Role of semi-quantitative bacteriological culture in management of chronic wound infections

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Abstract

Introduction: The role of quantitative microbiology in the management of wound infections is very crucial. In this study semi-quantitative swab technique is used as a simple procedure for assessing the bacterial load of chronic non healing wounds, thereby distinguishing the patients with infected ulcers from colonized one, thereby restricting the use of antibiotics only to the appropriate infected population.

Aim: To assess the role of semi-quantitative bacterial culture of chronic non healing wounds in differentiating between colonised and infected wounds based on bacterial load.

Setting and Design: This is a cross sectional study carried out in the Department of Microbiology in a tertiary care government medical College.

Materials and Methods: Two wound swabs were taken using sterile cotton swabs from 100 patients with chronic non healing ulcer wounds. Semi-quantitative aerobic bacterial culture was done.

Results: Among 100 patients for whom semi-quantitative culture was done, 72 patients showed growth, of which 21 patients had poly-microbial growth making a total of 93 isolates. Out of the total 93 isolates, 73 isolates had a significant count of $\geq 3+$, hence considered as pathogens. The most commonly isolated pathogens were *Staphylococcus aureus* (24.6%) and *Pseudomonas aeruginosa* (24.6%). 55.5% were *Methicillin resistant Staphylococcus aureus* (MRSA). 65% of Enterobacteriaceae were ESBL producers and 15% were AmpC β -lactamase producers.

Conclusion: This study emphasizes the importance and benefits of semi-quantitative bacterial culture technique in differentiating between pathogens and colonisers and enable refinement of antibiotic regimens and thereby curbing antibiotic resistance.

Keywords: Chronic ulcers, Semi-quantitative culture, Bacterial load, Antibiotic resistance, ESBL, MRSA.

Introduction

Chronic non healing wounds pose a significant burden to healthcare systems and also cause morbidity and mortality to patients. In India, the estimate of patients suffering from chronic wounds is around 2.8 million per year and the treatment cost accounts to billions of rupees. A community based epidemiological study of wounds conducted in India, estimated the prevalence rate of chronic wounds to be 4.5 per 1000 population.¹

Wound healing is a complex series of events that should have the capacity to overcome the obstacles and eventually result in wound closure. There are many factors that impede or prevent wound healing. One of the most serious extrinsic factors affecting wound healing is infection.

The quantitative definition of infection has defined as the presence of 10^5 colony forming units per square centimetre or gram of tissue.² It is reported that healing of ulcers were delayed when the bacterial load was more than 10^6 CFU/ml of wound.^{3,4} Considering the concept of critical colonisation,⁴ it is reported that a critical level of bacteria of more than or equal to 10^4 CFU/g of tissue should be achieved to cause wound infection. Hence the role of quantitative microbiology in the management of wound infections as suggested by many studies, is very

crucial, as it can predict the risk of wound infection and the capacity of wound healing.

Nevertheless, since quantification of bacteria by obtaining tissue biopsy is an invasive procedure, non-invasive techniques such as the semi-quantitative swab can be used to obtain reliable information on the bacterial load. And thus can assist the clinicians in differentiating between colonised and infected wounds.

Hence in this study semi-quantitative swab technique was used as a simple procedure for assessing the bacterial profile of wounds that are clinically infected or having a delayed wound healing, thereby distinguishing the patients with infected ulcers from colonized one. This would help the clinicians in the management of chronic wounds and also prevent inadvertent use of antibiotics and restricting the use only to the appropriate infected population.

Materials and Methods

This is a Cross sectional study carried out in the Department of Microbiology in a tertiary care government medical College, from September 2015 to February 2016. 100 patients with chronic non healing ulcer wounds were included in the study.

Two wound swabs were taken using sterile cotton swabs by Levine's technique.⁵ The wound was cleaned

thoroughly with sterile saline and all superficial exudates removed. A 1-cm² area of tissue around the centre of the wound base is sampled using sterile cotton swab for 5 seconds with sufficient pressure to express fluid from within the wound tissue. Direct Gram staining was done with one of the swabs and interpreted as per standard guidelines. The other swab was rolled over an area of 2cm diameter on the first quadrant of the blood agar plate followed by dilution streaking technique in the other 3

quadrants⁶ and also plated onto MacConkey agar plate. Plates were incubated for 18 to 24 hours at 37 °C under aerobic conditions. After 18 to 24 hours of incubation, MacConkey agar and Blood agar plates were observed for growth, colony count and colony morphology. Semi-quantisation of bacterial colonies was done according to the grading procedure based on the number of colonies in each quadrant.⁶

Semi-quantitative grading procedure for bacterial isolates on growth media:⁶

Score	Number of colonies visible in each quadrant			
	1st Quadrant	2nd Quadrant	3rd Quadrant	4th Quadrant
1+	<10			
2+	<10	<10		
3+	>10	>10	>10	
4+	>10	>10	>10	>5

In the present study, grading of 3+ and 4+ are considered as pathogens causing infection and grading ≤ 2+ are considered as either contaminants or colonisers.⁷ Antibiotic sensitivity testing of the pathogens and their resistance pattern was done as per CLSI document M 100-S26.⁸

Results

The results were analyzed statistically using SPSS version 17.0. In our study the age of the patients ranged from 18 to 75 years. The mean and Standard deviation of age (in years) of patients were 46.1 and 14.364 respectively. Majority of the patients were in the age group of 31-60 years. Out of 100 patients 77 were males and 23 were females. Majority of the patients had venous ulcers (56%), followed by diabetic ulcers (22%).

Out of 100 patients for whom semi-quantitative culture was done, 72 patients showed growth and 28 patients had no growth in culture (Table 1). Of the 72 patients who showed growth, 21 patients had polymicrobial growth making a total of 93 isolates. Out of the total 93 isolates, 73 isolates had a significant count of ≥3+, hence considered as pathogens and 20 isolates had semi-quantitative bacterial grading of 2+ and hence considered as either colonisers or contaminants (Table 2). The most commonly isolated pathogens in semi-quantitative culture were *Staphylococcus aureus* (24.6%) and *Pseudomonas aeruginosa* (24.6%) (Table 3).

The antibiotic susceptibility pattern of the *Staphylococcus aureus* in the present study, showed that the isolates were sensitive to Gentamicin (55.5%), Ciprofloxacin (55.5%), Tetracycline (88.8%) and least sensitive to Penicillin (22.2%), Cotrimoxazole (33.3%) and Erythromycin (44.4%). Enterobacteriaceae isolates in the present study, were sensitive to Ciprofloxacin (64%), Tetracycline (72%), Gentamicin (60%), and least susceptible to Cefotaxime 32%) and Cotrimoxazole (24%). All Enterobacteriaceae were highly susceptible to Imipenem (100%). *Pseudomonas aeruginosa* in the present study, were sensitive to Ceftazidime (55.5%), Gentamicin (44.4%), and Ciprofloxacin (33.3%). *Acinetobacter* were sensitive to Ciprofloxacin (33.3%) and Cotrimoxazole (33.3%) All isolates of *Pseudomonas aeruginosa* and *Acinetobacter* were highly susceptible to Imipenem (100%) and Piperacillin-Tazobactam (100%).

Out of 18 *Staphylococcus aureus* isolates, 10 (55.5%) were Methicillin resistant *Staphylococcus aureus* (MRSA) (Table 4). All the MRSA isolates (10) were susceptible to Vancomycin tested by determination of Minimum Inhibitory concentration by Macro broth dilution method. Out of the isolates tested for ESBL production 15 isolates (65%) were found to be ESBL producers and 8(15%) were AmpC β-lactamases producers. Test for MBL and carbapenamases production was not done as all Gram negative bacilli isolates were sensitive to Imipenem (Table 5).

Table 1: Interpretation of semi-quantitative culture results n=100

Semi-quantitative culture	Bacterial grading	No of isolates	No of patients
Growth	4+	32	72
	3+	41	
	2+	20	
No growth	-	-	28
Total	-	93	100

Table 2: Categorisation of ulcers based on semi-quantitative bacterial culture

Semi-quantitative culture	Bacterial grading	No of isolates	Total
Pathogens	4+	32	73
	3+	41	
Colonisers	2+	20	20
Total	-	93	93

Table 3: Bacteriological Profile of semi-quantitative bacterial culture n=93

Type of isolate	Organism	Total no of pathogens	Total no of colonisers	Total
Gram positive cocci	<i>Staphylococcus aureus</i>	18 (24.6%)	6 (30%)	24 (25%)
Gram negative bacilli	<i>Pseudomonas aeruginosa</i>	18 (24.6%)	5 (25%)	23 (24.7%)
	<i>Pseudomonas species</i>	6 (8.2%)	-	6 (6.4%)
	<i>Klebsiella pneumonia</i>	7 (9.5%)	2 (10%)	9 (9.7%)
	<i>Klebsiella oxytoca</i>	6 (8.2%)	1 (5%)	7 (7.5%)
	<i>Acinetobacter baumannii</i>	6 (8.2%)	-	6 (6.4%)
	<i>Escherichia coli</i>	5 (7%)	3 (15%)	8 (8.6%)
	<i>Proteus mirabilis</i>	5 (7%)	2 (10%)	7 (7.5%)
	<i>Proteus vulgaris</i>	2 (2.7%)	1 (5%)	3 (3.2%)
	Total	73 (100%)	20 (100%)	93 (100%)

Table 4: Methicillin resistance among *Staphylococcus aureus* isolates by cefoxitin screening method (n=18)

Gram positive cocci	Resistance pattern	Number of isolates	Percentage
Staphylococcus aureus n=18	MRSA	10	55.5%
	MSSA	8	44.4%

Table 5: Distribution of resistance pattern in Gram negative bacilli

Gram Negative Bacilli (n=55)	Extended-Spectrum Beta Lactamases (ESBL)		AmpC Beta-lactamases	
	N	%	N	%
<i>Escherichia coli</i> (n=5)	3	60	0	0
<i>Klebsiella oxytoca</i> (n=6)	4	66.6	2	3.3
<i>Klebsiella pneumonia</i> (n=7)	4	57.1	1	25
<i>Proteus vulgaris</i> (n=2)	-	-	0	0
<i>Proteus mirabilis</i> (n=5)	4	80	0	0
<i>Pseudomonas aeruginosa</i> (n=18)	-	-	3	16.6
<i>Pseudomonas species</i> (n=6)	-	-	0	0
<i>Acinetobacter baumannii</i> (n=6)	-	-	2	3.3
Total	15		8

Fig. 1: Semi-quantitative culture method

Discussion

Wounds or ulcers are classified as acute or chronic according to the duration they have persisted. However, there is no specific length of time to define chronicity.⁹ Chronic wounds or ulcers come with significant morbidity and cost for the patients and society as a whole. This study was carried out to assess the role of semi-quantitative bacterial culture in differentiating between pathogens and colonisers in chronic wounds thereby aid in the management.

In this study, out of 100 study population, 77 were males and 23 were females. Majority of the patients (25%) were in the age group of 51 to 60 years. According to many studies, it was stated that chronic leg ulcers affect 0.6–3% of patients aged over 60 years, increasing to over 5% of those aged over 80 years. The incidence of ulceration is also rising as a result of aging population and increased risk factors for atherosclerotic occlusion such as smoking, obesity and diabetes.¹⁰

In our study majority of the patients had venous ulcers (56%), followed by diabetic ulcers (22%). This is similar to studies from India that show that etiology of chronic wounds include causes like venous hypertension, arterial insufficiency, neuropathy like diabetes, pressure ulcers, vasculitis or trauma.^{10,11} In another study it had been reported that ulcers related to venous insufficiency constitute 70%, arterial disease 10%, and most of the others are due to either neuropathy (usually diabetic) or a combination of those diseases.⁶

Semi-quantitative Swab Culture: In our present study, 2 wound swabs were taken by Levine's technique from patients with chronic non healing ulcers and semi-quantitation of bacterial colonies were done according to the grading procedure based on the number of colonies in each quadrant. Out of 100 patients with chronic non healing ulcers, 72 patients showed growth and 28 patients had no growth in culture (Table 1). Of the 72 patients who showed growth, 21 patients had polymicrobial growth making a total of 93 isolates.

The interaction between the Wounds and Bacteria can be classified into four Levels: Contamination, colonization, critical colonization and infection. The line to differentiate between colonization and infection had always been difficult to define. The term 'critical

colonization' has been used to describe the stage at which bacteria begin to adversely affect wound healing.¹⁴ A critical threshold of 10^5 bacteria has been proposed as the delineation between colonization and a clinically relevant infection that may impede wound healing.^{15,16}

A correlation between semi-quantitative swab data and quantitative biopsy data has been reported in many studies.^{7,17,18} In semi-quantitative swab technique, 1+ growth correlates with a bacterial count of 10^2 to 10^3 CFU/g and 4+ correlates with a bacterial count of approximately 10^7 CFU/g.^{7,17} Swabs that yield more than 30 CFU indicate a tissue count of 10^5 CFU/g.^{7,18} According to studies semi-quantitative count of $\geq 3+$ was considered significant.⁷

Hence in our study grading of 3+ and 4+ are considered as significant growth causing infection and grade with 1+ & 2+ are considered either as contaminants or colonisers. In our study out of 93 isolates, 73 isolates had a significant count of $\geq 3+$, hence considered as pathogens. 20 isolates had semi-quantitative bacterial grading of 2+ and hence considered as colonisers or contaminants (Table 2).

Thus from the present study, the superficial swab sampling procedure becomes the simple, inexpensive, non-invasive and convenient method than a quantitative biopsy. From a quantitative perspective, this method is beneficial in providing a semi-quantitative estimation of the microbial load and thereby differentiating between pathogens and colonisers or contaminants and also determining the optimal time point for antibiotic treatment. From a qualitative perspective, this method provides the profile of causative bacteria.

Aerobic Bacteriological Profile of Chronic Wounds and their Antibiotic Resistance Pattern: In our study the predominant pathogens isolated by semi-quantitative culture were *Staphylococcus aureus* (24.6%) and *Pseudomonas aeruginosa* (24.6%) (Table 3). The other isolates were *Klebsiella pneumoniae* (9.5%), *Klebsiella oxytoca* (8.2%), *Escherichia coli* (7%), *Proteus mirabilis* (7%), *Proteus vulgaris* (2.7%), and *Acinetobacter baumannii* (8.2%). Six *Pseudomonas* species (8.2%) were isolated out of which 2 was *Pseudomonas fluorescens*, and 4 were *Pseudomonas putida*.

This is similar to the study conducted by Bowler et al, in which *Staphylococcus aureus* is reported to be the common bacterium isolated in traumatic ulcers, surgical site infections and burns ulcers.⁷ In review article by R. S. Howell-Jones et al, *Staphylococcus aureus* has been reported in frequencies varying from 43% of infected leg ulcers.¹⁴ *Pseudomonas aeruginosa* is another frequently identified organism and has been found in 7–33% of ulcers. A number of other aerobic species have also been reported, including *Escherichia coli*, *Klebsiella* species and *Proteus* species.^{14,21}

There are many studies which have investigated the role of microorganisms in wound healing. These studies

have shown that the primary pathogens that cause delayed wound healing and infection are *Staphylococcus aureus*, *Pseudomonas aeruginosa*, and *beta-hemolytic Streptococci* as they are notorious for their capacity to produce dangerous destructive virulence factors.^{7,19} Due to the chronicity of the wound, *Methicillin Resistant Staphylococcus aureus* (MRSA) like organisms may colonise the wound and cause wound infection⁽²⁰⁾ which is the most common pathogen in our study too.

Antibiotics are frequently used in the management of chronic wounds and these patients receive significantly more antibiotic prescriptions both as systemic and topical application. Antibiotic resistance in bacteria has thus become a concerning issue and the control of infections is extremely difficult. As a result, there has been a great interest to detect antibiotic resistance of the causative bacteria.¹⁴

The antibiotic susceptibility pattern of the *Staphylococcus aureus* in the present study, showed that the isolates were sensitive to Penicillin (22.2%), Erythromycin (44.4%), Gentamicin (55.5%), Ciprofloxacin (55.5%), and Tetracycline (88.8%). Enterobacteriaceae isolates in the present study, were sensitive to Ciprofloxacin (64%), Tetracycline (72%), Gentamicin (60%), and least susceptible to Cefotaxime (32%), Cotrimoxazole (24%). All Enterobacteriaceae were highly susceptible to Imipenem (100%). *Pseudomonas aeruginosa* in the present study, were sensitive to Ceftazidime (55.5%), Gentamicin (44.4%), and Ciprofloxacin (33.3%). *Acinetobacter* was sensitive to Ciprofloxacin (33.3%) and Cotrimoxazole (33.3%) All isolates of *Pseudomonas aeruginosa* and *Acinetobacter* were highly susceptible to Imipenem (100%) and Piperacillin-Tazobactam (100%).

In the present study out of 18 *Staphylococcus aureus* isolates, 10(55.5%) were *Methicillin resistant Staphylococcus aureus* (MRSA) (Table 4). All the MRSA isolates (10) were susceptible to Vancomycin tested by determination of Minimum Inhibitory concentration by Macro broth dilution method. Out of the isolates tested, 15 isolates (65%) were found to be ESBL producers and 8(15%) were AmpC β -lactamases producers. Test for MBL resistance and carbapenemase production was not done as all Gram negative bacilli isolates were sensitive to Imipenem (Table 5).

The above antimicrobial susceptibility pattern in this study is similar to many studies conducted to analyse the incidence of antimicrobial resistance of bacterial isolates causing wound infection and delayed wound healing.

According to Frankel et al,²² the incidence of MRSA was 45% among patients with chronic wounds. Colsky et al²³ found 50% of *Staphylococcus aureus* isolates from hospitalized patients with leg ulcers to be MRSA. A study by Tentolouris et al²⁴ in a diabetic foot clinic, found 40% of *Staphylococcus aureus* isolated from patients with infected foot ulcers to be MRSA. But in another study²⁵ it was found that only 12% of

Staphylococcus aureus to be MRSA in infected diabetic ulcer foot. They have also found high levels of resistance to erythromycin in *Staphylococcus aureus*.

In study conducted by Nyambura Moremi et al,²⁶ 41% were ESBL producers among the Enterobacteriaceae isolates. Mathur et al^{27,28} found out that 68.0% of Enterobacteriaceae were ESBL producers, whereas Kumar et al^{27,29} reported only 19.2% of *E. coli* isolates as ESBL producers. The resistance pattern of isolates in our study is also similar to the study conducted by Umadevi et al^{27,30} in which 65.5% of *S aureus* were MRSA positive and 56% of Enterobacteriaceae were ESBL producers, among which 62.5% of *Proteus* species, 60% of *Klebsiella pneumonia* and 56% of *Escherichia coli* were ESBL producers. It was also found that Amikacin, Piperacillin-tazobactam, Imipenem were sensitive against gram-negative bacilli, while Vancomycin was sensitive against gram-positive bacteria.

In the study by Citron et al^{27,31} all aerobic gram-positive organisms were susceptible to vancomycin like in our study and also *Pseudomonas aeruginosa* strains and the Enterobacteriaceae group were also largely susceptible to imipenem, piperacillin-tazobactam. The isolates were least susceptible to amoxicillin-clavulanate, doxycycline, and cephalexin. The study by Murugan et al^{27,32} also showed that *E. coli* exhibited 100% susceptibility to imipenem and meropenem and resistant to cephalexin, erythromycin, gentamycin. An Indian study on soft tissue infections documented 72.37% ESBL producers among *E. coli*. It was found that 36% of Gram negative bacilli were ESBL producers and AmpC was seen among enteric GNB (10%) and *Acinetobacter* species (18%).³³

Conclusion

Quantitative bacterial cultures become essential part of management of chronic wounds. According to our study semi quantitative bacterial culture using superficial swab technique can be used as a simple procedure for assessing the bacterial load and the bacterial flora of wounds that are clinically infected or having a delayed wound healing, thereby differentiate between pathogens and colonisers. Semi-quantitative swabs are thus beneficial for clinicians who want to adjust their wound care based on bacterial bio-burden and thus prevent inadvertent use of antibiotics causing increased antibiotic resistance.

References

1. Gupta N, Gupta SK, Shukla VK, Singh SP. An Indian community-based epidemiological study of wounds. *J Wound Care*. 2004;13(8): 323-5.
2. Terry Swanson, David Keast, Rose Cooper, Joyce Black, Donna Angel, Greg Schultz, Keryln Carville, Jacqui Fletcher. Ten top tips: identification of wound infection in a chronic wound. *Wounds International* 2015;6(2).
3. Bendy, R. H., P. A. Nuccio, E. Wolfe, B. Collins, C. Tamburro, W. Glass, and C. M. Martin. Relationship of

- quantitative wound bacterial counts to healing of decubiti. Effect of topical gentamicin. *Antimicrob. Agents Chemother.* 1964;10:147-55.
4. Breidenbach, W. C., S. Trager. Quantitative culture technique and infection in complex wounds of the extremities closed with free flaps. *Plast Reconstr Surg.* 1995(5):860-65.
 5. Paul Drinka, Phyllis Bonham, Christopher J. Crnich. Controversies in Long Term Care Swab Culture of Purulent Skin Infection to Detect Infection or Colonization with Antibiotic-Resistant Bacteria. *JAMDA.* 2012;(1):75-9.
 6. Patricia M. Tille. *Bailey and Scott's Diagnostic Microbiology*, 13 ed. USA:2014.
 7. P. G. Bowler, B. I. Duerden, D. G. Armstrong. Wound microbiology and associated approaches to wound management. *Clin Microbiol Rev.* 2001;14(2):244-69.
 8. Clinical and Laboratory Standards Institute. Performance standards for Antimicrobial susceptibility testing; CLSI document M100-S26. 26th ed. CLSI, Wayne. Pennsylvania USA. 2016;36(1).
 9. Manish Suthar, Saniya Gupta, Suhail Bukhari, Venkatesh Ponemone. Treatment of chronic non-healing ulcers using autologous platelet rich plasma: a case series. *J Biomed Sci.* 2017;24(16).
 10. Shubhangi Vinayak Agale. "Chronic leg ulcers; Epidemiology, Aetiopathogenesis, and Management. *Ulcers* 2013; 2013():
 11. V. K. Shukla, M A Anssari, S K Gupta. Wound healing research; a perspective from India. *Internationa Journal of Lower Extremity Wounds.* 2005;4(1):7-8.
 12. P K Sarkar, S Ballantyne. Management of leg ulcers. *Postgraduate Medical journal* 2000; 76(901): 674-682.
 13. M.C. Moloney, P. Grace. Undersatanding the underlying causes of chronic leg ulceration. *Journal of wound care.* 2004;13(6):215-218.
 14. R. S. Howell-Jones, M. J. Wilson, K. E. Hill, A. J. Howard, P. E. Price, D. W. Thomas. A review of the microbiology, antibiotic usage and resistance in chronic skin wound. *Journal of Antimicrobial Chemotherapy.* 2005;55(2):143-149.
 15. George Han, Roger Ceilley. Chronic Wound Healing: A Review of Current Management and Treatments. *Adv Ther.* 2017;34(3):599-610.
 16. Trengove NJ, Stacey MC, McGechie DF, Mata S. Qualitative bacteriology and leg ulcer healing. *J Wound Care.* 1996;5(6):277-80.
 17. Thomson, P. D, D. J. Smith. What is infection?. *Am. J. Surg.* 1994;167(1A):7S-11S.
 18. Lawrence, J. C. The bacteriology of burns. *J. Hosp. Infect* 1985; 6(2):3-17.
 19. Heggers J.P. Defining infection in chronic wounds: methodology. An historical review of the quantitative assessment of microbial flora in wounds. *J. Wound Care* 1998;7():452-456.
 20. Halcón L, Milkus K. Staphylococcus aureus and Wounds: a Review of Tea Tree Oil as a Promising Antimicrobial. *American Journal of Infection Control.* 2004; 32(7): 402-8.
 21. Hansson, C., Hoborn, J., Mo ¨ller, A. The microbial flora in venous leg ulcers without clinical signs of infection. *Acta Dermato-Venereologica.* 1995;75(24-30).
 22. Frankel YM, Melendez JH, Wang N, Price LB, Zenilman JM, Lazarus GS. Defining Wound Microbial Flora: Molecular Microbiology Opening New Horizons. *Arch Dermatol.* 2009;145(10):1193-1195
 23. Colsky, A. S., Kirsner, R. S. &Kerdel, F. A. Analysis of antibiotic susceptibilities of skin wound flora in hospitalized dermatology patients. The crisis of antibiotic resistance has come to the surface. *Archives of Dermatology* 1998; 134(): 1006-9.
 24. Tentolouris, N., Jude, E. B., Smirnof, I. Methicillin resistant Staphylococcus aureus: an increasing problem in a diabetic foot clinic. *Diabetic Medicine.* 1999; 16():767-71.
 25. Ge, Y., MacDonald, D., Hait, H. Microbiological profile of infected diabetic foot ulcers. *Diabetic Medicine* 2002; 19(): 1032-5.
 26. Moremi, N., Mushi, M. F., Fidelis, M., Chalya, P., Mirambo, M., &Mshana, S. E. Predominance of multi-resistant gram-negative bacteria colonizing chronic lower limb ulcers (CLLUs) at Bugando Medical Center. *BMC Research Notes.* 2014;7():211.
 27. Mohammad Zubair, Abida Malik, Jamal Ahmad. Microbiology of diabetic foot ulcer with special reference to ESBL infections. *American Journal of Clinical and Experimental Medicine.* 2015; 3(1):6-23.
 28. Mathur P, Tatman A, Das B, Dhavan B. Prevalence of ESBL gram negative bacteria in a tertiary care hospital. *Indian J Med Microbiol.* 2002;115:153-157.
 29. Kumar MS, Lakshmi V, Rajagopalan R. Occurrence of extended spectrum beta lactamases among Enterobacteriaceaespp isolated at a tertiary care institute. *Indian J Med Microbiol.* 2006;24(3):208-11.
 30. Umadevi S, Kumar S, Joseph NM, Easow JM, Kandhakumari G, Srirangaraj S, Raj S, Stephen S. Microbiological study of diabetic foot infections. *Indian Journal of Medical Specialities.* 2011;2(1):12-17.
 31. Citron DM, Goldstein EJC, Merriam CV, Lipsky BA, Abtamson MA. Bacteriology of moderate-to-severe Diabetic foot infections and in vitro activity of antimicrobial agents. *J. Clin Microbio.* 2006;45(9):2819-2829.
 32. Murugan. S, R. Bakkiya Lakshmi, P.Uma Devi and K.R. Mani (). Prevalence and Antimicrobial Susceptibility pattern of Metallo beta lactamase producing Pseudomonas aeruginosa in Diabetic Foot Infection. *International Journal of Microbiological Research.* 2010;1(3):123-128.
 33. Wilson R. Upward trend in acute anaphylaxis continued in 1998-9. *British Medical Journal.* 2000;321, 1021.
 34. Anusha Gopinathan, Chiranjay Mukhopadhyay, Vandana K. E. Characterization of antimicrobial resistance mechanisms of multidrug resistant Gram negative bacterial wound infections and their clinical epidemiology from a tertiary care hospital in Karnataka. *India Int J Res Med Sci.* 2017;5(3):824-828.
 35. Aerden D, Bosmans I, Skin grafting the contaminated wound bed; reassessing the role of pre-operative swab *J wound care.* 2013;22: 85-9.
 36. Ryan J. Donegan. An overview of factors maximizing successful split thickness skin grafting in diabetic wounds. *Diabetic Foot & Ankle.* 2014;5:24769.

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