Fatal Case of *Campylobacter lari* Prosthetic Joint Infection and Bacteremia in an Immunocompetent Patient

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*Campylobacter lari* is an infrequent cause of intestinal and extraintestinal infection in humans. We report a case of *C. lari* prosthetic joint infection and bacteremia in an 81-year-old immunocompetent man. The infection was associated with septic shock and fatal outcome. *C. lari* may cause severe disease, even in an immunocompetent host.

Bacteria of the genus *Campylobacter* are important causes of enteritis and extraintestinal infection in humans. Although the majority of documented *Campylobacter* infections are caused by *C. jejuni*, *C. coli*, and *C. fetus*, other species are being increasingly recognized as human pathogens.

*Campylobacter lari* is infrequently isolated from humans, but has been associated with enteritis (3, 5, 12, 15, 16), bacteremia (5, 6, 8–11, 14, 15, 16), permanent pacemaker infection (10), purulent pleurisy (4), and urinary tract infection (2). We report a case of *C. lari* prosthetic joint infection and bacteremia in an immunocompetent patient, which had a fatal outcome. To our knowledge, this is the first report to document infection of an orthopedic prosthesis with this microorganism.

An 81-year-old male was admitted to the hospital with a 1-day history of severe pain in his right prosthetic hip joint. He had experienced pain after flexion in the same joint 2 weeks earlier, but this had resolved completely following physiotherapy. The patient had also felt generally unwell for about 3 weeks prior to admission, with dizzy spells, feverishness, and chills. He did not experience diarrhea at any time. The right total hip prosthesis was inserted 4 years previously for osteoarthritis. His past medical history also included coronary artery disease, hypertension, and atrial fibrillation, for which he was being treated with cilazapril, amiodarone, and felodipine.

At the time of admission to hospital, he had a temperature of 38.1°C, was hemodynamically stable, but was unable to stand due to severe pain in the right hip, particularly on flexion and rotation. Blood tests at admission included a leukocyte count of 15.5 × 10⁹/liter, a neutrophil count of 13.9 × 10⁹/liter, and a C-reactive protein of 143 mg/liter (normal, <10 mg/liter). An aspirate of the right hip showed numerous leukocytes, but Gram staining revealed no organisms and was reported as negative.

A diagnosis of prosthetic joint infection was made, and the patient was commenced on intravenous penicillin, flucloxacillin, and gentamicin, while waiting for the results of blood and hip aspirate cultures. A washout of the affected joint was performed 1 day after admission, and aspirated purulent material and tissue were sent for microscopy and culture.

During surgery, the patient became hypotensive, developed ischemic electrocardiogram changes, and was transferred to the Intensive Care Unit for continued ventilation and hemodynamic support. In the Intensive Care Unit, his initial mean arterial pressure was 55 mmHg (low), the cardiac index was 3.5 liter/min/m² (elevated), and the pulmonary artery occlusion pressure was 20 mmHg (high). These measurements suggested significant myocardial dysfunction in the presence of distributive (septic) shock. Over the next few hours, he required escalating doses of norepinephrine and epinephrine. During this time, there was progressive acidosis and multiple organ dysfunction. Troponin T increased from 0.2 to 0.75 µg/liter (normal, <0.1 µg/liter) over the next day, and the creatinine kinase had increased from 105 to 278 IU/liter (normal, 25 to 175 IU/liter). The raised creatinine kinase was consistent with his postoperative state, and the rise in the troponin T level could be explained by acute renal failure. However, an acute myocardial infarction could not be excluded. In view of his failure to respond to these measures, supportive therapies were withdrawn. The patient died on hospital day 2, 1 day after his admission to the Intensive Care Unit.

Blood cultures taken on the day of admission were incubated in the automated BacT/ALERT system (Organon Teknika Corporation, Durham, N.C.). The anaerobic BacT/ALERT bottle became positive after 26 h of incubation with a faintly staining curved gram-negative bacillus. The aerobic bottle remained negative. The organism was subcultured onto 5% sheep blood agar, MacConkey agar, and chocolate agar and was incubated at 36°C in a 5% CO₂ environment. Inoculated sheep blood agar plates were also incubated in an anaerobic atmosphere and at 42°C in a microaerophilic environment; the latter was prompted by the Gram stain appearance suggestive of *Campylobacter* species. Growth appeared on the microaerophilic plate after 24 h and on the anaerobic plate after 2 days. In view of these findings, the original Gram stains of the hip joint aspirates were reviewed, and bacteria with similar Gram stain morphology were seen. The original report was amended appropriately. The same microorganism was subsequently cultured from both blood and joint aspirates and was presumptively identified as *C. lari* based on phenotypic characteristics.

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Key test results include growth at 42°C; growth in 1% glyce; and no growth in 3.5% NaCl; positive tests for catalase production and nitrate reduction; negative tests for hippurate hydrolysis, indoxyl acetate hydrolysis, and H2S production; and resistance to both nalidixic acid and cephalothin.

The identification of the isolates was also confirmed by molecular methods based on a multiplex PCR targeting of the _Campylobacter lpxA_ gene. Whole-cell lysates (WCLs) were prepared by suspending the _Campylobacter_ cells (48 h of growth) in 2 ml of distilled water to a McFarland equivalent of 1. A 1-ml aliquot was transferred to a sterile 1.5-ml Eppendorf tube and boiled for 10 min. Cellular debris was removed after centrifugation (10,000 × g for 10 min), and supernatants were transferred to a second sterile 1.5-ml Eppendorf tube and stored at −20°C until use.

Three primers, recognizing species-specific regions of the _Campylobacter jejuni_ (0121; 5'-ACAACCTGGTGACGATGT TGTA-3'), _C. coli_ (0120; 5'-AGACAATAAGAGAGAATC AG-3'), or _C. lari_ (0122; 5'-CTTACAAATGTTAAAATAG GC-3') _lpxA_ gene were used in combination with primer KK2 (5'-CAATCATGWNATAGRCAATANGCC-3'). KK2 binds to a conserved region of sequence identified for all of the thermotolerant _Campylobacter_ species. Fifty-microliter reaction mixtures for the _lpxA_ multiplex consisted of 10 pmol of the primers 0120, 0121, and 0122 per reaction; 30 pmol of KK2 per reaction; 200 μM each deoxynucleoside triphosphate (dNTP); 4 mM MgCl2; 1× PCR buffer (Roche); and 10 μl of WCLs. 

FIG. 1. _lpxA_ PCR analysis of _Campylobacter_ spp. isolated from the patient in this study. Lanes: 1 and 8, Gibco-BRL 1 kb Plus; 2, _Campylobacter coli_ control (isolate WA27); 3, _Campylobacter jejuni_ control (isolate ANR0697); 4, _Campylobacter lari_ control (isolate DR1879); 5, _Campylobacter_ spp., hip aspirate; 6, _Campylobacter_ spp., blood isolate; 7, no-DNA control.

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The conditions for thermocycling were as follows: DNA was initially denatured at 94°C for 3 min. Thirty-one cycles of the following steps were subsequently performed: denaturation at 94°C for 1 min followed by annealing at 64°C for 1 min and amplification at 72°C for 1 min. A final cycle was included in which denaturation and annealing times and temperatures remained the same, but the extension time was extended to 5 min to permit completion of all initiated products. PCR amplicons were analyzed by electrophoresis through 3% agarose, gels were stained with ethidium bromide, and amplicons were visualized under UV light (254 nm). Images were captured with a Kodak electrophoresis documentation and analysis system 120.

PCR amplicons to be sequenced were first passed through a Qiagen PCR cleanup spin column (Qiagen GmbH, Hilden, Germany) as per the manufacturer’s instructions. Amplicons were sent to the University of Waikato DNA Sequencing Centre, Hamilton, New Zealand, and were processed by fluorescence dye chemistry on an ABI3100 (Perkin-Elmer). Electronic readouts were systematically compared to electropherograms, and ambiguities were resolved by using a minimum of twofold sequence redundancy of the target DNA fragments. Edited nucleotide sequences were sent to the BLAST server (National Center for Biotechnology Information) for nucleotide sequence comparisons. Alignments of DNA sequences were performed with the program Clustal X.

This identification system was developed in the laboratory of one of the authors (J.D.K.) and was validated in the following manner. The complete, redundant nucleotide sequences of the _lpxA_ gene from five _C. jejuni_, three _C. coli_, and three _C. lari_ isolates, as well as one _Campylobacter upsaliensis_ isolate (the type strain of each species in addition to other phenotypically typical isolates), were determined by Sanger dideoxy DNA sequencing by fluorescent labeling technology. The edited sequence was aligned by the program Clustal X. Regions of maximal intraspecies identity and interspecies diversity were identified, and oligonucleotide sequence primers were developed for each of these species. The multiplex PCR primers were subsequently validated in two assays. In the first assay, purified genomic DNA from 17 _Campylobacter_ and 20 non-_Campylobacter_ species were used as a template in the multiplex PCR. With the exception of _Campylobacter jejuni_ subsp. Doyleyi (which was identified as a _C. jejuni_) and _Campylobacter hyoilei_ (which has been shown to be a strain of _C. coli_), none of the nontarget species resulted in an amplicon from this multiplex PCR. A second assay, in which over 100 environmental and clinical isolates of thermotolerant _Campylobacter_ were tested, also showed a 100% correlation with the multiplex PCR. These isolates were previously confirmed as either _C. jejuni_, _C. coli_, or _C. lari_ based on classic phenotypic tests as well as species-specific PCR and pulsed-field gel electrophoresis.

The WCLs of two independently collected _Campylobacter_ species isolates from the present case (hip aspirate and blood) subjected to _lpxA_ multiplex PCR were consistent with an identification of _C. lari_ (Fig. 1). In order to confirm this result, an approximately 700-bp fragment of the small subunit rRNA gene was amplified, sequenced by methods described elsewhere (7), and compared to similar sequences from other thermotolerant and nonthermotolerant _Campylobacter_ species. A comparison of the _C. lari_ nucleotide sequence from the patient to the nucleotide sequence contained in the GenBank database revealed less than 1% divergence.

_C. lari_ infection was first described in 1980 by Skirrow and...
Benjamin, who isolated the microorganism from seagulls of the genus *Larus* (13). The organism was named *Campylobacter laridis* in 1983 (1), and the name was changed to *C. lari* in 1990 (18). The main reservoir for *C. lari* is gulls, chickens, other birds, and some domestic mammals, although it does not appear to cause overt infection in these animals.

*C. lari* belongs to the group of thermophilic campylobacters that grow optimally at 42°C. Other members of this group are *C. jejuni* and *C. coli*, which are both well-recognized human pathogens. Useful tests to distinguish *C. lari* from other *Campylobacter* species include resistance to nalidixic acid, demonstration of anaerobic growth in the presence of trimethylamine-N-oxide, susceptibility to triphenyltetrazolium chloride, hydrolysis of indoxyl acetate, and the absence of hippurate hydrolysis. It may be difficult to distinguish *C. lari* from *C. jejuni* based on phenotypic properties alone, especially since both hippurate-negative and nalidixic acid-resistant *C. jejuni* strains have been described (1, 17). Molecular testing is useful in this situation to confirm identification.

Generally, *Campylobacter* species are not easily visualized with the safranin counterstain commonly used in Gram staining. In the present case, no microorganisms were seen when the Gram stains of the hip aspirates were initially examined. When reviewed after bacteria were detected in blood cultures, faintly staining curved gram-negative bacilli were seen in all hip aspirate samples and the tissue. This highlights the difficulty in identifying *Campylobacter* in clinical samples, especially from a site at which *Campylobacter* is infrequently or unexpectedly isolated.

Our case is unusual for several reasons. It is the first reported case of a prosthetic joint infection due to *C. lari*. *C. lari* had been the causative agent of a permanent pacemaker infection (10), raising the question as to whether *C. lari* could have a predilection for prosthetic devices. Second, our patient was not known to be immunocompromised. Most reported cases of *C. lari* bacteremia have occurred in people with significant medical conditions, usually associated with immune suppression (9). Third, our patient had a severe illness associated with septic shock and fatal outcome. Only two other fatal cases of *C. lari* bacteremia have been described before, both occurring in immunosuppressed patients: one patient with multiple myeloma and chronic renal failure, the other with AIDS (6, 11). Although our patient may have had a small perioperative myocardial infarction, his general condition and hemodynamic measurements were more in keeping with a picture of septic shock. Although we suspect bloodstream invasion arose from an intestinal focus, the patient experienced no antecedent gastrointestinal symptoms. This is in keeping with the findings from a review of eight other cases of *C. lari* bacteremia; only two cases were associated with gastroenteritis (9).

*C. lari* is an infrequent, possibly underrecognized, cause of human disease. Although usually having a favorable outcome, *C. lari* may cause severe disease, even in an immunocompetent host.

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**REFERENCES**