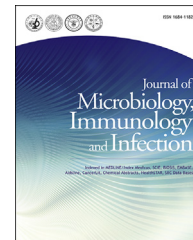




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## CASE REPORT

# Emergence of a strain of methicillin-resistant *Staphylococcus aureus* with decreased susceptibility to vancomycin 7 months after treatment with glycopeptide antibiotics



Cheng Len Sy<sup>a</sup>, Susan Shin-Jung Lee<sup>a,b</sup>, Kuan-Sheng Wu<sup>a,b</sup>, Hung-Chin Tsai<sup>a,b</sup>, Yung-Ching Liu<sup>c</sup>, Yao-Shen Chen<sup>a,b,\*</sup>

<sup>a</sup> Section of Infectious Diseases, Department of Medicine, Kaohsiung Veterans General Hospital, Kaohsiung, Taiwan

<sup>b</sup> National Yang-Ming University, Taipei, Taiwan

<sup>c</sup> Section of Infectious Diseases, Shuang-Ho Hospital, Taipei Medical University and School of Medicine, Taipei Medical University, Taipei, Taiwan

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*Staphylococcus aureus*;  
Vancomycin

This case report describes a methicillin-resistant *Staphylococcus aureus* isolated repeatedly from the blood of a patient with community-acquired endocarditis who developed a four-fold increase in the minimal inhibitory concentration of vancomycin and daptomycin 7 months after his last exposure to glycopeptide antibiotics. This is contrary to the expected situation in which antimicrobial resistance tends to decrease after a patient is no longer exposed to vancomycin. Copyright © 2013, Taiwan Society of Microbiology. Published by Elsevier Taiwan LLC. This is an open access article under the CC BY-NC-ND license (<http://creativecommons.org/licenses/by-nc-nd/4.0/>).

## Introduction

Methicillin-resistant *Staphylococcus aureus* (MRSA) emerged in the 1960s,<sup>1</sup> and gradually became an endemic pathogen in hospital settings in the 1980s. This led to the extensive use of vancomycin for the treatment of MRSA infections. MRSA with reduced vancomycin susceptibility

\* Corresponding author. Section of Infectious Diseases, Department of Medicine, Kaohsiung Veterans General Hospital, Kaohsiung, Taiwan.

E-mail address: [yschen@vghks.gov.tw](mailto:yschen@vghks.gov.tw) (Y.-S. Chen).

appeared in the 1990s, and tends to disappear following cessation of vancomycin therapy.<sup>2</sup> The current report describes a strain of MRSA, repeatedly isolated from the blood of a patient with community-acquired bacterial endocarditis, which exhibited a delayed increase in the minimum inhibition concentration (MIC) of vancomycin 200 days following treatment with vancomycin and teicoplanin.

## Case report

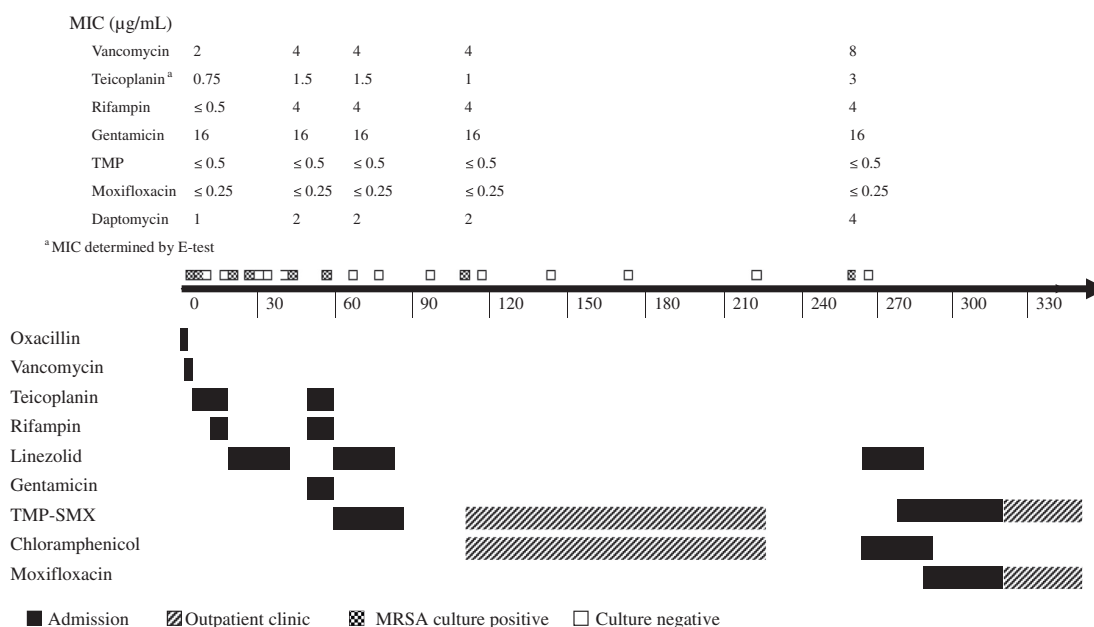
A 45-year-old male intravenous drug user was admitted to Kaohsiung Veterans General Hospital, Taiwan in March, 2007 with MRSA infective endocarditis and lumbar spine osteomyelitis. He was treated with vancomycin for 3 days followed by a 24-day, intermittent course of teicoplanin. During this period, the MIC of vancomycin rose from 2 mg/dL to 4 mg/dL. Teicoplanin was switched to linezolid on Day 19 due to persistence of fever. There were no glycopeptide exposures since then. Blood cultures were obtained as an outpatient at 1–3 week intervals. MRSA was re-isolated on the Day 110. He was treated for 110 days with oral trimethoprim–sulfamethoxazole and chloramphenicol. He was readmitted on the Day 257 because of fever. MRSA with an MIC of 8 mg/dL was isolated from his blood. There was no concurrent infection with vancomycin-resistant enterococci. He was treated with linezolid and chloramphenicol for 23 days and then switched to trimethoprim–sulfamethoxazole and moxifloxacin until Day 337. Blood cultures continued to remain sterile after Day 264. The patient was not screened for colonization with MRSA or vancomycin-resistant enterococci. A preliminary report of this case was published<sup>3</sup>

prior to noting the four-fold rise in the MIC of vancomycin and daptomycin on Day 257 and knowledge of the follow-up by Day 110 day.

## Laboratory studies

Blood culture isolates obtained at Day 0, Day 47, Day 55, Day 110, and Day 257 were stored at  $-80^{\circ}\text{C}$ . Following reconstitution antibiotic susceptibility and genetic profiles were performed on all the isolates together with *S. aureus* ATCC 29213. The MICs for vancomycin, daptomycin, rifampin, gentamicin, trimethoprim–sulfamethoxazole, and moxifloxacin were determined, in duplicate, by broth microdilution using Sensititre custom designed plates for staphylococci (Trek Diagnostics, East Grinstead, West Sussex, UK). The MIC for teicoplanin was determined by the E-test (AB Biodisk, bioMérieux SA, Marcy l'Etoile, France). Pulsed field gel electrophoresis (PFGE) was conducted using standard DNA extraction methods. Lambda ladder PFG marker (New England Biolabs, Schwalbach, Germany) was used as fragment size marker. The type IV staphylococcal cassette chromosome *mec* and Pantone–leukocidin genes were determined as previously described.<sup>3</sup> Multilocus sequence typing (MLST) was performed as described by Enright et al.<sup>4</sup> The MLST sequences were submitted to the MLST database (<http://www.mlst.net/>). Spa typing was performed using methods described by Harmsen et al.<sup>5</sup>

The antimicrobial susceptibility results are shown in Fig. 1. The key findings were a four-fold increase in the MICs of vancomycin, teicoplanin, and daptomycin without significant changes in the MICs of the other antibiotics during the study period. All isolates were found to be identical by



**Figure 1.** Course of treatment of a patient with bacterial endocarditis in Taiwan with the minimal inhibitory concentrations (MICs/dL) to 10 antibiotics of a methicillin-resistant strain of *Staphylococcus aureus* repeatedly isolated from his blood during March–November 2007. MIC determined by the E-test method. MRSA = methicillin-resistant *Staphylococcus aureus*; TMP-SMX = trimethoprim–sulfamethoxazole.

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