

Original Research Article

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Microbiological Profile of Chronic Suppurative Otitis Media and their Antibiogram in A Rural Tertiary Care Hospital

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ABSTRACT

Chronic suppurative otitis media (CSOM) is one of the major causes of deafness in India. The chronicity of the disease and poor response to routine antimicrobials prompted us to isolate and identify the causative organisms and study antimicrobial susceptibility pattern. The ear discharge was collected using two sterile cotton swabs. One swab was used for performing Gram's stain and KOH mount. The second swab was used to inoculate Blood agar, MacConkey agar and Sabouraud Dextrose agar (SDA) for the isolation of aerobic bacteria and fungal pathogens. The antibiotic susceptibility testing of the bacterial isolates was done by Kirby Bauer's Disc Diffusion method according to CLSI Guidelines. Among 104 samples included in the study, Single bacterial growth was obtained in 69 (66.3%) samples and single fungal growth was seen in 24 (23.1%) samples. Mixed growth was seen in 6 (5.8%) samples and no growth was observed in 5 (4.8%) samples. Among the bacteria *Pseudomonas aeruginosa* (30.5%) was the predominant isolate and *Aspergillus niger* (8.6%) was the predominant isolate among fungi. Majority of the Gram negative isolates were sensitive to imipenem and all the *Staphylococci* isolates were sensitive to vancomycin, teicoplanin and linezolid. Periodic monitoring of microbiological profile is essential for the effective management of CSOM cases in a particular geographical area.

Keywords

Chronic suppurative otitis media, Sabouraud Dextrose agar, antibiotic susceptibility testing, *Pseudomonas aeruginosa*

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Introduction

Chronic suppurative otitis media (CSOM) is a chronic inflammation of the middle ear cleft, with permanent abnormality of tympanic membrane which presents as recurrent otorrhea. Patients presenting with tympanic membrane perforations and discharging ear

for a period of 3 months or more, despite medical treatment are recognized as CSOM cases (Mehta, *et al.*, 2017). Incidence of this disease is higher in developing countries especially in low socio-economic group because of malnutrition, overcrowding, poor hygiene and inadequate health care (Rajesh, *et al.*, 2017).

It is a disease of multiple etiologies and is well known for its persistence and recurrence in spite of treatment. CSOM is the leading cause of preventable hearing loss in the developing world (Toleti, *et al.*, 2016).

Untreated cases of CSOM can result in various complications such as persistent otorrhea, mastoiditis, labyrinthitis and facial nerve paralysis to serious complications like meningitis, intracranial abscess and thrombosis.

So, timely management of CSOM cases is important (Harshika, *et al.*, 2015). Most common organisms found in CSOM are *Pseudomonas*, *Staphylococcus aureus*, *Proteus mirabilis*, *Klebsiella*, *E.coli*, *Aspergillus species* and *Candida*. These organisms vary among geographical area (Shreshta, *et al.*, 2011).

Microbial predominance and their antibiotic sensitivity pattern change over time. The knowledge of local pattern of infections is essential to enable efficacious treatment of this disease and thereby reducing risks of complications (Arvind, *et al.*, 2014).

So, the present study was undertaken to know the microbiological profile of CSOM and their antibiotic sensitivity pattern in our set up. This knowledge is very important for the clinicians in the appropriate management of cases and to prevent complications associated with it.

Materials and Methods

This prospective study was conducted for a period of 2 years from September 2018 to September 2020 in the department of Microbiology and in association with the department of ENT of a tertiary care hospital. The study includes 104 clinically diagnosed cases of CSOM attending the Outpatient department of ENT.

Inclusion criteria

Patients with persistent or intermittent ear discharge in one or both ears for more than 3 months duration, irrespective of age and gender

Exclusion criteria

Patients with ear discharge with intact tympanic membrane (Otitis externa) and Patients on antibiotics or antifungal drugs (Topical or systemic) at the time of presentation or within a week of presentation

Sample collection

The ear discharge was collected using two sterile cotton swabs under aseptic precautions with the help of an aural speculum. Samples were collected from each ear separately in case of bilateral infection and transported immediately to the Microbiology laboratory for further processing.

Sample processing

One swab was used for performing Gram's stain and KOH mount. A thin smear was prepared using a clean glass slide, Gram staining is done and examined under oil immersion objective of the microscope for the presence of pus cells, bacteria and budding yeast cells. KOH mount is prepared and examined under high power objective for presence of fungal elements. The second swab was inoculated onto Blood agar, MacConkey agar for the isolation of aerobic bacteria and onto Sabouraud Dextrose agar (SDA) for the isolation of fungal pathogens.

Aerobic culture

The inoculated aerobic bacterial cultures was incubated at 37°C for 18-24 hours. The bacterial growth was identified based on

morphology, cultural characteristics and biochemical reactions according to standard microbiological methods (Collee JG *et al.*, 1996). The antibiotic susceptibility testing of the bacterial isolates was performed as per Kirby Bauer's Disc Diffusion method on Mueller Hinton agar using the preferable antibiotics and interpretation is done based on CLSI Guidelines.

Fungal culture

The specimen inoculated onto SDA was incubated at room temperature. The growth was identified based on morphology and cultural characteristics. Tease mount was examined microscopically using Lactophenol Cotton Blue (LPCB) staining technique. Antifungal susceptibility testing is not done in the present study.

Results and Discussion

A total of 104 patients were included in the study. Ear swabs from all the patients were collected and sent to the microbiology laboratory for culture. Single bacterial growth was obtained in 69 (66.3%) samples and single fungal growth was seen in 24 (23.1%) samples. Mixed growth was seen in 6 (5.8%) samples and no growth of bacteria or fungi was observed in 5 (4.8%) samples (Table.1).

A total of 105 (76 bacterial & 29 fungal) isolates were obtained from 104 samples. Distribution of bacteria and fungi causing CSOM is shown in Table.2. Among the bacterial isolates, *Pseudomonas aeruginosa* (30.5%) was the predominant isolate obtained followed by *Staphylococcus aureus* (11.4%), *Klebsiella* (7.6%), MRCONS (4.8%), *Citrobacter* (4.8%), NFGNB (3.8%), MRSA (3.8%), CONS (2.9%), *Proteus* (1.9%) and *Escherichia coli* (0.9%). Among the fungal isolates, *Aspergillus niger* (8.6%) was the predominant organism followed by

Aspergillus fumigatus (6.6%), *Candida albicans* (5.7%), NAC (3.8%) and *Aspergillus flavus* (2.9%).

Out of the 104 cases, 6 showed mixed isolates. *P. aeruginosa* was seen in combination with *A. fumigatus*, *C. albicans*, Coagulase negative *Staphylococcus* and *Klebsiella*. *S. aureus* was seen with *A. niger* growth and Non albicans *Candida* with *A. fumigatus* which is depicted in Table.3.

Antibiotic sensitivity testing was done for all the bacterial isolates. All culture media and antibiotic discs were procured from Himedia Laboratories Private Limited. In the present study, majority of the *P. aeruginosa* isolates were sensitive to Imipenem (90.6%) followed by ciprofloxacin (87.5%), netilmicin (87.5%), tobramycin (84.4%), meropenem (84.4%), amikacin (81.2%), ofloxacin (81.2%), gentamicin (78.1%), piperacillin tazobactam (59.4%), cefipime (56.2%), ceftazidime (50%) and amoxiclav (37.5%) (Table.4).

Gram positive isolates like *S. aureus*, MRSA, CONS, MRCONS showed 100% sensitivity to vancomycin, linezolid and teicoplanin. They showed increased sensitivity to gentamicin (91.7%), tetracycline (87.5%), amikacin (87.5%), followed by cotrimoxazole (79.2%), azithromycin (79.2%), amoxiclav (58.3%), clindamycin (45.8%), erythromycin (45.8%) and less sensitive to penicillin (33.3%) shown in Table.5. Sensitivity pattern of other Gram negative bacilli is shown in table.6. Chronic suppurative otitis media (CSOM) is one of the most common childhood diseases worldwide.

Its prevalence is related to poor socioeconomic conditions and common in developing countries. Research into the microbial causes of CSOM has so far been reliant on culture-based techniques (Neeff, *et al.*, 2016). Poorly treated or untreated CSOM can lead to many complications.

Table.1 Growth seen in CSOM cases

Growth	Number	Percentage
Single bacterial culture	69	66.3
Single fungal culture	24	23.1
Mixed growth	6	5.8
No growth	5	4.8
Total	104	100

Table.2 Distribution of organisms causing CSOM

Organisms isolated	Number	Percentage
BACTERIA		
<i>Pseudomonas aeruginosa</i>	32	30.5
<i>Staphylococcus aureus</i>	12	11.4
<i>Klebsiella</i>	8	7.6
Methicillin Resistant Coagulase negative <i>Staphylococcus aureus</i> (MRCONS)	5	4.8
<i>Citrobacter species</i>	5	4.8
Methicillin Resistant <i>Staphylococcus aureus</i> (MRSA)	4	3.8
Non fermenting Gram negative Bacilli (NFGNB)	4	3.8
Coagulase Negative <i>Staphylococcus aureus</i> (CONS)	3	2.9
<i>Proteus species</i>	2	1.9
<i>Escherichia coli</i>	1	0.9
FUNGI		
<i>Aspergillus niger</i>	9	8.6
<i>Aspergillus fumigatus</i>	7	6.6
<i>Candida albicans</i>	6	5.7
Non albicans <i>Candida</i> (NAC)	4	3.8
<i>Aspergillus flavus</i>	3	2.9
Total	105	100

Table.3 Distribution of mixed isolates in CSOM

Mixed isolates	Number
<i>P. aeruginosa</i> + <i>A. fumigatus</i>	1
<i>P. aeruginosa</i> + <i>C. albicans</i>	1
<i>S. aureus</i> + <i>A. niger</i>	1
<i>P. aeruginosa</i> + CONS	1
<i>P. aeruginosa</i> + <i>Klebsiella</i>	1
NAC + <i>A. fumigatus</i>	1
Total	6

Table.4 Antibiotic sensitivity pattern of *Pseudomonas aeruginosa*

(n=32)

Antibiotic	Sensitive		Resistant	
	Number	percentage	Number	Percentage
Imepenem	29	90.6	3	9.4
Ciprofloxacin	28	87.5	4	12.5
Netilmicin	28	87.5	4	12.5
Tobramycin	27	84.4	5	15.6
Meropenem	27	84.4	5	15.6
Amikacin	26	81.2	6	18.8
Ofloxacin	26	81.2	6	18.8
Gentamicin	25	78.1	7	21.9
Piperacillin tazobactam	19	59.4	13	40.6
Cefipime	18	56.2	14	43.8
Ceftazidime	16	50	16	50
Amoxyclav	12	37.5	20	62.5

Table.5 Antibiotic sensitivity pattern of *Staphylococcus* species

(n=24)

Antibiotic	Sensitive		Resistant	
	Number	percentage	Number	Percentage
Vancomycin	24	100	00	00
Linolid	24	100	00	00
Teicoplanin	24	100	00	00
Gentamicin	22	91.7	2	8.3
Amikacin	21	87.5	3	12.5
Tetracycline	21	87.5	3	12.5
Azithromycin	19	79.2	5	20.8
Cotrimoxazole	19	79.2	5	20.8
Amoxyclav	14	58.3	10	41.7
Clindamycin	13	54.2	11	45.8
Erythromycin	11	45.8	13	54.2
Penicillin	8	33.3	16	66.7

Table.6 Antibiotic sensitivity pattern of other Gram negative bacilli

(n=20)

Antibiotic	Sensitive		Resistant	
	Number	percentage	Number	Percentage
Imepenem	18	90	2	20
Amikacin	18	90	2	20
Ciprofloxacin	18	90	2	20
Gentamicin	17	85	3	15
Tobramycin	17	85	3	15
Meropenem	17	85	3	15
Ofloxacin	17	85	3	15
Piperacillin tazobactam	16	80	4	20
Cefipime	15	75	5	25
Ceftazidime	14	70	6	30
Amoxyclav	11	55	9	45
Ampicillin	7	35	13	65

Hence, diagnosis of the causative organism is necessary for proper management of CSOM cases (Prakash M *et al.*, 2013). Topical and oral antimicrobials are prescribed to patients based on culture results when available, but the clinical benefit of antimicrobial therapy is not always clear. Surgery may prevent local, regional, or systemic complications, but some patients may continue to have ear discharge postoperatively (Neeff M *et al.*, 2016).

CSOM can be characterized by co-infections with polymicrobial growth. In our study, monomicrobial growth was seen in 89.4% of cases which is similar to the previous study by Prakash M *et al.*, and Agarwal *et al.*, Mixed growth was seen in 5.8% and no growth in 4.8% of cases.

The predominant organism isolated in our study is *Pseudomonas* (30.5%) followed by *S. aureus* which is in accordance with other studies (Kaur *et al.*, 2018, Saranya *et al.*, 2015, Al-Hilli Z B *et al.*, 2015, Arif D *et al.*, 2014). But many other studies have shown *S. aureus* as the predominant organism (Vaidya K *et al.*, 2015 & Bizimana A *et al.*, 2017).

Among fungi, the common organism isolated was *Aspergillus* species (18.1%), which is in accordance with the study by Agarwal R *et al.*, Next common fungi isolated was *Candida* species (9.5%).

The injudicious and indiscriminate use of antibiotics has changed the microbiological flora. The knowledge of the microbiological spectrum is important to deliver efficacious treatment of this disease and also to prevent emergence of resistant strains (Chavan A *et al.*, 2014). The important fact to be kept in mind is that the antibiotic susceptibility pattern of these CSOM causing organisms keeps changing. Hence, routine antibiotic susceptibility testing before treatment is recommended (Prakash, *et al.*, 2013).

In the present study, Gram negative bacilli showed increased sensitivity to imepenem, ciprofloxacin, amikacin, meropenem, gentamicin, ofloxacin and high resistance to amoxyclav and cephalosporins. All the Staphylococcal isolates were sensitive to vancomycin, linezolid, teicoplanin and high resistance was noted to penicillin, erythromycin, clindamycin, amoxyclav. Our

observations are similar with many other studies (Kumar *et al.*, 2013, 2015).

CSOM is a global problem and affects mainly children due to their horizontal, wider and short Eustachian tube but can persist during adulthood following poor management of acute otitis media. It is a massive health problem in developing countries like India. Situation is more critical in rural areas because of lack of hygiene and knowledge. The present study gives knowledge about local pathogens that can assist in selection of most appropriate treatment which in turn improves the management of CSOM patients. Continuous and periodic evaluation of microbial pattern and antibiotic sensitivity of CSOM also helps to decrease the potential risks of complications.

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