Research articles

HIGH RATES OF COMMUNITY-ACQUIRED, PANTON-VALENTINE LEUKOCIDIN (PVL)- POSITIVE METHICILLIN-RESISTANT S. AUREUS (MRSA) INFECTIONS IN ADULT OUTPATIENTS IN GREECE

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Staphylococcus aureus was isolated in 88 (30. 8%) of 286 adult patients suffering from various skin and soft-tissue infections examined in the outpatient department of a 650 bed tertiary-care hospital of Athens, Greece between January 2006 and December 2007. Twenty-seven (30.7%) of the S. aureus infections were caused by methicillin-resistant S. aureus (MRSA). All MRSA isolates were also resistant to tetracycline, fucidic acid and kanamycin, but were sensitive to gentamicin and tobramycin, as well as to to cotrimoxazole, chloramphenicol, guinolones, clindamycin and erythromycin. All isolates belonged to staphylococcal cassette chromosome mec elements (SCCmec) type IV, and were found to carry the *luk*F-PV and *luk*S genes coding for Panton-Valentine leukocidin (PVL). Pulsed-field gel electrophoresis (PFGE) and spatyping revealed high genetic similarity among all MRSA isolates and with the PFGE pattern of the well-described ST80 clone that seems to be spreading through Europe. The high prevalence of MRSA among S. aureus infections in the community signify that empiric therapy in Greece, when clinically indicated, should exclude β-lactam antibiotics. Moreover, the establishment of an active screening for PVL-positive community-acquired (CA)-MRSA carriage and the adoption of a search and destroy strategy for CA-MRSA in all patients admitted with purulent skin and soft-tissue is of high priority in Greece as well as in all European countries which face high rates of CA-MRSA infection.

Introduction

Methicillin-resistant *Staphylococcus aureus* (MRSA) is a wellrecognised major cause of healthcare-associated infections. Over the past 10 years the epidemiology of this pathogen has changed throughout the world and infections caused by it have emerged in the community [1,2]. First reports of MRSA infections in the community were described predominantly in children without established risk factors for MRSA acquisition and were defined as community-acquired MRSA (CA-MRSA) [3]. Further infections have been reported among selected populations, including sports teams and correctional facility inmates. Moreover, infections in outpatients, mainly healthy, non-immunocompromised adults without risk factors have also been documented [1,2]. CA-MRSA isolates primarily cause skin and soft-tissue infections but serious, life-threatening, invasive infections such as bacteraemia and necrotizing pneumonia have also been described [4]. In Greece, 20 to 40 % of all *S. aureus* skin and soft-tissue infections in paediatric outpatients are found to be due to CA-MRSA [5,6]. In the presented study we investigated the prevalence of Panton-Valentine leucidine (PVL)-positive CA-MRSA among *S. aureus* infections in adult outpatients in a tertiary-care hospital of Athens.

Materials and methods

Between January 2006 and December 2007, 286 patients suffering from various skin and soft-tissue infections and with no history of hospitalization or any contact with a hospital during the past twelve months were examined in the outpatient department of a 650 bed tertiary-care hospital of Athens.

Laboratory testing

Microbiological examination of the respective clinical specimens and the identification of the species were performed by standard procedures. Resistance to oxacillin in *S. aureus* was assessed by the disc diffusion method, through cefoxitin resistance, according to the Clinical and Laboratory Standards Institute (CLSI) criteria [7]. The same criteria were used to determine resistance levels to other antibiotics (tetracycline, kanamycin, tobramycin, gentamicin, fucidic acid, chloramphenicol, erythromycin, ciprofloxacin and cotrimoxazol).

Molecular testing

Staphylococcal cassette chromosome elements (SCC*mec*) typing as well as detection of the *mec*A gene was performed by PCR, as described by Oliveira and de Lencastre [8]. The *luk*F-PV and *luk*S-PV genes coding for the PVL toxin, were detected by PCR, as described by Lina et al. [9]. To determine the genetic relatedness of the isolates, Smal restriction fragments of genomic DNA were separated by PFGE as described previously [10] and analysed by BioNumerics software, version 4.6 (Applied Maths, Sint-Martens-Latem, Belgium), using Dice coefficients and the unweighted-pair group method by means of average linkages. *Spa*-typing was performed as described by Harmsen et al. [11] and *spa* types were determined using Ridom StaphType software version 1.4 (Ridom GmbH, Würzburg, Germany).

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Results

S. aureus was isolated from 88 (30.8%) of 286 patients presenting with skin infections without history of hospitalisation or any contact with a hospital during the last year. Upon sensitivity testing the infection was found to be caused by MRSA in 27 (30.7%). Fourteen of the affected were men and 13 women. The mean age of these patients was 43 years, ranging from 29 to 56 years. Abscesses (skin abscesses 7, soft-tissue abscesses 9) dominated the clinical presentations, followed by furuncles (6), wound infections (4) and folliculitis (1) (Table). No statistically significant difference was found between the rates of methicillinsensitive *S. aureus* (MSSA) and MRSA isolated from the various types of skin and soft-tissue infections. Moreover, there was no difference in age or sex between patients suffering from MSSA or MRSA infections (data not shown).

All MRSA isolates were resistant to tetracycline, fucidic acid and kanamycin, but they were sensitive to tobramycin and gentamicin, as well as to cotrimoxazole, chloramphenicol, quinolones, clindamycin and erythromycin.

All isolates belonged to SCC *mec* type IV and carried the *luk*F-PV and *luk*S-PV genes. PFGE revealed high genetic similarity among all MRSA strains (Table). The PFGE patterns of 18 isolates were identical and shared 100% similarity with the PFGE pattern of the well-described ST80 clone that seems to be spreading through Europe [12,13]. The remaining nine isolates revealed differences in one to three bands and were allocated into four subpatterns, comprising 5, 2, 1 and 1 isolates respectively. *Spa*-typing of the 27 strains, allocated 26 into *spa* type t044, a type closely associated with the ST80 clone and one to *spa* type t131 which is also associated with the ST80 clone (Table).

Discussion

MRSA has become a significant cause of community-acquired skin and soft-tissue infections in many parts of the world [12,13]. The worldwide spread of PVL-positive CA-MRSA clones that were initially described at the beginning of this decade to be continent specific [12] has been documented [12]. Furthermore, new lineages of PVL-positive CA-MRSA strains have also been detected [13].

It is well recognised that the high prevalence of MRSA among *S. aureus* skin and soft-tissue infections observed in the USA, is due to the spread of a single clone that can be identified on the basis of PFGE and other genotyping characteristics. This clone, a result of recent clonal expansion and diversification of a subset of isolates [14] is designated as the USA300 clone by the United States Centres for Disease Control and Prevention (CDC) in Atlanta. It belongs to MLST (ST8) and *spa* type (t008) which are different from the ones described in this study [15,16]

In Europe although CA-MRSA skin and soft-tissue infections have been reported from most countries, the prevalence of infections due to CA-MRSA appear to vary across the continent [17-27]. However, reports of prevalence rates of MRSA among *S. aureus* infections are, to the best of our knowledge, lacking. The currently prevailing genetic type among CA-MRSA in Europe is the PVL-positive, t044/ ST80-SCC*mec* type IV [12]. In a recent Danish study, travel to or residing in countries abroad, especially in the Mediterranean region, the Balkans (Serbia) and the Middle East, where there is a high prevalence of CA-MRSA infections caused by t044/ST80-SCC*mec* type IV, have been associated with infections with this type [17]. Moreover, in some European countries strains with USA300 genotype are starting to be isolated with increasing frequency: The emergence of clones that are related to the USA300 has been associated with increasing rates of CA-MRSA in Spain. These clones were primarily isolated from immigrants from South America [25]. Further increasing isolation rates of the USA300 clone have been reported in Germany [26].

Contrary to the high degree of molecular diversity among CA-MRSA that has been shown in various parts of Europe [17, 24, 27], our study documented high genetic relatedness among the PVL-positive CA-MRSA isolates, which might indicate a successful and rapid spread of this clone in Greece. The study has some limitations since it focuses on patients presenting at the outpatient department of a large hospital, a fact that might be a selective factor for more serious infections. Nevertheless, the high prevalence of PVL-positive CA-MRSA has implications for both antimicrobial

TABLE

Main characteristics of community-acquired methicillinresistant *Staphylococcus aureus* (CA-MRSA) strains* isolated in a tertiary-care hospital in Athens, Greece, January 2006 -December 2007 (n=27)

No	Sex**	Age	Disease	PFGE type	<i>spa</i> type	Resistance Phenotype***
1	М	45	Furuncle	A1	t044	Oxa Tet Km FA
2	М	43	Abscess (skin)	A	t044	Oxa Tet Km FA
3	F	45	Abscess (skin)	A	t044	Oxa Tet Km FA
4	м	34	Furuncle	A1	t044	Oxa Tet Km FA
5	F	43	Abscess (soft tissue)	Α	t044	Oxa Tet Km FA
6	F	46	Folliculitis	A	t131	Oxa Tet Km FA
7	М	51	Furuncle	A2	t044	Oxa Tet Km FA
8	F	32	Abscess (soft tissue)	A	t044	Oxa Tet Km FA
9	М	45	Abscess (soft tissue)	A	t044	Oxa Tet Km FA
10	м	32	Abscess (soft tissue)	A	t044	Oxa Tet Km FA
11	F	38	Abscess (skin)	A	t044	Oxa Tet Km FA
12	F	31	Wound infection	A1	t044	Oxa Tet Km FA
13	F	29	Abscess (skin)	A	t044	Oxa Tet Km FA
14	М	35	Wound infection	A	t044	Oxa Tet Km FA
15	F	48	Furuncle	A	t044	Oxa Tet Km FA
16	М	39	Abscess (skin)	A 3	t044	Oxa Tet Km FA
17	F	46	Abscess (skin)	A	t044	Oxa Tet Km FA
18	М	51	Abscess (soft tissue)	A2	t044	Oxa Tet Km FA
19	М	56	Abscess (soft tissue)	A1	t044	Oxa Tet Km FA
20	F	51	Furuncle	A	t044	Oxa Tet Km FA
21	М	45	Abscess (soft tissue)	A	t044	Oxa Tet Km FA
22	м	47	Abscess (soft tissue)	A	t044	Oxa Tet Km FA
23	М	43	Wound infection	A	t044	Oxa Tet Km FA
24	F	46	Abscess (soft tissue)	A	t044	Oxa Tet Km FA
25	F	51	Wound infection	A4	t044	Oxa Tet Km FA
26	М	34	Furuncle	A	t044	Oxa Tet Km FA
27	F	51	Abscess (skin)	A1	t044	Oxa Tet Km FA

* Note: All stains were sensitive to tobramycin and gentamicin, cotrimoxazole, chloramphenicol, quinolones, clindamycin, erythromycin. ** M=male; F=female

*** Oxa=Oxacyclin; Tet=Tetracyclin; Km=Kanamycin; FA=Fucidic acid

treatment and MRSA surveillance in Greece. Our results indicate that in this country empiric therapy when clinically indicated, should exclude β -lactam antibiotics. Moreover, empiric use of macrolides for purulent skin and soft-tissue infections should be monitored closely. Clindamycin, trimethoprim–sulfamethoxazole, or linezolid, because of their good activities against all *S. aureus* in general, are potential alternatives to β -lactams for oral application. However, routine microbiologic workup should be performed for all community-acquired skin and soft-tissue infections in this country.

In contrast to the well documented nosocomial spread of CA-MRSA in the USA, outbreaks of nosocomial infections due to CA-MRSA have so far been reported only sparsely in Europe, with eight cases in Germany in 2005 [28]. This might be due to an overall low prevalence of CA-MRSA in the European population, and thus a rare introduction of such strains to the hospitals by admission of colonised carriers on the one hand. On the other, the high clinical manifestation index of CA-MRSA might lead to an earlier detection of patients infected with CA-MRSA. The phenomenon may also indicate that ST80-MRSA type IV isolates are less well adapted to be sustained in hospital environments [17]. However, in Greece, PVL-positive ST80-MRSA type IV CA-MRSA have been introduced in at least one hospital since 2000 [29, 30], a fact of great public health significance. These strains are associated with increased disease severity mainly due to the presence of PVL genes, and a possible adaptation in the hospital environment would result in outbreaks of serious nosocomial infections. This perspective is of immense importance in a country already suffering from high rates of infections due to multidrug-resistant organisms (see Greek System for the Surveillance of Antimicrobial Resistance www. medne.gr/whonet and ERASS http://www.rivm.nl/earss/).

In conclusion we believe that the establishment of an active screening programme for PVL-positive CA-MRSA carriage and adopting a search and destroy strategy for CA-MRSA in all patients admitted with purulent skin and soft-tissue is of high priority Greece as well as in all European counties who face high rates of CA-MRSA infections.

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