Original Article

Prevalence of Gram positive bacteria in the affected individuals of Otitis media with effusion from the indigenous population of Southern Punjab, Pakistan: first report

Prevalência de bactérias Gram-positivas em andivíduos afetados por Otite média com efusão da população indígena do sul de Punjab, Paquistão: primeiro relatório

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Abstract

Otitis media with effusion (OME) is a type of otitis media (OM) characterized by the presence of fluid behind intact tympanic membrane and is one of the most common diseases of early childhood. It is an infectious disease associated with the presence of many pathogenic bacteria in the middle ear of affected individuals. This study was aimed to determine the prevalence of Gram-positive bacteria from the middle ear of OME patients in the population of Southern Punjab, Pakistan. The incidence of OME under comprehensive healthcare setting was investigated in patients who consulted at the department of ear, throat and nose, Bahawal Victoria Hospital (BVH), Bahawalpur, from December, 2019 to May, 2021. Ear swabs were taken from affected and normal individuals. After culturing bacteria from the ear swabs, microscopic analysis and biochemical tests were performed to characterize the cultured Gram-positive bacteria. Out of 352 patients examined, 109 (30.9%) patients had OME. Age of the participants ranged from 14 to 50 years; individuals between the ages of 14 and 22 years had the highest infection rates, while individuals between 40 and 50 years had the lowest rate of infection. Tympanic membrane perforation, fever, cough, sore throat, ear pain and hearing problem showed association with symptoms of OME. Microscopic analysis and biochemical characterization showed the presence of streptococci and staphylococci in all the studied samples. The most frequently isolated bacteria were Streptococcus pneumoniae, Streptococcus pyogenes, and Staphylococcus aureus with percentage of 53.3%, 20% and 13.3% respectively. Enterococcus faecalis (6.6%) and Staphylococcus epidermidis (6.6%) were also identified in the studied samples. This study will help in the better medical administration of OME affected individuals.

Keywords: Otitis media with effusion, Gram-positive bacteria, Tympanic membrane, *Streptococcus pneumoniae* and Pakistan.

Resumo

A otite média com efusão (OME) é um tipo de otite média (OM) caracterizada pela presença de líquido atrás da membrana timpânica intacta e é uma das doenças mais comuns durante a primeira infância. Trata-se de uma doença infecciosa associada à presença de muitas bactérias patogênicas na orelha média dos indivíduos afetados. Este estudo teve como objetivo determinar a prevalência de bactérias Gram-positivas do ouvido médio de pacientes com OME na população do sul de Punjab, Paquistão. A incidência de OME em ambiente de saúde abrangente foi investigada em pacientes que consultaram no departamento de ouvido, garganta e nariz, Hospital Bahawal Victoria (BVH), Bahawalpur, de dezembro de 2019 a maio de 2021. Cotonetes de orelha foram coletados de indivíduos afetados e normais. Após a cultura das bactérias dos swabs auriculares, análises microscópicas e testes bioquímicos foram realizados para caracterizar as bactérias Gram-positivas cultivadas. Dos 352 pacientes examinados, 109 (30,9%) apresentavam OME. A idade dos participantes variou de 14 a 50 anos; indivíduos entre 14 e 22 anos apresentaram as maiores taxas de infecção, enquanto indivíduos entre 40 e 50 anos apresentaram as menores taxas de infecção. Perfuração da membrana timpânica, febre, tosse, dor de garganta, dor de ouvido e problema de audição apresentaram associação com sintomas de OME. A análise microscópica e a caracterização bioquímica mostraram a presença de estreptococos e estafilococos em todas as amostras estudadas. As bactérias isoladas com maior frequência foram Streptococcus pneumoniae, Streptococcus pyogenes e Staphylococcus aureus com percentuais de 53,3%, 20% e 13,3%, respectivamente. Enterococcus faecalis (6,6%) e Staphylococcus epidermidis (6,6%) também foram identificados nas amostras estudadas. Este estudo ajudará na melhor administração médica de indivíduos afetados por OME.

Palavras-chave: Otite média com efusão, bactérias Gram-positivas, membrana timpânica, *Streptococcus pneumoniae*, Paquistão.

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1. Introduction

Otitis media (OM) is a clinical disorder that causes conductive hearing loss and involves the presence of fluid behind the intact tympanic membrane of the middle ear without any signs of acute inflammation (Sadé et al., 1989). Children who experience hearing loss because of OM may experience problems related to speaking and reading (Cai and McPherson, 2017; Carroll and Breadmore, 2018; Khavarghazalani et al., 2016; le Clercq et al., 2017). OM can lead to a number of problems, including mastoiditis, cholesteatoma, meningitis, brain abscess, and lateral sinus thrombosis. OM is caused by a number of factors, such as environmental factors, viruses, bacterial load, age, allergies, upper respiratory tract infections (URTIs), lack of breastfeeding, smoking, a low standard of living, a large number of siblings, clinical history of family and daycare attendance (Brennan-Jones et al., 2015; Zhang et al., 2014). OM may also result from immune system flaws. The multigenic nature of the genetic susceptibility factors is known to have broad effects on host immunological responses. The most prevalent infectious agents, such as Streptococcus pneumoniae, Haemophilus influenzae, Moraxella catarrhalis, Streptococcus pyogenes, Staphylococcus aureus and/ or respiratory viruses, colonize and multiply in the nasopharynx. These pathogens travel from the nasopharynx to the Eustachian tube and infect the middle ear which may result in OM (Coticchia et al., 2013; Mofatteh et al., 2018). It has also been reported that Streptococcus pneumoniae causes OM, acute respiratory infections, pneumonia, meningitis and sepsis (Silva et al., 2022). Depending on the type and severity, OM is treated differently, ranging from antibiotics to surgery (Lieberthal et al., 2013).

OM is majorly divided into three categories: acute OM (AOM), recurrent AOM (rAOM), and OM with effusion (OME). OME is also known as oto-tubaritis, OM serosa or OM with chronic exudate (Paradise et al., 2002; Mudry and Young, 2020). Approximately 80% of the children may face this disease before the age of 4 years which makes OME as one of the most common diseases of early childhood. The most common symptom of the disease is mild hearing loss. This hearing loss is so mild that it remains undetected for a large period of time. This mild hearing loss can only be detected with the help of tympanometry devices (Rosenfeld et al., 2016). Otoscopy is commonly performed to diagnose the disease but it is not very sensitive as the large number of OME cases remain undetected (Xu et al., 2020). OME is a disease of middle ear and its health is directly related to the health of upper respiratory tract (URT) (Ngo et al., 2016). Many bacteria have been isolated from the middle ear of OME patients which include Corynebacterium otitidis, Moraxella catarrhalis, Haemophilus influenzae, Streptococcus pneumoniae, Alloiococcus otitidis, Pseudomonas spp. and Staphylococcus spp. (Ari et al., 2019; Baek et al., 2018; Boers et al., 2018; Chan et al., 2016, 2017; Enoksson et al., 2020; Jervis-Bardy et al., 2015; Johnston et al., 2019; Kolbe et al., 2019; Krueger et al., 2017; Liu et al., 2011; Val et al., 2018; Xu et al., 2020).

Many of these bacteria are commonly present in healthy URT as well which is making difficult to associate the presence of these bacteria with OME (De Boeck et al., 2017; Frank et al., 2003; Stroman et al., 2001).

Current study was designed to study the prevalence of Gram-positive bacteria from the microbial flora of middle ear of OME patients who visited ear, nose, and throat (ENT) department of Bahawal Victoria Hospital (BVH) Bahawalpur, one of the biggest hospitals in the South Punjab region of Pakistan, from December 2019 to May 2021.

2. Materials and Methods

2.1. Enrollment of patients

We visited the ENT department of BVH from December 2019 to May 2021 to identify and enroll patients of OME from different areas of South Punjab, Pakistan. A detailed questionnaire was designed to record family history, age of onset, symptoms and the occurrence of OME. Endoscopy was performed to check for abnormalities in the internal structure of the middle ear. Written consent was obtained from the participants or their parents/legal guardians. The study was approved by the Institutional Bioethics Committee of the Islamia University of Bahawalpur, Pakistan.

2.2. Collection and culturing of Ear swab samples

Patients were enrolled for the study if they did not have any oral or systemic antibiotic treatment in the seven days prior to sample taken, had a tympanic perforation and ear drainage for more than two months. A fine, sterile cotton swab was used to obtain the middle ear discharge from the tympanic chamber. The ear was cleaned with normal saline before sample taking. The samples were delivered to the microbiology lab for additional identification procedures. Bacterial culturing was done according to the method described by (Kombade et al., 2021). Briefly, the method consists of inoculation of swabs on blood agar and MacConkey agar plates for 24–48 hours at 37°C for the optimal growth of bacteria.

2.3. Microscopic analysis

The simultaneous application of chemical reagents for a set amount of time was used to apply the Gram staining procedure. Crystal violet was used as the primary stain, followed by iodine as the mordant, ethanol or acetone as the decolorizer, and safranin as the counterstain. To distinguish between Gram-positive and Gram-negative isolates, stained slides were air-dried and examined under oil immersion (100X) through bright field microscope.

2.4. Biochemical identification of bacterial isolates

Identification of isolates was performed using Berge's Manual of determinative bacteriology. The dichotomous key was used to identify reported microbes causing OME.

We were interested in identification of Gram positive bacteria as there are many reports which showed the prevalence of Gram positive bacteria such as, *S. pneumoniae*, *S. aureus and S. pyogenes is* higher as compared to Gram negative bacteria in the bacterial flora of middle ear of OME patients (Afolabi et al., 2012; Al-Alousi and Mayas, 2012; Agha and Al-Delaimi, 2021; Brook and Frazier, 1996; Rosenblut et al., 2001; Silva et al., 2021).

A catalase test was performed to differentiate between *Streptococcus* and *Staphylococcus*. The positive results were further confirmed through a coagulase test to find the occurrence of widely distributed *Staphylococcus aureus* with coagulase activity while the Novobiocin test was performed to confirm the occurrence of *Staphylococcus epidermidis* (Dawodu and Akanbi, 2021). The samples with negative results were cultured on Blood Agar plates to observe the three types of hemolysis: α - hemolysis (partial hemolysis), β -hemolysis (complete hemolysis) and γ -hemolysis (no hemolysis). The results were recorded after the validation of the Optochin Sensitivity test, Bacitracin test and NaCl (6.5%) based test (Burckhardt et al., 2017; Facklam et al., 1982).

3. Results

Three hundred and fifty two patients were examined by the physicians of the ENT department during the period of the study. The age of the patients ranged from 14 to 50 years. Figure 1 provides a breakdown of the incidence of OM in the various age groups.

Using the criteria (Figure 2), 109 (30.9%) out of 352 patients were identified as having OME; Out of total



Figure 1. Prevalence of OME in different age groups.

109 selected individuals 58 (53.2%) males and 51 (46.8%) females had the symptom of ear discharge, which was the principal criterion for the selection. Some enrolled patients were belonging to same families and were close relatives of each other. Perforation of tympanic membrane was present as the second most commonly identified clinical problem with the incidence rate of 41.2% in the studied group along with the presence of fever with the same incidence rate (41.2%). Coughing, sore throat and hoarseness were the third most common problem with a percentage of 20.11%. The percentage of the studied individuals that had pain in their affected ear at least once from the presence of the disease was 23.7%. None of the studied individual had any speech problem whereas 6.4% of the totally studied individuals had some auditory problems.

3.1. Bacterial isolates cultured from ear swabs of OME patients

The distribution of bacterial isolates found in swab samples is displayed in Table 1 and Figure 3 while the results of biochemical identification are presented in Figure 4. Out of the total 45 bacterial isolates cultured from swab specimens, Gram-positive Cocci made up 100% of the population, out of these *streptococci* and *staphylococci* made up 80% and 20% of the population, respectively. 66.6% of the staphylococci were *Staphylococcus aureus*, whereas 33.3% were coagulase-negative staphylococci (*Staphylococcus epidermidis*). *Streptococcus pneumoniae* (66.6%), *Streptococcus pyogenes* (25%) and *Enterococcus faecalis* were the three most prevalent *streptococci* (8.3%) shown in Figure 3.

4. Discussion

Middle ear irritation linked to OM is a prevalent condition in both children and adults (Korona-Glowniak et al., 2020). In order to identify the presence of Gram-positive bacteria in the bacterial flora of middle ear of OME patients and to link them with infection, 109 patients from Southern Punjab region of Pakistan were enrolled. We also investigated how common OM was within the analyzed age groups. The results showed that men (53.2%) had a higher prevalence of OM than did women (46.8%). These findings are in line with the findings of (Kombade et al., 2021), who found that while men engage in more outdoor



Figure 2. Clinical and diagnostic profile of subjects with OME.

Isolates	Total no. of isolates n=45	Total (%)	Gram Staining Gram +ive	Catalase test	Coagulase test	Blood Agar test	Optochin test	Bacitracin test	Growth in 6.5% NaCl	Novobiocin test
Enterococcus faecalis	ε	6.6	Cocci (Single Diplo, chains Clusters)	-ive		γ -hemolysis			+ive	
Streptococcus pyogenes	6	20	Cocci (Single Diplo, chains Clusters)	-ive		β-hemolysis		+ive		
Staphylococcus aureus	9	13.3	Cocci in grape-like clusters	+ive	+ive					
Streptococcus pneumoniae	24	53.3	Single cocci, lanceolate diplococci, Short chains of cocci	-ive		α -hemolysis	+ive			
Staphylococcus epidermidis	3	6.6	Cocci in grape like clusters	+ive	-ive					+ive

Table 1. Profile of bacterial Isolates cultured from ear swabs of subjects.

activities and travel more frequently than women (47.7%), they were more likely to get OM (52.2%) and recurrent URT infections. The frequency of OME infections is significantly influenced by age. In the current study, the age group with the highest infection rates was 14-22 (37.6%), followed by 22-26 (22.6%). In individuals from 40 to 50 years, a low distribution of OME (4.5%) was seen. Children's propensity for upper respiratory tract infections makes them obviously more vulnerable to OM. Studies show children are more susceptible to OM as compared to adults as children's immune systems are more sensitive and they have to deal with a bigger number of infectious bacteria. Poor hygiene, overcrowding, inadequate housing, low socioeconomic level, decreased immunity, and repeated upper respiratory tract infections are the major risk factors for increasing susceptibility of young children for OM (Kombade et al., 2021; Rathi et al., 2018). In our study individuals from the age of 14-22 years had the maximum percentage of affected individuals from the group of studied people. We did not find many children having OME who were fulfilling the set criteria. This may be because of many



Figure 3. Percentage of bacterial strains isolated from cultured ear swabs from subjects.

reasons including the unawareness regarding the disease as the region has one of the high percentages of illiteracy, poor socioeconomic background and difficulties to visit the hospital (Kumar and Seth, 2011; Mazlan et al., 2002).

Tympanic membrane perforation is one of the major clinical phenotype in the patients of OME. This perforation may be present in one ear or in both ears. In our study 45 (41.2%) affected individuals had tympanic membrane perforation in at least one ear. These results are in line with the studies conducted by (Kamal et al., 2004), (Ibekwe et al., 2009) and (Shrikrishna et al., 2013). The phenotype of hearing loss is also one of major finding in association with OME but we found hearing loss in a fraction of studied group (6.4%). Fever, cough and sore throat are the other parameters in which changes have been recorded in the patients of OME. We also checked these parameters and found the presence of these symptoms in 41.2% and 20.1% of the studied population, respectively. The presence of these symptoms indicates the role of ENT associated bacteria in OME and in associated problems. We used biochemical analysis to identify bacterial isolates in order to confirm the dispersion of infection-causing organisms. 100% of the bacteria identified for this investigation were Gram-positive, with Streptococcus pneumoniae being the most frequent isolate (53.3%). With 20% and 13.3% respectively, *Staphylococcus aureus* and Streptococcus pyogenes made up the majority. Similar findings were observed by (Silva et al., 2021), who observed Streptococcus pneumoniae (50.9%) as the most dominant bacterial pathogen in AOM patients, while the Streptococcus pyogenes (30.9%) was also widely distributed. According to many studies, Staphylococcus aureus is the most prevalent bacterium in people with chronic OM but in our study Streptococcus pneumoniae is the most commonly found bacteria. The different patterns of bacterial dissemination could be attributed to geographic variability (Afolabi et al., 2012; Al-Alousi and Mayas, 2012; Brook and Frazier,



Figure 4. (A) Hemolysis on the blood agar media to differentiate the streptococci; (B) Microscopic observation of Gram positive *Streptococcus pneumoniae*; (C) Catalase test to differentiate the Streptococci and Staphylococci; (D) Coagulase test to differentiate the *Staphylococcus aureus* from *Staphylococcus epidermidis*.

1996; Rosenblut et al., 2001). Many studies showed the involvement of genetic factors in OME (Bartlett et al., 2015; Casselbrant et al., 1999; Crompton et al., 2017; Mira et al., 1999; Samuels et al., 2016; Santos-Cortez et al., 2016; Van Ingen et al., 2016). In our study, more than one individual affected with OME were present in some families. The presence of more than one affected individuals gives an indication about the association of genetic factors in these families and genetic screening can help to find out the involvement of any genetic factor in the enrolled patients.

5. Conclusion

We studied the population of Southern Punjab region of Pakistan having OME. This study indicates the OME to be associated with the presence of some or many symptoms liker tympanic membrane perforation, hearing problem, fever, cough and sore throat in certain individuals. We also identified the principal Gram-positive bacterial agents responsible for the illness in the Southern Punjab region of Pakistan. The most frequently isolated organisms were Streptococcus pneumoniae, Streptococcus pyogenes and Staphylococcus aureus. The findings of this study shed light on the Gram-positive harmful bacteria that may produce OM. Moreover, this is the first study, to the best of our knowledge, in which microbial flora of Gram-positive bacteria of the middle ear of OME patients from Southern Punjab region of Pakistan have been studied which will be helpful for the better management of the disease in the studied population.

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References

- AFOLABI, O.A., SALAUDEEN, A.G., OLOGE, F.E., NWABUISI, C. and NWAWOLO, C.C., 2012. Pattern of bacterial isolates in the middle ear discharge of patients with chronic suppurative otitis media in a tertiary hospital in North central Nigeria. *African Health Sciences*, vol. 12, no. 3, pp. 362-367. http://dx.doi.org/10.4314/ ahs.v12i3.18. PMid:23382753.
- AGHA, Z.H.M. and AL-DELAIMI, M.S., 2021. Prevalence of common bacterial etiology and antimicrobial susceptibility pattern in patients with otitis media in Duhok Province–Iraq. *Zanco Journal of Pure and Applied Sciences*, vol. 33, no. 4, pp. 11-25. http://dx.doi.org/10.21271/ZJPAS.33.4.2.
- AL-ALOUSI, M. and MAYAS, N., 2012. Bacteria causing otitis media in some private and public centers in Thamar governorate-Yemen. Yemeni Journal for Medical Sciences, vol. 6, no. 1, pp. 7-15. https://doi.org/10.20428/yjms.v6i1.29.
- ARI, O., KARABUDAK, S., KALCIOGLU, M.T., GUNDUZ, A.Y. and DURMAZ, R., 2019. The bacteriome of otitis media with effusion: does it originate from the adenoid? *International Journal of Pediatric Otorhinolaryngology*, vol. 126, pp. 109624. http://dx.doi.org/10.1016/j.ijporl.2019.109624. PMid:31408742.

- BAEK, I., KIM, M., LEE, I., NA, S.I., GOODFELLOW, M. and CHUN, J., 2018. Phylogeny trumps chemotaxonomy: a case study involving Turicella otitidis. *Frontiers in Microbiology*, vol. 9, pp. 834. http://dx.doi.org/10.3389/fmicb.2018.00834. PMid:29760685.
- BARTLETT, J.A., MEYERHOLZ, D.K., WOHLFORD-LENANE, C.L., NAUMANN, P.W., SALZMAN, N.H. and MCCRAY JUNIOR, P.B., 2015. Increased susceptibility to otitis media in a Splunc1deficient mouse model. *Disease Models & Mechanisms*, vol. 8, no. 5, pp. 501-508. http://dx.doi.org/10.1242/dmm.019646. PMid:25765466.
- BOERS, S.A., DE ZEEUW, M., JANSEN, R., VAN DER SCHROEFF, M.P., VAN ROSSUM, A., HAYS, J.P. and VERHAEGH, S.J., 2018. Characterization of the nasopharyngeal and middle ear microbiota in gastroesophageal reflux-prone versus gastroesophageal reflux non-prone children. *European Journal* of Clinical Microbiology & Infectious Diseases, vol. 37, no. 5, pp. 851-857. http://dx.doi.org/10.1007/s10096-017-3178-2. PMid:29404836.
- BRENNAN-JONES, C.G., WHITEHOUSE, A.J., PARK, J., HEGARTY, M., JACQUES, A., EIKELBOOM, R.H., SWANEPOEL, D.W., WHITE, J.D., and JAMIESON, S.E., 2015. Prevalence and risk factors for parent-reported recurrent otitis media during early childhood in the Western Australian Pregnancy Cohort (Raine) study. *Journal of Pediatrics and Child Health*, vol. 51, no. 4, pp. 403-419. https://doi.org/10.1111/jpc.12741.
- BROOK, I. and FRAZIER, E.H., 1996. Microbial dynamics of persistent purulent otitis media in children. *The Journal of Pediatrics*, vol. 128, no. 2, pp. 237-240. http://dx.doi.org/10.1016/S0022-3476(96)70397-9. PMid:8636819.
- BURCKHARDT, I., PANITZ, J., BURCKHARDT, F. and ZIMMERMANN, S., 2017. Identification of Streptococcus pneumoniae: development of a standardized protocol for optochin susceptibility testing using total lab automation. *BioMed Research International*, vol. 2017, pp. 4174168. http://dx.doi.org/10.1155/2017/4174168. PMid:28401155.
- CAI, T. and MCPHERSON, B., 2017. Hearing loss in children with otitis media with effusion: a systematic review. *International Journal of Audiology*, vol. 56, no. 2, pp. 65–76. http://dx.doi.org /10.1080/14992027.2016.1250960. PMid:27841699.
- CARROLL, J.M. and BREADMORE, H.L., 2018. Not all phonological awareness deficits are created equal: vidence from a comparison between children with otitis media and poor readers. *Developmental Science*, vol. 21, no. 3, pp. e12588. http://dx.doi. org/10.1111/desc.12588. PMid:28880490.
- CASSELBRANT, M.L., MANDEL, E.M., FALL, P.A., ROCKETTE, H.E., KURS-LASKY, M., BLUESTONE, C.D. and FERRELL, R.E., 1999. The heritability of otitis media: a twin and triplet study. *Journal of the American Medical Association*, vol. 282, no. 22, pp. 2125-2130. http://dx.doi.org/10.1001/jama.282.22.2125. PMid:10591333.
- CHAN, C.L., WABNITZ, D., BARDY, J.J., BASSIOUNI, A., WORMALD, P.J., VREUGDE, S. and PSALTIS, A.J., 2016. The microbiome of otitis media with effusion. *The Laryngoscope*, vol. 126, no. 12, pp. 2844– 2851. http://dx.doi.org/10.1002/lary.26128. PMid:27335217.
- CHAN, C.L., WABNITZ, D., BASSIOUNI, A., WORMALD, P.J., VREUGDE, S. and PSALTIS, A.J., 2017. Identification of the bacterial reservoirs for the middle ear using phylogenic analysis. JAMA Otolaryngology-Head & Neck Surgery, vol. 143, no. 2, pp. 155-161. http://dx.doi.org/10.1001/jamaoto.2016.3105. PMid:27812691.
- COTICCHIA, J.M., CHEN, M., SACHDEVA, L. and MUTCHNICK, S., 2013. New paradigms in the pathogenesis of otitis media in children. *Frontiers in Pediatrics*, vol. 1, pp. 52. http://dx.doi. org/10.3389/fped.2013.00052. PMid:24400296.

- CROMPTON, M., PURNELL, T., TYRER, H.E., PARKER, A., BALL, G., HARDISTY-HUGHES, R.E., GALE, R., WILLIAMS, D., DEAN, C.H., SIMON, M.M., MALLON, A.M., WELLS, S., BHUTTA, M.F., BURTON, M.J., TATEOSSIAN, H. and BROWN, S.D.M., 2017. A mutation in Nischarin causes otitis media via LIMK1 and NF-kB pathways. *PLOS Genetics*, vol. 13, no. 8, pp. e1006969. http:// dx.doi.org/10.1371/journal.pgen.1006969. PMid:28806779.
- DAWODU, O. and AKANBI, R., 2021. Isolation and identification of microorganisms associated with automated teller machines on Federal Polytechnic Ede campus. *PLoS One*, vol. 16, no. 8, pp. e0254658. http://dx.doi.org/10.1371/journal.pone.0254658. PMid:34351934.
- DE BOECK, I., WITTOUCK, S., WUYTS, S., OERLEMANS, E.F., VAN DEN BROEK, M.F., VANDENHEUVEL, D., VANDERVEKEN, O. and LEBEER, S., 2017. Comparing the healthy nose and nasopharynx microbiota reveals continuity as well as niche-specificity. *Frontiers in Microbiology*, vol. 8, pp. 2372. http://dx.doi. org/10.3389/fmicb.2017.02372. PMid:29238339.
- ENOKSSON, F., RUIZ RODRIGUEZ, A., PENO, C., BALCAZAR LOPEZ, C., TJERNSTROM, F., BOGAERT, D., HAKANSSON, A.P. and BERGENFELZ, C., 2020. Niche-and gender-dependent immune reactions in relation to the microbiota profile in pediatric patients with otitis media with effusion. *Infection and Immunity*, vol. 88, no. 10, pp. e00147-20. http://dx.doi.org/10.1128/ IAI.00147-20. PMid:32661126.
- FACKLAM, R.R., THACKER, L.G., FOX, B. and ERIQUEZ, L., 1982. Presumptive identification of streptococci with a new test system. *Journal of Clinical Microbiology*, vol. 15, no. 6, pp. 987-990. http://dx.doi.org/10.1128/jcm.15.6.987-990.1982. PMid:7050157.
- FRANK, D.N., SPIEGELMAN, G.B., DAVIS, W., WAGNER, E., LYONS, E. and PACE, N.R., 2003. Culture-independent molecular analysis of microbial constituents of the healthy human outer ear. *Journal* of Clinical Microbiology, vol. 41, no. 1, pp. 295-303. http://dx.doi. org/10.1128/JCM.41.1.295-303.2003. PMid:12517864.
- IBEKWE, T.S., NWAORGU, O.G. and IJADUOLA, T.G., 2009. Correlating the site of tympanic membrane perforation with hearing loss. *BMC Ear, Nose, and Throat Disorders*, vol. 9, no. 1, pp. 1. http:// dx.doi.org/10.1186/1472-6815-9-1. PMid: 19121227.
- JERVIS-BARDY, J., ROGERS, G.B., MORRIS, P.S., SMITH-VAUGHAN, H.C., NOSWORTHY, E., LEONG, L.E., SMITH, R.J., WEYRICH, L.S., DE HAAN, J., CARNEY, A.S., LEACH, A.J., O'LEARY, S. and MARSH, R.L., 2015. The microbiome of otitis media with effusion in Indigenous Australian children. *International Journal of Pediatric Otorhinolaryngology*, vol. 79, no. 9, pp. 1548-1555. http://dx.doi. org/10.1016/j.ijporl.2015.07.013. PMid:26228497.
- JOHNSTON, J., HOGGARD, M., BISWAS, K., ASTUDILLO-GARCIA, C., RADCLIFF, F.J., MAHADEVAN, M. and DOUGLAS, R.G., 2019. Pathogen reservoir hypothesis investigated by analyses of the adenotonsillar and middle ear microbiota. *International Journal of Pediatric Otorhinolaryngology*, vol. 118, pp. 103-109. http://dx.doi.org/10.1016/j.ijporl.2018.12.030. PMid:30599284.
- KAMAL, N., JOARDER, A.H., CHOWDHURY, A.A. and KHAN, A.W., 2004. Prevalence of chronic suppurative otitis media among the children living in two selected slums of Dhaka city. *Bangladesh Medical Research Council Bulletin*, vol. 30, no. 3, pp. 95-104. PMid:16240980.
- KHAVARGHAZALANI, B., FARAHANI, F., EMADI, M. and HOSSENI DASTGERDI, Z., 2016. Auditory processing abilities in children with chronic otitis media with effusion. *Acta Oto-Laryngologica*, vol. 136, no. 5, pp. 456-459. http://dx.doi.org/10.3109/000164 89.2015.1129552. PMid:26881324.
- KOLBE, A.R., CASTRO-NALLAR, E., PRECIADO, D. and PEREZ-LOSADA, M., 2019. Altered middle ear microbiome in children with chronic otitis media with effusion and respiratory illnesses.

Frontiers in Cellular and Infection Microbiology, vol. 9, pp. 339. http://dx.doi.org/10.3389/fcimb.2019.00339. PMid:31637220.

- KOMBADE, S.P., KAUR, N., PATRO, S.K. and NAG, V.L., 2021. Clinicobacteriological and antibiotic drug resistance profile of chronic suppurative otitis media at a tertiary care hospital in Western Rajasthan. *Journal of Family Medicine and Primary Care*, vol. 10, no. 7, pp. 2572-2579. http://dx.doi.org/10.4103/jfmpc. jfmpc_2480_20. PMid:34568138.
- KORONA-GLOWNIAK, I., WISNIEWSKA, A., JUDA, M., KIELBIK, K., NIEDZIELSKA, G. and MALM, A., 2020. Bacterial etiology of chronic otitis media with effusion in children-risk factors. *Journal of Otolaryngology - Head & Neck Surgery*, vol. 49, pp. 24. http://dx.doi.org/10.1186/s40463-020-00418-5. PMid:31898554.
- KRUEGER, A., VAL, S., PEREZ-LOSADA, M., PANCHAPAKESAN, K., DEVANEY, J., DUAH, V., DEMASON, C., POLEY, M., ROSE, M. and PRECIADO, D., 2017. Relationship of the middle ear effusion microbiome to secretory mucin production in pediatric patients with chronic otitis media. *The Pediatric Infectious Disease Journal*, vol. 36, no. 7, pp. 635-640. http://dx.doi.org/10.1097/ INF.000000000001493. PMid:28027286.
- KUMAR, H. and SETH, S., 2011. Bacterial and fungal study of 100 cases of chronic suppurative otitis media. *International Journal* of Scientific Research, vol. 5, no. 6, pp. 1224-1227. http://dx.doi. org/10.36106/ijsr.
- LE CLERCQ, C.M.P., VAN INGEN, G., RUYTJENS, L., GOEDEGEBURE, A., MOLL, H.A., RAAT, H., JADDOE, V.W.V., BAATENBURG DE JONG, R.J. and VAN DER SCHROEFF, M.P., 2017. Prevalence of hearing loss among children 9 to 11 years old: the generation R study. JAMA Otolaryngology-Head & Neck Surgery, vol. 143, no. 9, pp. 928-934. http://dx.doi.org/10.1001/jamaoto.2017.1068. PMid:28750130.
- LIEBERTHAL, A.S., CARROLL, A.E., CHONMAITREE, T., GANIATS, T.G., HOBERMAN, A., JACKSON, M.A., JOFFE, M.D., MILLER, D.T., ROSENFELD, R.M., SEVILLA, X.D., SCHWARTZ, R.H., THOMAS, P.A. and TUNKEL, D.E., 2013. The diagnosis and management of acute otitis media. *Pediatrics*, vol. 131, no. 3, pp. e964-e99. http://dx.doi.org/10.1542/peds.2012-3488. PMid:23439909.
- LIU, C.M., COSETTI, M.K., AZIZ, M., BUCHHAGEN, J.L., CONTENTE-CUOMO, T.L., PRICE, L.B., KEIM, P.S. and LALWANI, A.K., 2011. The otologic microbiome: a study of the bacterial microbiota in a pediatric patient with chronic serous otitis media using 16SrRNA gene-based pyrosequencing. *Archives of Otolaryngology*-*Head & Neck Surgery*, vol. 137, no. 7, pp. 664–668. http://dx.doi. org/10.1001/archoto.2011.116. PMid:21768410.
- MAZLAN, R., SAIM, L., THOMAS, A., SAID, R. and LIYAB, B., 2002. Ear infection and hearing loss amongst headphone users. *The Malaysian Journal of Medical Sciences : MJMS*, vol. 9, no. 2, pp. 17-22. PMid:22844220.
- MIRA, J.P., CARIOU, A.F., GRALL, C., DELCLAUX, M.R., LOSSER, F., HESHMATI, C., CHEVAL, M., MONCHI, J.L., TEBOUL, J.L., RICHÉ, F., LELEU, G., ARBIBE, L., MIGNON, A., DELPECH, M. and DHAINAUT, J.F., 1999. Association of TNF2, a TNF-α promoter polymorphism, with septic shock susceptibility and mortality: a multicenter study. *Journal of the American Medical Association*, vol. 282, no. 6, pp. 561-568. http://dx.doi.org/10.1001/jama.282.6.561. PMid:10450718.
- MOFATTEH, M.R., SHAHABIAN MOGHADDAM, F., YOUSEFI, M. and NAMAEI, M.H., 2018. A study of bacterial pathogens and antibiotic susceptibility patterns in chronic suppurative otitis media. *The Journal of Laryngology and Otology*, vol. 132, no. 1, pp. 41-45. http://dx.doi.org/10.1017/S0022215117002249. PMid:29151379.
- MUDRY, A. and YOUNG, J.R., 2020. Otitis media with effusion: Politzer's 100 year legacy. International Journal of Pediatric

Otorhinolaryngology, vol. 136, pp. 110160. http://dx.doi. org/10.1016/j.ijporl.2020.110160. PMid:32544637.

- NGO, C.C., MASSA, H.M., THORNTON, R.B. and CRIPPS, A.W., 2016. Predominant bacteria detected from the middle ear fluid of children experiencing otitis media: a systematic review. *PLoS One*, vol. 11, no. 3, pp. e0150949. http://dx.doi.org/10.1371/ journal.pone.0150949. PMid:26953891.
- PARADISE, J.L., BLUESTONE, C.D., COLBORN, D.K., BERNARD, B.S., ROCKETTE, H.E. and KURU-LASKY, M., 2002. Tonsillectomy and adenotonsillectomy for recurrent throat infection in moderately affected children. *Pediatrics*, vol. 110, no. 1, pp. 7-15. http:// dx.doi.org/10.1542/peds.110.1.7. PMid:12093941.
- RATHI, S., JAISWAL, A.A., SHARMA, N., BANERJEE, P. and GARG, A., 2018. Bacteriological profile and drug sensitivity patterns in chronic suppurative otitis media patients at JLN Hospital and Research Centre, Bhilai, Chhattisgarh State India. IP Indian Journal of Anatomy and Surgery of Head. *Neck and Brain*, vol. 4, no. 2, pp. 27-37. http://dx.doi.org/10.18231/2581-5229.2018.0009.
- ROSENBLÜT, A., SANTOLAYA, M.E., GONZALEZ, P., CORBALAN, V., AVENDANO, L.F., MARTÍNEZ, M.A. and HORMAZABAL, J.C., 2001. Bacterial and viral etiology of acute otitis media in Chilean children. *The Pediatric Infectious Disease Journal*, vol. 20, no. 5, pp. 501-507. http://dx.doi.org/10.1097/00006454-200105000-00006. PMid:11368107.
- ROSENFELD, R.M., SHIN, J.J., SCHWARTZ, S.R., COGGINS, R., GAGNON, L., HACKELL, J.M., HOELTING, D., HUNTER, L.L., KUMMER, A.W. and PAYNE, S.C., 2016. Clinical practice guideline: otitis media with effusion (update). *Otolaryngology - Head and Neck Surgery*, vol. 154, no. 1, suppl., pp. S1-S41. http://dx.doi. org/10.1177/0194599815623467. PMid:26832942.
- SADÉ, J., LUNTZ, M. and PITASHNY, R., 1989. Diagnosis and treatment of secretory otitis media. Otolaryngologic Clinics of North America, vol. 22, no. 1, pp. 1-14. http://dx.doi.org/10.1016/ S0030-6665(20)31462-6. PMid:2649851.
- SAMUELS, T.L., YAN, J.P., KHAMPANG, A., MACKINNON, W., HONG, N., JOHNSTON, N. and KERSCHNER, J.E., 2016. Association of microRNA 146 with middle ear hyperplasia in pediatric otitis media. *International Journal of Pediatric Otorhinolaryngology*, vol. 88, pp. 104–108. http://dx.doi.org/10.1016/j.ijporl.2016.06.056. PMid:27497395.
- SANTOS-CORTEZ, R.L.P., HUTCHINSON, D.S., AJAMI, N.J., REYES-QUINTOS, M.R.T., TANTOCO, M.L.C., LABRA, P.J., LAGRANA, S.M., PEDRO, M., LLANES, E.G.V., GLORIA-CRUZ, T.L., CHAN, A.L., CUTIONGCO-DE LA PAZ, E.M., BELMONT, J.W., CHONMAITREE, T., ABES, G.T., PETROSINO, J.F., LEAL, S.M. and CHIONG, C.M., 2016. Middle ear microbiome differences in indigenous Filipinos with chronic otitis media due to a duplication in the A2ML1 gene.

Infectious Diseases of Poverty, vol. 5, no. 1, pp. 97. http://dx.doi. org/10.1186/s40249-016-0189-7. PMid:27799062.

- SHRIKRISHNA, B.H., JYOTHI, A.C., SANJAY, G. and SAMSON, S.G., 2013. Age and gender differences in the incidence eof non-cholesteatomatous chronic suppurative otitis media. *International Journal of Research in Pharmaceutical and Biomedical Sciences*, vol. 4, no. 4, pp. 1172-1174.
- SILVA, M., OLIVEIRA, J., CARVALHO, A., SANTOS, D., LIMA, N., SANTOS, F., TABORDA, R., RODRIGUES, R., DALL'ACQUA, D. and MATOS, N., 2022. Colonization by Streptococcus pneumoniae among children in Porto Velho, Rondônia, Western Brazilian Amazon. *Brazilian Journal of Biology*, vol. 82, pp. e260617. https://doi. org/10.1590/1519-6984.260617.
- SILVA, M.D., LIMA, A., MARCAL, N., DIAS, L., GAMA, M. and SILLANKORVA, S., 2021. Identification of the bacterial pathogens in children with otitis media: a study in the Northwestern Portuguese district of Braga. *Microorganisms*, vol. 10, no. 1, pp. 54. http://dx.doi. org/10.3390/microorganisms10010054. PMid:35056502.
- STROMAN, D.W., ROLAND, P.S., DOHAR, J. and BURT, W., 2001. Microbiology of normal external auditory canal. *The Laryngoscope*, vol. 111, no. 11 Pt 1, pp. 2054-2059. http://dx.doi. org/10.1097/00005537-200111000-00035. PMid:11801996.
- VAL, S., POLEY, M., ANNA, K., NINO, G., BROWN, K., PEREZ-LOSADA, M., GORDISH-DRESSMAN, H. and PRECIADO, D., 2018. Characterization of mucoid and serous middle ear effusions from patients with chronic otitis media: implication of different biological mechanisms? *Pediatric Research*, vol. 84, no. 2, pp. 296-305. http://dx.doi.org/10.1038/s41390-018-0060-6. PMid:29915406.
- VAN INGEN, G., LI, J., GOEDEGEBURE, A., PANDEY, R., LI, Y.R., MARCH, M.E., JADDOE, V.W., BAKAY, M., MENTCH, F.D., THOMAS, K., WEI, Z., CHANG, X., HAIN, H.S., UITTERLINDEN, A.G., MOLL, H.A., VAN DUIJN, C.M., RIVADENEIRA, F., RAAT, H., BAATENBURG DE JONG, R.J., SLEIMAN, P.M., VAN DER SCHROEFF, M.P. and HAKONARSON, H., 2016. Genome-wide association study for acute otitis media in children identifies FNDC1 as disease contributing gene. *Nature Communications*, vol. 7, no. 1, pp. 12792. http://dx.doi. org/10.1038/ncomms12792. PMid:27677580.
- XU, J., DAI, W., LIANG, Q. and REN, D., 2020. The microbiomes of adenoid and middle ear in children with otitis media with effusion and hypertrophy from a tertiary hospital in China. *International Journal of Pediatric Otorhinolaryngology*, vol. 134, pp. 110058. http://dx.doi.org/10.1016/j.ijporl.2020.110058. PMid:32388082.
- ZHANG, Y., XU, M., ZHANG, J., ZENG, L., WANG, Y. and ZHENG, Q.Y., 2014. Risk factors for chronic and recurrent otitis media–a meta-analysis. *PLoS One*, vol. 9, no. 1, pp. e86397. http://dx.doi. org/10.1371/journal.pone.0086397. PMid:24466073.