Surveillance and outbreak reports

SURVEILLANCE OF LISTERIOSIS IN NAVARRE, SPAIN, 1995-2005 – EPIDEMIOLOGICAL PATTERNS AND CHARACTERISATION OF CLINICAL AND FOOD ISOLATES

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We monitored the incidence of human listeriosis in Navarre, a region in north of Spain between 1995 and 2005, and carried out the characterisation of *Listeria monocytogenes* isolates obtained from clinical samples and ready-to-eat products (sliced cooked meat, smoked salmon and liver pate). The active surveillance requesting hospitals to notify all listeriosis cases (n=40) yielded higher incidence rates (average annual rate 0.65/100,000 inhabitants, range 0.18-1.18/100,000 inhabitants) than expected. Pregnant women were the largest group affected (n=13, 32.5% of the cases), with a peak in incidence during the last three years of the study period. From the 40 human cases we obtained 33 *Listeria* isolates. Serological and molecular characterisation by PFGE identified 20 different pulsotypes, which on three occasions enabled us to link sporadic cases into clusters. Although we could not identify the incriminated food product we found two clinical pulsotypes among smoked salmon and cooked meat isolates. Surveillance of listeriosis in Spain should be improved and coordinated with other European Union Member States in order to better estimate the burden of disease and to prevent foodborne outbreaks.

Introduction

Listeria monocytogenes has been recognised as a serious foodborne pathogen, with a case-fatality rate between 20% and 50% [1-3]. However, the important impact that this disease has on public health is not always recognised, particularly since listeriosis is a relatively rare disease compared with other common foodborne infections such as salmonellosis. Listeriosis is likely to be under-reported due to its status as a non-notifiable disease in many countries, including Spain, and because of the absence of adequate surveillance programs. The Community Summary Report on Trends and Sources of Zoonoses, Zoonotic Agents, Antimicrobial resistance and Foodborne outbreaks in the European Union produced jointly by the European Food Safety Authority (EFSA) [4] reported an incidence between 0 and 10 cases per million in 2006 among the 24 countries that submitted listeriosis information. It is significant that incidence rates above 5 cases per million were reported in countries where listeriosis is statutorily notifiable, such as France, Germany, Finland and Switzerland. However, even in countries with obligatory notification of listeriosis and efficient surveillance systems, such as PulseNet in the United States [5], the number of cases could be greater than reported due to the occurrence of

many sporadic cases and spontaneous miscarriages which are not investigated.

As pointed out by Kiss et al. and MacKenzie et al. [6,7], integrated food chain surveillance is necessary for all national and international authorities in order to achieve adequate information regarding the true impact of listeriosis in the population. Serotyping of *L. monocytogenes* has a low discriminatory power for subtype differentiation [8], however, when combined with molecular methods based on DNA macrorestriction pattern analysis [9] it becomes a useful tool in epidemiological investigation. Among the molecular typing methods, pulsed-field gel electrophoresis (PFGE) has been one of the most frequently used in epidemiological investigation of listeriosis because of its excellent discriminatory power and reproducibility [10-15].

In Spain, the current surveillance of listeriosis is based on voluntary reporting of cases to the Microbiological Information System of the National Reference Laboratory. The present study set out to evaluate the incidence of listeriosis in Navarre, a region in northern Spain, over a period of 11 years from 1995 to 2005, by implementing active surveillance in this geographical area. The objectives of this study were:

- 1) to obtain epidemiological data on cases of listeriosis reported by the three main hospitals of Navarre;
- 2) to compare *L. monocytogenes* isolates recovered from food products and human cases of listeriosis in Navarre over the same period of time, by serological and PFGE characterisation.

Methods

Listeriosis surveillance and epidemiological data

In order to determine the incidence of listeriosis, we asked the three main hospitals in Navarre to report cases of listeriosis. The case definition was based on the isolation of *L. monocytogenes* from a hospitalised patient with a clinically compatible illness. A case was considered perinatal in the following cases: infected pregnant woman, miscarriage, stillbirth or newborn less than one month old. When the pathogen was isolated from both the pregnant woman and her newborn child, this was considered to represent a single case. Information regarding sex, age, clinical symptoms, immunosuppressive treatment or underlying disease, and death or recovery of the patients, was reported when available.

In addition, patients diagnosed in 2005 were interviewed about their consumption habits with regard to high risk foodstuffs. The questionnaire (available upon request from the corresponding author) covered different aspects relating to the consumption of ready-to-eat (RTE) products (type, brand and store where purchased) during the two months preceding disease onset. Specific questions regarding high-risk RTE products sampled in the study were also included.

Collection of *L. monocytogenes* strains

A total of 87 *L. monocytogenes* isolates were obtained from food samples in a study which we performed in 2003-2005 [16]. Of these, 45 were isolated from a market sampling pool of 783 RTE high-risk food products that included sliced cooked meat products (pork, chicken and turkey), sliced smoked fish products (salmon and trout) and liver pate. Isolation and identification of *L. monocytogenes* was carried out using aseptic techniques following the NF EN ISO 11290-1 [17]. The remaining 42 food strains were obtained as a result of our earlier study on the occurrence of *L. monocytogenes* in the same type of RTE food products carried out in Navarre between 1995 and 2002 [18].

With respect to clinical strains, from the 40 human cases of listeriosis reported between 1995 and 2005, we were able to obtain only 33 isolates. They were isolated in the hospital microbiology laboratories and most of them originated from either blood or cerebral spinal fluid or placenta, while stool cultures for *Listeria* were not available. These isolates were subsequently submitted to our laboratory at the University of Navarre, where identification of strains was carried out by biochemical and serological methods.

All strains were stored at -80°C in sterilised skimmed milk.

Serological characterisation

Serotyping was carried out using commercial specific antisera (Denka Seiken Co., Ltd., Tokyo, Japan) following the manufacturer's instructions. Both polyclonal anti-O antisera (O-I/II, O-V/VI, O-I, O-II, O-VI, O-VII, O-VIII, O-IX) and anti-H (H-A, H-AB, H-C, H-D) were used in the determination of somatic and flagellar antigen, respectively. Interpretation of the results was carried out according to the serotyping scheme established by Seeliger and Höhne [19].

Molecular characterisation by pulsed-field gel electrophoresis (PFGE)

PFGE was performed according to Graves and Swaminathan [8], with minor modifications. Before performing PFGE, strains were revitalised by plating onto blood agar (Biomérieux, Marcy L'Etoile, France) and incubated at 37°C for 18 h. DNA from a single *Listeria* colony was digested with *Apal* (Roche Diagnostics, Barcelona, Spain) and separated at 6 V/cm for 19.5 h on a CHEF-DR II PFGE apparatus (Bio-Rad, Hercules, California, US) with switch time from 4 to 40 seconds at 14°C. *Staphylococcus aureus* ATCC 29213 was used as a control for digestion. The obtained images were digitised and analysed using Gel Compar II® software (Applied Maths, Kortrijk, Belgium). Restriction patterns were analysed using the criteria described by Tenover et al. [20]. Similarity values of the patterns were calculated using the Dice correlation coefficient with a 1.0% band position tolerance and unweighted pair group method using arithmetic average (UPGMA). Clinical and food isolates were compared, and pulsotypes were numbered consecutively.

Statistical analysis

The statistical package used was SPSS v13.0. The contingency table analysis was based on the chi square distribution (Pearson's chi square test).

Results

Clinical and epidemiological data

A total of 40 cases of listeriosis were documented in Navarre during the 11-year surveillance study period (Figure 1). Between 1995 and 2005, the mean annual incidence was 0.65/100,000 inhabitants, ranging from 0.18/100,000 in 1998 (n=1) to 1.18/100,000 in 2005 (n=7). Table 1 shows the epidemiological data of reported cases. From the available information, 26 cases (65%) correspond to non-perinatal infections, while perinatal infections (pregnant women and newborns) were described in 13 cases.

In total, 21 deaths were reported resulting in the average case fatality rate of 52.5%. These included eight foetal deaths in

FIGURE 1
Cases of listeriosis reported in Navarre, Spain, from 1995 to 2005 (n=40)

TABLE 1
Principal characteristics of listeriosis cases reported in Navarre, Spain, from 1995 to 2005 (n = 40)

Epidemiological data ^a	Number of cases (% of the total)	Number of deaths	Case-fatality rate (%) by group
Clinical form			
Perinatal	13 (32.5)	8 ^b	61.5 ^b
Non-perinatal	26 (65.0)	13	50.0
Risk factor (non-perinatal)			
No detectable pathology	5 (12.5)	1	20.0
Known risk ^c	15 (37.5)	8	53.3
Other conditions ^d	6 (15.0)	4	66.7
Age group (in years)			
<1	1	1	100
1-19	2 (5.0)	1	50.0
20-39	12 (32.5)	0	0
40-59	8 (20.0)	2	25.0
>60	12 (30.0)	8	66.7

^a Information not available in all cases

^b Only foetal death (all women recovered)

^c Aged >60 or immunocompromised patients (cancer, HIV, organ transplantation).

^d Chronic diseases

perinatal cases (all women recovered) and 13 deaths among the non-perinatal cases (case-fatality rate of 50%). Among this latter group, 15 patients (37.5% of the total) had an underlying listeriosis risk factor defined as age >60 years and/or immunosuppressive conditions such as cancer, HIV or organ transplantation. Further six cases had underlying chronic conditions, such as diabetes (n=1), addiction to alcohol (n=2) and other (n=3). The remaining five non-perinatal cases (12.5% of the total) were healthy people aged between 2 and 59 years.

Clinical symptoms most frequently reported were septicaemia (37.5%) and meningitis (15.0%) (Table 2). No cases of acute gastroenteritis caused by listeriosis were detected in the course of the study.

The most affected group at risk of listeriosis was pregnant women (n=13; two of them with underlying diseases). The number

of perinatal cases was significantly higher in the years 2003-2005 compared with the previous period of 1995-2002 (10 vs. 3) (Figure 2). The clinical information available showed that all of the infected mothers recovered (one of them was diagnosed with meningitis). However, 61.5% of all pregnancy-associated cases resulted in miscarriage (n=5), stillbirth (n=2) or infant death within 24-48 hours of birth (n=1) (Table 2).

Serotyping results

Four serovars were determined among the 33 clinical isolates as shown in Table 3. In all of the different risk groups most of the isolates belonged to serogroup 4 (78.8%). Serotype 4b was the predominant (75.8%, n=25), followed by serotype 1/2a (18.2%, n = 6). With respect to the food isolates, the predominance was for serogroup 1 (77.0%). In contrast with the results obtained from clinical strains, serotype 1/2a was the most common (51.7%, n=45), followed by serotype 4b (23.0%, n=20), serotype 1/2c

TABLE 2

Clinical symptoms of listeriosis patients reported in Navarre, Spain, 1995-2005 (n=40)

Clinical symptoms ^a	Number of cases	Proportion of total (%)
Septicaemia ^b	15	37.5
Meningitis ^c	6	15.0
Miscarriage	5	12.5
Stillbirth	2	5.0
Premature	1	2.5
Endocarditis	2	5.0
Other ^d	5	12.5
Unkown	5	12.5

^a Multiple responses were possible

^b One case also with encephalitis

^c Classified as meningitis (n = 4), meningoencephalitis (n = 1) and cerebral abscess (n = 1)

^d Hormonal disorders, arthritis, osteomyelitis, kidney failure, heart attack

FIGURE 2

Incidence of listeriosis in Navarre, Spain in 1995-2005 and pregnancy-associated cases per 1,000 births in the same time period

TABLE 3

Serotype distribution of *Listeria monocytogenes* isolated from ready-to-eat (RTE) food and clinical cases of listeriosis in Navarre, Spain, during the period 1995-2005

Source	Proportion of serogroup 1 (%)	Proportion of serogroup 4 (%)	Number of isolates of each serotype					Total number of isolates
			1/2a	1/2b	1/2c	4b	3b	
Clinical isolates	21.2	78.8	6	1	- ^a	25	1	33
Non condition associated with risk of listeriosis	20.0	80.0	1	-	-	4	-	5
Pregnancy	20.0	80.0	2	-	-	8	-	10
Transplantation	28.6	71.4	1	1	-	4	1	7
Cirrhosis/Alcoholism	0	100	-	-	-	4	-	4
Cancer	33.3	66.7	1	-	-	2	-	3
Others	25.0	75.0	1	-	-	3	-	4
RTE foods	77.0	23.0	45	3	19	20	-	87
Sliced cooked meat	94.6	5.4	32	3	18	3	-	56
Sliced smoked fish	43.3	56.7	13	-	-	17	-	30
Pate	100 ^b	0	-	-	1	-	-	1

^a No clinical isolates of this serotype detected

^b Only one isolate obtained

(21.8%, n=19) and finally by serotype 1/2b (3.5%, n=3). When food categories were examined according to serotype, we found that serotype 4b was the predominant in smoked fish (56.7%, n=17), while serotype 1/2a was the most frequent among sliced meat products (57.1%, n=32). The unique strain isolated in liver pate belonged to serotype 1/2c.

PFGE results

PFGE revealed a total of 20 different pulsotypes among clinical isolates, distinguished by one or more band differences ranging in size from 50 to 500 kb (Figure 3). Pulsotypes 1, 5, 8 and 9 contained two or more strains which remained indistinguishable from each other. While strains of pulsotype 5 (n=3) were recovered from different years, strains of pulsotype 1 (n=10) and pulsotype 9 (n=2) were related in time and geographical distribution, showing that possible outbreaks could have occurred. Among isolates with pulsotype 1, three strains corresponded to listeriosis cases diagnosed between November and December 2003, and four were isolated from four pregnant women affected between November and December 2005. However, the oral interviews about food intake in these patients did not give us information about a possible common food source. With respect to the 87 food isolates, we found 19 different pulsotypes (data not shown), but only two of them were similar to the previous clinical pulsotypes characterised. We found a cluster of seven strains isolated from smoked salmon showing pulsotype 1. Two of these were isolated in October 2003, 1-2 months before the isolation of three clinical strains with identical

pulsotype. In addition, pulsotype 16 was shared by a clinical strain and seven food isolates (three isolates from sliced meat and four from smoked salmon).

Discussion

A collaborative surveillance study of *Listeria* infections in Europe led by the French Institut de Veille Sanitaire and Institut Pasteur [21] described an incidence of 0.15 cases per 100,000 inhabitants in the year 2000 for Spain (based on information from voluntary reporting), whilst for that same year the incidence detected for Navarre in our study was 0.91 (five cases reported in our active surveillance). Similarly, for 2004, the EFSA report [22] described an incidence of 0.20 in Spain whereas the rate obtained in our study was 0.51 (three cases). Considering that the likelihood of listeriosis in Navarre was similar to the whole of Spain (similar proportion of people at risk and food intake habits), we think that the active surveillance carried out in our study was the reason for the higher incidence obtained in Navarre compared to that reported for the whole of Spain in the mentioned publications. Similar incidence rates to those we found in Navarre were described in countries with mandatory notification of listeriosis, such as Denmark, where Gerner-Smidt et al. [23] reported incidences ranging from 0.45 to 0.75 cases per 100,000 between 1994 and 2003, and increasing to 0.8. during 2004 [22]. It is interesting to note that the sensitivity of the Danish system for *Listeria* infection is thought to be almost 100% and is based on immediate notification to the National Surveillance Centre (Staten Serum Institute) of all patients from whom *Listeria* has been isolated. This corresponds to the methods used in our study. Likewise, active surveillance of *L. monocytogenes* infections in the Netherlands revealed an increase in incidence from 0.26 cases per 100,000 inhabitants in 2001 to 0.62 per 100,000 in 2005 [24].

It should be noted that the 52% case-fatality rate (foetal death included) obtained in our study is similar to that described in other Spanish reports [25] but higher than the average reported over recent years in the EFSA publication [4]. However, the EFSA report admits that the lower than expected reported case-fatality rate might be due to a lack of data on patient outcomes after the initial notification, which indicates the importance of clinical data recovery to assess the impact of listeriosis.

With respect to listeriosis associated with pregnancy, several authors have reported high incidences with case-fatality rates above 45% [6,26]. The high numbers of listeriosis cases we detected among this group in Navarre is due, in part, to the fact that we included investigation of tissue samples obtained from hospitalised women after miscarriage. However, it should be noted that testing for *L. monocytogenes* was performed in a similar way during the entire study period, without a systematic analysis of all miscarriage tissues. So, the increase of the number of pregnancy-associated cases in recent years can not be considered a surveillance artefact. Taking into account that bacterial cultures are not routinely obtained from spontaneously aborted foetus or stillborn neonates in a wide range of Spanish hospitals, we believe that the true incidence of listeriosis in this risk group may still be underestimated. In order to assess the rates more accurately, we recommend the routine investigation of *L. monocytogenes* in tissue after miscarriage and stillbirth in the whole of Spain and also at European level.

To assess the real impact of listeriosis in Spain and in the whole of EU, better harmonisation of data collection systems at national

FIGURE 3
Dendrogram for *Listeria monocytogenes* pulsotypes of all 33 isolates obtained from clinical cases and some isolates from food samples, Navarre, Spain, 1995-2005

level is required. Validated clinical and food questionnaires would be valuable in all diagnosed or suspected cases of listeriosis, providing more precise and complete information about symptoms, outcomes and food consumption habits of affected people. This way we would be able to conduct epidemiological studies (useful for outbreak investigations) and to provide dietary advice to high-risk individuals in avoiding specific foods. In our study, suspecting a possible outbreak in December 2005 after four cases had been detected in a short period of time, we decided to interview the patients using a food questionnaire but this oral survey failed to identify a possible common food source. Nevertheless, this does not exclude the possibility of a common source but rather reflects the limitations of this preliminary survey and the need for validation. Active surveillance in Italy [27] which involved distributing clinical and food questionnaires to the hospitals and the characterisation of all strains resulted in a higher number of cases of listeriosis than reported by mandatory notifications.

In addition to the accurate recovery of epidemiological information, there is a need to isolate and characterise *L. monocytogenes* clinical strains. Although most cases of listeriosis occur sporadically [28], serological and molecular analysis could help us to relate sporadic cases that are geographically and time-related, allowing the detection of possible outbreaks that perhaps go unnoticed if few people are involved. Serological characterisation is useful for rapid screening of strains during suspected outbreaks. In line with the findings of several authors [6,11,15,25,29] our results confirm the fact that most cases of human listeriosis are caused by serotypes 4b (75.8%). However, recent studies observed a variation of this classical distribution with an increase of serotype 1/2a [27,30].

The combination of serology and PFGE has provided us with the opportunity to link sporadic cases on three different occasions during the period 2002-2005. It should be noted that pulsotype 1 was the most frequent among the clinical strains isolated in the study (10 out of 33), and also one of the predominant profiles in the food isolates (seven isolates from smoked salmon). The profile of three patient isolates in November-December 2003 (pulsotype 1) was indistinguishable from that of two isolates obtained from smoked salmon a month before, but we had no available information about the food intake of these patients as the food questionnaire was not carried out at that time. The repeated isolation of pulsotype 1 in smoked salmon (2003-2004), and pulsotype 16 in smoked salmon and sliced cooked meat (1995-1998) suggests the persistence of specific subtypes in food processing plants. Considering that food products at risk of containing *Listeria* are often commercially available over a wide area, characterisation of food and clinical strains should be managed at a national level in order to trace probable sources of infection and to detect related cases occurring in other regions of Spain.

In conclusion, the present study shows that harmonised and active surveillance of listeriosis is needed in Spain in order to increase knowledge about real impact of this serious health problem. Accurate national surveillance should be based on the obligatory notification of listeriosis, the collection of epidemiological information by the application of a standardised food and clinical questionnaire and the sending of isolated strains to a reference laboratory for a serological and molecular characterisation. This active surveillance could be extended at European level, improving the available information to detect compatible cases and to trace probable sources of infection.

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