Emerging drug resistance in *Salmonella* Typhi and *S. Paratyphi* is a substantial public health concern. We report what appears to be the first case and isolation of multidrug resistant *S. Paratyphi* A carrying CTX-M-15-type extended-spectrum beta-lactamase from a Japanese traveller returning from India.

Here, we report the isolation of multidrug resistant *S. Paratyphi* A producing CTX-M-15-type extended-spectrum beta-lactamase (ESBL) from a traveller returning from India. To our knowledge, this is the first report of *S. Paratyphi* A with CTX-M-15-type ESBL isolated from a human. Enteric fever, including typhoid fever caused by *S. enterica* serotype Typhi (*S. Typhi*) and paratyphoid fever caused by *S. enterica* serotype Paratyphi A (*S. Paratyphi* A), is one of the most important febrile illnesses in tropical and subtropical countries, with high rates of morbidity and mortality [1]. In industrialised countries, enteric fever is a common cause of fever in returned travellers [2]. The emergence of drug-resistance in *S. Typhi* and *S. Paratyphi* A is an emerging public health problem. Owing to the recent increase of fluoroquinolone resistance, third-generation cephalosporins, such as ceftriaxone or cefotaxime, have become the primary drugs for treatment of enteric fever [3].

No further details on the treatment regimen in India were available. The patient returned to Japan on day 4 of illness owing to sustained fever, diarrhoea, and anorexia. During her travel from India to Japan (days 5–7), the patient took oral combined ofloxacin-cefixime. On day 7, she was admitted to our hospital with diarrhoea and anorexia but without fever. On examination, deep palpation of the lower abdominal area caused mild discomfort, otherwise, physical examination was normal. She did not have any underlying illness, and was not on any regular medication. Eight months prior to the admission at the hospital in India she had been vaccinated against *Salmonella* Typhi (Vi polysaccharide).

**Laboratory analyses**

On admission, stool and blood samples were taken for culture. Antimicrobial susceptibility test was performed by broth microdilution in accordance with the Clinical and Laboratory Standards Institutes [4].

The patient’s stool was screened on admission for meticillin-resistant *Staphylococcus aureus*, vancomycin-resistant Enterococci, and Gram-negative rods resistant to one or multiple agents in the extended-spectrum cephalosporin class and/or that demonstrated elevated minimum inhibitory concentrations (MICs) (≥1 mg/L) to imipenem and/or meropenem. Screening culture of stool was positive only for drug-sensitive *Escherichia coli* and *Enterobacteriaceae* spp.

Results from blood cultures were positive for *S. Paratyphi* A two days after admission and *S. Paratyphi* A was found to be resistant to cefotaxime (MIC = 64 mg/L) and ceftazidime (MIC > 16 mg/L). Addition of clavulanic acid to each cephalosporin lowered the MIC to 0.25 mg/L and 1 mg/L, respectively. This isolate was also resistant to nalidixic acid, and categorised as...
isolate.

quinolone resistance genes were not detected in this described elsewhere [6-9]. However plasmid-mediated qnrC, qnrS quinolone resistance genes including qnrA, qnrB, for 2 www.eurosurveillance.org

tial antibiotic treatment was changed on day 9 from the final susceptibility results were available, the ini-
toms resolved after eight days of admission. After one (2 g per day for 8 days), and the patient’s symp-
treatment and outcome

Patient treatment and outcome

Upon admission, we treated the patient with ceftriax-
one (2 g per day for 8 days), and the patient’s symp-
toxinemia (orally, 500 mg per day), which was administered for seven days. A stool culture obtained on admission was negative for S. Paratyphi A. Stool cultures repeated before changing antibiotic treatment as well as eight, nine, and 10 days after completing azithromycin treatment also remained negative for S. Paratyphi A. The patient was discharged on 10 days after admission.

Discussion and conclusion

Although ESBL-producing organisms are emerging among Enterobacteriaceae, ESBL-producing S. Typhi or S. Paratyphi A have been reported rarely in Nepal and the Middle East [10,11]; CTX-M-15 type ESBL-producing S. Typhi were previously reported in patients in Middle Eastern countries [11,12].

bla_{CTX-M-15} genes have been identified worldwide in Enterobacteriaceae; further, isolation of these genes from Escherichia coli has been reported to be increasing [13]. The acquisition of bla_{CTX-M-15} as a transferable plasmid from another enteric bacterium has previously been reported in S. Typhi [5].

Of particular concern are reports of increasing isolation rates of S. Paratyphi A from India and other parts of Asia [14], which might be attributed to use of protective vaccines (Vi polysaccharide and live oral Ty21a) effective against S. Typhi, but not S. Paratyphi A [14].

The number of reports on ESBL in S. Typhi and S. Paratyphi A is still limited, and thus, extended-spectrum cephalosporins would still be a reasonable empiric treatment for a suspected case of enteric fever in the current situation. The patient improved on day 8 before antibiotics were changed. The oral treatment by a fluoroquinolone (ofloxacin) prior to the admission might have contributed to her clinical improvement. Other possible explanations include clinical recovery by natural history of S. Paratyphi A, or infection of two different antimicrobial patterns of S. Paratyphi A, i.e. ESBL-S. Paratyphi A and non-ESBL-S. Paratyphi A, with non-ESBL mainly contributing to clinical symptoms.

Potential increase in the plasmid-mediated spread of ESBL in S. Paratyphi A in the future would pose a threat to public health. Judicious use of antibiotics to avoid unnecessary selective pressure on intestinal bacterial flora, and careful microbiological analysis of patients with typhoid fever, especially those returning from the Middle East or south Asia are approaches critical for prevention of the potential spread of multidrug resistant S. Paratyphi A.

Acknowledgements

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### Table

Minimum inhibitory concentrations of various antibiotics on Salmonella enterica serotype Paratyphi A isolated from a Japanese traveller returning from India, Japan, July 2013

<table>
<thead>
<tr>
<th>Antibiotics</th>
<th>MIC (mg/L), interpretation result</th>
</tr>
</thead>
<tbody>
<tr>
<td>Ampicillin</td>
<td>&gt;16, R</td>
</tr>
<tr>
<td>Amoxicillin/clavulanic acid</td>
<td>=16, I</td>
</tr>
<tr>
<td>Cefotaxime</td>
<td>&gt;64, R</td>
</tr>
<tr>
<td>Cefotaxime/clavulanic acid</td>
<td>=0.25, S</td>
</tr>
<tr>
<td>Ceftazidime</td>
<td>&gt;16, R</td>
</tr>
<tr>
<td>Ceftazidime/clavulanic acid</td>
<td>=1, S</td>
</tr>
<tr>
<td>Ceftriaxone</td>
<td>&gt;16, R</td>
</tr>
<tr>
<td>Nalidixic acid</td>
<td>&gt;16, R</td>
</tr>
<tr>
<td>Ciprofloxacin</td>
<td>=0.5, I</td>
</tr>
<tr>
<td>Levofloxacin</td>
<td>=1, I</td>
</tr>
<tr>
<td>Azithromycin</td>
<td>=16', I</td>
</tr>
<tr>
<td>Aztreonam</td>
<td>&gt;16, R</td>
</tr>
<tr>
<td>Trimethoprim/sulfamethoxazole</td>
<td>=2, R</td>
</tr>
<tr>
<td>Chloramphenicol</td>
<td>=4, S</td>
</tr>
<tr>
<td>Tetracycline</td>
<td>=2, S</td>
</tr>
</tbody>
</table>

MIC: Minimum inhibitory concentration; R: resistant; S: susceptible; I: intermediate.

* The MIC breakpoint for azithromycin is not provided by the Clinical and Laboratory Standards Institutes. We thus refer to the breakpoints table by the European Committee on Antimicrobial Susceptibility Testing, which mentions that azithromycin has been used in the treatment of infections with Salmonella Typhi (MIC <16 mg/L for wild type isolates) [15].

intermediate to ciprofloxacin (Table). Due to the atypical susceptibility pattern and the requirement for the verification susceptibility test process, it took longer than one week for the final susceptibility result to be available.

The presence of the gene conferring the ESBL phenotype was verified by polymerase chain reaction (PCR) as described elsewhere [5]. Sequencing of the PCR products revealed that S. Paratyphi A was positive for bla_{CTX-M-15}'. The presence of the plasmid-mediated quinolone resistance genes including qnrA, qnrB, qnrC, qnrS, and aac(6')-Ib was investigated using PCR described elsewhere [6-9]. However plasmid-mediated quinolone resistance genes were not detected in this isolate.
Conflict of interest

None declared.

Authors’ contributions

M. Mawatari collected the data and drafted the manuscript; YK and KH participated in the coordination and concept of the manuscript and edited the manuscript and helped with the draft of the manuscript; M. Morita, MO and HI performed the molecular tests; TK, YF, S. Kutsuna and NT collected the data and participated in the concept of the manuscript; S. Kanagawa and NO revised the article for intellectual content. All authors read and critically revised the first as well as the subsequent and final drafts of this manuscript.

References