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Title	First outbreak of methicillin-resistant Staphylococcus aureus USA300 harboring the Panton-Valentine leukocidin genes among Japanese health care workers and hospitalized patients.
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First outbreak of methicillin-resistant *Staphylococcus aureus* USA300 harboring the
 Panton-Valentine leukocidin genes among Japanese healthcare workers and hospitalized
 patients
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- 9 Abstract
- 10 This report describes the first outbreak of methicillin-resistant Staphylococcus aureus

11 USA 300 in a general hospital ward in Japan, involving six healthcare workers and four

- 12 patients. This report emphasizes the need for healthcare personnel to be alert that MRSA
- 13 harboring *pvl* poses a threat for both nosocomial and occupational infection.
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- 15

16	Accumulating evidence indicates that community-associated methicillin resistant
17	Staphylococcus aureus (CA-MRSA) isolates can readily produce outbreaks in hospitals,
18	adding to the threat posed by these organisms (1-3). CA- MRSA is genetically
19	heterogeneous, and includes a variety of clones such as the multilocus sequence (ST) 1
20	(USA400) and ST8 (USA300) types that emerged as major clones in the United States.
21	The USA300 clone can replace preexisting MRSA clones, and it now represents the
22	predominant CA-MRSA clone in the United States (4,5). Until now, CA-MRSA
23	infections reported in Japan have been sporadic, and most strains did not harbor the
24	Panton-Valentine leukocidin (pvl) genes (6).
25	In September 2009, we were notified that a cluster of skin infections had broken out
26	among healthcare workers (HCWs) and hospitalized patients in a general ward. We
27	document herein the first outbreak of MRSA harboring pvl genes belonging to the
28	USA300 clone in a healthcare setting in Japan.
29	At the time of notification, 65 patients were being cared for by 96 HCWs in a general
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	ward at the tertiary care, 1240-bed Kyoto University Hospital (Japan), where
31	ward at the tertiary care, 1240-bed Kyoto University Hospital (Japan), where dermatological disorders were quite prevalent. We were notified that a cluster of skin
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32	dermatological disorders were quite prevalent. We were notified that a cluster of skin infections had developed among four healthcare workers (HCW 1-4) and it seemed
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32 33 34	dermatological disorders were quite prevalent. We were notified that a cluster of skin infections had developed among four healthcare workers (HCW 1-4) and it seemed compatible with <i>S. aureus</i> infection. Two weeks later, one patient (Patient 4) developed a skin abscess in the left arm from which MRSA was isolated. The isolate
32 33 34 35	dermatological disorders were quite prevalent. We were notified that a cluster of skin infections had developed among four healthcare workers (HCW 1-4) and it seemed compatible with <i>S. aureus</i> infection. Two weeks later, one patient (Patient 4) developed a skin abscess in the left arm from which MRSA was isolated. The isolate was susceptible to erythromycin, clindamycin and gentamicin. The antimicrobial
32 33 34 35 36	dermatological disorders were quite prevalent. We were notified that a cluster of skin infections had developed among four healthcare workers (HCW 1-4) and it seemed compatible with <i>S. aureus</i> infection. Two weeks later, one patient (Patient 4) developed a skin abscess in the left arm from which MRSA was isolated. The isolate was susceptible to erythromycin, clindamycin and gentamicin. The antimicrobial susceptibility pattern was distinct from that of healthcare-associated MRSA

40 Eventually MRSA isolates were recovered from HCWs.

41

42which MRSA with a specific antibiogram was recovered from a clinical specimen. The case-defined antibiogram was susceptible to erythromycin, clindamycin and gentamicin, 43but resistant to levofloxacin and β -lactams with MIC levels below those of HA-MRSA. 44 We reviewed the antimicrobial susceptibility profiles of MRSA strains from all adult 45and pediatric hospitalized patients who were under care at Kyoto University Hospital 46 during 2009 to detect any unidentified MRSA. HCWs were screened in the ward 47using nasal swabs to identify MRSA carriers. 48Clinical specimens were inoculated onto mannitol salt agar plates and examined after 495048h. Susceptibility testing proceeded according to the Clinical and Laboratory 51Standards Institute. The mecA gene, PVL determinants and arcA gene on the arginine catabolite mobile element (ACME) were detected and SCCmec typing was performed 5253by PCR (7-9). The typing procedure included PFGE using the restriction enzyme SmaI as described (5). Multilocus sequence typing proceeded as described and the 54nomenclature was specified as previously described. (www.MLST.net) 55Patients 1 to 3 who became infected with MRSA were newly identified based on the 56 57antimicrobial susceptibility profile described in the case definition (Table 1). A review 58of the medical records revealed that the first MRSA infection occurred in March 2009.

Based on information derived from these cultures, we developed a case definition in

59 Patient 1 developed catheter-related bloodstream infection followed by pneumonia and 60 required intravenous anti-MRSA drug administration. Six of nine skin and soft tissue 61 infections (skin abscesses, folliculitis) were treated with antibiotics, whereas three were 62 cured by drainage alone. Patient 1 as well as HCW 1 and 5 developed recurrent 63 infections. No case patient had a history of visiting abroad recently. All MRSA

isolates recovered from the case patients contained SCC *mec* type IV, the *pvl* gene and
ACME-associated *arcA* gene. PFGE-based findings identified all isolates as being
identical to and indistinguishable from the USA 300 clone. MLST defined all of them
as ST8.

Excluding the isolates recovered from the case patients, four of 825 strains of MRSA isolates at our institution in 2009 had the same antimicrobial susceptibility profile as the outbreak strain. Those isolates were recovered from swab specimens and the patients did not have a symptom of infection when the specimens were taken. Screening nasal swabs of HCWs did not identify any carriers of CA-MRSA USA 300 other than the case patients.

74 Discussion

To our knowledge, this is the first report to document an outbreak of 75healthcare-associated and -transmitted CA-MRSA USA300 in Japan. 76 To date. 77 outbreaks of PVL-positive CA-MRSA have been reported, especially in neonatal intensive care units and long term care facilities in the United states and European 78countries (1-3). However, no outbreaks of CA-MRSA in either community- or 79 nosocomial settings in Japan have been described. Only one sporadic infection with 80 81 the USA300 clone in Japan has been documented (10). We speculate that infected HCWs or unidentified PVL-positive MRSA carriers served as the source of infection for 82 Patients 1 to 4, because all of them became infected while in hospital. In addition, we 83 considered that the causative agent was community-associated because antibiograms of 84 85 the outbreak strain were distinct from those of HA-MRSA strains. This was supported by a review of the antibiotic susceptibility of MRSA strains isolated at our institution 86 87 during 2009; the case-defined antibiogram occurred in only 0.5% of isolates.

The findings of this investigation have considerable public health implications. 88 Although HA-MRSA remains a serious threat to hospitalized patients, the introduction 89 90 of CA-MRSA strains into tertiary care hospitals like our hospital represents an especially serious challenge. Many of the infections caused by these strains have been 9192reported to cause serious infections among healthy adults and can be severer and more life-threatening to those who are highly immunocompromised. In this study, healthy 93 HCWs suffered from skin abscesses, although most infections were mild and cured 94 95 without parenteral anti-MRSA drugs. Considering that infection relapsed in some case patients, further investigations are needed to establish the management of PVL-positive 96 97 MRSA carriers, especially when they are caregivers. Systematic studies involving 98 healthcare settings are needed to reveal the transmission of such CA-MRSA isolates within the healthcare system. These would provide not only an accurate estimate of 99 100 CA-MRSA prevalence, but would help monitor the emergence of more resistant and/or 101 virulent clones and help with therapeutic infection control and patient management 102 policies.

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105 Potential conflict of interest: All authors; no conflict

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146 Table 1 Characteristics of patients with CA-MRSA infections and their treatment, Kyoto

Case	Age,	Underlying	Onset of	Type of	Site of	Antimicrobial	Drainage	
	Sex	disease	infection	infection	infection	drug treatment	Drainage	
Pt 1	60,F	Inflammatory bowel disease	3/29	CRBSI	Blood- stream	TEIC	None	
			8/27	Pneumonia	Lung	LZD	None	
Pt 2	65,F	Decubitas ulcer	8/22	Skin abscess	Thigh	GEN	Spontaneou	
Pt 3	42,M	Polyarteritis nodosa	10/19	Folliculitis	Legs	ST	Spontaneou	
Pt 4	14,F	Decubitus ulcer	11/2	Skin abscess	Arm	GEN	Surgical,	
HCW	31,F	,F none _	9/19	Skin abscess	Arm	MINO	Surgical	
1			11/24	Skin abscess	Arm	CLI	None	
HCW 2	27,F	none	9/21	Skin abscess	Arm	GEN	Surgical	
HCW 3	27,F	none	9/21	Skin abscess	Leg	MINO	Surgical	
HCW 4	25,F	none	9/21	Skin abscess	Leg	None	Surgical	
HCW	HCW 27,F 5	27 F	none	11/20	Skin abscesses	Leg, Chest	CLI	Spontaneou
5			12/14	Folliculitis	Arm, Finger	ST MUP	None	
HCW 6	31,M	none	11/24	Skin abscess	Thigh	CLI	Spontaneou	

147 University Hospital, 2009

- 148 TEIC, teicoplanin; LZD, linezolid; GEN, gentamicin(topical); ST,
- 149 Trimethoprim-sulfamethoxazole; MINO, minocyclin; CLI, clindamycin; MUP,
- 150 mupirocin (topical), CRBSI; catheter-related bloodstream infection