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Nasopharyngeal Carriage of Potential Pathogenic Bacteria in Healthy Children Living in İstanbul

İstanbul'da Yaşayan Sağlıklı Çocuklarda Potansiyel Patojen Bakterilerin Nazofarengeal Taşıyıcılığın Araştırılması

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Abstract –

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Aim: The purpose of this study was to determine the prevalence of nasopharyngeal carriage of potentially pathogenic bacteria (especially *Streptococcus pneumoniae*, *Haemophilus influenzae*, *Moraxella catarrhalis*) in healthy 1-7 years old children and risk factors affecting it.

Methods: A total of four hundred forty-five healthy children (1-7 years old) from the European side of Istanbul (10 different units) were included in this cross-sectional study. Isolated microorganisms were identified by standard laboratory methods and the results were evaluated. Risk factors affecting the prevalence of nasopharyngeal carriage were also evaluated.

Results: In the study, 139 (31.2%) of the samples were positive. *S. pneumoniae* two (0.4%), *H. influenzae* 50 (11.2%); of them type b 10 (2.2%), *H. parainfluenzae* 10 (2.2%), *M. catarrhalis* 14 (3.1%), *S. aureus* 35 (7.9%), Group A β hemolytic streptococci (*Streptococcus pyogenes*) 13 (2.9%), Group B streptococci (*S. agalactiae*) 18 (4.0%), Non A Non B streptococci 16 (3.6%) were detected. The results obtained from different sites (places) were found to be variable in terms of pathogen density.

Conclusion: Nasopharyngeal carriage rates found in our study were generally lower than in some studies on this subject. However, when the samples taken from different places were evaluated one by one, it was seen that the colonization rate reached 45.9% in crowded and poor building properties (small, poorly ventilated, unhealthy buildings). The highest rate of *H. influenzae* carriage in the 2-3 age group was 33.3%. In this study, crowded and unhealthy school environments and low maternal education level have been determined as risk factors for increased nasopharyngeal carriage rate. Results of such studies vary depending on the region, sample frequency, individual and

Amaç: Bu çalışmada sağlıklı 1-7 yaş çocuklarda (*Streptococcus* pneumoniae, Haemophilus influenzae, Moraxella catarrhalis başta olmak üzere) potansiyel patojen bakterilerin nazofaringeal taşıyıcılık prevalansının ve bunu etkileyen risk faktörlerinin belirlenmesi amaçlanmıştır.

Yöntemler: Bu kesitsel çalışmaya İstanbul Avrupa yakasından (10 farklı birimden) 1-7 yaş grubu toplam dört yüz kırk beş sağlıklı çocuk dahil edildi. İzole edilen mikroorganizmalar standart laboratuvar yöntemleri ile tanımlanarak elde edilen sonuçlar değerlendirildi. Ayrıca, nazofaringeal taşıyıcılık prevalansını etkileyen risk faktörleri değerlendirildi.

Bulgular: Çalışılan örneklerin 139'u (%31,2) pozitifti. *S. pneumoniae* iki (%0,4), *H. influenzae* 50 (%11,2); bunlardan tip b 10 (%2,2), Haemophilus parainfluenzae 10 (%2,2), *M. catarrhalis* 14 (%3,1), *S. aureus* 35 (%7,9), A grubu β hemolitik streptokok (*Streptococcus pyogenes*) 13 (%2,9), B Grubu streptokok (*S. agalactiae*) 18 (%4,0), Non A Non B streptokok 16 (%3,6) oranlarında tespit edildi. Farklı yerlerden elde edilen sonuçlarda, patojen yoğunluğunun ve çeşitliliğinin değişken olduğu görüldü.

Sonuç: Çalışmamızda tespit edilen nazofaringeal taşıyıcılık oranları bu konuda yapılmış bazı çalışmalara göre düşük bulundu. Ancak örnek alınan farklı kurumlar tek tek değerlendirildiğinde, mevcudu fazla ve bina özellikleri kötü olan yerlerde (kalabalık, küçük, havalandırılması iyi yapılamayan, sağlıksız binalar) oranın %45,9'a kadar ulaştığı görüldü. H. influenzae taşıyıcılığında ise 2-3 yaş grubunda en yüksek oranı %33,3 olarak tespit edildi. Bu tür çalışmaların sonuçları bölgeye, örnekleme sıklığına, bireysel ve sosyal faktörlere göre değişiklik göstermektedir. Daha sağlıklı sonuçlara ulaşmak için daha fazla çalışmaya ihtiyaç vardır.

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social factors. Further studies are needed in order to achieve healthier results.

Keywords: Nasopharyngeal carriage, potential pathogenic bacteria, *S. pneumoniae*, *H. influenzae*, *M. catarrhalis*, vaccinated healthy children, risk factors

Anahtar Sözcükler: Nazofaringeal taşıyıcılık, potansiyel patojen bakteriler, *S. pneumoniae, H. influenzae, M. catarrhalis*, aşılanmış sağlıklı çocuklar, risk faktörleri

Introduction

Nasopharynx is heavily colonized by commensal bacteria as well as by a variety of microorganisms, including potential pathogens such as Streptococcus pneumoniae, Haemophilus influenzae (essentially non-typeable strains) and Moraxella catarrhalis. Nasopharyngeal microbiota occurs in the first year after birth, and changes throughout life due to different factors. Carriage rates generally decrease with age. However, these rates are also high in adults with chronic respiratory diseases. Reported nasopharyngeal carriage rates of potentially pathoge, nic bacteria vary widely between different studies and geographic regions. There are a lot of parameters including genetic factors and socio-economic conditions, access to health services, poor hygiene, number of family members, household crowding, access to health services, number of siblings, vaccination, age, type of child care, acute respiratory disease, and sleeping position (1). In Turkey, pneumococcal conjugate vaccine pneumococcal vaccine (PCV) 7 was introduced into the national immunization program in 2008, with a switch to PCV13 in 2011: the vaccination rate is estimated to be higher than 95% for children <2 years of age and it begins at 2 months of age with the other doses at 4 and a booster dose at 12 months. H. influenzae type b (Hib) vaccination is performed also at 2, 4 and 6 months of age with a booster at 18 months. Different results have been reported from different countries, and the different regions of Turkey about the rate of nasopharyngeal carriage of potentially pathogenic bacteria (1-3). In this study, we aimed to determine the rates of nasopharyngeal colonization of potential respiratory pathogens isolated from healthy school children vaccinated in Istanbul and some risk factors for nasopharyngeal carriage.

Methods

This cross-sectional study was conducted between 2014 and 2017 (sample collection, February-April 2016). The study involved four hundred forty-five children aged 1-7 years, asymptomatic, who had not taken antibiotics in the past 15 days, and were vaccinated in ten different institutions in the European side of Istanbul. Sociodemographic characteristics of the children (age,

gender, number of household members, average income, etc.) and situations that may be risk factors for carriage (antibiotic use in the past month, number of people sharing the same room in the family, presence of siblings under the age of 5, exposure to smoke at home, etc.) were questioned.

Nasopharyngeal samples were collected using cotton tipped swabs and immediately transferred into transport medium and delivered to the laboratory. On the same day, 5% sheep blood agar and chocolate agar (Becton Dickinson, USA) were cultured by single colony method. After 24 hours of incubation at 37 °C, Gram staining was performed from suspicious colonies that grew. Standard tests were applied for identification. For the detection of Streptococcus, bacteria displaying beta hemolysis (β-hemolysis) were first identified by classical methods. Group identifications were performed using bacitracin (0.04 units), trimethoprim (1.25 µg) -sulfamethoxazole (23.75 µg) discs and commercial latex agglutination kits (Dryspot Streptococcal Grouping Kit-Oxoid, England). Bacitracin-sensitive and trimethoprim-sulfamethoxazoleresistant ones were accepted as Group A beta-hemolytic streptococci (GABHS). Catalase test was applied for differentiation between streptococcus and staphylococcal microorganisms with similar hemolysis properties. In vitro coagulase test to determine the pathogenicity of growing staphylococci, optochin susceptibility test and bile dissolution tests to distinguish pneumococci were performed. H. influenzae needs both X and V factors simultaneously. With this feature, diagnosis was made. Typing was done by slide agglutination using antiserum specific for Hib (Denka Seiken™, Japan). Biochemical properties of the growing Gram-negative bacilli were also examined and detected. Identification of S. aureus isolates was performed according to standard microbiological procedures (morphology, Gram stain, catalase test, coagulase test, mannitol salt agar fermentation).

This research study was approved by the İstanbul University-Cerrahpaşa Faculty of Medicine Ethics Committee of Clinical Research (approval no: A-12/07 Jan 2014). Written informed consent was obtained from all of the parents of children who participated in the study.

Necessary permissions were obtained from the İstanbul Governorship, the Provincial Health Directorate and the Provincial Directorate of National Education for the study.

Results

Four hundred forty-five healthy children, including two hundred three girls (45.6%) and two hundred forty-two boys (54.4%), aged 1-7 years were included in the study. The number of positive specimens was a hundred thirty-nine (31.2%), and the carriage rate was 48% in girls and 52% in boys. Eleven bacteria species were isolated in the study. Potential pathogenic bacteria isolated from nasopharyngeal specimens are given below (Table 1, 2).

In this study, samples from 10 different educational institutions (10 different groups) were collected. The results obtained for five types of bacteria from each of these are given in Figure 1. It was observed that the obtained positivity rates and microorganism distributions differ between institutions. The results show that the carriage rates have increased significantly in crowded places. The positivity rate reached 45.8% in a crowded school. The carriage rates were between 12.5% and 45.8% in different institutions. *H. influenzae* was the most commonly detected microorganism with 33.3% in children aged 2-3 years in a nursery. Gender, exposure to

Table 1. Distribution of microorganisms isolated from nasopharyngeal specimens							
Distribution types	Microorganisms	n (%)					
1	H. influenzae	36 (8.9)					
2	H. influenzae + S. aureus	3 (0.7)					
3	H. influenzae + S. pyogenes	4 (0.9)					
4	H. influenzae + non-A non-B Streptococci	4 (0.9)					
5	H. influenzae + group B Streptococci	1 (0.2)					
6	H. influenzae + non-A non-B Streptococci + S. aureus	2 (0.4)					
7	H. parainfluenzae	8 (1.8)					
8	H. parainfluenzae + S. aureus	2 (0.4)					
9	S. aureus	25 (5.6)					
10	S. aureus + Pseudomonas	1 (0.2)					
11	M. catarrhalis	12 (2.7)					
12	M. catarrhalis + S. aureus	2 (0.4)					
13	S. pyogenes	9 (2.0)					
14	non-A non-B Streptococci	10 (2.2)					
15	Group B Streptococci	16 (3.6)					
16	Group B Streptococci + S. aureus	1 (0.2)					
17	S. pneumoniae	2 (0.4)					
18	Acinetobacter	1 (0.2)					
19	E. coli	1 (0.2)					
n: Number							

the negative effects of cigarette smoke (passive smoking), crowded household and recent respiratory infections do not constitute important risk factors for carriage; however, low maternal education level, and crowded and unsuitable school environments were found to be risk factors that increase carriage. There was no significant difference between the socio-economic levels of the groups. The characteristic features of the study group are given in Table 3.

Table 2. Microorganisms isolated from nasopharyngeal samples					
Bacteria types	Microorganisms	n (%)			
1	H. influenzae	50 (11.2)			
2	H. parainfluenzae	10 (2.2)			
3	M. catarrhalis	14 (3.1)			
4	S. pneumoniae	2 (0.4)			
5	S. aureus	35 (7.9)			
6	S. pyogenes	13 (2.9)			
7	Group B Streptococci	18 (4.0)			
8	Non-A Non-B Streptococci	16 (3.6)			
9	Acinetobacter	1 (0.2)			
10	Pseudomonas	1 (0.2)			
11	E. coli	1 (0.2)			
n: Number					

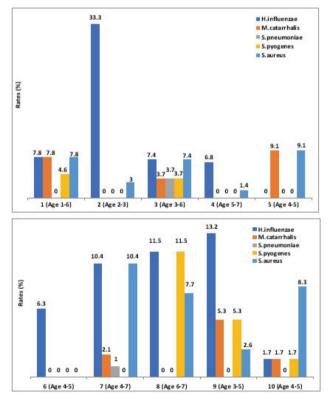


Figure 1. Carriage percentages of five bacteria in ten different educational institutions

Discussion

For hundred and forty-five healthy children aged 1-7 years from 10 different places (nursery, kindergarten and primary school 1st grade) were included in the study. According to the results we have achieved; the total nasopharyngeal carriage rate for 11 types of potential respiratory pathogens was 31.23%: *S. pneumoniae* 0.44%, *H. influenzae* 11.23%; type b 2.24%, *H. parainfluenzae* 2.24%, *M. catarrhalis* 3.14%, *S. aureus* 7.86%, *S. pyogenes* 2.92%. It is seen that the results

Table 3. Characteristics of 445 children						
Characteristics (n=445)	Number (%)					
Sex						
Male	242 (54.4)					
Female	203 (45.6)					
Age						
1-7	445 (100)					
Education of mother						
High	98 (22)					
Low	347 (78)					
Number of people at home						
≤3	136 (30.6)					
≥3	309 (69.4)					
Exposure to smoke at home						
Yes	254 (57.1)					
No	191 (42.9)					
Family economic status						
Low	-					
Medium	445 (100)					
High	-					
n: Number						

about multiple bacteria colonization were obtained only in a small number of publications and they are different from each other (2,4-6). This situation can be attributed to the effect of many factors on nasopharyngeal carriage (1).

The distribution of the pathogens are shown in Table 4 and compared with other studies' findings.

Torun et al. (2) conducted a study with 330 healthy children in the 6-10 age group in istanbul and they found *H. influenzae* in 48.7%, type b 7.2%, *S. pneumoniae* in 29.1%, and *M. catarrhalis* in 23.9%. This study is most suitable for comparing our results with the situation in previous years. However, given the possible differences in risk factors, healthy comparisons cannot be made.

Principi et al. (4) conducted a study with a thousand seven hundred twentythree children aged 1-7 years from eighteen schools and day care centers in eighteen Italian cities. They found that the rate of carriage respiratory pathogens was 17.9% (*S. pneumoniae* 3.5%, *H. influenzae* 11.9%, *M. catarrhalis* 4.1%). Especially in children aged 1-5, siblings under 3 years of age, living in rural areas and staying in a full-day care home significantly affect carrier rates; however, they found that breast-feeding, gender, passive smoking, and recent respiratory infections were not significant variables. In this study, it was reported that antibiotic treatment in the previous three months did not affect nasopharyngeal carriage and *S. pneumoniae* carriage was associated with recurrent macrolide treatment.

In Belgium, a study with three hundred thirthy three healthy children of different socio-economic levels aged 3-6 point out that 50% of children were colonized at least once with *S. aureus*, 69% with *S. pneumoniae*, 67% with *M. catarrhalis* and it was found that 83% of them were colonized with *H. influenzae* (5).

Table 4. Distribution of pathogens in our study and compared with other studies from different countries.									
Study (Reference)	Country	n	S. pneumoniae (%)	H. influenza (%)	H. influenza type B (%)	H. parainfluenza (%)	M. catarrhalis (%)	S. aureus (%)	S. pyogenes (%)
Our Study	Turkey	445	0.44	11.23	2.24	2.24	3.14	7.86	2.92
Torun et al. (2)	Turkey	330	29.1	48.7	7.2	_	23.9	_	_
Principi et al. (4)	Italy	1723	3.5	11.9	_	_	4.1	_	_
Jourdain et al. (5)	Belgium	333	67	_	_	_	83	69	_
Dunne et al. (6)	Indonesia	302	49.5	27.5	_	_	42.7	7.3	_
Abaye et al. (8)	Ethiopia	_	18.4	_	_	_	_	_	_
Wada et al. (9)	Ethiopia	_	43.8	_	_	_	_	_	_
Claire et al. (11)	Kenya	323	19.4-20.0	_	_	_	_	_	_
Birindwa et al. (12)	Congo	794	21	_	_	_	_	_	_
Koliou et al. (13)	Cyprus	402	35.3	_	_	_	_	_	_
Yang et al. (16)	China	17338	_	17.1	_	_	_	_	_
Ortiz-Romero et al (17)	Spain	404	_	_	_	_	_	_	_
Toprak et al. (21)	Turkey	1129	_	_	_	_	_	_	6.5

In a study conducted by Dunne et al. (6) with three hundred two healthy children aged 12-24 months, the carriage prevalence was 49.5% for *S. pneumoniae*, 27.5% for *H. influenzae*, 42.7% for *M. catarrhalis* and 7.3% for *S. aureus*.

In our country, studies conducted in Istanbul in recent years showed that the rate of *S. pneumoniae* carriage varies depending on age and the rates decreased after vaccination (3,7).

Arvas et al. (3) found 14% *S. pneumoniae* carriage in their study with a hundred fifty vaccinated healthy children (in 2014) aged 0-6 in İstanbul. However, the rate of *S. pneumoniae* carriage in young children of 24 months has been reported to be 76.2%. The low, 0.44%, *S. pneumoniae* rate detected in our study suggests that other factors may also be effective. In our study, this can be attributed to the fact that the number of children under 2 was quite low.

In a study by Soysal et al. (7), the low prevalence of pneumococcal carriage was remarkable. Soysal et al. (7) determined the prevalence of pneumococcal carriage as 6.4% 5 years after PCV-7 vaccination and 2 years after PCV-13 childhood vaccination in 2,165 healthy children in the 0-18 age group in Istanbul between 2011 and 2013.

In recent years, there are important findings related to pneumococcal carriage risk factors in studies conducted abroad: the number of siblings under certain ages, sleeping in the same room, crowded family environment, young age, mothers with low education level and low socioeconomic status were found to be related to carriage (8-10).

In their study including children aged ≤6 years, Abaye et al. (8) found a general pneumococcal carriage rate of 18.4%. It was determined that there was no significant variation in colonization according to gender and age of children. Children living with younger siblings (1-2<6 years) and number of persons >5 in the household were associated with higher rates of *S. pneumoniae* carriage.

Wada et al. (9) found a *S. pneumoniae* carriage rate of 43.8% in seven hundred fourteen children aged 3-13 years. The rate was determined as 62.5% in 3-5 years old children and 38.6% in 6-13 years old children. Sleeping with siblings and age below five years were significantly associated with *S. pneumoniae* carriage. Findings of Fadlyana et al. (10) emphasized the association of the level of maternal education and socio-economic factors with nasopharyngeal carriage. Having more than one small child in the household has been identified as a risk factor for transport of potential pathogenic bacteria. There was no difference in the frequency of carriage of *S. pneumoniae*, *H. influenzae*, *M. catarrhalis* and *S. aureus* among urban and semi-rural children. Low maternal

education level is also valid for our study group as a risk factor that increases carriage.

Some publications suggest that vaccination affects the overall prevalence of pneumococcal carriage, and among other known risk factors, the prevalence of serotypes included in the vaccine (vaccine strains) is lower in vaccinated children and the prevalence is higher in children without vaccination and in poor living conditions (11-13). It has been reported that pneumococcal conjugate vaccines could have a protective effect against colonization by drug-resistant pneumococcal strains (11-13).

Claire et al. (11) determined in a study with nasopharyngeal swab samples from 323 children aged 4-7 years from coastal Kenya before and after vaccination with 10-valent pneumococcal vaccine, it was found that vaccination did not decrease the overall prevalence of pneumococcal carriage; the number of colonized children before vaccination was 65 (20%) and the number of colonized children after vaccination was 63 (19.4%). However, they determined that the prevalence of serotypes (vaccine strains) included in the vaccine decreased from 43% to 19%.

In 2014 and 2015 Birindwa et al. (12) investigated, pneumococcal carriage in seven hundred ninety fdur healthy children aged 1 to 60 months who attended health centers in the east part of Congo and who were vaccinated with a 13-valent conjugated PCV13 in 2013. According to their results, the rate of pneumococcal carriage was 21%. They found that the rate of carriage was higher in children who had not received PCV13 vaccination, lived in rural areas, in a house without an enclosed kitchen, malnourished and had high fever. It is stated that the results show that in addition to PCV13 vaccination, better living conditions are needed to reduce pneumococcal burden. In the study, worryingly decreased susceptibility to commonly used antibiotics for pneumococci infections and need for more appropriate use of antibiotics in the region were highlighted.

Koliou et al. (13) found that 35.3% of the children were colonized with *S. pneumoniae* in a study they conducted with four hunred two healthy children aged 6 months-5 years in Cyprus. Attending a day-care center has been identified as a risk factor that increases colonization approximately by 3 times. They stated that factors such as age, gender, breastfeeding history and passive exposure to cigarette smoke were not statistically significantly related to the risk of *S. pneumoniae* colonization. On the other hand, complete vaccination with PCV7 has been shown to have a protective effect against colonization compared to unvaccinated children (the risk decreased two times).

Conclusion

It has been stated that the careful use of antibiotics for upper respiratory tract infections, especially in children, and increasing the coverage of pneumococcal conjugate vaccines can be effective in reducing levels of colonization by drug-resistant pneumococcal strains.

In our study, the *H. influenzae* carriage rate was 11.23%, while in Italy, it was 14.1% (14) in a study involving seven hundred seventeen healthy children <6 years of age, 15 years after vaccination of Hib. On the other hand, in a study conducted in Spain, it was found that 42% of 960 healthy <5 years old children were colonized with *H. influenzae* and the rate varied between centers (12-83%) (15). In our study, it was determined that there was variability among institutions (0-33.3%).

H. influenzae carriage rate was determined to be 17% in children in a study in which a total of 42 studies with 17,388 participants were included in China. It was reported that the rate was relatively low in healthy population, but showed great differences between provinces (16). This study also explains the differences in carriage rates.

Ortiz-Romero et al. (17) performed a study with four hundred four healthy children <5 years of age in Spain and determined the rate of *H. influenzae* carriage as 73.2% in winter and 26.8% in summer.

As a result, according to our findings, *Haemophilus influenzae* carriage rates vary depending on factors such as study region, centers, seasons (15-17).

Staphylococcus aureus carriage rates are known to vary by country, demographic group and occupation (5,18).

Streptococcus agalactiae, known as group B Streptococcus GBS, is an important cause of neonatal sepsis and meningitis (19). In our study, 4.04% of carriers were detected to have this bacterium. The natural reservoir for GABHS is humans. They are the most common cause of streptococcal pharyngitis. It is generally seen in school age children at the ages of 5-15 (20). Toprak et al. (21) detected 6.5% S. pyogenes carriage in 6.5% of 1,129 primary school children in the 7-15 age group in the city center of Afyonkarahisar. Tonsillopharyngitis developed in 5.5% of the children within a year. In our study, S. pyogenes carriage rate was found to be 2.9%. However, in one group (6-7 years) the rate was 11.5%.

When looking at the results obtained from different places in different studies, the differences between them (carriage rates and microorganism diversity, etc.) are thought-provoking. In order to minimize the health problems associated with carriage, there is a need for new studies in which the determination of carriage rates will be handled together with risk factors.

It is an advantage of our study that this study was not performed in a single center. We observed that

the carriage prevalence varies considerably between institutions. At the same time, the change in the diversity of microorganisms has emerged strikingly. Low maternal education level and crowded classroom environments were found to increase the prevalence of nasopharyngeal carriage. Especially *S. pneumoniae* carriage rate was very low. Among the possible reasons for this result; the age of the children in our study (scarcely any number of children under the age of 2) and their vaccination are important. Besides, the leading factors affecting the results are: there was no significant difference in the socio-economic status which was at a good level, between the families, and the children could benefit from health services adequately and correctly.

Authorship Contributions

Concept: N.G., S.Y., O.A., Z.T., S.Ö., S.A., H.T., F.K.Ç. Design: N.G., S.Y., O.A., Z.T., S.Ö., S.A., H.T., F.K.Ç. Data Collection or Processing: N.G., S.Y., O.A., Z.T., S.Ö., S.A., H.T., F.K.Ç. Analysis or Interpretation: N.G., S.Y., O.A., Z.T., S.Ö., S.A., H.T., F.K.Ç. Literature Search: N.G., S.Y., O.A., Z.T., S.Ö., S.A., H.T., F.K.Ç. Writing: N.G., S.Y., O.A., Z.T., S.Ö., S.A., H.T., F.K.Ç.

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