results of strain Manara in EMJH and Fletcher media diluted with seawater.

day	EMJH Medium					Fletcher Medium				
	1%	3%	5%	7%	10%	1%	3%	5%	7%	10%
1	-	-	-	-	-	-	-	-	-	-
2	-	-	-	-	-	-	-	+	-	-
3	-	-	-	-	-	+	+	+	-	-
4	-	-	-	-	-	+	+	+	+	+
5	-	-	-	-	-	+	+	+	+	+
6	-	-	-	-	-	+	+	+	+	+
7	+	-	-	-	-	+	+	+	+	+
8	+	-	-	-	-	+	+	+	+	+
9	+	-	-	-	-	+	+	+	+	+
10	+	-	-	-	-	+	+	+	+	+
11	+	-	-	-	-	+	+	+	+	+
12	+	-	-	-	-	+	+	+	+	+

Growth results of strain Manara in EMJH and Fletcher media diluted with seawater at percentages 1%, 3%, 5%, 7% and 10% (v/v). + and–correspond to detection and no detection of growth, respectively; Subcultures (marked with an asterisk) under the same conditions were started at day 8.

doi:10.1371/journal.pone.0144974.t001

sequence similarities within the range 97–100% (results not shown), and closely related to *L. interrogans* (Fig 3). Further molecular characterization must be pursued to confirm this possibly new species, such as sequencing secY gene. The sequence obtained is available in GenBank under Accession number KP901270.

Detection of new strains in the environment and carrier animals is vital to the understanding of enzootic and epizootic leptospirosis in marine mammals. Since pathogenic *Leptospira* spp. serovars are extremely difficult to cultivate *in vitro* [29,30], the mode of transmission of this organism in marine species is not understood [10–13]. Detection by PCR of leptospiral DNA in sand was observed by Cameron et al. [10], however this study could not determine if *Leptospira* sp. was dead or alive in sand. In this study, no samples were taken at site, where the Southern Right Whale stranded dead, but the possibility exists that leptospiral DNA could be also in sand. Potential environmental sources of pathogen exposure in the marine environment could increase the zoonotic potential of Leptospirosis in marine mammals, however the way of transmission is yet unclear [5,7,8,10,11,12,14].

Table 2. Growth results of subcultures of strain Manara in Fletcher media diluted with seawater after 8 days.

day	Subculture from Fletcher Medium after 8 days				
	1%	3%	5%	7%	10%
1	-	-	-	-	-
2	-	-	-	-	-
3	+	+	+	+	+
4	+	+	+	+	+
5	+	+	+	+	+

Growth results of subcultures strain Manara started at day 8 of the first culture assay, in Fletcher media diluted with seawater at percentages 1%, 3%, 5%, 7% and 10% (v/v). + and–correspond to detection and no detection of growth, respectively.

doi:10.1371/journal.pone.0144974.t002

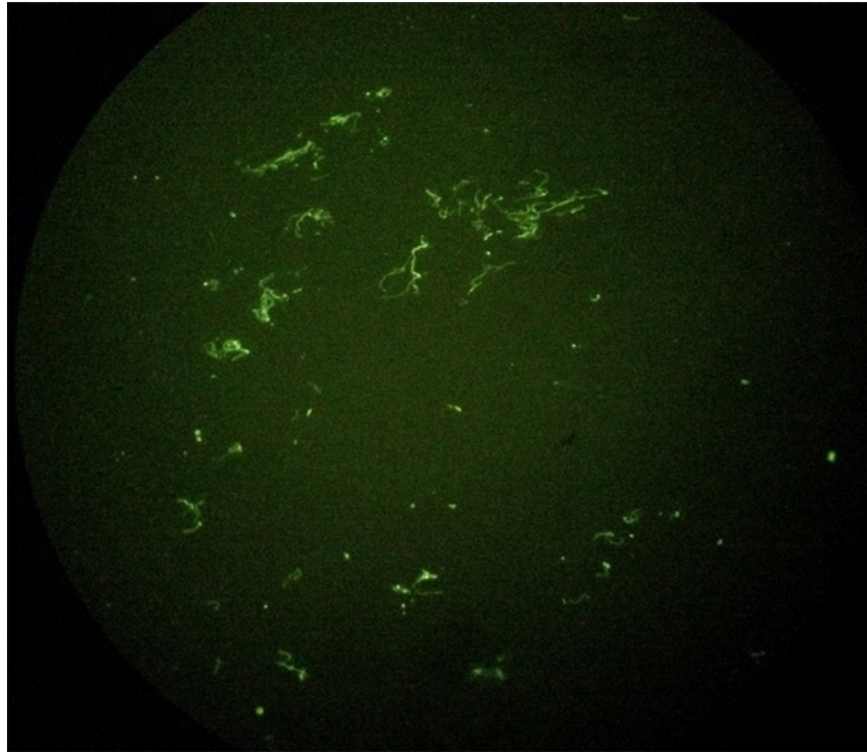


Fig 2. Immunofluorescence of strain Manara. Immunofluorescence of strain Manara isolated from a kidney sample of a Southern right whale (*Eubalaena australis*) 40X.

doi:10.1371/journal.pone.0144974.g002

Confirmation that this isolate contained the gene LigB, a gene associated with virulence and present in all pathogenic *Leptospira* spp., suggests this strain is pathogenic and is consistent with our sequencing results which place strain Manara within the radiation of other pathogenic *Leptospira* spp. Further studies in animal models must be conducted to confirm this possibility. In addition, determination of the serogroup and serovar of this strain are the object of ongoing research.

Due to the phenotypic findings in this study we propose that strain Manara belongs to a new species of *Leptospira* spp., phylogenetically closely related to *L. interrogans*, but with the particular features of being able to survive during a long period of time in seawater, and to colonize unreported hosts, as the Southern right whale.

Ethiology of *Leptospira brihuegai* sp. Nov. The proposed name for strain Manara is *Leptospira brihuegai*, after professor Bibiana Brihuega, an Argentinean veterinarian and bacteriologist who has made significant contributions to the study of Leptospirosis.

Conclusion

This study reports the first isolation of a *Leptospira* sp. from a cetacean. Phylogenetic analysis of a sequence obtained by targeting the 16S rRNA as well as its host specificity and growth requirements identified this as a novel halophilic *Leptospira* sp. strain. These findings contribute highly relevant information to current knowledge on leptospirosis epidemiology and ecology.

New findings concerning the biology of leptospires is a challenge and more extensive studies are required to monitor the presence of *Leptospira* spp. in the environment and different

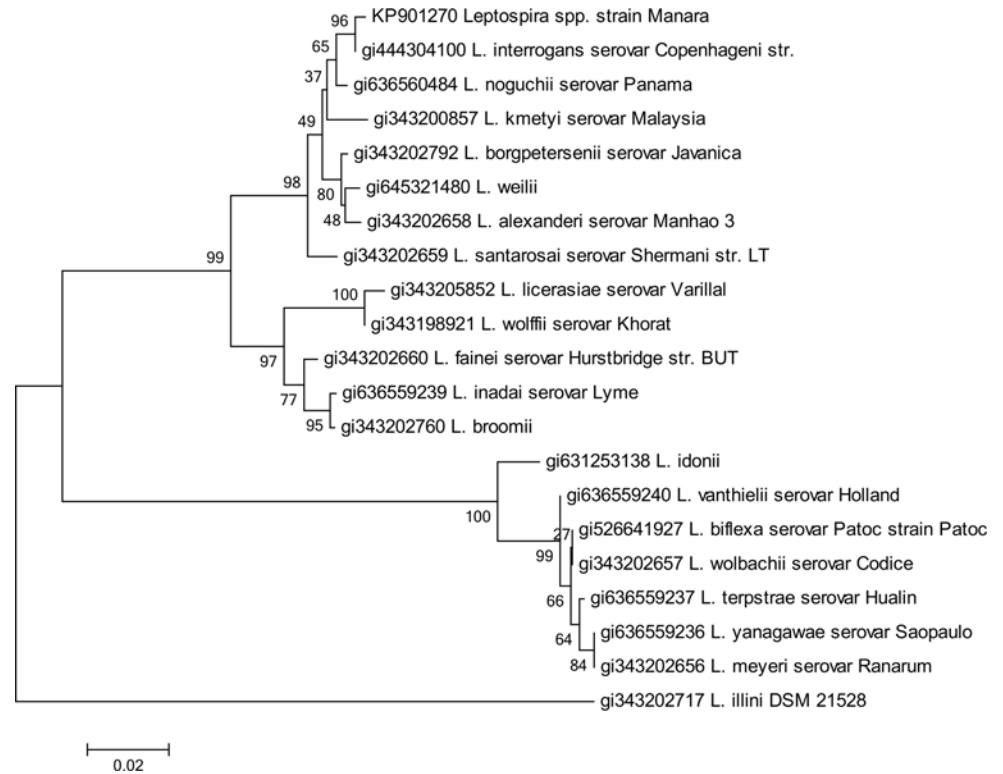


Fig 3. Neighbor-joining analysis of the sequence 16S rRNA of strain Manara. Phylogenetic analysis based of 16S rRNA including 19 representative species of *Leptospira* spp. The corresponding sequence of *Leptonema illini* strain DSM 21528 was used as outgroup. The dendrogram was constructed using Neighbor-joining. Bootstrap values are displayed as percentages.

doi:10.1371/journal.pone.0144974.g003

mammalian hosts. Our study contributes valuable information to our knowledge on leptospires overall, and on marine mammals in particular.

Acknowledgments

We thank members, researchers and collaborators of the SRWHMP: M. Sironi, V. Rowntree, La Sala, L. M. Pozzi, L. Musmeci, N. Mohamed, J. E. Sala, J. Andrejuk, L. Bandieri, A. Chirife, M. Di Martino, L. Beltramino, D. Taboada, R. Schteinbarg, L. Valenzuela, M. Ricciardi, P. de Diego, C. Marón, and the many volunteers who make fieldwork possible. The Instituto de Conservación de Ballenas, Wildlife Conservation Society, Ocean Alliance, Fundación Patagonia Natural, Ecocentro Puerto Madryn, Armada Argentina, Prefectura Naval Argentina and Aluar also provided invaluable support of various kinds. Research permits for this work were issued annually by the Dirección de Fauna y Flora Silvestre and the Subsecretaría de Turismo y Áreas Protegidas of Chubut Province, Argentina.

Author Contributions

Conceived and designed the experiments: SGL BB GR MFC MM. Performed the experiments: SGL MM. Analyzed the data: SGL BB MFC GR MM. Contributed reagents/materials/analysis tools: VR MU BB SGL MM. Wrote the paper: SGL BB MFC MM MU VR.

References

1. Sterling CR, Thiermann AB. Urban rats as chronic carriers of leptospirosis: an ultrastructural investigation. 1981. *Vet Pathol*, 18:628–637. PMID: [7281461](#)
2. Faine S. Guidelines for the control of leptospirosis, WHO offset publication N° 67. 1982. World Health Organization, Geneva, 161 pg.
3. Levett PN. Leptospirosis. 2001. *Clin Microbiol Rev*, 14:296–326. PMID: [11292640](#)
4. Gulland FMD, Koski M, Lowenstine LJ, Colagross A, Morgan L, Spraker T. Leptospirosis in California sea lions (*Zalophus californianus*) stranded along the central California coast, 1981–1994. 1996. *J. Wildl. Dis.* 32; 572–80. PMID: [9359054](#)
5. Smith AW, Brown RJ, Skilling DE, Bray HL, Keyes MC. Naturally occurring Leptospirosis in Northern fur seals (*Callorhinus ursinus*). 1977. *J. Wildl. Dis.* 13;144–8. PMID: [864847](#)
6. Colegrove KM, Lowenstine JL, Gulland FMD. Leptospirosis in northern elephant seals (*Mirounga angustirostris*) stranded along the California coast. 2005. *J. Wildl. Dis.* 41; 426–30. PMID: [16107678](#)
7. Stamper MA, Gulland FMD, Spraker T. Leptospirosis in rehabilitated Pacific harbor seals from California. 1998. *J. Wildl. Dis.* 34; 407–10. PMID: [9577797](#)
8. Greig DJ, Gulland FM, Smith WA, Conrad PA, Field CL, Fleetwood M, et al. Surveillance for zoonotic and selected pathogens in harbor seals *Phoca vitulina* from central California. 2014. *Dis Aquat Org* 111;93–106. doi: [10.3354/dao02762](#) PMID: [25266897](#)
9. Colagross-Schouthern A, Mazet JA, Gulland FMD, Miller MA, Hietala S. Diagnosis and seroprevalence of leptospirosis in California sea lions from coastal California. 2002. *Journal of Wildlife Diseases* 38 (1);7–17. PMID: [11838231](#)
10. Cameron C, Zuerner RL, Raverty S, Colegrove KM, Norman S, Lambourn DM, et al. Detection of Pathogenic *Leptospira* bacteria in Pinniped Population via PCR and Identification of a source of Transmission for zoonotic leptospirosis in the Marine Environment. 2008. *Journal of Clinical Microbiology*, 46 (5);1728–33 doi: [10.1128/JCM.02022-07](#) PMID: [18367568](#)
11. Zuerner RL, Cameron CE, Raverty S, Robinson J, Colegrove KM, Norman SA, et al. Geographical dissemination of *Leptospira interrogans* serovar Pomona during seasonal migration of California sea lions. 2009. *Vet. Microbiol.* 137; 105–10. doi: [10.1016/j.vetmic.2008.12.017](#) PMID: [19186009](#)
12. Waltzek TB, Cortés-Hinojosa G, Wellehan JFX Jr, Gray GC. Marine Mammal Zoonoses: A review of Disease Manifestations. 2012. *Zoonoses and Public Health* Dec 59(8); 521–35.
13. Norman SA, DiGiacomo RF, Gulland FMD, Meschke JS, Lowry MS. Risk factors for an outbreak of leptospirosis in California sea lions (*Zalophus californianus*) in California, 2004. 2008. *J. Wildl. Dis.* 44; 837–44. PMID: [18957639](#)
14. Lloyd-Smith J, Greig DJ, Hietala S, Ghneim G, Palmer L, Leger J, et al. Cyclical changes in seroprevalence of leptospirosis in California sea lions: endemic and epidemic disease in one host species? 2007. *BMF Infectious Diseases* 7; 125–146.
15. Delgado PM, Sanchez Perea N, Biffi Garcia C, Garcia Davila CR. Detection of infection with *Leptospira* spp. in manatees (*Trichechus inunguis*) of the Peruvian Amazon. 2015. *Latin American Journal of Aquatic Mammals* 10(1); 58–61.
16. Sepulveda MA, Seguel M, Alvarado-Rybak M, Verdugo C, Muñoz-Zanzi C, Tamayo R. Postmortem Findings in Four South American Sea Lions (*Otaria byronia*) from a Urban Colony in Valdivia, Chile. *Journal of Wildlife Disease* 51(1);279–282.
17. Cerqueria GM, Picardeau M. A century of *Leptospira* strain typing. 2009. *Infect Genet Evol* 9 (5); 760–8. doi: [10.1016/j.meegid.2009.06.009](#) PMID: [19540362](#)
18. Cerqueira GM, McBride AJ, Picardeau M, Ribeiro SG, Moreira AN, Morel V, et al. Distribution of the leptospiral immunoglobulin-like (lig) genes in pathogenic *Leptospira* species and application of ligB to typing leptospiral isolates. 2009. *Journal of Medical Microbiology* 58,1173–1181. doi: [10.1099/jmm.0.009175-0](#) PMID: [19528180](#)
19. Nalam L, Ahmed A, Majulata S, Francalacci P, Baig M, Sechi L, et al. Genetic Affinities within a Large Global Collection of Pathogenic *Leptospira*: Implications for Strain Identification and Molecular Epidemiology. 2010. *PLOS ONE* 5(8);e12637. doi: [10.1371/journal.pone.0012637](#) PMID: [20805987](#)
20. Brihuega B, Pavan M, Cairo F, Auteri C, Funes D, Romero G, et al. Leptospira patógena en riñón de *Didelphis albiventris* (comadreja). 2007. *Rev Arg Microb*; 39:19.
21. Grune Löffler S, Pavan ME, Vanasco B, Samartino L, Suarez O, Auteri C. et al. Pathogenic *Leptospira* spp. from urban and periurban rodents in Argentina. 2014. *Memorias do Instituto Oswaldo Cruz*. Rio de Janeiro: Fundaco Oswaldo Cruz (2:);1–5.
22. Grune S. Aislamiento y caracterización genotípica de leptospiras provenientes de animales silvestres en tres ecoregiones argentinas mediante la técnica del Múltiple-Locus Variable-number tandem repeat

- Análisis (MLVA): Coincidencia con genotipos provenientes de animales de producción. 2014. PhD Thesis. Universidad de Buenos Aires, 135pg.
23. Pavan M, Cairo F, Brihuega B, Samartino L. Multiple-locus variable-number tandem repeats analysis (MLVA) of *Leptospira interrogans* serovar Pomona from Argentina reveals four new genotypes. 2008. *Comp Immun Microbiol & Infec Diseases*, 31 (1); 37–45.
 24. Pavan ME, Cairó F, Pettinari MJ, Samartino L, Brihuega B. Genotyping of *Leptospira interrogans* strains from Argentina by Multiple-Locus Variable-number tandem repeat Analysis (MLVA). 2011. *Comp Immunol Microbiol Infect Dis*. 34(2); 135–41. doi: [10.1016/j.cimid.2010.06.002](https://doi.org/10.1016/j.cimid.2010.06.002) PMID: [20674025](https://pubmed.ncbi.nlm.nih.gov/20674025/)
 25. Grune Loffler S, Passaro D, Samartino L, Soncini A, Romero G, Brihuega B. Genotypes of *Leptospira* spp. strains isolated from dogs in Buenos Aires, Argentina. 2014. *Rev Argent Microbiol*. 46(3);1–13.
 26. François S, Brihuega B, Grune S, Gattarello V, Correa D, Petrakovsky J, et al. Aislamiento de *Leptospira borgpetersenii* de fuentes de agua en Argentina. 2013. *Revista Cubana de Medicina Tropical*. Nr. 2.
 27. Cinco M, Tamaro M, Cociancich L. Taxonomical, cultural and metabolic characteristics of halophilic leptospirae. *Zentralblatt für Bakteriologie, Parasitenkunde, Infektionskrankheiten und Hygiene*. 1975. *Erste Abteilung Originale Reihe A: Medizinische Mikrobiologie und Parasitologie* 233(3);400–57.
 28. Cinco M, Tamaro M, Rottini GD, Monti-Bragadin C. Comparative Serological Studies Between a newly Isolated Halophilic *Leptospira* and two other *Leptospiras* isolated from Brackish Water. 1974. *International Journal of Systematic Bacteriology* 24(1); 131–135.
 29. Khairani-Bejo S, Bahaman AR, Zamri-Saad M, Mutalib AR. The survival of *Leptospira interrogans* serovar Hardjo in the Malaysian environment. 2004. *J. Anim. Vet. Adv.* 3;123–129
 30. Trueba G, Zapata S, Madrid K, Cullen P, Haake D. Cell aggregation: a mechanism of pathogenic *Leptospira* to survive in fresh water. 2004. *International microbiology* 7; 35–40. PMID: [15179605](https://pubmed.ncbi.nlm.nih.gov/15179605/)
 31. Saito M, Miyahara S, Villanueva SYAM, Aramaki N, Ikejiri M, Kobayashi Y, et al. PCR and Culture Identification of Pathogenic *Leptospira* spp. from Coastal Soil in Leyte, Philippines, after a Storm Surge during Super Typhoon Haiyan (Yolanda). 2014. *Appl. Environ. Microbiol* 80(22);6926–32.
 32. McClellan WA, Rommel SA, Moore MJ, Pabst DA. Right whale necropsy protocol. 2004. Contract report to NOAA Fisheries.
 33. Geraci JR, Lounsbury VJ. Marine mammals ashore: a field guide for strandings (Geraci JR and Lounsbury VJ, Eds. 2005. 2nd edn. National Aquarium of Baltimore, Inc., Baltimore, MD. 344pg.
 34. Gravekamp C, Van de Kemp H, Franzen M, Carrington D, Schoone GJ, Van Eys GJ, et al. Detection of seven species of pathogenic leptospirae by PCR using two sets of primers. 1993. *J. Gen. Microbiol*. 139(8);1691–700 PMID: [8409911](https://pubmed.ncbi.nlm.nih.gov/8409911/)
 35. Martínez M, Grune Loffler S, Samartino L, Romero G, De la Fuente I, Brihuega B. Detección de leptospiras patógenas mediante PCR usando los iniciadores LigBF y LigBR. 2014. *Revista Argentina de Zoonosis y Enfermedades Infecciosas Emergentes* 9(2);41.
 36. Djadid ND, Ganji ZF, Gouya MM, Rezvani M, Zakeri S. A simple and rapid nested polymerase chain reaction-restriction fragment length polymorphism technique for differentiation of pathogenic and non-pathogenic *Leptospira* spp. 2009. *Diagn. Microbiol. Infect Dis*. 63(3); 251–6. doi: [10.1016/j.diagmicrobio.2008.10.017](https://doi.org/10.1016/j.diagmicrobio.2008.10.017) PMID: [19097839](https://pubmed.ncbi.nlm.nih.gov/19097839/)
 37. Tamura K, Peterson D, Peterson N, Stecher G, Nei M, Kumar S. MEGA 5: Molecular Evolutionary Genetics Analysis using maximum Likelihood, Evolutionary Distance, and Maximum Parsimony Methods. 2011. *Molecular Biology and Evolution* 28; 2731–2739. doi: [10.1093/molbev/msr121](https://doi.org/10.1093/molbev/msr121) PMID: [21546353](https://pubmed.ncbi.nlm.nih.gov/21546353/)