Epidemiologic application of pulsed-field gel electrophoresis to an outbreak of *Campylobacter jejuni* in an Austrian youth centre

A. LEHNER^{1*}, C. SCHNECK¹, G. FEIERL², P. PLESS³, A. DEUTZ³, E. BRANDL¹ AND M. WAGNER¹

(Accepted 5 April 2000)

SUMMARY

We report the first documented *Campylobacter jejuni* outbreak in an Austrian youth centre. Sixty-four children were involved of which 38 showed classical signs of campylobacter gastroenteritis. Since unpasteurized milk distributed by a local dairy was suspected to be the source of infection, stool samples were collected from 20 cows providing the milk. Five of the cows tested positive for *C. jejuni*. These isolates together with 37 clinical samples were compared by pulsed-field-gel electrophoresis (PFGE). The PFGE patterns, using the restriction endonucleases *SmaI* and *SaII*, were identical for the human and bovine isolates. This finding confirmed that the outbreak was caused by the consumption of unpasteurized milk contaminated with *C. jejuni*.

INTRODUCTION

Campylobacter jejuni and C. coli are well-recognized causes of human gastroenteritis. Accurate identification of the sources of infections is especially difficult as the organism is very common in nature and most cases of campylobacter infections in humans occur sporadically. The National Campylobacter Surveillance Centre has determined the incidence of reported cases of campylobacter gastroenteritis in the USA as approximately 5-6 cases per 100000 population. However, if estimates of the number of milder and/or unreported infections are correct, and considering the difficulties in cultivation and identification of the bacteria from faeces, the true incidence may be 200 times or more the nationally reported rate, or 1 % of the population [1]. In Austria, the incidence of campylobacter infections, based on laboratory confirmed cases was 68.8/100000 inhabitants as determined in a national survey in the province Styria

During the last decade, traditional methods of strain differentiation, namely serological and phage typing [6–8], have been supplemented with molecular methods, such as plasmid fingerprinting, [9] ribotyping [10], PCR-based methods [11] and pulsed-field gel electrophoresis (PFGE) [12, 13]. PFGE has been shown to be a highly discriminative method for many species and in *Campylobacter* spp. several genotypes can be found within a serotype. The aim of the present study was to apply PFGE to trace-back the source of *C. jejuni* in an outbreak of gastroenteritis in an Austrian youth centre in September 1998.

¹ Institute of Milk Hygiene, Milk Technology and Food Science, University of Veterinary Medicine, Veterinärplatz 1, A-1210 Vienna, Austria

² Institute of Hygiene, University Graz, Universitätsplatz 1, A-8010 Graz, Austria

³ Veterinary Administration, Styrian Government, Zimmerplatzgasse, 15, A-8010 Graz, Austria

^{(1.2} million inhabitants) in 1998 [2], making it the second most common bacterial pathogen after *Salmonella* spp. causing diarrhoea. Contaminated drinking water, consumption of unpasteurized milk and poultry have been shown to be risk factors [3–5]. However, there have been no reports so far in Austria, where unpasteurized milk was involved in transmission of the pathogen.

^{*} Author for correspondence.

MATERIALS AND METHODS

Stool samples

The *C. jejuni* strains from human (n = 37) and bovine (n = 5) faeces were cultured on selective medium (mCCDA; Oxoid, UK) at 42 °C under microaerobic conditions for 48 h. Species identity was confirmed by Gram stain, cytochrome oxidase production, catalase activity and hydrolysis of hippurate.

Typing of isolates by PFGE

Isolates were cultured on 5% sheep blood agar plates (Oxoid, UK) in a Gas-Pak jar at 42 °C for 48 h under microaerobic conditions produced with a gas generator kit (GENbox microaer, bioMérieux, France). Bacterial cells were harvested from plates with a swab into 2 ml phosphate buffered saline, pH 7.3 (PBS; 130 mm NaCl, 10 mm NaH₂PO₄). The optical density of the suspension was adjusted to 1.2 at 600 nm [14] and 1.5 ml was centrifuged and the pellet resuspended in 300 μ l PBS. The cells were incubated with 33 μ l of 37% formaldehyde solution (Merck, BRD) for 1 h at room temperature to inhibit DNAse activity [15], and subsequently washed three times with PBS. $300 \mu l$ aliquots of each isolate were embedded in 1 % agarose (pulsed-field certified, Bio-Rad Laboratories, USA) and the agarose plugs were incubated in 3 ml ESP lysis buffer (1% sodium dodecyl sulphate, 0.25 m EDTA. pH 8.0, 0.5 mg/ml proteinase K) at 55 °C for 48 h [16]. Digestion of the chromosomal DNA using restriction enzymes SmaI and SalI (New England Biolabs, USA) was performed according to the manufacturer's instructions. The DNA fragments were separated in 1 % agarose gels using a CHEF DR-III apparatus under the following conditions: SmaI: 5-40 s for 20 h at 6 V/cm, 14 °C; SalI: 10-30 s for 20 h at 6 V/cm, 14 °C. The gels were stained with ethidium bromide and the profiles were photographed under UV-light.

RESULTS AND DISCUSSION

In autumn 1998, an outbreak of gastroenteritis occurred in a youth centre. Of the 64 persons attending the centre (aged 10–18 years), 38 showed classical signs of campylobacter infection such as stomach ache (100%), fever $(72\cdot2\%)$, headache (63%), sickness $(54\cdot5\%)$, nausea $(22\cdot7\%)$ and rheumatism $(18\cdot2\%)$. In most cases stools were highly liquid $(54\cdot5\%)$ or soft liquid $(36\cdot4\%)$. Blood and mucous was present in

19·1% and 23·8% respectively. The average duration of the diarrhoea as well as that of the illness was 7 days. Twenty-eight of 38 patients with diarrhoea, 8 children without clinical signs of infection and 1 healthy member of the staff were tested positive for *C. jejuni*.

Evaluation of the questionnaires regarding the possible source of infection revealed two possible risk factors: poultry meat, and unpasteurized milk together with the breakfast cereal. Sixty-three (98%) of the children had consumed poultry meat, but only 32 (50%) had unpasteurized milk. Twenty-three (72%) of the milk consumers were ill, whereas 9 (2%) stayed healthy. In the group of children that had not consumed raw milk 15 (47%) were ill and 17 (53%) were healthy. The poultry meat served 2 days before onset of the outbreak was not available for testing but milk samples from the local dairy, that distributed the milk directly to the youth centre, were collected by the Austrian Food Surveillance authority. C. jejuni was not detected in any of them. This was not surprising as cross contamination from faeces to raw milk, as a result of inadequate hygiene measures was suspected as the route of infection. Therefore, rectal swabs of all cows on the dairy farm (n = 20) were taken and C. jejuni was recovered from five of these.

By PFGE, 37 human and 5 bovine isolates showed identical macrorestriction patterns using enzymes SmaI (Fig. 1) and SalI. Additionally, ten isolates sent to the Institute of Hygiene in Hamburg (Professor Aleksic) were typed as serotype 062. It was therefore concluded that the consumption of contaminated milk provided by the local dairy, caused the outbreak in the centre. However, there were clearly other routes of infection as some of the children developed illness without having consumed the milk in the centre. In these cases human-to-human transmission probably occurred, suggesting that the infective dose of organisms was low as the hygiene standards of the centre were sufficient. Moreover, a member of staff who cleaned the toilets contracted the infection but had no direct contact with the affected children or contaminated milk. This suggests possible person-toperson spread of campylobacter infection resulting from contact with contaminated human faeces.

The value of PFGE as a discriminant method of DNA fingerprinting isolates for epidemiological studies has been commented on by other writers [13, 16]. It has also been shown previously, that *SmaI* defined macrorestriction types of *C. jejuni* can be further differentiated by the use of a second enzyme such

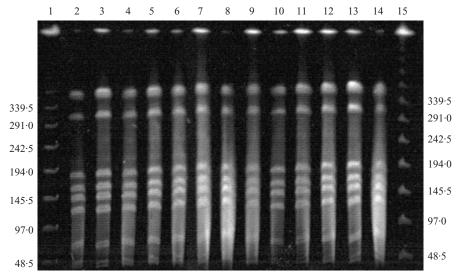


Fig. 1. PFGE patterns of DNA from 13 human (lanes 2–14) and 5 bovine (15–19) *C. jejuni* isolates digested with *Sma*I. Lanes 1 and 20: molecular weight standard in kb (lambda concatemers).

as Sall [17]. It is noteworthy that all isolates showed the same susceptibility pattern to seven antimicrobial compounds (data not shown) and this corroborates the finding of a single DNA pattern for all isolates examined.

During the evaluation of a parallel study comparing the PFGE profiles of poultry and human campylobacter isolates, four additional clinical isolates, drawn from the same region in Austria, from persons originally not connected to the youth centre could be linked to this outbreak strain. Further questioning of these individuals revealed that all of them had had contact with the farm, either as consumers or visitors. This finding underlines the importance of establishing national DNA fingerprint databases based on PFGE.

We have documented the first outbreak of *C. jejuni* attributed to unpasteurized milk in Austria. Apart from other sources and routes of contamination the handling and sale of unpasteurized milk should be considered a potential risk for gastroenteritis and clear guidelines on hygienic measures should be given to minimize the risk of infection. Because many of the reported campylobacter infection cases are among children, individuals involved in youth activities must be alert to the danger of consumption of raw milk.

ACKNOWLEDGEMENT

This project was financially supported by the Veterinary Administration, Styrian Government, Zimmerplatzgasse, 15, A-8010 Graz, Austria, grant number: VV-293 L2/113-98 ad.

REFERENCES

- 1. Tauxe RV. Epidemiology of *Campylobacter jejuni* infections in the United States and other industrialized nations. In: Nachamkin I, Blaser MJ, Tompkins LS, eds. *Campylobacter jejuni*, current status and future trends. Washington: ASM Press, 1992: 9–19.
- 2. Feierl G, Sixl B, Berghold C, et al. Epidemiologie in der Steiermark. Mitteilungen der Sanitätsverwaltung, 1999; **6**: 3–7.
- Kapperud GE, Skjerve LH, Bean NH, Ostroff SM, Lassen J. Risk factors for sporadic *Campylobacter* infections: results of a case-control study in southeastern Norway. J Clin Microbiol 1996; 30: 3117–21.
- 4. Beutling D. Vorkommen und Überleben von *Campylobacter* spp. in Lebensmittel. Arch Lebensmittelhyg 1998; **49**: 1–24.
- 5. Altekruse SF, Stern NJ, Fields PI et al. *Campylobacter jejuni* an emerging foodborne pathogen. Emerg Infect Dis 1999; **5**: 28–35.
- Lior H. New, extended biotyping scheme for Campylobacter jejuni, Campylobacter coli and 'Campylobacter laridis'. J Clin Microbiol 1984; 20: 636–40.
- Penner JL, Henessy JN, Congi RV. Serotyping of Campylobacter jejuni and Campylobacter coli on the basis of thermostable antigens. Eur J Clin Microbiol 1983; 2: 378–83.
- 8. Patton CM, Wachsmut IJ. Typing schemes: are the current methods useful? In: Nachamkin I, Blaser MJ, Tompkins L, eds. *Campylobacter jejuni*, current status and future trends. Washington: American Society for Microbiology Press, 1992: 110–28.
- Tenover FC. Plasmid fingerprinting. A tool for bacterial strain identification and surveillance of nosocomial and community-acquired infections. Clin Lab Med 1985; 5: 413–36.
- 10. Stull TL, LiPuma JJ, Eding TD. A broad spectrum

- probe for molecular epidemiology of bacteria: ribosomal RNA. J Infect Dis 1988; **157**: 280–6.
- 11. van Belkum A. DNA fingerprinting of medically important microorganisms by use of PCR. J Clin Microbiol 1994; **31**: 2697–701.
- 12. Finney M. Pulsed-field gel electrophoresis. In: Ausubel FM, Brent R, Kingston RE, et al, eds. Current protocols in molecular biology, Vol 1. New York: Greene-Wiley, 1993:2.5.9–2.5.14.
- 13. Maslow JN, Slutsky AM, Arbeit RD. Application of pulsed-field gel electrophoresis to molecular epidemiology. In: Persing DH, Smith T, Tenover FC, White TJ, eds. Diagnostic molecular microbiology: principles and applications. Washington, D.C.: American Society for Microbiology Press, 1993: 563–72.
- 14. Greimel SE. Campylobacter spp.: Isolation and mol-

- ecular characterization (dissertation). Karl-Franzens University, Graz, Austria, 1997.
- Gibson JR, Sutherland K, Owen RJ. Inhibition of DNAse activity in PFGE analysis of DNA from Campylobacter jejuni. Lett Appl Microbiol 1994; 19: 357–8.
- Morooka T, Umeda A, Fujita M, et al. Epidemiologic application of pulsed-field gel electrophoresis to an outbreak of *Campylobacter fetus* meningitis in a neonatal intensive care unit. Scan J Infect Dis 1996; 28: 269–70.
- 17. On SLW, Nielsen EM, Engberg J, Madsen M. Validity of *Sma*I-defined genotypes of *Campylobacter jejuni* examined by *Sal*I, *Kpn*I, and *Bam*HI polymorphisms: evidence of identical clones infecting humans, poultry, and cattle. Epidemiol Infect 1998; **120**: 231–7.