

In Vitro Susceptibility to Glycopeptides, Linezolid, Tigecycline, and Ceftaroline against Methicillin Resistant *Staphylococcus aureus* Isolates in a Tunisian University Hospital

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ABSTRACT

Background: Vancomycin remains the drug of choice for treatment of serious methicillin-resistant *Staphylococcus aureus* (MRSA) infections. However, Strains with reduced susceptibility to vancomycin have been reported in several countries around the world. Therefore, new antibiotics are introduced into the treatment.

Objectives: The present study aimed to determine the vancomycin, teicoplanin, linezolid, tigecycline and ceftaroline susceptibility pattern and to investigate the presence and the frequency of heterogeneous vancomycin intermediate *S. aureus* (hVISA) among clinical isolates of MRSA in Tunisia.

Methods: A total of 162 non duplicate MRSA strains isolated between 2017 and 2018 were investigated. Vancomycin, teicoplanin, tigecycline, linezolid and ceftaroline minimum inhibitory concentrations (MIC) values were detected by broth microdilution method and interpreted according to the European Committee on Antimicrobial Susceptibility testing criteria. Etest GRD, Etestmacromethod, Mueller-Hinton screen agar, and population analysis profile-area under the curve (PAP-AUC) methods were used to detect hVISA.

Results: The MIC₅₀, MIC₉₀, and MIC ranges were respectively 1, 1, and 0.5-2 µg/ml for vancomycin; 1, 2, and 0.125-4 µg/ml for teicoplanin; 2, 2, and 0.5-4 µg/ml for linezolid; and 0.25, 0.5, and 0.064-0.5 µg/ml for tigecycline. Twelve strains were suspected as hVISA by EtestGRD, 28 by screen agar, and one by Etestmacromethod, but only one strain was confirmed hVISA by PAP-AUC. No vancomycin, tigecycline and linezolid resistance was found among MRSA isolates. Four strains were teicoplanin resistant, and four were intermediate to ceftaroline.

Conclusion: The prevalence of hVISA, teicoplanin and ceftaroline resistance are low. Linezolid and tigecycline were found to be highly active against MRSA isolates. Therefore, they could be considered as alternative agents for the treatment of serious infections. Continuous and regular monitoring of MICs at local and regional level is necessary to guide clinician in their empiric antibiotic selection.

Keywords: Methicillin resistant *Staphylococcus aureus*, glycopeptide, linezolid, tigecycline, ceftaroline, MIC.

INTRODUCTION

Glycopeptides, vancomycin and teicoplanin are considered the treatment of choice for severe methicillin-resistant *Staphylococcus aureus* (MRSA) infections for decades. However, strains with reduced susceptibility to vancomycin, including vancomycin intermediate *S. aureus* (VISA) and heterogeneous VISA (hVISA) and even vancomycin resistant *S. aureus* (VRSA) have been reported since 1997 in many parts of the world [1-3]. Reduced susceptibility to vancomycin is frequently accompanied by acquisition of teicoplanin resistance [1,4]. hVISA is an *S. aureus* isolate with a minimum

inhibitory concentration (MIC) for vancomycin within the susceptible range but contain a resistant subpopulation to vancomycin at a frequency of 10⁻⁵ to 10⁻⁶ [5]. Although VRSA strains are rare, the prevalence of hVISA/VISA strains is increasing [2, 6].

For testing susceptibility to glycopeptides, the broth microdilution method (BMD) should be used. The European Committee on Antimicrobial Susceptibility Testing (EUCAST) defines the vancomycin and teicoplanin MIC breakpoint of >2 µg/ml for resistant, and ≤ 2 µg/ml for susceptible [7]. However, *S. aureus* strains with an MIC vancomycin and / or teicoplanin > 1 µg/ml should be tested for

hVISA. In fact, vancomycin treatment failure has been reported even in susceptible strains, which may be attributable to the presence of resistant hVISA subpopulation [5]. The gold standard technique to detect hVISA strains is the population analysis profile area under the curve (PAP-AUC) method, which is time-consuming, expensive, and is unsuitable for routine use in the clinical microbiology laboratories [1,8,9].

Infections caused by multidrug resistant MRSA strains with reduced susceptibility to vancomycin are frequently associated with therapeutic failure and persistent infections justifying the use of alternatives such as linezolid, tigecycline, daptomycin or new generation cephalosporin which have shown potent activity against MRSA in previous publications [1, 10]. In Tunisia, ceftaroline is not commercially available; add to this, there have been no studies focused on prevalence of hVISA among MRSA strains using PAP-AUC method. Therefore, we aimed to determine vancomycin, teicoplanin, linezolid, tigecycline and ceftaroline susceptibility pattern and to investigate the presence and the frequency of hVISA isolates among clinical isolates of MRSA.

MATERIAL AND METHODS

Bacterial Strains

A total of 162 consecutive and non-duplicate MRSA isolates collected between 2017 and 2018 at Sfax university hospital were included in this study. Isolates were collected from various clinical samples, including blood (n=52), skin and soft tissue (n=62), respiratory tract (n=40), and catheter (n=8).

Identification of *S. aureus* isolates was performed using conventional methods. Methicillin resistance was identified by cefoxitin disc according to the EUCAST guidelines [7], and confirmed by the detection of *mecA* gene by PCR [11]. *S. aureus* ATCC 43300 was used as positive control.

Antimicrobial Susceptibility Testing

Antimicrobial susceptibilities were performed by a disc diffusion method according to the EUCAST criteria [7]. MICs of vancomycin, teicoplanin, linezolid and tigecycline were determined for all of isolates by reference BMD [7]. Ceftaroline MICs were determined for intermediate or resistant strains by the disc method. *S. aureus* ATCC 29213 was used as a control.

Detection of hVISA

Screening for hVISA was performed by various tests on all isolates. *S. aureus* ATCC 29213 (vancomycin-susceptible), *S. aureus* Mu3 (ATCC 700698; hVISA), and *S. aureus* Mu50 (ATCC 700699; VISA) were included as controls.

MHA5T screening agar

Ten microliters of a 2 McFarland inoculum of each strain was inoculated as spot onto the surface of the Mueller-Hinton agar plates with 5 µg/ml teicoplanin. Growth of ≥ 4 colonies after 48 h of incubation at 35°C indicated a positive result [1].

Etestmacromethod

A 2 McFarland inoculum (200µl) was swabbed on Brain-Heart Infusion (BHI) agar and allowed to dry. Vancomycin and teicoplanin Etest strips (BioMérieux) were applied. After incubation for 48 h at 37°C, the MICs were read at complete inhibition. The criteria used to detect hVISA were MICs of ≥ 8 µg/ml for both vancomycin and teicoplanin or a teicoplanin MIC of ≥ 12 µg/ml [8].

Etestglycopeptide-resistance detection (GRD)

Etest GRD was performed according to the manufacturer instructions (BioMérieux) using a double-ended Etest strip with vancomycin and teicoplanin. A standard inoculum (0.5 McFarland) was swabbed onto a Mueller-Hinton agar plates with 5% Blood; next a GRD strip was applied. The elliptical zone was read at 24 and 48 h after incubation at 35°C. If either teicoplanin or vancomycin Etest GRD was ≥ 8 µg/ml, the isolate was considered hVISA[8].

PAP-AUC

PAP-AUC was done as described previously [8, 9]. Briefly, after 24 h incubation in BHI broth, cultures were diluted in saline to 10^{-3} and 10^{-6} , and plated on to BHI plates containing 1, 2,3,4,5 and 6 µg/ml vancomycin. Colonies were calculated after 48 h incubation at 37°C and plotted against vancomycin concentration using GraphPad prism. The ratio of AUC of the test isolate/ AUC of Mu3 was calculated. If ratio < 0.9 , the isolate was considered vancomycin susceptible. Ratios of 0.9-1.3 and ≥ 1.3 were considered positive for hVISA and VRSA respectively [12].

RESULTS

All MRSA isolates were susceptible to linezolid, tigecycline, vancomycin and quinupristin-dalfopristin (Table 1).

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Table1. Antimicrobial susceptibility of MRSA isolates against evaluated antimicrobial agents.

Antimicrobial agents	%R	%I	%S
Gentamicin	46.9	0	53.1
Erythromycin	38.3	0	61.7
Clindamycin	32.7	0	67.3
Quinupristin-dalopristin	0	0	100
Ceftaroline ^a	0	2.5	97.5
Chloramphenicol	2.5	0	97.5
Tetracycline	69.2	0	30.8
Tigecycline ^a	0	0	100
Ofloxacin	44.5	0	55.5
Rifampicin	37.7	5.5	56.8
Trimethoprim-sulfamethoxazole	1.2	0	98.8
Fusidic acid	50.6	0	49.4
Vancomycin ^a	0	0	100
Teicoplanin ^a	2.5	0	97.5
Linezolid ^a	0	0	100

S, susceptible; I, intermediate; R, resistant.

^aVancomycin, teicoplanin, linezolid, tigecycline and ceftaroline susceptibilities were determined by broth microdilution method.

Of the eight strains classified as resistant or intermediate to ceftaroline by disc diffusion method, four had ceftaroline MIC value of 1

Table2. MIC distributions and activities of vancomycin, teicoplanin, linezolid and tigecycline against MRSA strains.

	No. of isolates with MIC (µg/ml)								MIC ₅₀	MIC ₉₀	Geometric mean
	0.064	0.125	0.25	0.5	1	2	4	8			
Vancomycin	0	0	0	30	126	6	0	0	1	1	0.90
Teicoplanin	0	2	2	61	48	45	4	0	1	2	0.92
Linezolid	0	0	0	5	70	82	5	0	2	2	1.45
Tigecycline	9	70	58	25	0	0	0	0	0.25	0.5	0.19

MIC₅₀: minimum inhibitory concentration which 50% of the strains were inhibited.

MIC₉₀: minimum inhibitory concentration which 90% of the strains were inhibited.

By Etestmacromethod, vancomycin MICs ranged from 2 to 6 µg/ml, teicoplanin MICs ranged from 2 to 12 µg/ml, and one isolate met the criteria of hVISA. By Etest GRD, vancomycin MICs ranged from 1 to 16 µg/ml, teicoplanin MICs ranged from 4 to 32 µg/ml,

µg/ml (susceptible) and four were in the intermediate category (MIC = 2 µg/ml). The distribution of vancomycin, teicoplanin, linezolid and tigecycline MICs is shown in table 2. The four strains resistant to teicoplanin were susceptible to ceftazolin (MIC of 1 µg/ml) and to vancomycin (MIC of 2 µg/ml).

and 12 isolates met the criteria of hVISA. By MHA5T, 28 isolates met the criteria of hVISA.

The PAP-AUC ratios of the isolates were between 0.32 and 0.98. Only one isolate was hVISA, with a PAP-AUC ratio to Mu3 of 0.98 (figure 1). This strain was also detected by Etest GRD, EtestMacromethod and MHA5T. The hVISA isolate had a vancomycin MIC of 2 µg/ml and teicoplanin MIC of 4 µg/ml.

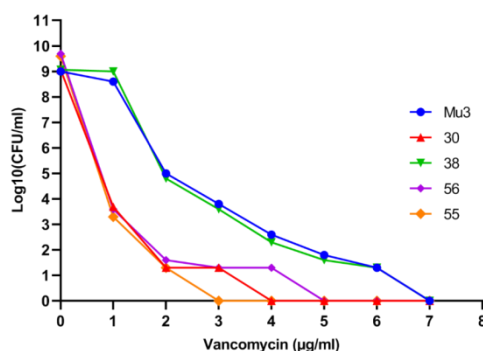


Figure1. Population analysis profile curves of four isolates.

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One isolate (38) was identified to be hVISA and three isolates were defined vancomycin-susceptible (30, 55, 56) compared to the susceptibility of the Mu3 reference strain.

DISCUSSION

Vancomycin remains as the only widespread therapeutic preference of serious MRSA infections although new anti-staphylococcal antibiotics such as linezolid and tigecycline have been developed. However, treatment failure may occur even when MRSA is susceptible to vancomycin [5,13,14]. Recently a phenomenon of gradual increase in the value of vancomycin MIC over time was reported in literature as MIC creep. It was described as one of the suspected causes of vancomycin treatment failure [15]. The studies reporting vancomycin creep have shown conflicting results. There are reports of increased MIC over the time [16,17], but other studies did not confirm these findings in MRSA [18,19]. A systematic review and meta-analysis did not report an increase in vancomycin MIC, suggesting that vancomycin continues to be the treatment of choice of MRSA infections [20]. The proportion of MRSA isolates with vancomycin MIC $>1 \mu\text{g/ml}$ was 26% in the USA, 18% in Asia, and 17% in Europe. The pooled means of vancomycin MIC were 1.12 $\mu\text{g/ml}$ in Europe, 1.17% in Asia and 1.37% in USA [20]. In Tunisia, only one multicenter study, conducted between 2011 and 2012, evaluated the activity of glycopeptides on MRSA by determination of MICs by BMD [21]. By comparing the results of this multicenter study with the present study, we observed an increase in the geometric means for vancomycin MIC (0.73 to 0.90 $\mu\text{g/ml}$) and for teicoplanin MIC (0.49 to 0.92 $\mu\text{g/ml}$), accompanied by an increase in the percentage of strains with vancomycin MIC $> 1 \mu\text{g/ml}$ (from 1.5% to 3.7%).

Our study documented the presence of 0.6% hVISA and 2.5% teicoplanin resistance amongst MRSA isolates. The previously study in Tunisia reported hVISA and teicoplanin resistance prevalence of 0.8% [21]. No resistance to vancomycin was noted in our study and in Tunisia [21]. VRSA due to the acquisition of the *vanA* gene from enterococci are currently very low. To date, few cases of VRSA have been reported from different countries such as the United States, India and Iran [6]. The prevalence of hVISA/VISA varied geographically. The

differences between studies may be explained by the use of different screening methods of hVISA and VISA strains. Add to this, confirmation of hVISA strains by the reference method is not performed in many studies [3,19,22,23]. A systematic review and meta-analysis [2] showed that the prevalence of hVISA/VISA isolates increased gradually from 4.68/2.05 % before 2006 to 7.01/7.93 in 2010-2014.

The PAP-AUC is the reference method to detect hVISA. However, this method is time-consuming, expensive and is not applicable in routine. Various screening strategies have been investigated for detection of hVISA. Several studies showed low sensitivity but good specificity ($> 92\%$) with EtestGRD and Etestmacromethod. However, agar screening plates with vancomycin or teicoplanin were highly sensitive but less specific [3,8,12,19,22-24]. In our study, false positive results have been found with Etest GRD and MHA5T. This result could be related to the rarity of strains with vancomycin MIC $\geq 2 \mu\text{g/ml}$, since there are studies in which hVISA strains were more commonly found among the isolates having vancomycin MICs of 2 $\mu\text{g/ml}$ [22].

Heteroresistance to vancomycin should be considered and investigated in case of clinical failure while using vancomycin to treat severe MRSA infection, and newer agents can be used as alternative if available. Linezolid and tigecycline are popular choices for the treatment of MRSA infections [10]. Li et al reported that the efficacy of linezolid should be better than that of vancomycin in the treatment of MRSA infections [25]. A systematic review and meta-analysis showed that linezolid and tigecycline have the best effect on MRSA with very low resistance ($<1\%$) [10]. In our study, all the MRSA isolates were susceptible to linezolid and tigecycline. The MIC₅₀, MIC₉₀ and the mean MIC were similar to those reported in the previously study from Tunisia [21].

Ceftaroline is a fifth-generation broad-spectrum cephalosporin that has activity against MRSA. It is reported to be non-inferior to vancomycin against MRSA [26,27]. In several studies around the globe, ceftaroline has an excellent in vitro activity against *S. aureus* isolates. Although MIC₅₀ and MIC₉₀ are significantly higher for MRSA. Susceptibility of MRSA to ceftaroline was 99.5% in the United States, 94% in Europe, 92.3% in Africa/West Asia, 84.4% in

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South America and 75.9% in Asia-Pacific [27]. In our study, 97.5% of the MRSA isolates were susceptible to ceftaroline. This was expected finding for us, as ceftaroline is not commercially available in Tunisia.

CONCLUSION

This is the first study in Tunisia investigating the prevalence of hVISA among MRSA strains by PAP-AUC method. We have demonstrated that the prevalence of hVISA and teicoplanin resistance are low. However, it is essential to test for hVISA especially for strains having teicoplanin or vancomycin MIC ≥ 2 $\mu\text{g/ml}$. Susceptibility to linezolid and tigecycline was higher than that of ceftaroline. These antibiotics should be kept as alternative therapy for critical cases of MRSA infections. Continuous and regular monitoring of MICs at local and regional level is necessary to guide clinician in their empiric antibiotic selection.

REFERENCES

- [1] Howden BP, Davies JK, Johnson PD, Stinear TP, Grayson ML. Reduced vancomycin susceptibility in *Staphylococcus aureus*, including vancomycin-intermediate and heterogeneous vancomycin-intermediate strains: resistance mechanisms, laboratory detection, and clinical implications. *Clin Microbiol Rev*. 2010;23(1):99-139.
- [2] Zhang S, Sun X, Chang W, Dai Y, Ma X. Systematic Review and Meta-Analysis of the Epidemiology of Vancomycin-Intermediate and Heterogeneous Vancomycin-Intermediate *Staphylococcus aureus* Isolates. *PLoS One* 2015;10(8):e0136082. Available from: doi: 10.1371/journal.pone.0136082.
- [3] Huang SH, Chen YC, Chuang YC, Chiu SK, Fung CP, Lu PL, et al. Prevalence of vancomycin-intermediate *Staphylococcus aureus* (VISA) and heterogeneous VISA among methicillin-resistant *S. aureus* with high vancomycin minimal inhibitory concentrations in Taiwan: A multicenter surveillance study, 2012–2013. *J Microbiol Immunol Infect*. 2016;49(5):701-7.
- [4] Bakthavatchalam YD, Babu P, Munusamy E, Dwarakanathan HT, Rupali P, Zervos M, et al. Genomic insights on heterogeneous resistance to vancomycin and teicoplanin in Methicillin-resistant *Staphylococcus aureus*: A first report from South India. *PLoS One* 2019;14(12):e0227009. Available from: doi: 10.1371/journal.pone.0227009.
- [5] Van Hal SJ, Paterson DL. Systematic review and meta-analysis of the significance of heterogeneous vancomycin-intermediate *Staphylococcus aureus* isolates. *Antimicrob Agents Chemother*. 2011;55(1):405-10.
- [6] Cong Y, Yang S, Rao X. Vancomycin resistant *Staphylococcus aureus* infections: A review of case updating and clinical features. *J Adv Res*. 2019;12(21):169-76.
- [7] The European Committee on Antimicrobial Susceptibility Testing. Breakpoint tables for interpretation of MICs and zone diameters. Version 8.1, 2018. <http://www.eucast.org>.
- [8] Satola SW, Farley MM, Anderson KF, Patel JB. Comparison of detection methods for heteroresistant vancomycin-intermediate *Staphylococcus aureus*, with the population analysis profile method as the reference method. *J Clin Microbiol*. 2011;49(1):177-83.
- [9] Wootton M, Howe RA, Hillman R, Walsh TR, Bennett PM, Mac Gowan AP. A modified population analysis profile (PAP) method to detect hetero-resistance to vancomycin in *Staphylococcus aureus* in a UK hospital. *J Antimicrob Chemother*. 2001; 47(4):399-403.
- [10] Shariati A, Dadashi M, Chegini Z, Van Belkum A, Mirzaii M, Khoramrooz SS, et al. The global prevalence of Daptomycin, Tigecycline, Quinupristin/Dalfopristin, and Linezolid-resistant *Staphylococcus aureus* and coagulase-negative staphylococci strains: a systematic review and meta-analysis. *Antimicrob Resist Infect Control*. 2020; 9(1):56. Available from: doi: 10.1186/s13756-020-00714-9.
- [11] Murakami K, Minamide W, Wada K, Nakamura E, Teraoka H, Watanabe S. Identification of methicillin-resistant strains of staphylococci by polymerase chain reaction. *J Clin Microbiol*. 1991;29(10):2240-4.
- [12] Walsh TR, Bolmström A, Qwärnström A, Ho P, Wootton M, Howe RA, et al. Evaluation of current methods for detection of staphylococci with reduced susceptibility to glycopeptides. *J Clin Microbiol*. 2001;39(7):2439-44.
- [13] Howden BP, Ward PB, Charles PG, Korman TM, Fuller A, du Cros P, et al. Treatment outcomes for serious infections caused by methicillin-resistant *Staphylococcus aureus* with reduced vancomycin susceptibility. *Clin Infect Dis*. 2004; 38(4):521-8.
- [14] Takesue Y, Nakajima K, Takahashi Y, Ichiki K, Ishihara M, Wada Y, et al. Clinical characteristics of vancomycin minimum inhibitory concentration of 2 $\mu\text{g/ml}$ methicillin-resistant *Staphylococcus aureus* strains isolated from patients with bacteremia. *J Infect Chemother*. 2011; 17(1):52-7.
- [15] Zarakolu P, Metan G, Ünal S, Karahan ZC, Tekeli A. Bacterial factors influencing the mortality for methicillin-resistant *Staphylococcus aureus* (MRSA) bacteremia. *Infect Dis (Lond)*. 2016;48(8):649-50.

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- [16] Yeh YC, Yeh KM, Lin TY, Chiu SK, Yang YS, Wang YC, et al. Impact of vancomycin MIC creep on patients with methicillin-resistant *Staphylococcus aureus* bacteremia. *J Microbiol Immunol Infect.* 2012;45(3):214-20.
- [17] Chang W, Ma X, Gao P, Lv X, Lu H, Chen F. Vancomycin MIC creep in methicillin-resistant *Staphylococcus aureus* (MRSA) isolates from 2006 to 2010 in a hospital in China. *Indian J Med Microbiol.* 2015;33(2):262-6.
- [18] Joana S, Pedro P, Elsa G, Filomena M. Is vancomycin MIC creep a worldwide phenomenon? Assessment of *S. aureus* vancomycin MIC in a tertiary university hospital. *BMC Res Notes.* 2013;6:65. Available from: doi: 10.1186/1756-0500-6-65.
- [19] OzmenCapin BB, Tekeli A, Karahan ZC. Evaluation of the Presence and Characterization of Vancomycin-Intermediate and Heterogeneous Vancomycin-Intermediate Level Resistance among Bloodstream Isolates of Methicillin-Resistant *Staphylococcus aureus*. *Microb Drug Resist.* 2020; 26(3):238-244.
- [20] Diaz R, Afreixo V, Ramalheira E, Rodrigues C, Gago B. Evaluation of vancomycin MIC creep in methicillin-resistant *Staphylococcus aureus* infections-a systematic review and meta-analysis. *Clin Microbiol Infect.* 2018;24(2):97-104.
- [21] MezghaniMaalej S, JdidiTrabelsi J, Claudealexandre G, Boutiba I, Mastouri M, Besbes S, et al. Antimicrobial Susceptibility and Molecular Epidemiology of Methicillin-Resistant *Staphylococcus aureus* in Tunisia: Results of a Multicenter Study. *J Infect Dis Epidemiol.* 2019, 5:071. Available from: doi: 10.23937/2474-3658/1510071.
- [22] Sancak B, Yagci S, Gür D, Gülay Z, Ogunc D, Söyletir G, et al. Vancomycin and daptomycin minimum inhibitory concentration distribution and occurrence of heteroresistance among methicillin-resistant *Staphylococcus aureus* blood isolates in Turkey. *BMC Infect Dis.* 2013;13: 583. Available from: doi: 10.1186/1471-2334-13-583.
- [23] Mirza HC, Sancak B, Gür D. The Prevalence of Vancomycin-Intermediate *Staphylococcus aureus* and Heterogeneous VISA Among Methicillin-Resistant Strains Isolated from Pediatric Population in a Turkish University Hospital. *Microb Drug Resist.* 2015;21(5):537-44.
- [24] Van Hal SJ, Wehrhahn MC, Barbagiannakos T, Mercer J, Chen D, Paterson DL, et al. Performance of various testing methodologies for detection of heteroresistant vancomycin-intermediate *Staphylococcus aureus* in bloodstream isolates. *J Clin Microbiol.* 2011;49(4):1489-94.
- [25] Li J, Zhao QH, Huang KC, Li ZQ, Zhang LY, Qin DY, et al. Linezolid vs. vancomycin in treatment of methicillin-resistant *Staphylococcus aureus* infections: a meta-analysis. *Eur Rev Med Pharmacol Sci.* 2017;21(17):3974-79.
- [26] Arshad S, Huang V, Hartman P, Perri MB, Moreno D, Zervos MJ. Ceftarolinefosamil monotherapy for methicillin-resistant *Staphylococcus aureus* bacteremia: a comparative clinical outcomes study. *Int J Infect Dis.* 2017;57:27-31.
- [27] Zhang Z, Chen M, Yu Y, Liu B, Liu Y. In Vitro Activity of Ceftaroline And Comparators against *Staphylococcus aureus* Isolates: Results From 6 Years Of The ATLAS Program (2012 To 2017). *Infect Drug Resist.* 2019; 12:3349-58.

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