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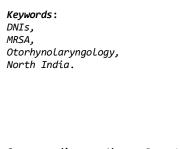
An Observational Study of Community-Acquired MRSA from Head and Neck Space Infections in Northern India

Binod Kumar Pati^{*3}, S Krishna Prakash¹, A K Agarwal², B Uppal¹ and A Gulati²

¹Department of Microbiology, Maulana Azad Medical College and Lok Nayak Hospital, New Delhi, 110002, India ²Department of Otorhinolaryngology, Maulana Azad Medical College and Lok Nayak Hospital, New Delhi, 110002, India ³Department of Microbiology³ Sanjay Gandhi Post Graduate Institute of Medical Sciences, Lucknow, 226014 U.P. India

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Corresponding author: Department of Microbiology, Sanjay Gandhi Post graduate institute of medical sciences, Rae bareli Road, Lucknow, India, PIN - 226014.

E-mail address: drbkpati@gmail.com

ABSTRACT

Background: Deep neck infections (DNIs) continues to be encountered in daily clinical practice even in the settings of widespread antibiotic use. We investigated the microbiology of deep neck infections and antibiogram profiles of *Staphylococcus aureus* isolates.

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Methods: Isolation and characterization of *Staphylococcus aureus* strains were performed from pus samples aspirated from clinically suspected patients of DNIs attending the Otorhinolaryngology department of a tertiary care health setting in northern India.

Results: *Staphylococcus aureus* was the most commonly recovered agent, out of which 11.5% were methicillin-resistant (MRSA).

Conclusion: Community-acquired MRSA (CA-MRSA) from deep neck infections is an issue of grave concern and emphasizes the need for culture-directed clinical decisions and appropriate choice of antibiotics.

Introduction

Deep neck infection is defined as a suppurative infectious process of the neck that often starts as a soft tissue cellulitis and eventually leads to abscesss¹. It could be lymphadenitis, cellulitis, necrotic node or abscess in nature². *Staphylococcus aureus* (*S. aureus*) is one of the commonest gram positive aerobic bacteria in DNIs besides the Streptococci. Antibiotic resistance is a matter of grave concern. While it is well-known that MRSA infections have become

endemic in hospitals, and usually afflict patients with established risk factors, more recently MRSA infections have also been described in patients without established risk factors and these infections involving the community are referred to as communityacquired MRSA (CA-MRSA) infections³. A significant morbidity and mortality is attributed to it. In one study⁴ a mortality of 5.9% and 6.2% was demonstrated in adults and children respectively. Monitoring the



British Biomedical Bulletin spread of MRSA by epidemiological surveillance, especially in DNIs should be considered seriously to initiate aggressive treatment strategies for efficient management^{3,5}.

Whilst the microbiology of DNI varies depending on certain underlying factors and geographic locations, the of aetiology DNIs generally is polymicrobial⁶. While aerobic bacteria are commonly isolated, the presence of anaerobes may be underestimated because of the difficulty in culturing them⁷. We investigated the antimicrobial susceptibility pattern of S. aureus isolated from DNIs from a tertiary care clinical setting in northern India

Materials and Methods

After taking informed consent, pus collected specimens were from 56 consecutive clinically suspected cases of DNIs in a tertiary care hospital. Patients with 48 hours prior history of antibiotic treatment and cases of tuberculosis were excluded. Samples were promptly transported to Microbiology department and processed within 30 minutes of collection. Both aerobic and anaerobic cultures were attempted. Plates incubated aerobically were read the next day and cultures reporting no growth, were incubated further with subcultures made from BHI broth. Anaerobic cultures were examined after 48 hours and in case of no growth on plates, subcultures were made from cooked meat broth and subsequently incubated anaerobically for another 48 hours. Bacterial isolates from cultures were identified and characterized following standard techniques. Grouping of the β -haemolytic streptococci isolates performed bv were latex agglutination kit of Plasmatec ltd (UK).

All the isolates were subjected to antimicrobial susceptibility testing to a wide battery of antibiotics by disc diffusion method using modified Stokes' technique⁸. Isolates of *E. coli* and *K. pneumoniae* resistant to one or more of the third generation cephalosporins were subjected to detection of extended-spectrum β -lactamase (ESBL) using three established methods, namely double-disk synergy, phenotypic confirmatory combined disc test and ESBL E-test⁹. All the *S. aureus* isolates were tested for methicillin resistance by standard oxacillin screening agar test and cefoxitin disc method^{10,11}.

The *S. aureus* isolates were phage typed employing the conventional set of phages described by Blair and Williams¹². The MRSA isolates were further biotyped based on Tween-80 hydrolysis, urease production, pigmentation and gentamicin susceptibility and phenotyped based on their antibiogram^{13,14}.

Results

A total of 53 bacteria were isolated. S. aureus was the most common isolate accounting for 49% of all the bacterial isolates. 11.5% of these isolates were resistant to methicillin. While methicillinsusceptibile isolates were recovered from all the six neck spaces investigated, the MRSA isolates primarily confined to the sites of infection namely, the submandibular, submental and peritonsillar spaces. Of the 12 isolates of β -hemolytic streptococci, eight were isolated in pure growth while four occurred in conjunction with S. aureus. Among them 09 belonged to Group A, two to Group F and one belonged to Group G streptococcus. Other aerobic pathogens isolated were Enterococcus fecalis (1), Gemella morbillorum (1), Streptococcus mitior oralis (1), Escherichia coli (2), Klebsiella pneumoniae (2), Pseudomonas Anaerobic aeruginosa (1). spectrum constituted 13.2% of the bacterial aetiology and of the seven anaerobes recovered, Peptostreptococcus species were



predominant (85.7%) followed by *Bacteroides* species.

D-test performed with a single clinical isolate of MRSA tested positive confirming the presence of erm-mediated resistance to clindamycin. MIC testing by Etest revealed that, all the methicillinsusceptible S. aureus (MSSA) isolates were uniformly sensitive vancomycin, to teicoplanin, mupirocin and fusidic acid. All the three MRSA isolates were sensitive to vancomycin, teicoplanin and mupirocin. Whilst two isolates were sensitive to fusidic acid, one showed high level resistance at $>256 \mu g/mL$. The comparative antibacterial susceptibility profile of MRSA and MSSA is listed in table1.

The 26 *S. aureus* isolates subjected to phage typing by the conventional (basic) set of phages, the three MRSA isolates were found to be non-typeable, whereas 14 (60.9%) MSSA isolates were typeable. An overall typeablilty of 53.9% was observed for the staphylococci isolates. Maximum typeablity was observed in the mixed group of phages (39.1%) followed by Group I (13.1%). None of the isolates were typeable by phages belonging to group II and the non-allocated group. The various phage patterns seen within each group for the typeable isolates of MSSA are shown in Table 2.

Biotyping of MRSA isolates showed that all the 3 MRSA isolates belonged to biotype A. Based on the antibiogram pattern, the 3 different isolates of MRSA showed 3 distinct mnemonic codes, the resistant phenotypes namely Bc (33.3%), Da (33.3%) and Ag (33.3%).

All the 12 isolates of β -hemolytic streptococci isolated irrespective of their serogroups and both the α -hemolytic isolates were susceptible to all the antibiotics tested. All the *E. coli* and *K. pneumoniae* showed resistance to all third-generation cephalosporins tested and were also tested positive for the presence of extendedspectrum β -lactamases (ESBL) by all the methods employed herein. Both the *E. coli* and *Klebsiella* isolates, that were ESBL positive were subjected to MIC testing and the results showed that all the isolates were sensitive to Imipenem and Meropenem.

Discussion

As with previous reports, Gram positive cocci (GPC) were the most commonly recovered bacteria followed by GNB^{3,5,15,16} in our study. Recovery of ESBL producing E. coli and K. pneumoniae from aspirated pus underscores the magnitude of resistance demonstrated by these isolates and the significance of prescribing ESBL inhibitor group of antibiotics to the afflicted population. Evidence that MRSA is a growing clinical problem has been very well documented in the literature with increasing frequency over the past decade and more recently the increasing incidence of CA-MRSA infections has been the source of grave concern. All of the MRSA cases reported in our study were communityacquired. The increasing incidence of CA-MRSA infections has important clinical implications while approaching a patient presenting with head and neck space infections since empirical treatment is advised before culture results are available and now it is clear that careful consideration should be given to cover MRSA^{3,5}. Thus, it becomes the responsibility of physicians to be aware of the prevalence of CA-MRSA in communities and its antimicrobial susceptibility patterns prescribe and empirical treatment accordingly in cases of deep neck infections. Hence, we suggest that CA-MRSA be considered a potential challenge in head and neck space infections and a high index of suspicion and aggressive treatment is believed to be the prime requisite to prevent untoward complications.



An essential issue surrounding the use of clindamycin for the treatment of MRSA infections is the probable risk of treatment failure if the infection is by erythromycin-resistant *S. aureus* with the potential for selecting for clindamycin resistance⁵. Thus, determination of presence of erythromycin-inducible clindamycin resistance by D-test becomes arterial and one of the MRSA isolates tested positive for D-test emphasizes the need for constant monitoring of the clinical isolates.

Conclusion

This study supports the notion that CA-MRSA infections are an important part diagnosis of the differential when approaching a patient with head and neck space infection. It is essential to consider the rising incidence of MRSA when choosing the antibiotic while waiting for culture and susceptibility results. These days many alternative antibacterial strategies are also under trial leeding to a re-evaluation of the therapeutic use of ancient remedies, such as plants and plant-based products like Nigella sativa¹⁷. Cultures are therefore very critical in the diagnosis and management of head and neck space infections.

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Conflict of interest

No financial support was received for this work. There is no conflict of interest to be declared by the authors.

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	MSSA isolates (n=23)	MRSA isolates (n=3)	
Antibiotics	Resistance (%)	Resistance (%)	
Penicillin	100	100	
Cefazolin	0	100	
Cephalexin	4.3	100	
Gentamicin	4.3	0	
Tobramycin	8.6	0	
Netilmycin	0	0	
Amikacin	0	0	
Chloramphenicol	0	0	
Rifampicin	0	33.3	
Tetracycline	0	33.3	
Ofloxacin	26	66.6	
Ciprofloxacin	56.5	66.6	
Cotrimoxazole	69.6	100	
Erythromycin	0 33.3		
Clindamycin	0	0	
Fosfomycin	0 33.3		
Fusidic Acid	0	0	
Dalfopristine / Quinupristine	0 0		
Vancomycin	0	0	
Teicoplanin	0	0	
Linezolid	0	0	

Table 1. Antimicrobial resistance phenotypes of isolates of methicillin-sensitive and resistant staphylococci from pus aspirates of Deep Neck Infections

Table 2. Phage patterns among the 14 typeable isolates of methicillin-sensitive staphylococci.

Phage group	No. of isolates	Phage pattern *	
I	3	29/52/52A/79/80	(3)
II	0	-	-
III	2	6/42E/47/53/54	(2)
Non -allocated group	0	-	-
Mixed group	9	29/52/52A/79/80/84/96 29/52/52A/79/80/6/42E/47 29/52/52A/79/80/6/47/53 29/52/79/80/6/42E/47/53 29/52/52A/79/80/6/42E/47/53/54 3A/3c/6/42E/47/53/54/77/84/94/96	(1) (1) (1) (1) (4) (1)
Total		14	

* Numbers in parenthesis indicate the number of isolates demonstrating a particular phage Pattern.

