Cheese-related listeriosis outbreak, Portugal, March 2009 to February 2012

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In Portugal, listeriosis has been notifiable since April 2014, but there is no active surveillance programme for the disease. A retrospective study involving 25 national hospitals led to the detection of an outbreak that occurred between March 2009 and February 2012. The amount of time between the start of the outbreak and its detection was 16 months. Of the 30 cases of listeriosis reported, 27 were in the Lisbon and Vale do Tejo region. Two cases were maternal/neonatal infections and one resulted in fetal loss. The mean age of the non-maternal/neonatal cases was 59 years (standard deviation: 17); 13 cases were more than 65 years-old. The case fatality rate was 36.7%. All cases were caused by molecular serogroup IVb isolates indistinguishable by pulsed-field gel electrophoresis and ribotype profiles. Collaborative investigations with the national health and food safety authorities identified cheese as the probable source of infection, traced to a processing plant. The magnitude of this outbreak, the first reported food-borne listeriosis outbreak in Portugal, highlights the importance of having an effective listeriosis surveillance system in place for early detection and resolution of outbreaks, as well as the need for a process for the prompt submission of *Listeria monocytogenes* isolates for routine laboratory typing.

Introduction

Listeria monocytogenes is an intracellular bacterial pathogen of humans and a variety of animal species. In humans, *L. monocytogenes* infections are mainly food-borne and can cause an invasive and often fatal disease in pregnant women and their fetuses, newborns, elderly people and immunocompromised individuals, with a case fatality rate of up to 30% [1]. The incidence of listeriosis increased in several European countries between 2009 and 2013 (such as Germany, the Netherlands, Spain and the United Kingdom [1,2]) and, was the most frequent cause of hospitalisation and death (15.6%) due to the consumption of contaminated food in Europe in 2013 [2]. This increase reinforces the

need for each country to establish enhanced molecular surveillance of listeriosis for efficient outbreak detection, investigation and control, as carried out by PulseNet USA or the Centre National de Référence des *Listeria*, Institut Pasteur, Paris, for example [3,4]. A similar programme for listeriosis surveillance at European Union level by harmonising methodological variables such as case definition, laboratory procedures and reporting systems is crucial. A pilot project was conducted by the European Centre for Disease Prevention and Control (ECDC) between January and March 2013 aiming to evaluate a *Listeria* external quality assurance scheme for the typing of *L. monocytogenes* that covered pulsed-field gel electrophoresis (PFGE) method and serological typing (both as a phenotypic and a multiplex polymerase chain reaction (PCR)-based method) [5]. Results demonstrated that the majority (59%) of the participating laboratories were able to produce a PFGE gel of sufficiently high quality and the average score for serotyping among the participants was 94% and 97% for traditional and multiplex PCR based methods, respectively; however, higher quality could be achieved through trouble-shooting assistance and training.

In the absence of an active surveillance system for listeriosis at a national level, a collaborative study between the *Listeria* Research Centre of Escola Superior de Biotecnologia (LRCEB) and 25 of the major national hospitals (on a voluntary basis), covering about 90% of the population, was established in 2003 with the aim of obtaining epidemiological data on human listeriosis cases in Portugal and characterising clinical isolates of *L. monocytogenes* both phenotypically and genetically. In 2003, the incidence of listeriosis was 0.14 cases per 100,000 population [6]. An increase was reported between 2003 and 2007, i.e. it was 0.23 cases per 100,000 inhabitants for the year 2007 [7]. As a result of this study, an increase in the number of listeriosis cases was detected between January and July 2010, particularly in the Lisbon and Vale do Tejo region

that corresponds to 13% of the total area of mainland Portugal and 34% of the total population (3.6 million inhabitants) [8], representing the first detected outbreak of listeriosis in Portugal. Here we describe the outbreak, as well as give details of the investigations carried out in order to determine the source of infection.

Methods

Case definition

A listeriosis case was defined as a non-maternal/neonatal (non-MN) patient who met the laboratory criteria or a mother with a laboratory-confirmed listeriosis infection in her fetus, stillborn or newborn, as described in the Commission Decision of 28/IV/2008 [9]. Cases (laboratory confirmed with unknown clinical criteria) were detected through voluntary reporting by hospitals to the LRCESB in Porto.

If the pathogen was isolated from a pregnant woman and her newborn, stillborn or fetus, this was counted as a single case. Information regarding the sex and age of the patient, underlying pathology (if present), the tissue or fluid from which the bacteria were isolated and the year of isolation was reported.

Culture collection

Hospitals sent isolates of *L. monocytogenes* to LRCESB for species confirmation and typing. Species confirmation was performed by carbohydrate fermentation (rhamnose, xylose and mannitol) and Christie Atkins Munch-Petersen (CAMP) test [10]. Confirmed isolates of *L. monocytogenes* were stored in tryptic soy broth with 30% (v/v) glycerol at -80°C in the culture collection of the LRCESB.

Molecular-serotyping

Molecular serotype of *L. monocytogenes* isolates was determined by multiplex PCR according to Doumith et al. [11]. This assay differentiates five major subtypes, each representing more than one serotype: geno-serogroup IVb (serotypes 4b, 4d and 4e), geno-serogroup IIa (serotypes 1/2a and 3a), geno-serogroup IIb (serotypes 1/2b, 3b and 7), geno-serogroup IIc (serotypes 1/2c and 3c) and geno-serogroup IVa (serotypes 4a and 4c).

Pulsed-field gel electrophoresis

PFGE typing was performed according to the standard CDC PulseNet protocol [12] using the restriction enzymes *Ascl* and *Apal* and gel run in CHEF III DR System (Bio-Rad, Laboratories, Hercules, CA, United States). *Salmonella enterica* serovar Braenderup H9812 (ATCC) DNA digested with *XbaI* was used as a reference size standard. Cluster analysis of the PFGE types was performed with the GelCompar software (Applied Maths, Sint-Martens-Latem, Belgium) by the unweighted pair group method with average linkages (UPGMA), using the Dice coefficient, and visually validated.

Ribotyping

Automated ribotyping was performed using the restriction enzyme *EcoRI* and the RiboPrinter microbial characterisation system (Qualicon Inc., Wilmington, DE, United States), as previously described [13,14].

Outbreak investigation

The outbreak was investigated by the national health (Direção Geral de Saúde and Administração Regional de Saúde de Lisboa e Vale do Tejo) and food safety (Autoridade de Segurança Alimentar e Económica) authorities in collaboration with LRCESB.

A standardised questionnaire (adapted from a Canadian listeriosis outbreak, kindly supplied by Dr Jeff Farber of the Public Health Agency of Canada) was administered by the national health authority to patients diagnosed with listeriosis or their families (face-to-face interview) concerning their diet histories in the two months before symptom onset, with reference to the type of food consumed and household shopping patterns.

FIGURE 1

Distribution of human listeriosis cases in Portugal with the outbreak *Listeria monocytogenes* strain, March 2009–February 2012 (n = 30)

Map adapted from www.d-maps.com/m/europa/portugal/portugal19.pdf.

FIGURE 2Human listeriosis cases with the 2010 *Listeria monocytogenes* outbreak strain, Portugal, March 2009–February 2012 (n = 30)

All except three cases were in the Lisbon and Vale do Tejo region (two were in Centre region, one case occurred in the Algarve region). Information on the number of listeriosis cases caused by *Listeria monocytogenes* strains with non-outbreak pulsed-field gel electrophoresis types for each year is available from Magalhães et al. [15].

Analysis of food products and environmental samples was conducted by the food safety authority. *L. monocytogenes* isolates from food and environmental samples were sent to LRCESB for typing.

International enquiry

To determine if the outbreak-associated strain of *L. monocytogenes* had been recovered from clinical or food samples from other countries, the PFGE type was communicated and compared with those of *L. monocytogenes* isolates in databases in France (Centre National de Référence des Listeria, Institut Pasteur), Canada (Listeriosis Reference Centre, Health Canada) and United States (Food Microbe Tracker, Food Safety Laboratory, Cornell University).

Results

Recognition of the outbreak

Between January and July 2010, a high number of listeriosis cases was observed (40 cases compared with 20 cases observed during all of 2009) [15], particularly in the Lisbon and Vale do Tejo region, where the majority of the cases were reported. Molecular typing of the 40 *L. monocytogenes* clinical isolates revealed that 18 serotype IVb isolates presented the same PFGE type and ribotype, which had been observed for five isolates recovered in 2009, four of which were in the Lisbon and Vale do Tejo region (in March, April and September) and one in the Centre region (in July) (Figures 1 and 2). This PFGE type was not found in the databases searched.

In July 2010, the national health and food safety authorities were alerted to the increased number of cases and an outbreak investigation was initiated. A public health alert was issued to national hospitals requesting prompt notification and reporting of cases.

LRCESB continued to receive clinical isolates for typing. Continued monitoring detected two more cases with the outbreak strain in November 2010, and three more cases in January, February and March 2011 (two in the Lisbon and Vale do Tejo region and one in the Algarve). Thereafter, in February 2012, there were two new cases with the same strain in the Lisbon and Vale do Tejo region. The total number of outbreak cases between March 2009 and February 2012 was 30.

Trace-back and investigation of the food source

Analysis of the epidemiological questionnaires pointed to different types and sizes of food retailers and identified the following as possible sources of infection: cheeses (cured cheese and queijo fresco, made from pasteurised cow and goat milk), ice cream, ham and fermented sausages. On the basis of data gathered concerning the type of establishments where the food products were purchased, as well as the geographical location of the cases, suspected foods and foods commonly associated with listeriosis, the food safety authority inspected 42 food retailers and collected 103 samples for analysis (51 meat products, 24 dairy products, 13 ready-to-eat foods and 15 environmental swabs). *L. monocytogenes* was detected in four samples collected at a retailer: three from queijo fresco and one from a swab taken from a ham slicing-machine; one queijo fresco sample contained counts of *L. monocytogenes* greater than 100 colony-forming units/g.

PFGE typing revealed that isolates recovered from two queijo fresco samples of different brands from the same retailer showed the same PFGE type as the clinical isolates with the outbreak strain. Further investigation of the processing plants where these cheeses had been produced involved collecting and testing environmental and cheese samples. The outbreak strain was

FIGURE 3

Pulsed-field gel electrophoresis type of the *Listeria monocytogenes* outbreak strain from 2010 in Portugal, using *ApaI* and *AscI* restriction enzymes

A: *L. monocytogenes* isolate from a listeriosis case from a hospital in the Lisbon and Vale do Tejo region (February 2010); B: *L. monocytogenes* isolate from a cheese sample collected at a retailer selected on the basis of the results of the epidemiological questionnaires (October 2010); C: *L. monocytogenes* isolate from a contaminated cheese sample collected at a processing plant located in the Alentejo region and identified by trace-back investigations (March 2011); S: *Salmonella* Braenderup size control.

detected in *L. monocytogenes* isolates from cheese samples from one of the two processing plants investigated (located in the Alentejo region) (Figure 3). Thus, cheeses produced by this plant were considered the probable source of the outbreak; cross-contamination between products or contamination from the environment, or both, may have occurred at retail level, as both suspected brands of queijo fresco were sold in the same market. As a result of these findings, the food safety authority recalled both products and more samples from the processing plant were analysed. Cheeses made with pasteurised cow and goat milk collected at the processing plant tested positive for *L. monocytogenes* and the collected isolates had the same PFGE pattern as the outbreak strain. Subsequently, in March 2011 the processing plant voluntarily suspended its activities during 15 days. After appropriate cleaning and disinfection measures, intensified product and environmental sampling was carried out. No positive samples were detected and products were allowed to be sold in the marketplace. Samples were then collected monthly by the food safety authority and no further positive samples have been detected.

Characteristics of listeriosis outbreak-associated cases

Of the 30 cases, two were MN cases, both of which occurred in 2010 (Table).

One MN case resulted in stillbirth and the other MN case involved a newborn with unknown outcome. For the 28 non-MN cases, isolates were collected from blood (n=16), cerebrospinal fluid (n=10) and from both blood and cerebrospinal fluid (n=2). The mean age of the 27 non-MN cases with a reported age was 58.9 years (standard deviation: 17); the median was 64 years (range: 15–83); 13 non-MN cases were older than 65 years. The ratio of male:female non-MN cases was 22:6. Information was available for 20 non-MN patients with underlying conditions (e.g. diabetes mellitus, cancer, hepatitis, human immunodeficiency syndrome (HIV) infection/acquired immunodeficiency syndrome

(AIDS)). For seven non-MN cases, no such information was available. The absence of known predisposing condition was reported for one 15 year-old patient. The overall case fatality rate, for MN and non-MN cases, was 37% (11/30).

Discussion

As there is no active surveillance programme for listeriosis in Portugal, outbreak detection is extremely difficult. The incubation period of the infection can be very long, up to 70 days, which makes it difficult to find a link between cases [1]. Detection of the outbreak

TABLE

Listeriosis outbreak-associated cases, Portugal, March 2009–February 2012 (n = 30)

Data	Number of cases				
	2009	2010	2011	2012	Totals
Clinical form					
Non-MN	5	18	3	2	28
MN	0	2	0	0	2
Sex					
Female	2	5	1	0	8
Male	3	15	2	2	22
Age of non-MN cases (years)					
<65	3	8	2	1	14
≥65	2	10	1	0	13
Age unknown	0	0	0	1	1
Clinical sample of non-MN cases					
Blood	4	10	1	1	16
CSF	1	6	2	1	10
Blood and CSF	0	2	0	0	2
Fatal outcome					
Non-MN	3	7	NA	NA	10
MN	NA	1	NA	NA	1

CSF: cerebrospinal fluid; MN: maternal/neonatal cases; NA: not available; non-MN: non-maternal/neonatal cases.

reported here was due mainly to retrospective investigations. The amount of time between the presumed onset of the outbreak (March 2009) and its recognition was extremely long (16 months). This long delay amplified the magnitude of the outbreak, leading to a high number of cases ($n=30$) and a high case-fatality rate (36.7%). Underestimation of the number of cases is likely, as many cases usually go unreported and unrecognised, since patient data and strains are voluntarily reported and listeriosis is an infrequent disease [2], with clinical symptoms that are difficult to identify [16].

Typing of clinical and food isolates of *L. monocytogenes* by molecular techniques, such as PFGE, was essential for the identification of cheese of a specific brand as being the most probable source of contamination. Although a cheese from another producer was contaminated at retail by a strain with the outbreak-associated PFGE type, this was probably a result of cross-contamination since no positive samples were detected in the processing plant. Increased risk of cross-contamination of ready-to-eat foods by *L. monocytogenes* in a retail environment has been demonstrated in several studies [17-20]. For example, a quantitative risk assessment conducted by Endrikat et al. suggested that ready-to-eat deli meats sliced at a retailer are five times more likely to cause listeriosis than pre-packaged products (per annum basis) [21].

Additional information is needed for a better understanding of the risk factors and for the development of improved strategies for controlling *L. monocytogenes* in these environments.

The long duration of this outbreak (March 2009 to February 2012) is noteworthy and reinforces the importance of setting up an effective multidisciplinary team able to help ensure rapid notification of cases and the prompt submission of *L. monocytogenes* isolates for routine laboratory typing.

Of the 28 non-MN cases, 13 were 65 years of age or older and at least 20 cases presented an underlying condition. In European countries with established surveillance programmes, such as France, Germany and the United Kingdom, the incidence of listeriosis is reported to be increasing and the distribution of cases is shifting, primarily affecting elderly persons and those with predisposing medical conditions, leading to a high case fatality rate [1,2]. This is of concern as life expectancy increases, including for those who are immunocompromised (e.g. those with AIDS, under immunosuppressive therapy for cancer) [22]. In addition, food habits are changing worldwide, with an increasing demand for processed ready-to-eat foods [23]. Therefore, it is likely that there will be an increased risk of food-borne listeriosis.

Data gathered from the surveillance of human disease and also from all stages in the food production chain should be continuously collected and analysed

to understand the ecology of *L. monocytogenes* and its routes of transmission. This will be crucial for developing enhanced strategies to control this organism and contribute to a decrease in the incidence of food-borne listeriosis in Portugal.

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Conflict of interest

None declared.

Authors' contributions

Maria Manuela Sousa was responsible for the collection of epidemiological data and for the link between hospitals and Centro de Biotecnologia e Química Fina (CBQF). Autoridade de Segurança Alimentar e Económica (ASAE) team was in charge of collection and analysis of food products and environmental samples. CBQF team was responsible for collecting clinical isolates supplied by the hospitals and food isolates supplied by ASAE. CBQF and Listeriosis Reference Centre for Canada teams were responsible for characterisation of the isolates. All authors participated in the analysis and interpretation of data. Rui Magalhães, Gonçalo Almeida, Vânia Ferreira and Paula Teixeira drafted the manuscript. Jeffrey Farber and Franco Pagotto critically reviewed the draft manuscript and provided substantive input. All authors approved the final version.

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