

# Results Of Microbiological Research In Patients With Acute Odontogenic Purulent Inflammatory Diseases

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## Introduction

**Relevance of the study.** In children with odontogenic inflammatory diseases, a sharp change in inflammatory processes is observed due to the relative maturity of organs and tissues, imperfection of immunity, incomplete formation of lymphatic tissues, the presence of anatomical and physiological features of the structure of teeth and jaws, susceptibility to injury and its complications. The results of numerous epidemiological studies show that an average of 3-5 acute respiratory viral infectious diseases (ORVI) have been found to have a tendency to be diagnosed per child per year. The greatest prevalence of acute respiratory infections is seen in young children, preschool children and primary school children. Children in the first 3 years of life are diagnosed with infectious diseases 10-10 times more often than children aged 2-2.5 years and older in one year. Severe respiratory infections lead to a violation of the functional state of the body, lead to a violation of adaptation and the development of chronic pathology [1.3.5.7.9].

Diseases of the affected infection are often viruses (mainly respiratory syncytial, influenza and parainfluenza, adenoviruses), pathogens of the Chlamydia and Mycoplasma family (especially Chlamydia pneumoniae and Mycoplasma pneumoniae), Haemophilus influenzae bacteria (often Type B), Streptococcus pneumoniae, S. pyogenes, Staphylococcus aureus, Moraxella [2.4.6]. The main bacterial agents are beta-hemolytic streptococci Group a (BGSG), Staphylococcus aureus (S. aureus), which are able to form mixed infections with respiratory viruses [10.12.21.22]. Low immune system, metabolic disorders, decreased pain sensitivity change the clinical picture and the course of many surgical diseases, which can lead to serious diagnostic errors and negatively affect the results of surgical pathology.

It was concluded that odontogenic inflammatory diseases and their complications in children with frequent diseases are precisely microcirculation disorders, the presence of microtrombs, dystrophic and necrotic processes, the predominance of the inflammatory component over reparative, the formation of cell proliferation, the formation of phagocytic activity of leukocytes, incomplete phagocytosis, microbiological damage to their tissues with certain

characteristics characterized by high-level microcirculation diseases, a decrease in general and local immunological reactivity. An analysis of studies conducted in Uzbekistan showed that scientific research was carried out on the prevalence, clinical picture, etiopathogenesis, treatment methods, preventive measures of jawperiostitis, abscess and phlegmona, odontogenic osteomyelitis in children with odontogenic inflammatory diseases. Among dental diseases, the number of people with odontogenic inflammatory diseases accounts for 10-11% of total applicants, which is the 2nd most common after caries and its complications [11.13.15.17.19].

A number of studies have been carried out among preschool and school-age children with odontogenic inflammatory diseases, focusing on the methods of treatment of jawperiostitis, abscess and phlegmona, prevalence, clinical manifestations of odontogenic osteomyelitis, etiopathogenesis. There has also been no development of a perspective-setting system for the Prevention of odontogenic inflammatory diseases among school-age children. Therefore, the urgent solution to the need for a complex approach in determining the problem of odontogenic inflammatory diseases in preschool and school-aged people, to identify the advantages and disadvantages of various programs based on comparative analysis, and to offer optimal solutions is one of the necessary problems.

Microbiological analysis was carried out in order to determine the microflora composition in patients with acute odontogenic purulent-inflammatory diseases, as well as correlations in research groups of patients. Acute odontogenic purulent-inflammatory diseases in 151 children, when a purulent exudate obtained during surgery was studied, it was determined that 62.5% of the 8 species studied from colony-forming microorganisms were obligate-anaerobic bacteria, among them *Neisseria* spp., *Bacillus* sp., *E. cloacae*, *Proteus* spp., *Cl. perfringens* defined. The concentration of microorganisms listed in the pus exudate varied from  $3.71 \cdot 10^3$  to  $2.71 \cdot 10^5$  Koe/ml (colony-forming unit in 1 ml). Facultative-anaerobic microorganisms were formed from more gram-positive species (staphylococci, streptococci, corinebacteria). The concentration of these microorganisms in the pus furnace changed from  $2 \cdot 10^3$  to  $1 \cdot 10^6$  Koe/MLD, that is, their amount is 10-100 times less than the amount of obligate anaerobes [12.14.16.18]. This data confirms the opinion of most researchers about the predominance of anaerobic microflora in diseases of the face-jaw odontogenic inflammation.

**The purpose of the study.** Patients with acute odontogenic purulent-inflammatory diseases improve methods of studying the microflora of the oral cavity. The analysis we conducted showed that the composition and structure of microflora in different age groups is faqr [18.19.20.21.22]. The least number of species were identified in children between 2 and 3 years of age. This age group was dominated by streptococci and corinebacteria. It is noteworthy that the composition of the complications included 2-3 species, in which the quantitative ratio facultative and obligate anaerobes became almost identical (from  $2 \cdot 10^4$  to  $1 \cdot 10^6$  Koe/ml). The overall concentration of microorganisms was one to two times lower compared to other groups. A similar rate was observed in children aged 4-6 years. But in the composition of the assosias appeared anaerobic microorganisms that do not form spores (bacteroids, fuzobacteria). Between 2 and 4 species were identified in the associations. In the case where anaerobes were present, they dominated the Assyrians and were expressed in very

high concentrations ( $1.8 \times 10^8$  Koe/ml). In the age group of 7-11, both a sharp increase in the number of microbial complications and the predominance of anaerobes that do not form spores were observed in them, the frequency of bacteriological separation of Streptococci, staphylococci, corinebacteria decreased. Among anaerobes, a lot of Fusobacteria, peptostreptococci, bacteroids were identified. That being said, the amount of staphylococci isolated in this age group was only 7.6%, at which time the literature recorded a significant number of observations of these microorganisms in the observation group of inflammatory diseases of the face-jaw in children.

An expressive trend towards greater dominance in strict anaerobic Association was also found in the 12-17 age group. There was an increase in both Gram-negative (bacteroids, fuzobacteria) and Gram-positive bacteria. In this case, the separation frequency of anaerobes was higher than 60.0%, that is, the microbial "landscape" approached the appearance in adults [17.19.21.22]. At the same time, the percentage of Streptococci remained high enough, which is not observed in adults. In this age group, there was a further decrease in the percentage of corinebacteria and staphylococci, the other specific side of which was isolated (4.0%). In this group, the concentration of microorganisms ranged from  $2 \times 10^6$  to  $4 \times 10^8$  Koe/ml, in which obligate anaerobes were separated in higher concentrations than facultative ones. There are acute odontogenic purulent inflammatory diseases the fact of a large difference in the composition of ascocias in children of different groups, in our opinion, most of all, patients are associated with the peculiarities of oral microbiocenosis, which, undoubtedly, is associated with a weakening of the child's body at the base of accompanying diseases in the observation group.

The above makes it possible to conclude that in the formation of acute odontogenic purulent inflammatory diseases in patients of the observation group, there is not only an odontogenic pathway of entry into the microflora, but also an endogenous pathway [20.21.22]. On the day of application of patients with acute odontogenic purulent inflammatory diseases, an emergency surgical purulent inflammatory ulcer was opened, and an ECMA was taken for microbiological research from an exudate detached from the purulent ulcer. The results of the microbiological study are presented.

In an analysis of research materials from Observation Group (N=122) patients, *S. aureus* has been observed in 75 (61.47%) of total patients, with an average of  $4.83 \pm 0.17 \times 10^5$  KHQB Log/ml detected. *pyogenes* was found to average  $4.21 \pm 0.26 \times 10^5$  KHQB Log/ml in 49 (40.16%) patients. The next place in the occurrence of the bacterial environment in the patients of the observation group is *E. coli* was occupied, averaging  $2.33 \pm 0.14 \times 10^4$  KHQB Log/ml per 42 (34.43%) of total patients. *Cl. perfringens* is found in 32 (26.23%) patients with an average of  $2.27 \pm 0.21 \times 10^4$  KHQB Log/ml, *Proteus* spp. An average of  $2.71 \pm 0.21 \times 10^5$  KHQB Log/ml was found in 25 (20.49%) patients. With similar rates, 23 (18.85%) patients had *E. cloacae* averages  $2.34 \pm 0.19 \times 10^4$  KHQB Log/ml and *Neisseria* spp. it averaged  $3.71 \pm 0.21 \times 10^3$  khqb from Log/ml. *Bacillus* sp, identified only in the minimum 6 (4.91%) patients in the bacterial environment. showed an average of  $1.08 \pm 0.19 \times 10^5$  KHQB Log/ml.

**Conclusion.** In patients with acute odontogenic purulent inflammatory diseases existing control group (n=29), a large difference in the composition of the complications was observed. Bunda *S. aureus* averaged  $3.32 \pm 0.18 \times 10^5$  KHQB Log/ml, s in 22 (75.86%) patients. *pyogenes* averaged  $2.87 \pm 0.42 \times 10^5$  KHQB Log/ml, *E. coli* was found to

average  $6.75 \pm 0.21 \times 10^3$  KHQB Log/ml in 4 (13.79%) patients. Only 1 (3.45%) of control group patients had *Cl. perfringens* have been observed at  $8.91 \times 10^2$  KHQB Log/ml whereas 2 patients have *Proteus* spp. as well as *E. cloacae* were identified at  $4.28 \pm 0.19 \times 10^3$  KHQB Log/ml and  $4.54 \pm 0.22 \times 10^3$  KHQB Log/ml, respectively. In 3 (10.34%) patients of the group, *Neisseria* spp. The average was found at  $8.20 \pm 0.28 \times 10^2$  KHQB Log/ml. In control group patients, *Bacillus* sp. was not detected.

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