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Bacteremia Associated with Pressure Ulcers at Alyarmuk Teaching Hospital in Baghdad

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Abstract

Fifty patients(24 female and 26 male)with pressure ulcersassociated with different diseases and attending AL-yarmouk Teaching Hospital in Baghdad were selected in this study. The duration of sample collection was from March to December 2018. All blood samples collected from patients were submitted to a blood culturing technique to examine bacteremia. The results showed that12 blood bacterial isolates were obtained. The isolated bacteria were subjected to Vitek-2, which is an accurate identification technique. The results of the blood culturing technique revealed that 33.3% were Gram negative bacteria, while 66.6% were Gram positive. Diagnosis by Vitek-2 showed that 33.3% were Staphylococcus spp., 33.3% were Enterococcus spp., 25.1% wereSerratiamarcescens and 8.3% comprised Acinetobacterbaumannii. The results of minimum inhibitory concentration (MIC)by Vitek-2showed that Trimethoprime -Sulfamethazole concentration at 320 µg/ml was the MIC for Acinetobacterbaumanni, while piperacilin, Ticarcillin, and Ticarcillin-Clavulanic acidat 128 µg\ml were the MIC for Serratia marcescens . Acinetobacterbaumanniishowed 100% resistance to all antimicrobial agents, while for the Serratiamarescenceresistancevalues were 54.55%, 54.55%, and 45.45% for isolate numbers 1, 2, and 3, respectively. Gram positive bacteria recorded NitrofurantionMIC of 256 µg/ml against Staphylococcus epidermidisand Enterococcus spp., withboth species showinghigh resistance compared with the others which had a value of 87.50%.

Keywords: Pressure ulcers, bacteremia, Blood culture.

تجرثم الدم المرتبط بمرضى قرح الفراش في مستشفى اليرموك التعليمي في بغداد نهى صلاح جاسم*, سمير عبد الامير علش قسم علوم الحياة, كلية العلوم, جامعة بغداد، بغداد، العراق

الخلاصة

خمسون مريضا تم اختيارهم في هذه الدراسة يعانون من قرح الفراش ومن امراض مختلفة اخرى (24 انثى ذكر و 26) المقيمين في مستشفى اليرموك التعليمي في بغداد. كل عينات الدم التي جمعت من المرضى خضعت لتقنية زراعة الدم للتحري عن تجرثم الدم. اظهرت النتائج الحصول على 12 عزلة بكتيرية. تم تشخيص هذه العزل بواسطة جهاز Vitek-2 وهو جهاز دقيق في التشخيص. كشفت نتائج تقنية زراعة الدم 33.3% بكتريا سالبة لصبغة كرام بينما 66.6% تعود الى البكتريا الموجبة لصبغة كرام. أظهر التشخيص بواسطة Vitek-2. ان نسبة . Staphylococcus بلغت 33.3% وكانت Enterococcus 33.3% كانت 33.3% كانت 25.1% واخيرا سجلت بكتريا Acinetobacter Serratia marcescensوليكترياspp.

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MIC) ان تركيز المثبطة (MIC) ان تركيز المثبطة (MIC) ان تركيز عن تركيز من تركيز المثبطة (Trimethoprime –Sulfamethazole ليكيريا Trimethoprime –Sulfamethazole ليكتيريا تركيز مضادات 128piperacilin, Ticarcillin, Ticarcillin–Clavulanic acidµg\ml ليكتيريا *تركيز مضادات Acinetobacter baumannii* كان انواع المضادات *Serratia marcescens* مقاومة ليكتريا *Acinetobacter baumannii* معاومة لكل انواع المضادات المستخدمة في هذه الدراسة . بينما نسبة المقاومة ليكتريا *Serratia marcescens* كانت ,84.55% *Staphylococcus epidermidis* على التوالي.البكتريا الموجبة لصبغة كرام اظهرت *MIC* 87.50 على الأنواع الأخرى التي كانت ,87.50 *و. Enterococcus spp.*هذان النوعان يظهران مقاومة عالية مقارنة مع الأنواع الأخرى التي كانت .87.50 *ز.*

Introduction

Pressure ulcers (PUs) are an injury to the skin or underlying tissue due to unrelieved pressure [1]. It is a serious health problem for the world, specifically toweakened geriatric or bed-bound patients in hospital [2]. The symptomstypically range from skin redness to serious injuries to thebones or attached tissues, raising a significant threat to patients with restricted mobility [3]. The prevalence of developed pressure sore is high in elderly people, appearing within those between the 70s and 80s decades. These ulcers appear in community setting, nursing homes and hospitals, with an incidence varying from 1.2% to 11.2% [4]. Pressure ulcers are generally followed by an inflammatory response andmostly bylocal bacterial colonization or systemic disease[5].

Risksof pressureulcers are correlated with remarkable morbidity and mortality with bacteria, which are the most prevalent complicating factors related with pressure ulcers[6].

PUs can serve as foci for blood infection as the most prevalent considerable Infected PUs complication. Patients are often more probable to develop bacteremia[7].

The association between PUand bacteremia was related with 50 percent mortality rate in hospitalize d patients[8].Septicaemiaor secondary bacteremia can represent complications of the pressure ulcer where both of these situations are correlated with increased death[9]. Precise identification of bacterial isolates from blood at species level as well as accurate identification of portal of entry and/or thesource of infection are essential for the ideal management of such infections[10].

From 1995 to 2002, a database of a hospital in the United Statesrecognizedcoagulase negative staphylococcus (CoNS) as the most prevalent cause, responsible for 31 percent of cases[11].In recent years, the prevalence of *Acinetobacterbaumannii*bacteremia has increased significantly, particularly in immunocompromised populations and intensive care units[12]. Enterococci have recently become one of the most prevalent nosocomial pathogens, with an elevated mortality rate of up to 61% [13].*Serratiamarcescens* is considered as an opportunistic bacterium that causes a variety of human infections, including keratitis ,bacteremia, as well as urinary tract and wound infections [14].

The aim of this study is to detect bacteremia associated with pressure ulcers, along withtesting MIC values of several antibacterial agents.

Materials and methods

Patients

Fifty patients (24 female and 26 male). 40% of patients between 70 and 80 years were included in this study suffering from pressure ulcer and another disease (30% heart disease, 18% lung disease, 16% kidney disease, 16% diabetic patients and the remaining percent for another disease) all these disease with a pressure ulcer were made the patients bed redden at department of medicine\ AL-yarmouk teaching hospital and the patients diagnosed clinically by a physician for pressure ulcer and bacteremia. The duration of study from the march 2018 to December 2018.

Blood sample collection and bacteremia

The following guidelines were implemented rigidly when samples of blood were obtained for blood culture [15]:

Whenever possible, blood sampleswere taken for culture before antimicrobial therapy was administered.ninemillilitres of blood was injected into a sterile bottle containing brain heart infusion broth culture. The same method was repeated to another blood sampletaken from separate sites over a duration of 10 min. Then, the bottles were incubated for 18-24 hours at 37 ° C. The presence of macroscopic alterations such as haemolysis, turbidity, cotton ball like colonies, and gas bubbles were

screened during the next days.Gram staining was performed irrespective to the macroscopic indications of growth, whileblind subcultures of blood and Macconkyagar were performed after 1,3, and 7 days.

Identification of bacterial isolates

Morphological identification was performed by examining the colonieson different mediaand by gram staining. The precise identification achieved through diagnosis by vitek-2 system.

Antimicrobial screening of bacterial isolates

Antimicrobial screening test wasperformed by using vitek-2 system, with the susceptibility card for Gram positive bacteria was AST-P580 and that for Gram negative bacteria was AST-222. Interpretation of the results wascarried out using the criteria of the Clinical Laboratory Standards Institute (CLSI, 2018) [16].

Statistical Analysis:

The Statistical Analysis System- SAS (2012) program was used to detect the effects of differentfactors onstudy parameters[17]. Least significant difference –LSD- test was used to compare significant differences between means and Chi-square test was used to compare significant differences between percentages.

Resultsand Discussion

From 50 patients with pressure ulcer, 12 samples (24%) were blood culture positive and different types of bacterial isolates were isolated and stained (33.3% Gram negative bacteria and Gram positive 66.6%), as shown in Table-1. Accurate diagnosis by vitek-2 revealed 33.3% *Staphylococcus sp.*, 33.3% *Enterococcus sp.*, 25.1% *Serratia maresscence*, and8.3% *Acinetobacterbaumannii*, as shown in Table-1. Antimicrobial test was performed for bacterial isolates.

	The results of stood culture and recharication s	j viten 2 compact system			
NO.	Blood culture	Vitek-2			
1	33.3% of Staphylococcus sp.	25 % of S.heamolyticus 25 % of S.epidermidis 50 % of S.aureus			
2	33.3% of Enterococcus sp.	50% of <i>E.faecalis</i> 50% of <i>E.gallinarum</i>			
3	25.1% Serratia maresscence				
4	8.3% of Acinetobacterbaumannii				

Table 1-The results of blood culture and identification by Vitek- 2 compact system

The results demonstrated a significant difference between bacterial isolates (P<0.05). The highest minimum inhibitory concentration (MIC) achieved using Trimethoprime-Sulfamethazolewas against *Acinetobacterbaumannii* was 320 µg ml, whereas for *Serratia marescence*, the piperacilin, Ticarcillin and Ticarcillin-Clavulanic acid showed highest MIC (128 µg ml), as shown in Table-2 and Figure-1.

Table 2-Theminimum inhibitory concentration for Gram negative bacteria

Bacterial isolate	Serratia marescence	Serratia marescence	Serratia marescence	Acinetobacter baumannii	
Antimicrobial	MIC µg\ml	MIC µg\ml	MIC µg∖ml	MICµg\ml	
Cefepime	64 (R)	8 (S)	64 (R)	32 (R)	
Ceftazidime	64 (R)	16 (R)	64(R)	64 (R) 4 (R)	
Ciprofloxacine	0.25 (S)	0.25 (S)	0.25 (S)		
Gentamicin	4 (S)	16 (R)	1 (S)	16 (R)	
Meropenem	16 (R)	0.25 (S)	0.25 (S)	16(R)	
Minocycline	8 (I)	4 (S)	8 (I)	16 (R)	
piperacilin	128 (R)	128 (R)	128 (R)	128(R)	
Ticarcillin	128	128	128	128 (R)	
Ticarcillin-Clavulanic acid	128 (R)	128 (R)	128 (R)	128(R)	

Tobramycin	8 (I)	16 (R)	1 (S)	16 (R)			
Trimethoprime- Sulfamethazole	20 (S)	20 (S)	20 (S)	320 (R)			
LSD value	16.38 *	`18.24 *	16.55 **	27.93 *			
* (P<0.05).							

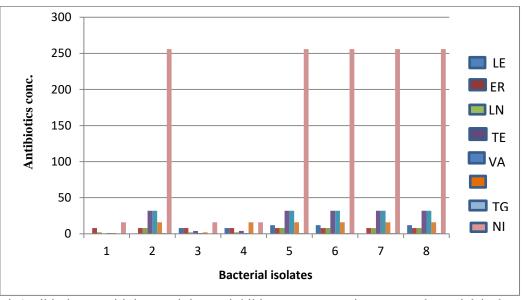


Figure 1-Antibiotics sensitivity to minimum inhibitory concentrations amongbacterial isolates (Gram positive).

1 (*S.aureus*); 2 (S. epidermidis); 3 (*S.heamolyticus* 1); 4 (*S.heamolyticus* 2); 5 (*E.fecalis* 1); 6 (*E.fecalis* 2); 7 (*E.gallinarum*1); 8 (*E.gallinarum*2).Abbervation : Levofloxcin(LEV),Erythromycin(ERY),Linezolid(LNZ), Ticoplanin(TEC), Vancomycin(VAN), Tetracycline(TET), Tigecycline(TGC),Nitrofurantion(NIT).

Acinetobacterbaumanniishowed 100% resistance to all antimicrobial agents used in this study, while the resistance values for the *Serratiamarescence*were54.55%, 54.55%, and 45.45% for isolates number 1,2, and 3, respectively, as shown in Table-3.

	Acinetobacter	Serratia	Serratia	Serratia			
	baumannii	marescence(1)	marescence (2)	marescence(3)			
Resistant (R)	(100%)	(54.55%)	(54.55%)	(45.45%)			
Resistant (R)	(11\11)	(6\11)	(6\11)	(5\11)			
Sensitive (S)	(0.00%)	(27.27%)	(45.45%)	(45.45%)			
Sensitive (S)	(0\11)	(3\11)	(5\11)	(5\11)			
Intermidate (I)	(0.00%)	(18.18%)	(0.00%)	(9.10%)			
Intermidate (I)	(0\11)	(2\11)	(0\11)	(1\11)			
Total	11*	11*	11*	11*			
Chi-Square (χ^2)	15.00 **	9.73 **	12.39 **	11.03 **			
	** (P<0.01).						

Table 3-<u>The</u> resistance percentages for gram negative bacteria

*11 : Is the number of antimicrobial agent used in this study

There was also asignificant difference between bacterial isolates (P<0.05), with the highest MIC usingNitrofurantion against *Staphylococcus epidermidis* being 256 μ g/ml while that for all *Enterococcus spp*.was 256 μ g/ml, as shown in Table-4 and Figure-2.

Bacterial	S.	S.	S.	S.	E.	E.	E.	E.
isolate	aureus	epidermidis	heamolyticus	heamolyticus	faecalis	faecalis	gallinarum	gallinarum
Antimicrobial	MIC µg∖ml	$MIC \ \mu g \backslash ml$	MIC µg∖ml	MIC µg\ml	MIC µg∖ml	MIC µg∖ml	MIC µg∖ml	MIC <u>µg∖ml</u>
Levofloxcin	0.12	0.12	8	8	12	12	0.25	12
	(S)	(S)	(R)	(R)	(S)	(S)	(S)	(S)
Erythromycin	8	8	8	8	8	8	8	8
	(R)	(R)	(R)	(R)	(R)	(R)	(R)	(R)
Linezolid	2	8	2	2	8	8	8	8
	(S)	(R)	(S)	(S)	(R)	(R)	(R)	(R)
Ticoplanin	0.5	32	4	4	32	32	32	32
	(S)	(R)	(S)	(S)	(R)	(R)	(R)	(R)
Vancomycin	1	32	1	1	32	32	32	32
	(S)	(R)	(S)	(S)	(R)	(R)	(R)	(R)
Tetracycline	1	16	2	16	16	16	16	16
	(S)	(R)	(S)	(S)	(R)	(R)	(R)	(R)
Tigecycline	0.12	1	0.5	0.5	1	1	1	1
	(S)	(R)	(S)	(S)	(R)	(R)	(R)	(R)
Nitrofurantion	16	256	16	16	256	256	256	256
	(S)	(R)	(S)	(S)	(R)	(R)	(R)	(R)
LSD value	6.19 *	17.94 *	6.55 *	6.09 *	20.44 *	20.44 *	17.52 *	20.44 *
* (P<0.05).								

 Table 4-The MIC values for Gram positive bacteria

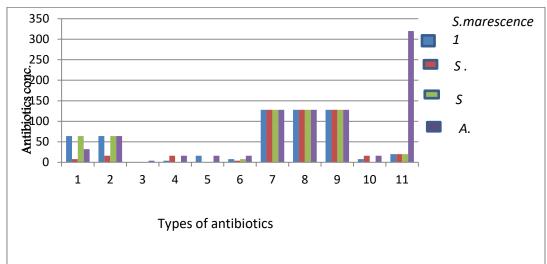


Figure 2-Antibiotic sensitivity variations among Gram negative bacterial isolates (series1 *Serratia marescence*38; series 2 *Serratia marescence*15; series3 *Serratia marescence*7; series4 *Acinetobacterbumannii*). 1-Cefepime;2-Ceftazidime;3-Ciprofloxacine;4-Gentamicin;5-Meropenem;6-Minocycline;7-piperacilin;8-Ticarcillin;9-Ticarcillin-Clavulanic acid;10-Tobramycin;11-Trimethoprime-Sulfamethazole

The bacterial isolate that had the highestbacterial resistance was *S.epidermidis*87.50%, with the same percentage being recorded for *Enterococcus spp.*, as shown in Table-5.

	S. aureus	S. epidermidi s	S. heamolyticu s	S. heamolyticu s	E. faecalis	E. faecalis	E. gallinarum	E. gallinarum	
R	(12.50%) (1\8)	(87.50%) (7\8)	(25.00%) (2\8)	(37.50%) (3\8)	(87.50%) (7\8)	(87.50%) (7\8)	(87.50%) (7\8)	(87.50%) (7\8)	
S	(87.50%) (7\8)	(12.50%) (1\8)	(75.00%) (6\8)	(62.50%) (5\8)	(12.5%) (1\8)	(12.50%) (1\8)	(12.50%) (1\8)	(12.50%) (1\8)	
Ι	(0.00%) (0\8)	(0.00%) (0\8)	(0.00%) (0\8)	(0.00%) (0\8)	(0.00%) (0\8)	(0.00%) (0\8)	(0.00%) (0\8)	(0.00%) (0\8)	
To tal	8	8*	8*	8*	8*	8*	8*	8*	
Chi - Squ are (χ^2)	12.56 **	12.56 **	10.47 **	9.85 **	12.56 **	12.56 **	12.56 **	12.56**	
	** (P<0.01).								

Table 5-The resistance percentage for Gram negative bacteria

*8: Is the numbers of antimicrobial agents used in this study.

Discussion

A study published by Thomas (2006) on 21 sepsis syndrome with a attributable to pressure ulcers revealed that 76% had bacteremia resulted from pressure ulcer [18]. Another study by Braga*et al.*(2017) revealed that, amongsixteen patients with infected pressure ulcer, 62.5% developed bacteremia [19]. These results demonstrated higher proportions of bacteriemic patients than that recorded in our study, which was24%.

The Gram negative bacteria associated with bacteremia:

(Acinetobacterbaumannii andS. marescence)

In our study, *A. baumannii* showed resistance to all antimicrobial agents. This result is corresponding with other studies. A study by Yang *et al.*, (2018) showed that 77.8% of the patients were multidrugresistant [20]. In addition, China's antimicrobial resistance monitoring program has wide ly identified extensive drug resistance to *A. baumannii* (XDRAB)[21].

In another study done by Xuet al., (2016) state that87.7% isolates from bacteremic patients were considered to be XDR[22]. In the same study of Xuet al., (2016) Acinetobacterbaumanniiisolated from blood was resistant to Cefepime, Ceftazidime, Ciprofloxacin, Gentamicin and Tobramycin this results correspond the current study results only the different in the isolatewas intermediately resistant to Meropenem, while in a present study was resistant to it [22].

The abilityof*A*. *baumannii*for the acquisition of genetic resistance determinants is responsible for the development of MDR strains. Other resistance mechanisms include Beta-lactamases, changes in porin canals, efflux pump (responsible for resistance to beta lactam antibiotics), mutations in deoxyribonucleic acid topoisomerase (mediated resistance to quinolone), and genes coding amino-glycoside-modifying enzymes .In addition, oxacillinases and metallo-blactamases (e.g., blaOXA58andblaOXA24\40, blaOXA23,) contribute to the resistance of carbapenem [23].

A retrospective cohort study was also previously performed, were 10 patients with one or more positive blood cultures for *S. marcescens* were recorded in a tertiary care hospital in Seoul, South Korea, from January 2006 to December 2012 [24]. While in the present study, 3 patients with positive blood cultures of *S. marcescens* were recorded for the period from March 2018 to December 2018.

The majority of the isolates in a study by Kim *et al.*, (2015) were susceptible tomeropenem, cefepime, and ceftazidime [24].While in this study, only 2 isolates weresensitive to meropenemsensitive and one isolate was sensitive to cefepime, whereasall isolates were resistant to ceftazidime.

Recent epidemiological analysis demonstrated an increase in the rate of antimicrobial resistance among isolates of *S*. Marcescens. In contrast, the multidrug-resistant (MDR) strains of *S*. *Marcescens*were linked with severe outcomes [25].

The Gram positive bacteria associated with bacteraemia

The most common bacteria associated with pressure ulcers were reported to be *Enterococcus faecalis* and *Staphylococcus aureus*[26].

The incidence of MRSA infections, particularly bacteremia, varies worldwide. In 2014, the proportio n of MRSA isolates in Europe ranged from 0.9% in the Netherlands to 56% in Romania[27].In a study on patients with coagulase-negative staphylococci(CoNS), three cases out of 56 (5.4%) of bacteremia were associated with pressure ulcers [28], whereas the proportion of those withCoNS was 10.09% [29]

Enterococcus spp. was shown to be responsible for 3.6% of bacteremia associated with pressure ulcers[30].Enterococci have recently become one of the most prevalent nosocomial pathogens, with an elevated death rate of up to 61%.Enterococci are reported as the second most cause of urinary tract and wound infections and the third common cause of bacteraemia[31].In the UK,there were 7066 cases reported of bacteremia by *Enterococcus* species in 2005, reflecting an increase of 8% from 2004. *E. Faecalis*was responsible for 63% of these cases,whereas28% were caused by *E. Faecium*. In addition, 80% of all cases were resistant to antibiotics. Also, it was reported that approximately 12 percent of nosocomial infections in the USA are caused by *Enterococcus* species [32].

Conclusions

Pressure ulcer is a serious health problem andbacteremiacould certainly be one of its dangerous complications. Appropriate antibiotic treatment should be selected in order to eradicate the infection associated with pressure ulcer and avoid bacteremia.

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