Three cases of community-associated meticillin-resistant Staphylococcus aureus infection in China

The spread of meticillin-resistant Staphylococcus aureus (MRSA) in the community has challenged infection control recently since it can be acquired by patients without established risk factors for MRSA infection and can lead to a fatal outcome. Such community-associated MRSA (CA-MRSA) has been observed in many parts of the world (Hsu et al., 2005; Aires de Sousa et al., 2003; Ho et al., 2004). Three cases of CA-MRSA in mainland China and the genetic characteristics of the strains are described here in detail.

Case 1: In September 2006, a 36-year-old woman was admitted to the West China Hospital with a 4-day history of a carbuncle on the back of her foot. Pus from the lesion site was sent for culture immediately. MRSA was isolated; it was susceptible to ciprofloxacin, gentamicin, rifampicin, co-trimoxazole and vancomycin. The patient was treated with levofloxacin and metronidazole intravenously. Gentamicin was also used on the lesion area. The patient was previously healthy without any fundamental diseases or recent hospitalization.

Case 2: In February 2007, a 39-year-old Japanese man was brought to our hospital because of an abscess in the centre of his left armpit. About 1 week before admission, he got the infection in his armpit after swimming in Thailand but was not treated there. The bacterial culture of pus was performed in West China Hospital. Results showed a pure growth of MRSA, which was susceptible to gentamicin, rifampicin, clindamycin, co-trimoxazole, tetracycline and vancomycin. He was treated with vancomycin. This patient was strong and healthy without any history of hospitalization, surgery or residence in medical settings for the previous year.

Case 3: In April 2007, a 43-year-old man was admitted to our hospital because of a progressive pain and abscess in his finger. The tip of the finger was necrotic. Secretion from the lesion site was cultured and amputation was performed. MRSA was isolated and found to be susceptible to ciprofloxacin, gentamicin, rifampicin, co-trimoxazole and vancomycin. The patient was treated with ciprofloxacin intravenously. He was previously healthy without any established risk factors for health-care-associated MRSA infection.

The MRSA strains from these three cases were named s35301, s19165 and s29635 sequentially. They were confirmed to be S. aureus by colony morphology, production of coagulase, as well as catalase tests. Resistance to meticillin was verified by both the cefoxitin disc diffusion test recommended by CLSI and multi-PCR of meca/femB genes as described previously (CLSI, 2005; Jonas et al., 1999). Genes encoding Panton–Valentine leukocidin (PVL) were co-amplified with primers described elsewhere (Lina et al., 1999). Staphylococcus chromosomal cassette mec (SCCmec) types were investigated by the novel multiplex PCR strategy described previously (Zhang et al., 2005). Genetic backgrounds of the three organisms were characterized further by PFGE, multilocus sequence typing (MLST) and staphylococcus protein A (spa) typing as described previously (Murchan et al., 2003; Enright et al., 2000; Aires de Sousa & de Lencastre, 2003).

Two strains (s35301 and s19165) out of the three were positive for PFGE. All three strains carried the type IVa SCCmec. Patterns with four or more different bands were designated separate PFGE types. They showed three different PFGE patterns, and clustered in two clones by MLST and spa typing (Fig. 1). Strains from case 1 and case 3 (s35301 and s29635) were ST39-t437. Both were isolated from the local Chinese people in Chengdu, a city in west China. The MRSA strain of case 2 was typed as ST8-4008. This strain had a unique PFGE pattern which was similar to that of the USA300 clone. ST59 was considered to be a community clone in Taiwan and has spread to San Francisco, USA (Aires de Sousa et al., 2003; Huang et al., 2006). It is very different in genotype from the present five MRSA clone complex (CC5, CC8, CC22, CC30 and CC45) (Enright et al., 2002). Five meticillin-susceptible S. aureus (MSSA) strains from the UK and the USA and six MRSA strains from the USA, Taiwan, Sweden and Singapore also belong to ST59. Two ST59 MRSA strains from Singapore have been specified as community-acquired possessing the type V SCCmec (from http://www.mlst.net). S. aureus of t437 has spread in some European countries such as Sweden, Norway, Germany, Denmark and others, and is also common in Taiwan. So far, 11 strains of t437 from Norway have been typed as ST59 (from http://www.ridom.de/spaserver). The two ST59 MRSA strains in our study may be from the local ST59 MSSA which acquired the type IVa SCCmec horizontally. However, the genetic background of MSSA in our region has not been surveyed. The possibility of ST59 MRSA input by foreign carriers can not be excluded.

Strain s19165 in our study was ST8-4008, possessing the type IVa SCCmec, with positive detection of the PVL gene, which showed the same genetic characteristics as those of the USA300 clone (King et al., 2006; Moran et al., 2006). This clone is mainly prevalent in the USA and some European countries, such as the UK, Netherlands, Germany, Austria and others, and it has also been found in Japan (from http://spa.ridom.de/spaserver), but it has not been found in mainland China before. This strain was isolated from a Japanese tourist who got his armpit infection when travelling in Thailand, which suggests that USA300 may have been prevalent there and was carried to west China.

All three strains can be considered community-associated by referring to the clinical information. Laboratory results
also characterized them as CA-MRSA: susceptible to several non-β-lactams, carriage of SCC\textsuperscript{mec} type IVa, positive for PVL genes (two of them), and shown to be the same as CA-MRSA clones prevalent in other parts of the world by MLST and spa typing (Davis \textit{et al.}, 2007). Conclusively, CA-MRSA (SCC\text{mec} IVa-ST59-t0437 and SCC\text{mec} IVa-ST8-1008) has been present in mainland China. It should be paid much attention because of the probable fatal results and potential for outbreak. Screening patients with risk factors for CA-MRSA might be useful for controlling its domestic spread in communities. Patients who have a history of travelling abroad should be screened for MRSA carriage before admittance to hospitals.

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![Fig. 1. Genetic types of the three CA-MRSA isolates. 1, s35301 (case 1); 2, s19165 (case 2); 3, s29635 (case 3); M, marker.](image-url)