

Post-Prostate Biopsy Infection with *Escherichia coli* ST131 Leading to Epididymo-Orchitis and Meningitis Caused by Gram-Negative Bacilli

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A 57-year-old man who had recently undergone a transrectal prostate biopsy for a rising prostate-specific antigen level developed postbiopsy necrotizing epididymo-orchitis (requiring orchiectomy) and then Gram-negative meningitis, despite fluoroquinolone administration for periprocedural prophylaxis and subsequent therapy. The causative organism proved to be a fluoroquinolone-resistant *Escherichia coli* strain from sequence type ST131.

CASE REPORT

A 57-year-old man was admitted to the hospital in 2007 for fever, chills, headache, low back pain, nausea, and vomiting.

Eight weeks previously, he had undergone transrectal ultrasound-guided prostate (TRUSP) biopsy due to a rising prostate-specific antigen (PSA) level despite long-term dutasteride therapy (2+ years) and a trial of oral ciprofloxacin (500 mg orally twice daily for 20 days before biopsy). Transrectal ultrasound showed suspicious hypoechoic areas. Perioperative oral ciprofloxacin (500 mg twice daily) was given for 72 h, beginning just prior to the biopsy, which revealed moderate chronic prostatitis, without malignancy. No culture was done.

Five days postbiopsy, bilateral scrotal swelling developed. Sequential oral levofloxacin and amoxicillin-clavulanate therapy were tried, without benefit. By day 14 postbiopsy, worsening left testicular pain and swelling prompted hospital admission and intravenous levofloxacin and gentamicin therapy. Scrotal ultrasound showed absent blood flow to the left testicle. Urgent left orchiectomy revealed testicular ischemia from severe epididymo-orchitis. The patient was discharged on hospital day 3 with 5 days of oral levofloxacin, 500 mg daily. Intraoperative cultures yielded *Escherichia coli* resistant to ampicillin, fluoroquinolones, and gentamicin; intermediate to tobramycin and ampicillin-sulbactam; and susceptible to aztreonam, cephalosporins, carbapenems, piperacillin-tazobactam, and trimethoprim-sulfamethoxazole (Table 1).

Five days before the index admission (i.e., approximately 7 weeks postbiopsy and 5 weeks postorchiectomy), the patient developed fever, chills, anorexia, nausea, emesis, low back pain, and severe headache. Upon presentation and admission, he was in distress from the headache and mild photophobia but otherwise appeared fairly well. Vital signs, physical examination, and screening blood tests were normal. Urine was cloudy, with ≥ 100 white blood cells (WBC), 11 to 20 red blood cells (RBC), and many bacteria per high-power field. Blood and urine cultures were collected. Intravenous levofloxacin, 750 mg daily, was begun.

On day 2, lumbar puncture yielded cerebrospinal fluid (CSF) containing 1,000 leukocytes (100% neutrophils) and 2,000 erythrocytes per microliter, 140 mg/dl protein, 9 mg/dl glucose, and Gram-negative bacilli on Gram stain. Admission blood cultures grew Gram-negative bacilli.

On day 3, the CSF and urine cultures also yielded Gram-negative

bacilli. Therapy was changed from levofloxacin to ceftriaxone, 2 mg intravenously daily, which was continued for 14 days. The blood, urine, and CSF isolates all proved to be *E. coli*, with the same susceptibility pattern and MICs as the orchiectomy isolate from 6 weeks previously (Table 1). The patient recovered uneventfully.

Context. Infectious complications following TRUSP biopsy are increasingly frequent, due largely to the widespread emergence of fluoroquinolone-resistant *Escherichia coli* (FQREC) strains (3, 14). Such strains may be selected for by any previous fluoroquinolone therapy, e.g., as used here, and by the fluoroquinolone prophylaxis that is conventionally used prior to TRUSP biopsy (3, 14). Presentations of post-TRUSP biopsy infection include urosepsis, prostatitis, and, rarely, meningitis, which can be fatal (12).

The main contributor to the recent upsurge in post-TRUSP biopsy infections and FQREC is *E. coli* sequence type ST131, a clonal group within *E. coli* phylogenetic group B2 that has emerged globally over the past decade as a disseminated cause of multidrug-resistant extraintestinal infections (14). Although best known for its association with extended-spectrum β -lactamases (ESBLs), especially CTX-M-15, in many regions, ST131 is encountered more frequently as an ESBL-negative but fluoroquinolone-resistant pathogen, accounting for two-thirds or more of FQREC clinical isolates (8).

ST131 has been documented to cause distinctive, severe infections in specific host groups, including pediatric septic arthritis and osteomyelitis (7), pyomyositis in neutropenic patients (13), emphysematous pyelonephritis and renal abscesses in diabetic patients (5), and neonatal meningitis (2). To date, however, neither has ST131 been reported to cause meningitis in an adult nor have the several reported post-TRUSP biopsy *E. coli* meningitis isolates (as reviewed in reference 12) been characterized beyond their antimicrobial susceptibility profile.

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TABLE 1 Susceptibility profile of the patient's four *Escherichia coli* clinical isolates^a

Agent	MIC ($\mu\text{g/ml}$)	Interpretation
Amikacin	4	Susceptible
Ampicillin	≥ 32	Resistant
Ampicillin-sulbactam	16	Intermediate
Aztreonam	≤ 1	Susceptible
Cefepime	≤ 1	Susceptible
Ceftazidime	≤ 1	Susceptible
Ceftriaxone	≤ 1	Susceptible
Ciprofloxacin	≥ 4	Resistant
Ertapenem	≤ 0.5	Susceptible
Gentamicin	≥ 16	Resistant
Imipenem	≤ 1	Susceptible
Levofloxacin	≥ 8	Resistant
Piperacillin-tazobactam	≤ 4	Susceptible
Tobramycin	8	Intermediate
Trimethoprim-sulfamethoxazole	≤ 20	Susceptible

^a Isolates included (i) the initial orchiectomy isolate and, 6 weeks later, the (ii) urine, (iii) blood, and (iv) cerebrospinal fluid isolates. All isolates gave identical susceptibility testing results.

Given the prominence of ST131 in post-TRUSP biopsy infections, ST131's association with multidrug resistance, and the multidrug-resistant nature of the reported post-TRUSP biopsy *E. coli* meningitis isolates, such isolates might be predicted to be ST131. The case patient, who developed post-TRUSP biopsy *E. coli* meningitis, allowed us to test this hypothesis and to identify ways to better prevent and manage this problem.

Laboratory analysis. Major *E. coli* phylogenetic group (A, B1, B2, D), presence of 62 virulence genes associated with extraintestinal pathogenic *E. coli*, and O and H types were defined by using established multiplex PCR-based assays (8). ST131 status was defined based on ST131-specific single nucleotide polymorphisms in *gyrB* and *mdh* and 7-locus multilocus sequence typing (8). Presence of genes encoding group 1 CTX-M ESBLs and CTX-M-15 was determined by PCR (10). Antimicrobial susceptibility testing was by Vitek II (in the clinical laboratory). XbaI pulsed-field gel electrophoresis (PFGE) was done according to a standardized protocol (9). Using BioNumerics (Applied Maths), the PFGE profile of the patient's isolate was compared to a large private PFGE

library (J.R.J.) within which isolates were assigned to a particular pulsotype if $\geq 94\%$ similar to that pulsotype's index isolate (9).

Findings. The patient's CSF isolate derived from phylogenetic group B2 and represented *E. coli* ST131. It exhibited the O25b *rfb* (O lipopolysaccharide) variant and the H4 *fliC* (flagellin) variant. Its virulence gene profile, which was typical of ST131 isolates generally (8), included the F10 *papA* allele (P fimbriae structural subunit) without *papCEFG*, *iha* (adhesin-siderophore receptor), *fimH* (type 1 fimbriae), *sat* (secreted autotransporter toxin), *fyuA* (yersiniabactin system), *iutA* (aerobactin system), *kpsM* II (group 2 capsule synthesis), *kfiC* (K5 capsular antigen), *traT* (serum resistance associated), *usp* (uropathogenic-specific protein), *ompT* (outer membrane protease), and *malX* (pathogenicity island marker). Its PFGE profile placed it within pulsotype 968 (Fig. 1), the most prevalent ST131-associated pulsotype in the private PFGE library (9), which is globally distributed and is associated with severe and distinctive infections (5, 7, 9, 11, 13). The strain's 10 nearest neighbors in the PFGE library were from multiple different locales, from humans and animals, and from diverse specimen types (Fig. 1), reflecting the broad geographical, host, and clinical range of this ST131-derived clonal lineage (9). No CTX-M group 1 or CTX-M-15 *bla* gene was detected in the patient's isolate, consistent with its susceptibility to extended-spectrum cephalosporins. Likewise, all but two of the nearest-neighbor comparator pulsotype 968 isolates were ESBL negative and lacked the corresponding genes (Fig. 1).

Comment. The case patient experienced a series of unfortunate but avoidable events, including a questionably indicated 20-day course of ciprofloxacin therapy and TRUSP biopsy (both done because of a rising screening PSA level), subsequent necrotizing epididymo-orchitis due to FQREC, ineffective empirical therapy for this, subsequent orchiectomy, ineffective postoperative therapy, subsequent meningitis due to FQREC ST131, and delayed effective therapy for this. This illustrates how the widespread emergence of a new multidrug-resistant and virulent *E. coli* clonal group, ST131 (8), can interact with established practice patterns regarding antimicrobial therapy (e.g., routine use of fluoroquinolones regardless of recent and current culture results and the patient's antimicrobial history) and delivery of preventive ser-

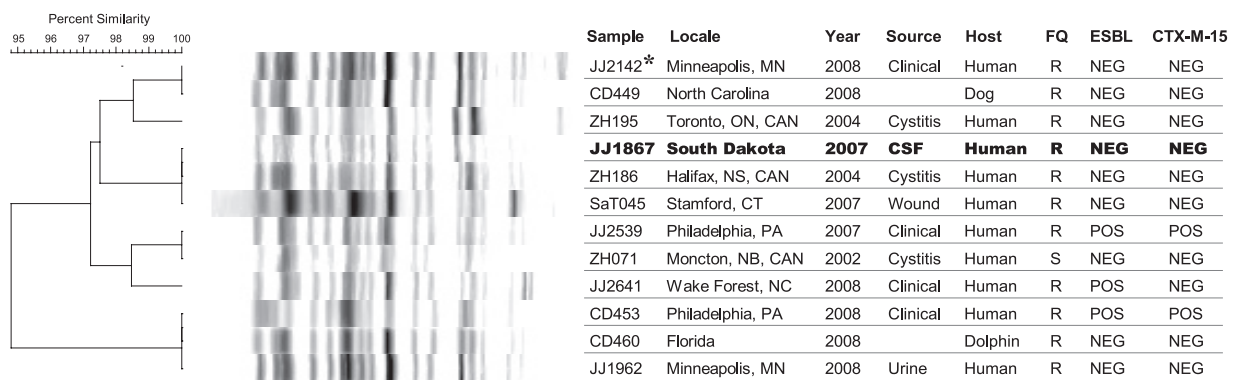


FIG 1 Pulsed-field gel electrophoresis profiles of the patient's *Escherichia coli* meningitis isolate and closely related strains from sequence type ST131. Abbreviations: FQ, fluoroquinolone phenotype; R, resistant; S, susceptible; ESBL, extended-spectrum β -lactamase; NEG, negative; POS, positive. Shown are the patient's cerebrospinal fluid isolate (JJ1867; boldface), the index isolate for pulsotype 968 (JJ2142; asterisk), and the 10 next most similar isolates (relative to the patient's isolate) in a large private pulsotype library (J.R.J.). All isolates represent pulsotype 968. Note the broad geographic range of the isolates (which spans multiple widely distributed locales in the United States and Canada), the diversity of sources (blank = unknown), and the broad host range (human, dog, and dolphin).

vices (e.g., routine PSA screening and TRUSP biopsy), resulting in highly morbid iatrogenic complications (3, 14).

E. coli ST131 caused this case patient's post-TRUSP biopsy meningitis and, presumably (based on chronology and antibiogram), also his epididymo-orchitis, bacteremia, and bacteriuria. ST131 also was the single most common cause of *E. coli* bacteremia post-TRUSP biopsy in a recent study from Australia (14), with pulsotype 968 predominating (J. R. Johnson, unpublished data). Although these observations might be interpreted as indicating enhanced virulence for ST131, and especially pulsotype 968, in the post-TRUSP biopsy context, other factors may be operative. For example, ST131 is also the predominant *E. coli* lineage among FQREC fecal surveillance isolates from men about to undergo prostate biopsy, where pulsotype 968 also predominates (Johnson, unpublished), suggesting that ST131's predominance among postbiopsy urinary tract infection (UTI) isolates may simply reflect its high prevalence within the fecal reservoir, the presumed source for post-TRUSP biopsy infections (4), rather than enhanced pathogenicity. Additionally, our patient's progression to extreme clinical manifestations may represent the natural history of ineffectively treated *E. coli* genitourinary infection, with ineffective treatment in this instance resulting from repeated reliance on agents to which the organism exhibited *in vitro* resistance. This is a predictable consequence of unrecognized resistance to the chosen antimicrobial regimen, particularly if a patient's recent antimicrobial exposure history and susceptibility data are not factored into drug selection.

This case represents a cautionary tale regarding key management principles that should be incorporated into routine clinical practice by providers, with support from the clinical microbiology laboratory, to avoid future such mishaps. First, in most locales, fluoroquinolone monotherapy is no longer acceptable for prophylaxis prior to TRUSP biopsy (14). Substitution or addition of a more broadly active empirical agent (selected based on current local cumulative susceptibility data) (1, 6, 15), or culture-guided therapy (4), is mandatory. Second, a patient who presents with suspected infection after receiving fluoroquinolone therapy for either prophylaxis or treatment should not receive empirical monotherapy with a fluoroquinolone or another agent (e.g., gentamicin) to which coselected resistance is likely. Third, discovery of *in vitro* resistance to an empirically selected antimicrobial regimen should prompt, at minimum, reassessment of the patient's clinical status, if not regimen change. Finally, according to the best available evidence (as assessed by the U.S. Preventive Services Task Force, <http://www.uspreventiveservicestaskforce.org/prostatecancerscreening.htm>), routine PSA screening should be avoided, which would avert many unnecessary TRUSP biopsies, with their attendant infectious and noninfectious risks.

Novel aspects of this report warrant mention. To our knowledge, it describes the first case of post-TRUSP biopsy meningitis from the United States, the first case of adult meningitis or epididymo-orchitis due to ST131, and the first molecular characterization of a post-TRUSP *E. coli* meningitis isolate. Limitations include the unavailability of the orchiectomy, blood, and urine isolates for molecular analysis and the inherent uncertainty regarding causal relationships.

In summary, this patient developed post-TRUSP biopsy epididymo-orchitis (leading to orchiectomy), bacteremia, bacteriuria, and Gram-negative meningitis, all presumably due to the same FQREC ST131 strain, in part because of overreliance on

fluoroquinolones for prophylaxis and empirical therapy. The widespread prevalence of ST131 in the general population, its ability to cause serious disease, and its typically multidrug resistance profile oblige rethinking of many standard approaches to antimicrobial prophylaxis and therapy, especially involving the genitourinary tract. Close coordination between clinicians and the clinical microbiology laboratory is essential to optimal management.

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