



MICROBIOLOGICAL SPECTRUM OF DACROCYSTITIS IN A TERTIARY CARE TEACHING HOSPITAL.

microbiology

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ABSTRACT

Introduction: The aim of the study is to determine microbiological spectrum of acute and chronic dacryocystitis and their antibiotic susceptibility pattern in a tertiary care teaching hospital.

Materials and methods: This prospective observational study was carried out in the department of microbiology and ophthalmology of a tertiary care teaching hospital. 205 were patients diagnosed with dacryocystitis and categorised into acute and chronic dacryocystitis. Specimens for microbiological analysis were collected under strict aseptic precautions from dacryocystitis. All specimens were subjected to standard microbiological procedures. methicillin resistant Staphylococcus aureus (MRSA) and extended spectrum of beta lactamases (ESBL) strains were also detected as per CLSI guidelines.

Results: A total of 205 specimens were collected from cases of dacryocystitis. 55(26.83%) specimens were from acute dacryocystitis and 150 (73.17%) specimens were from chronic dacryocystitis. Culture positivity was high in the specimens collected from chronic dacryocystitis (110/150; 73.33%). Overall Staphylococcus aureus was the predominant pathogen found in both acute 9(29.03%) and chronic dacryocystitis 27(24.55%). Among Gram negative bacilli, H. influenza 4(12.90%) was found to be the predominant pathogen in specimens collected from acute dacryocystitis and preponderance Pseudomonas aeruginosa 9(8.19%) was revealed in specimens collected from chronic dacryocystitis. 33.33% were found to be methicillin resistant Staphylococcus aureus (MRSA) in acute dacryocystitis. In chronic dacryocystitis 40.74% strains of S. aureus were found to be MRSA. Out of 8 Gram negative bacilli, E. coli (25%) was found to be ESBL producers from cases of acute dacryocystitis. Among 39 isolates of Gram negative bacilli, E. coli (7.62%) and Klebsiella species (10.26%) were found to be ESBL producers in chronic dacryocystitis.

Conclusion: Staphylococcus aureus was the predominant organism isolated from acute and chronic dacryocystitis. Gram positive cocci from both acute and chronic dacryocystitis were highly susceptible to cefazolin and vancomycin. Efficacy of amikacin was high towards Gram positive and Gram negative bacteria.

KEYWORDS

Acute dacryocystitis, Chronic dacryocystitis, MRSA.

Introduction:

Dacryocystitis is an inflammation of the lacrimal sac, which usually occurs because of obstruction of the nasolacrimal duct. 1 Dacryocystitis might present in two forms, acute and chronic dacryocystitis. Acute dacryocystitis is an acute inflammation of the lacrimal sac secondary to nasolacrimal duct obstruction and possibly bacterial overgrowth in the stagnant fluid in the lacrimal sac. There is a varied spectrum of its clinical presentations ranging from tenderness and erythema of the overlying tissues to a frank lacrimal abscess. 2 Chronic dacryocystitis is more common than acute dacryocystitis and has several stages of presentation like epiphora, mucoid discharge, conjunctival hyperaemia and chronic conjunctivitis. 2

It is an important cause of ocular morbidity in India. It is the most common cause of epiphora and may present with or without mucopurulent discharge. 3 Untreated lacrimal abscess can progress to orbital cellulitis, superior orbital vein thrombosis, and cavernous sinus thrombosis. 4 The studies have shown that microbial spectrum differs in chronic and acute dacryocystitis. Most articles in the literature have discussed the microbiologic spectrum of chronic dacryocystitis and very few dealt with acute dacryocystitis. 5

There are distinct patterns of geographical variation in terms of etiology. Knowledge on microbial spectrum of dacryocystitis in a particular region is essential. In recent years, there has been an increase in the incidence and the prevalence of multi drug resistant bacteria such as methicillin resistant Staphylococcus aureus (MRSA) and extended spectrum of beta lactamases (ESBLs). Currently, there is a paucity of data on multi drug resistant bacteria involved in dacryocystitis, especially in this region.

Hence, we aimed to focus light on microbial spectrum of acute and

chronic dacryocystitis along with their antibiotic susceptibility pattern.

Materials and methods:

This prospective observational study was conducted in the department of microbiology and ophthalmology, Vinayaka missions medical college, Karaikal from March 2015 to June 2017. All the patients diagnosed with dacryocystitis were categorised into acute and chronic dacryocystitis. Acute dacryocystitis was diagnosed in patients with pain, redness, and swelling in the lacrimal sac area. Patients with persistent epiphora and regurgitation of mucopurulent material on pressure over the sac area or during irrigation of the lacrimal drainage system are diagnosed as chronic dacryocystitis.

Specimens for microbiological analysis were collected under strict aseptic precautions. In cases of acute dacryocystitis, along with conjunctival swab, pus discharge following spontaneous bursting of abscess and/or following incision and drainage was also taken. In cases of chronic dacryocystitis, those cases with mucoid or mucopurulent discharge on syringing of the lacrimal sac. 6

Inclusion criteria:

Clinically diagnosed cases of acute and chronic dacryocystitis were included in the study.

Exclusion criteria:

Patients who had antibiotics in past seven days were excluded.

All the specimens collected from patients were inoculated on aerobic and anaerobic bacteriological media (5% sheep blood agar, chocolate agar, MacConkey agar and thioglycollate broth). And also inoculated on Sabouraud's dextrose agar and processed according to standard microbiological procedures. 7

MRSA and ESBL strains were also detected as per CLSI guidelines, 20148.

MRSA detection:

The phenotypic test for the detection of MRSA was done by using a cefoxitin (30 µg) disc. A zone of inhibition which was equal to or more than 22 mm was considered as susceptible to Cefoxitin and the organism was reported as Methicillin Sensitive *Staphylococcus aureus*. Those isolates which produced a zone of inhibition less than or equal to 21 mm were considered as Methicillin Resistant *Staphylococcus aureus* (MRSA).

Quality control strains:

Methicillin sensitive *S. aureus* (MSSA) ATCC 25923
Methicillin resistant *S. aureus* (MRSA) ATCC 43300

ESBL detection:

ESBL production was confirmed by using discs of Ceftazidime (30 µg) and Ceftazidime Clavulanic acid (30/10µg) respectively. The test organism was inoculated as a lawn on a Mueller Hinton agar plate and the above mentioned discs were placed on the plate. The plates were incubated at 37°C overnight and they were examined next day. An increase in the zone diameter, which was equal to or more than 5 mm for the antimicrobial agent which was tested in combination with clavulanic acid, in comparison to the antimicrobial which was tested alone, indicated that the strain was an ESBL producer.

Quality control strains:

K. pneumoniae ATCC 700603 (ESBL positive control)
E. coli ATCC 25922 (ESBL negative control)

Results:

A total of 205 specimens were collected from cases of dacryocystitis. 55 (26.83%) specimens were from acute dacryocystitis and 150 (73.17%) specimens were from chronic dacryocystitis. 121(85.82%) specimens yielded monomicrobial growth and 20(14.18%) specimens yielded polymicrobial growth. Culture positivity was high in the specimens collected from chronic dacryocystitis(110/150;73.33%) compared to acute dacryocystitis(31/55;56.36%). Out of 205, female patients were 167(81.46%) and male patients were only 38(18.54%) in both acute and chronic dacryocystitis. Majority of patients in our study were between the age group of 30-55 years (87%). Overall 31 organisms isolated from cases of acute dacryocystitis. *Staphylococcus aureus* 9(29.03%) was the predominant bacteria isolated. *Streptococcus pneumoniae* 7(22.58%) and *S.epidermidis*(22.58%) were the second commonest pathogens grown. *H.influenzae* 4 (12.90%) was the common pathogen isolated among Gram negative bacilli. *Escherichia coli* 2 and *Pseudomonas aeruginosa* 2 both together accounted for (12.90%).Table.1. In acute dacryocystitis no anerobic bacteria and no fungi were isolated.

Table 1: Organisms isolated from acute dacryocystitis

Organism	Number (%)
<i>Stahylococcus aureus</i>	9(29.03%)
<i>Streptococcus pneumonia</i>	7(22.58%)
<i>Stahylococcus epidermidis</i>	7(22.58%)
<i>Haemophilus influenzae</i>	4(12.90%)
<i>Escherichia coli</i>	2(6.45%)
<i>Pseudomonas aeruginosa</i>	2(6.45%)
TOTAL	31(100%)

Again *S.aureus* 27(24.55%) was the predominant pathogen isolated from chronic dacryocystitis.This was followed by *Stahylococcus epidermidis* 21(19.09%) and *Streptococcus pneumoniae*17(15.45%). In chronic dacryocystitis, among Gram negative bacilli *Pseudomonas aeruginosa* 9(8.19%) was frequently isolated pathogen. Three specimens collected from chronic dacryocystitis, anaerobic bacteria *Bacteroides*(2.73%) were isolated. Fungal pathogens, *Candida* 2(1.82%) and *Aspergillus* 1(0.10%) were also isolated from chronic dacryocystitis. Table.2

Table 2: Organisms isolated from chronic dacryocystitis

Organism	Number(%)
<i>Stahylococcus aureus</i>	27(24.55%)
<i>Stahylococcus epidermidis</i>	21(19.09%)
<i>Streptococcus pneumonia</i>	17(15.45%)

<i>Pseudomonas aeruginosa</i>	9(8.19%)
<i>Escherichia coli</i>	7(6.36%)
<i>Haemophilus influenza</i>	7(6.36%)
<i>Klebsiella species</i>	6(5.45%)
<i>Enterobacter</i>	5(4.54%)
<i>Proteus</i>	3(2.73%)
<i>Citrobacter</i>	2(1.82%)
<i>Bacteroides</i>	3(2.73%)
<i>Candida</i>	2(1.82%)
<i>Aspergillus</i>	1(0.10%)

Table 3: Antibiotic susceptibility pattern of isolates from acute dacryocystitis

Bacterial isolate(n=31)	Cefazolin	Vancomycin	Ciprofloxacin	Moxifloxacin	Gentamicin	Amikacin	Tobramycin
<i>S.aureus</i> (9)	7(77.79%)	9(100%)	5(55.56%)	6(66.67%)	4(44.44%)	8(88.89%)	3(33.33%)
<i>S.pneumoniae</i> (7)	7(100%)	7(100%)	4(57.14%)	7(100%)	4(57.14%)	6(85.71%)	4(57.14%)
<i>S.pidermidis</i> (7)	7(100%)	7(100%)	4(57.14%)	4(57.14%)	1(14.28%)	5(71.43%)	2(28.57%)
<i>H.influenza</i> (4)	2(50%)	NT	1(25%)	3(75%)	1(25%)	3(75%)	2(50%)
<i>E. coli</i> (2)	1(50%)	NT	0(0%)	1(50%)	1(50%)	2(100%)	1(100%)
<i>P.aeruginosa</i> (2)	1(50%)	NT	0(0%)	0(0%)	1(100%)	2(100%)	2(100%)

Table 4: Antibiotic susceptibility pattern of isolates from chronic dacryocystitis

Bacterial isolate(n=110)	Cefazolin	Vanco mycin	Ciprofl oxacin	Moxiflo xacin	Gentam ycin	Amikac in	Tobram ycin
<i>S.aureus</i> (27)	21(77.8%)	24(88.9%)	15(55.6%)	17(62.6%)	8(29.6%)	22(81.4%)	17(62.9%)
<i>S.epider midis</i> (21)	20(95.2%)	21(100%)	18(85.7%)	19(90.4%)	10(47.6%)	20(95.2%)	9(42.6%)
<i>S.pneumo niae</i> (17)	16(94.1%)	16(94.1%)	15(88.2%)	15(88.2%)	7(41.1%)	15(88.2%)	12(70.5%)
<i>P.aerugin osa</i> (9)	1(11.1%)	NT	4(44.4%)	4(44.4%)	2(22.2%)	6(66.6%)	3(33.3%)
<i>E. coli</i> (7)	4(57.14%)	NT	3(42.85%)	5(71.43%)	1(14.29%)	7(100%)	5(71.43%)
<i>H.influen zae</i> (7)	2(28.57%)	NT	7(100%)	6(85.71%)	2(28.57%)	6(85.71%)	5(71.43%)
<i>Klebsiella</i> (6)	3(50%)	NT	6(100%)	6(100%)	3(50%)	6(100%)	5(83.3%)
<i>Enterobac ter</i> (5)	3(60%)	NT	5(100%)	5(100%)	3(60%)	5(100%)	4(80%)
<i>Proteus</i> (3)	1(33.33%)	NT	2(66.67%)	3(100%)	2(66.67%)	3(100%)	1(33.33%)
<i>Citrobact er</i> (2)	1(50%)	NT	1(50%)	2(100%)	1(50%)	2(100%)	1(50%)
<i>Bacteroid es</i> (3)	1(33.33%)	2(66.67%)	3(100%)	3(100%)	2(66.67%)	3(100%)	1(33.33%)

Antibiotic susceptibility pattern of isolates were displayed in tables. (Table.3)(Table.4).

Among 9 isolates of *Staphylococcus aureus*, 3(33.33%) were found to be methicillin resistant *Staphylococcus aureus*(MRSA) in acute dacryocystitis. All MRSA strains were sensitive to vancomycin. Among 8 Gram negative bacilli from acute dacryocystitis,2(*E.coli*) were identified as extended spectrum of beta lactamase(ESBL) producing strains. Both ESBL strains of *E.coli* were 100% resistant to tested quinolones except to moxifloxacin(50%).

In chronic dacryocystitis 11(40.74%) strains of *S.aureus* were found to be methicillin resistant *Staphylococcus aureus*(MRSA). Among 39 isolates of Gram negative bacilli, 3 strains of *E.coli* and 4 strains of *Klebsiella* species were found to be ESBL producers. All ESBL producing strains were susceptible to amikacin.

Discussion:

The most common inflammation of the lacrimal apparatus is

dacryocystitis. Lacrimal abscess is a very painful condition. Pathogenesis is believed to be microbial overgrowth in the stagnant fluid of the lacrimal sac in the background of nasolacrimal duct obstruction.⁹ Obstructed nasolacrimal duct may occur in any age group. Congenital blockage occurs in 3–6% of term infants. In majority of cases this is due to blockage of nasal end by epithelial debris or an imperforate mucosal membraneresulting from incomplete canalization of embryonic duct.¹⁰ Primary and secondary acquired nasolacrimal duct obstruction usually occurs mainly in middle-aged or older people. In our study 77% of cases were between the age group 30-60 years. According to Shakya et al , it was observed that the maximum numbers of dacryocystitis cases were found to be most common (54%) between 30 and 60 years of age.¹¹ Similar studies conducted by by Diggle, Duke-Elder , Reddy and Reddy, also showed maximum incidence in 35-65 years.^{12,13,14}

In our study, overall females were predominantly affected in acute and chronic dacryocystitis which accounted for 81%. These results are similar to other studies conducted by Shakya et al ¹¹ and Mallik and Chatterjee ¹⁵ in which the incidence of dacryocystitis was 81% and 71% respectively among females. Most probably cause for the preponderance of females could be due to excessive lacrimation, long stay in the kitchen with consequent greater exposure to soot, smoke, and heat. However, the anatomical factor of the narrow bony nasolacrimal canal appears to be more acceptable.¹¹

In our study, Out of 55 specimens collected from cases of acute dacryocystitis, 31(56.36%) specimens yielded bacterial growth. No anaerobic bacteria and fungi were isolated from acute dacryocystitis. Gram positive cocci was predominantly isolated and accounted for 74.19%. Among the gram-positive isolates, *S. aureus* was the most common organism noted, accounting for 39.13% of all the gram-positive isolates. Among Gram negative organisms *H. influenzae* (50%) was the predominant followed by *E. coli* (25%) and *P. aeruginosa* (25%). As per the study conducted by American Society of Ophthalmic Plastic and Reconstructive Surgery (ASOPRS)¹⁶ dacryocystitis group, it was found that 78.3% of the isolates were gram-positive and 21.7% were gram negative. *S. aureus* was the most common organism revealed and accounted for 50% of all the gram-positive isolates. In contrast, Briscoe et al. exclusively studied acute dacryocystitis in 39 patients and found the preponderance of gram-negative isolates (61%) with *P. aeruginosa* being the commonest isolate.¹⁷

In our study, out of 150 specimens collected from cases of chronic dacryocystitis, 150(73.33%) specimens yielded microbial growth. In chronic dacryocystitis, *S. aureus* (24.55%) was found to be the predominant bacterial pathogen followed by *S. epidermidis* and *S. pneumoniae*, accounting for, 19.09% and 15.45% of the isolates, respectively. According to the study conducted by Sun et al ¹⁸, in chronic dacryocystitis, CoNS was found to be the predominant bacterial pathogen followed by *S. aureus* and *S. pneumoniae*, accounting for 44.2, 10.8, and 8.7% of the isolates, respectively. Among Gram negative bacilli, *P. aeruginosa* was the predominant bacteria isolated. In our study, 20 specimens from chronic dacryocystitis revealed more than one organism.

In our study *Bacteroides* was the only anaerobic bacteria isolated and accounted for 2.73%. However, anaerobic microorganisms have been isolated in as many as 15.7% of the positive cultures, in some studies, the most common genus being *Bacteroides* (5.7%).¹⁸ Other commonly isolated anaerobic bacteria from dacryocystitis were *Peptostreptococcus* species, *Propionibacterium* species, *Prevotella* species and *Fusobacterium* species.¹⁹ According to the study by Sun et al,¹⁸ the most frequently isolated fungi being *Candida*, although *Aspergillus* and *Mucor* may also be found.¹⁸ In our study, *Candida* and *Aspergillus* was isolated and accounted for 1.82% and 0.10% respectively.

Antibiotic susceptibility pattern revealed , cefazolin and vancomycin has a higher efficacy towards Gram positive cocci. Higher susceptibility rates were also found with amikacin compared to gentamycin and tobramycin towards Gram positive and Gram negative bacteria. Pathogens isolated from acute dacryocystitis are more susceptible than pathogens recovered from chronic dacryocystitis. In acute dacryocystitis, MRSA was found to be 33% whereas in chronic dacryocystitis, MRSA was 40%. All strains of MRSA from acute dacryocystitis were susceptible to vancomycin. But, 3 strains of MRSA isolated from chronic dacryocystitis showed resistance to all tested

antibiotics including vancomycin. But, as per the study conducted by Mills et al, 15 methicillin-resistant *Staphylococcus aureus* (MRSA) have been isolated more frequently as etiologic agents in acute dacryocystitis than in chronic dacryocystitis.

Among Gram negative isolates from acute dacryocystitis, only two strains of *E. coli* were found to be extended spectrum of beta lactamases (ESBLs). Among 39 isolates of Gram negative bacilli, 3 strains of *E. coli* and 4 strains of *Klebsiella* species were found to be ESBL producers. All ESBL producing strains isolated from acute and chronic dacryocystitis were susceptible to amikacin.

Conclusion:

Staphylococcus aureus was the predominant organism isolated from acute and chronic dacryocystitis. Among Gram negative bacilli, *H. influenzae* was found to be the predominant pathogen in specimens collected from acute dacryocystitis and preponderance *Pseudomonas aeruginosa* was revealed in specimens collected from chronic dacryocystitis. Bacterial strains isolated from chronic dacryocystitis showed higher resistance than those from acute cases. Gram positive cocci from both acute and chronic dacryocystitis were highly susceptible to cefazolin and vancomycin. Efficacy of amikacin was high towards Gram positive and Gram negative bacteria.

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