

АЛЛЕРГОЛОГИЯ И ИММУНОЛОГИЯ В ПЕДИАТРИИ ALLERGOLOGY AND IMMUNOLOGY IN PEDIATRICS

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TOPICS IN THE ISSUE:

ROLE OF NONSPECIFIC RISK
FACTORS IN ATOPIC DERMATITIS

THE EFFECT OF VITAMIN D
PROVISION ON BIOMARKERS OF
INFLAMMATION
IN BRONCHIAL ASTHMA IN
CHILDREN

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REPORT

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The aim of this journal is to promote and maintain professional contacts and interactions between basically and clinically oriented allergologists and immunologists. This journal is the official organ of the Association of Pediatric Allergists and Immunologists of Russia (APAIR). «Allergology and Immunology in Pediatrics», founded in 2003, is devoted to the wide spectrum of interests of the pediatricians, allergists and immunologists in clinical practice and related research. As the journal intends promoting productive communication between scientists engaged in the basic research and clinicians working with children, both experimental and clinical research related findings are accepted for publication. The regular format of the Journal includes original articles, concise communications, case reports, discussions, comprehensive reviews, book reviews, correspondence, news, recent advances in clinical research, and selected APAIR proceedings. The Journal also presents Selected Abstracts from other periodicals in related disciplines. Areas of interest also includes but not limited to the evaluation, management and prevention of allergic and other immune-mediated diseases with a special attention to the pediatric allergy and asthma. Furthermore, new sections and activities focusing on the continuing medical education will be introduced shortly. «Allergology and Immunology in Pediatrics» is published quarterly (4 volumes per annum). The journal was founded in 2003. From 2003–2004 it was called Scientific and Practical Journal of Allergology and Immunology in Pediatrics. From 2004 to the present time it is called «Allergology and Immunology in Pediatrics». The journal is published 4 times a year.

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Postal address:

«Allergology and Immunology in Pediatrics» 6 Ostrovityanova st., 117513, Moscow, Russia, phone: 8(495) 225-71-04
Fax: 8(495) 225-7107. E-mail: adair@adair.ru

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АЛЛЕРГОЛОГИЯ И ИММУНОЛОГИЯ В ПЕДИАТРИИ

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Журнал «Аллергология и иммунология в педиатрии» — рецензируемое научно-практическое периодическое издание, предназначенное для педиатров, аллергологов-иммунологов, а также специалистов разного профиля, работа которых связана с областью педиатрической аллергологии и иммунологии. Журнал является официальным печатным органом Ассоциации детских аллергологов и иммунологов России (АДАИР); издается при участии ведущих специалистов страны — педиатров, аллергологов, клинических иммунологов. На страницах издания — оригинальные статьи, образовательные программы для врачей, клинические наблюдения, дискуссии, информация о последних достижениях отечественной, зарубежной науки и практики. Все публикации журнала связаны с вопросами диагностики, лечения, профилактики аллергических и других иммуноопосредованных заболеваний у детей с акцентом на детскую аллергологию. Журнал основан в 2003 году. С 2003–2004 гг. носил название «Научно-практический журнал Аллергология и иммунология в педиатрии». В 2004 году переименован и носит название «Аллергология и иммунология в педиатрии».

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Role of nonspecific risk factors in atopic dermatitis

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Dali Sh. Macharadze¹, Ekaterina A. Rassanova², Tatyana A. Ruzhentsova¹, Alena V. Galanina³, Vladimir S. Malyshev⁴

¹ *Research Institute of Epidemiology and Microbiology named after G. N. Gabrichevsky Rospotrebnadzor, 10 Admiral Makarov st., Moscow, 125212, Russia.*

² *Federal State Budgetary Educational Institution of Higher Education «Kirov State Medical University» of the Ministry of Healthcare of the Russian Federation, 112 K. Marx st., Kirov, 610998, Russia*

³ *Federal State Autonomous Educational Institution of Higher Education «N. I. Pirogov Russian National Research Medical University» of the Ministry of Health of the Russian Federation, 1 Ostrovityanova st., Moscow, 117997, Russia*

⁴ *«FIDES Lab», Limited Liability Company, 9 Altufyevskoe shosse, Moscow, 127106, Russia*

Dali Shotaevna Macharadze — Doc. Sci., Leading Researcher of Research Institute of Epidemiology and Microbiology named after G. N. Gabrichevsky Rospotrebnadzor, ORCID ID: 0000-0001-5999-7085, e-mail: dalim_a@mail.ru.

Ekaterina Andreyevna Rassanova — assistant of the Department of pediatrics Federal State Budgetary Educational Institution of Higher Education “Kirov State Medical University” of the Ministry of Healthcare of the Russian Federation, ORCID ID: 0009-0005-5298-056X, e-mail: ekaterinarassanova@yandex.ru.

Tatyana Alexandrovna Ruzhentsova — Doc. Sci., Deputy Director of Research Institute of Epidemiology and Microbiology named after G.N. Gabrichevsky Rospotrebnadzor, ORCID ID: 0000-0002-6945-2019, e-mail: ruzhencova@gmail.com.

Alena Vasilevna Galanina — Doc. Sci., Professor of Department of Federal State Autonomous Educational Institution of Higher Education «N. I. Pirogov Russian National Research Medical University» of the Ministry of Health of the Russian Federation, ORCID ID: 0000-0003-1670-0506, e-mail: alenagalanina@yandex.ru.

Vladimir Sergeevich Malyshev — Doc. Sci. (Biol.), Head of the Lab Department of «FIDES Lab», Limited Liability Company, ORCID ID: 0009-0009-5351-4893, e-mail: com.delafere@mail.ru.

Annotation

The increasing prevalence of atopic dermatitis (AD) over recent decades suggests that environmental factors play an important role in the etiology and pathogenesis of the disease. Nonspecific factors refer to external (or exposomal) factors and include human and natural factors that influence the health of a population: for example, the socioeconomic status of the patient; climate, including air temperature, exposure to ultraviolet radiation, air pollution; and living in a city or rural area. Although studies have shown the influence of these factors on the course of AD, in general, none of them significantly increases or decreases the risk of developing the disease. This review briefly discusses studies on the role of nonspecific environmental risk factors and their impact on the course of AD in children and adults.

Keywords: atopic dermatitis, nonspecific risk factors, climate, socio-economic conditions, urban/rural environment.

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Роль неспецифических факторов риска при атопическом дерматите

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For correspondence:

Dali Sh. Macharadze, Research Institute of Epidemiology and Microbiology named after G. N. Gabrichevsky Rospotrebnadzor.

Address: 10 Admiral Makarov st., Moscow, 125212, Russia.

E-mail: dalim_a@mail.ru.

Для корреспонденции:

Мачарадзе Дали Шотаевна, д. м. н., в. н. с. клинического отдела ФБУН МНИИЭМ им. Г. Н. Габричевского Роспотребнадзора.

Адрес: 125212, Москва, ул. Адмирала Макарова, д. 10.

E-mail: dalim_a@mail.ru.

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Мачарадзе Д. Ш.¹, Рассанова Е. А.², Руженцова Т. А.¹, Галанина А. В.³, Малышев В. С.⁴

¹ ФБУН МНИИЭМ им. Г. Н. Габричевского Роспотребнадзора, 125212, Москва, ул. Адмирала Макарова, д. 10, Россия

² ФГБОУ ВО Кировский ГМУ Минздрава России, 610998, г. Киров, ул. К. Маркса, д. 112, Россия

³ ФГАОУ ВО РНИМУ им. Н. И. Пирогова Минздрава России, 117997, г. Москва, ул. Островитянова, д. 1, Россия

⁴ ООО «Фидес Лаб», 127106, г. Москва, Алтуфьевское ш., д. 9, Россия

Мачарадзе Дали Шотаевна — д. м. н., в. н. с. клинического отдела ФБУН МНИИЭМ им. Г. Н. Габричевского Роспотребнадзора, ORCID ID: 0000-0001-5999-7085, e-mail: dalim_a@mail.ru.

Рассанова Екатерина Андреевна — ассистент кафедры педиатрии ФГБОУ ВО Кировский ГМУ Минздрава России, ORCID ID: 0009-0005-5298-056X, e-mail: ekaterinarassanova@yandex.ru.

Руженцова Татьяна Александровна — д. м. н., зам. директора по клинической работе ФБУН МНИИЭМ им. Г. Н. Габричевского Роспотребнадзора, ORCID ID: 0000-0002-6945-2019, e-mail: ruzhencova@gmail.com.

Галанина Алена Васильевна — д. м. н., профессор кафедры пропедевтики детских болезней ФГАОУ ВО РНИМУ им. Н. И. Пирогова Минздрава России, ORCID ID: 0000-0003-1670-0506, e-mail: alenagalanina@yandex.ru.

Малышев Владимир Сергеевич — д. б. н., зав. Лабораторией ООО «Фидес Лаб», ORCID ID: 0009-0009-5351-4893, e-mail: com.delaferre@mail.ru.

Аннотация

Увеличение распространенности атопического дерматита (АтД) за последние десятилетия свидетельствует о том, что факторы окружающей среды играют важную роль в этиологии и патогенезе заболевания. Неспецифические факторы относятся к внешним (или экспозомным) факторам и включают человеческие и природные факторы, влияющие на здоровье популяции: например, социально-экономический статус больного; климат, в том числе температуру воздуха, воздействие ультрафиолетового излучения, загрязнение воздуха; а также проживание в городе или сельской местности. Несмотря на то, что в исследованиях показано влияние этих факторов на течение АтД, в целом ни один из них достоверно не увеличивает или не снижает риск развития заболевания. В этом обзоре кратко обсуждаются исследования, посвященные роли неспецифических внешних факторов риска и их влиянию на развитие и течение АтД у детей и взрослых.

Ключевые слова: атопический дерматит, неспецифические факторы риска, климат, социально-экономические условия, городская/сельская среда.

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The pathogenesis of atopic dermatitis (AtD) is affected by a number of environmental factors in different ways (i.e. exposomes) [1–3]. External (exposome) factors are divided into *nonspecific*, or a *general external environment*, (human and natural factors, affecting the health of the population, — climate, urban environment, social and economic conditions); *specific* (air humidity, ultraviolet (UV) radiation, allergens, microbes, diet, tobacco and other pollutants) and *internal environment, dependent on the host* (interaction between body cells, for example, cutaneous and intestinal microbiota; inflammation and oxidative stress) [2, 3].

According to the latest data, mechanisms, responsible for the onset and exacerbation of AtD, involve interactions between genes and epithelial barrier, immune disorders, skin dysbiosis as well as the effect of various environmental factors [4, 5]. Due to impaired

skin barrier function, in particular due to a genetic defect (including filaggrin mutations), penetration of allergens and infectious agents into the skin becomes easier [6].

According to the new concept — “*hypothesis of epithelial barrier*” — inflammation in the epithelial layer, which covers the skin surface as well as respiratory, urogenital and gastrointestinal tract, develops, first of all, when patients live in the urban environment [1]. Epithelial cell activation and release of such cytokines as IL-33, IL-25, thymic stromal lymphopoietin (TSLP), etc., upon allergen exposure, infectious agents or tissue damage due to skin itching, cause immune responses of Th2-type [7, 8]. Although Th2 axis is considered major in AtD pathogenesis, recent studies confirm the involvement of additional immune pathways, in particular Th1, Th17 and Th22-lymphocytes [7, 8]. As a result of impaired epi-

thelial barrier function, due to external factors, there is also change in the microbiome structure; in conjunction with disorders of immune regulation, it affects the maintenance of chronic inflammation in the skin [2,4–6].

Any of exposome factors might cause exacerbation and trigger AtD. The most significant role of external factors (e.g., air pollution and climate) in children with variants of filaggrin mutation, having lesions on exposed skin (face, arms, neck) [6]. The famous American scientist Leung

D. believes that other equally important triggers with AtD are *S. aureus*, herpes simplex virus, stress and allergens [4].

Although the impact of various risk factors on AtD development has been shown in numerous studies and confirmed under experimental conditions, their role in the start of the atopic march has not been determined yet [7].

Meanwhile, a wide variety of risk factors with AtD has been studied mainly in children: starting from nutrition and ending with certain external and internal, in particular, in different periods (pre- and postnatal) child development [8].

In many cases patients use exposure to the specific triggers to control their disease activity. Among the many environmental factors that affect the course of AtD, some can even play a protective role (e.g., consumption of unpasteurized milk) [2, 3].

NONSPECIFIC FACTORS

Urbanization, air pollution and climate change are the global factors, which affect public health worldwide, in contrast to the local ones, one of which is, in particular, the healthcare system in specific regions.

HEALTHCARE

Availability and quality of healthcare differ, depending on economic development of the country of residence, social status, race and ethnicity, including AtD patients [9]. It is clear that obstacles to obtaining highly qualified medical help are a lack of expert physicians and even medical institutions with a modern infrastructure [2]. Also, if a patient belongs to an economically poor family or lives in conditions with limited access to education, this directly affects the effectiveness of treatment, including the opportunity to purchase necessary medicine. Particularly, in pa-

tients with bronchial asthma the above-mentioned socio-economic conditions lead to an increase in exposure to allergens and the development of more frequent and severe attacks of the disease [10]. Also, dilapidated and old housing as well as living in ecologically unfavourable areas were associated with increased prevalence and a more severe course of AtD in children [9, 11]. On the other hand, a reduction in the risk of AtD in the children, living in rural areas was revealed in the population (black population of South Africa), homogeneous by ethnicity and geographical origin [12]. It has long been known that colonization of the skin *S. aureus* is a risk factor, with which researches associate severity of AtD, allergic sensitization and impaired barrier function of the epidermis [4–6].

It is interesting to note that even among children with bronchial asthma the highest indicators of positive skin culture on *S. aureus* were more common in those who lived in the city or families with a low income [13]. The level of air pollution, especially inside the premises is also associated with a patient's low socioeconomic status [14]. In turn, impediments to access to quality health care due to socioeconomic problems can contribute to the development of stress in patients [8].

URBAN/RURAL LIFESTYLE

Hygiene hypothesis has long been debated as a possible explanation for growing allergic diseases: in particular, it has been observed that the youngest child among brothers and sisters has the lowest risk of developing AtD; or this risk decreases in infants, visiting kindergarten in the first year of life [15].

Other evidence, supporting the importance of environmental factors with AtD, include ecological differences between a city and a village.

It is believed that AtD is more common among patients, living in the city, compared to rural and suburban areas [3]. However, according to meta-analysis, conducted in 2010 by Schram M. et al, out of 26 studies, included in the analysis, 11 show a significantly higher prevalence of AtD in patients, living in the city; 14 — no such association is found, and only 1 study identifies a lower incidence of AtD in patients, living in the city [16]. Thereby, this comparative analysis underlines the potential role of various environmental factors in developing AtD.

Although exact factors are not quite clear, one explanation is related to the effect of farm animals on human health. Moreover, a link is shown between certain elements of a rural way of life (especially, the consumption of unpasteurized milk) and a reduction in prevalence of AtD [17, 18]. There was an interesting study, conducted jointly by Finnish and Russian scientists in a population of children, living in a genetically homogenous, but economically different region of Karelia at the Russian-Finnish border [19]. The comparative analysis showed 3–10 times higher prevalence of allergic diseases (bronchial asthma, hay fever, AtD, rhinitis as well as atopic sensitization) in Finland than in Russian Karelia, and, moreover, these patterns persisted for 10 years of observation [19]. Skin microbiome and bacterial and fungal content in the nasal mucus are also contrastingly different with a predominance of the genus *Acinetobacter* in children, living in Russia.

In addition, the disruption of the gut microbiome, especially at an early age, might affect human immunity and atopy pathogenesis (development of tolerance or sensitization). The change of microbiocenosis in urban environments, taking antibiotics, reducing exposure to farm animals and (or) their absence lead to a low effect of endotoxins and increased Th2-cells in epithelial and mucous membranes in patients with allergic diseases [20].

The urban way of living is also associated with progression of AtD and atopic multimorbidity in children with atopy in their parents, filaggrin mutations and allergic polysensitization in the anamnesis [21].

CLIMATE

One of the reasons for rising prevalence of AtD in industrialized countries is considered global climate change: in particular, rise in air temperature and increased greenhouse gas emissions that cause changes in atmospheric UV radiation and air humidity [22]. Furthermore, climate changes (droughts; mass displacement of people from new uninhabitable areas,

etc.) pose the greatest threat to health of people, living in low- and middle-income countries.

Climate factors (temperature and air humidity, precipitation, UV radiation, etc.) are characterized by severe seasonality, and all of them certainly affect the skin. Particularly, both local and global changes in the outside temperature might be associated with the severity of skin manifestations in patients with AtD (although we cannot exclude simultaneous relationship between these factor and others).

Meanwhile, evidence of the association between air temperature and prevalence of AtD contradicts each other, perhaps because of various analytical methods or criteria for determining cold or hot weather. Several studies showed that both high and low air temperature is associated with exacerbation of AtD in observed patients [23–28]. Thus, in the USA, where there is one of the most diverse climates in the world, the frequency of out-patient visits was the highest in adults and children with AtD, living in the east of the country, where there is the lowest temperature, which reached its peak in winter [23]. According to other data, only cold or, on the contrary, hot weather is associated with exacerbation of AtD [24, 25].

It is more logical to expect a decrease in the prevalence of AtD in areas with high air temperature [22, 24]. Usually people, living in warmer climates, spend more time outdoors, and, therefore, exposure to UV rays might have a protective effect on the skin [23]. On the other hand, patients with an established diagnosis of AtD do not tolerate high air temperature as heat can cause sweating. As a rule, with sweat, which may develop the irritation action on the skin and contribute to Th2-type inflammation, itching of the skin becomes worse. Children with AtD reacts most to changes in temperature, who have more severe symptoms in spring, autumn and winter [25].

Despite the fact that sun exposure has a positive effect on the skin (UV rays contribute to raising the level of antimicrobial peptides in the skin, modulate the composition of the microbiota, etc.), some studies have found an association between poorly con-

trolled AtD and higher air temperature, including children [26–28].

Patients with AtD do not also tolerate extreme cold weather that might cause dry and itchy skin. Low air temperature may contribute to production of Th2-cytokines and affect the activity of mast cells in the skin that is strongly correlated with inflammation and impaired epidermal barrier of the skin [29].

Two studies show a very significant impact of climate on the course of AtD. Thus, displacement of children aged 4–13 with severe AtD, living in Norway (subarctic/moderate climate), to the subtropical climate of Gran Canaria for 4 weeks led to a significant improvement in skin symptoms on SCORAD scale and quality of life as well as reduced need for local steroids in 1 and 3 months [30]. A rapid decline in the intensity of itching is also noted in adult patients with AtD, who underwent sanatorium treatment in the high mountain region of Davos, Switzerland [24].

It is important to consider the influence of other effects of climate change on AtD (cold and dry weather conditions, flood, etc.) [31, 32]. Scientists believe that global warming will be accompanied by flood due to melting polar ice, sea level rise and longer rains. Thus, during flood events, occurred in Thailand, there was a significant increase in emergency department visits of children aged 0–12, suffering from AtD [25].

Climate changes and global warming will probably contribute to an increase in the concentration of such aeroallergens as pollen and fungi in the air, and also in

the duration of pollination of plants and allergenicity of pollen itself [30–33]. It is shown that higher air temperature is associated with the extension of flowering season, and higher levels of CO₂ contribute to an increase in biomass of pollen produced and its allergens. In particular, raising level of pollutants in the air causes changes in allergenicity of ambrosia pollen [33]. This, ultimately, exacerbates allergenic load in patients with seasonal rhinoconjunctivitis. The study by Krämer U. et al confirms deterioration of skin symptoms in children with AtD and sensitization to grass pollen in summer [30].

CONCLUSION

The impact of different exposome factors on the development and course of AtD has been found in many studies. However, according to the literature, there are the most studied issues related to specific exposome factors, including their exposure in the prenatal and early period of a child's life (type of feeding, exposure to air pollutants, in particular tobacco smoke, vitamin D level, contact with pets, etc.). Although studies show the effect of nonspecific exposome factors on clinical manifestations of AtD, generally, none of them reliably increase or, on the contrary, decrease the risk of disease in children.

Undoubtedly, other factors have a modulating effect on above-mentioned nonspecific ones. Particularly, viruses, allergens, antioxidants, different pollutants, etc., which are closely associated with environment. These and other questions require further study.

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THE AUTHORS' CONTRIBUTION TO THE WORK

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The effect of vitamin D provision on biomarkers of inflammation in bronchial asthma in children

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Natalia A. Belykh, Inna V. Pisnyur, Aleksandr A. Nikiforov, Larisa V. Nikiforova

Ryazan State Medical University, 390026, Ryazan, Vysokovoltynaya str., 9, Russia

Natalia Anatolyevna Belykh — Dr. Sci., Professor, Head of the Department of Faculty and Polyclinic Pediatrics with the Course of Pediatrics of Ryazan State Medical University, ORCID ID: 0000-0002-5533-0205, e-mail: nbelyh68@mail.ru.

Inna Vladimirovna Pisnyur — Assistant of the Department of Faculty and Polyclinic Pediatrics with the Course of Pediatrics of Ryazan State Medical University, ORCID ID: 0000-0002-9267-439X, e-mail: innaabramova@yandex.ru.

Alexander Alekseevich Nikiforov — Cand. Sci., Associate Professor, Head of the Central Research Laboratory, Ryazan State Medical University, ORCID ID: 0000-0001-9742-4528, e-mail: a.nikiforov@rzgmu.ru.

Larisa Vladimirovna Nikiforova — Senior Researcher at the Central Research Laboratory, Ryazan State Medical University, ORCID ID: 0000-0001-6296-9034, e-mail: a.nikiforov@rzgmu.ru.

Annotation

Introduction. Asthma is a widespread disease in childhood and has a persistent tendency to increase. Therefore, the search for factors influencing this process, as well as biomarkers reflecting the degree of asthma control, is an urgent problem.

Objective. To study the relationship of vitamin D levels with the serum periostin and TGF- β 1 concentration in children with asthma.

Materials and methods. The cross-sectional (one-stage) study included 80 children aged 6 to 17 years (average age — 12 ± 3.2 g). The subjects were divided into 2 groups: children with asthma — group 1 ($n=40$); group 2 — the control group ($n=40$). In all children, the assessment of the concentration of 25(OH)D, periostin and TGF- β 1 in the blood serum was studied.

Results. Median (Me) 25(OH) in patients with asthma was statistically significantly lower than in children of the comparison group (16.7 ng/ml, versus 25.7 ng/ml, $p=0.017$), and did not depend on the severity of the disease, corresponded to a deficiency condition in both mild (16.2 ng/ml) and with an average severity of asthma (16.8 ng/ml) ($p=0.041$). Me of periostin in 1st group was within the normal range (730.2 ng/ml), but statistically significantly exceeded the indicator of 2nd group (539.7 ng/ml, $p<0.05$) and did not depend on the age and duration of asthma. High rates of periostin were observed in children with moderate severity of asthma with a disease experience of 4–6 years (617.2 ng/ml). Me of TGF- β 1 in both groups corresponded to normal values (309.0 and 369.6 pg/ml, respectively, $p>0.05$) and did not depend on the age and duration of asthma.

Conclusions. VD deficiency is registered in children with asthma 2 times more often than in healthy children in Ryazan region. The serum concentration of periostin increased in proportion to the severity of asthma. Vitamin D deficiency can be one of the risk factors for the development of asthma and lead to an imbalance in the periostin and TGF- β system.

Keywords: asthma, vitamin D, periostin, transforming growth factor β 1, children.

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Влияние обеспеченности организма витамином D на биомаркеры воспаления при бронхиальной астме у детей

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For correspondence:

Inna Vladimirovna Pisnyur, Assistant of the Department of Faculty and Polyclinic Pediatrics with the Course of Pediatrics of Ryazan State Medical University.

Address: 390026, Ryazan, Vysokovoltynaya str., 9, Russia.

E-mail: innaabramova@yandex.ru.

Для корреспонденции:

Пизнюр Инна Владимировна, ассистент кафедры факультетской и поликлинической педиатрии с курсом педиатрии ФДПО, ФГБОУ ВО «Рязанский государственный медицинский университет имени академика И. П. Павлова».

Адрес: 390026, г. Рязань, ул. Высоковольная, д. 9, Россия.

E-mail: innaabramova@yandex.ru.

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Белых Н. А., Пизнор И. В., Никифоров А. А., Никифорова Л. В.

Федеральное государственное бюджетное образовательное учреждение высшего образования «Рязанский государственный медицинский университет имени академика И. П. Павлова» Министерства здравоохранения Российской Федерации, 390026, г. Рязань, ул. Высоковольная, д. 9, Россия

Белых Наталья Анатольевна — д. м. н., доцент, заведующая кафедрой факультетской и поликлинической педиатрии с курсом педиатрии ФДПО, ФГБОУ ВО «Рязанский государственный медицинский университет имени академика И. П. Павлова», ORCID ID: 0000-0002-5533-0205, e-mail: nbelyh68@mail.ru.

Пизнор Инна Владимировна — ассистент кафедры факультетской и поликлинической педиатрии с курсом педиатрии ФДПО, ФГБОУ ВО «Рязанский государственный медицинский университет имени академика И. П. Павлова», ORCID ID: 0000-0002-9267-439X, e-mail: innaabramova@yandex.ru.

Никифоров Александр Алексеевич — к. м. н., доцент, заведующий Центральной научно-исследовательской лабораторией ФГБОУ ВО «Рязанский государственный медицинский университет имени академика И. П. Павлова», ORCID ID: 0000-0001-9742-4528, e-mail: a.nikiforov@rzgmu.ru.

Никифорова Лариса Владимировна — старший научный сотрудник Центральной научно-исследовательской лаборатории, ФГБОУ ВО «Рязанский государственный медицинский университет имени академика И. П. Павлова», ORCID ID: 0000-0001-6296-9034, e-mail: a.nikiforov@rzgmu.ru.

Аннотация

Актуальность. Бронхиальная астма (БА) является широко распространенным в детском возрасте заболеванием и имеет стойкую тенденцию к росту. Поэтому поиск факторов, влияющих на этот процесс, а также биомаркеров, отражающих степень контроля БА, является актуальной проблемой.

Цель. Изучить взаимосвязь уровня витамина D с концентрацией периостина и TGF- β 1 в сыворотке крови у детей с БА.

Материалы и методы. В поперечное (одномоментное) исследование были включены 80 детей в возрасте от 6 до 17 лет (средний возраст — $12 \pm 3,2$ г.). Обследуемые были распределены на 2 группы: дети с БА — 1-я группа ($n = 40$); 2-я группа — контрольная группа ($n = 40$). У всех детей проводили оценку концентрации 25(OH)D, периостина и TGF- β 1 в сыворотке крови.

Результаты. Медиана (Me) 25(OH)D витамина D у пациентов с БА была статистически значимо ниже, чем у детей группы сравнения (16,7 нг/мл, против 25,7 нг/мл, $p = 0,017$), и не зависела от тяжести течения заболевания, соответствовала дефицитному состоянию как при легкой степени (16,2 нг/мл), так и при средней степени тяжести БА (16,8 нг/мл) ($p = 0,041$). Ме периостина в 1-й группе была в пределах нормы (730,2 нг/мл), но статистически значимо превышала показатель 2-й группы (539,7 нг/мл, $p < 0,05$) и не зависела от возраста и длительности БА. Высокие показатели периостина имели дети со средней степенью тяжести БА при длительности заболевания 4–6 лет (617,2 нг/мл). Ме TGF- β 1 в обеих группах соответствовала нормальным значениям (309,0 и 369,6 пг/мл соответственно, $p > 0,05$) и не зависела от возраста и длительности БА.

Заключение. В г. Рязани у детей с БА дефицит VD регистрируется в 2 раза чаще, чем у здоровых детей. Показатели периостина в сыворотке крови возрастали пропорционально степени тяжести БА. Дефицит витамина D может выступать одним из факторов риска развития БА и приводить к дисбалансу в системе периостина и TGF- β .

Ключевые слова: бронхиальная астма, витамин D, периостин, трансформирующего фактора роста β 1, дети.

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INTRODUCTION

Bronchial asthma (BA) is one of the most common chronic diseases in children population and characterized by the presence of respiratory symptoms such as wheezing, shortness of breath, chest tightness, cough that are varied in time and intensity [1]. Internal (predominantly genetic) and external factors (allergens, tobacco smoke, industrial dust, atmospheric pollution) are involved in the development of asthma. The prevalence of BA increases from year to year and a rapid growth is particularly noted in children population that prompts a further search for factors, contributing to this trend as well

as biomarkers, demonstrating the level of asthma control [2–4].

Periostin is one of markers of allergic inflammation, which is a protein of the epithelium extracellular matrix with a molecular weight of 90 kDa, belonging to the Fascioline family. Periostin is expressed by osteoblasts, localized in fetal tissues, embryonic periosteum, placenta, heart valves, adrenal tissue, lungs and thyroid [5–6]. In BA the synthesis of periostin is carried out by fibroblasts, epithelial and endothelial cells as well as bronchial smooth muscle cells [7]. Periostin is involved in the acute phase of inflammation in BA, providing synthesis and secretion by eosino-

phils of IL-6 and IL-8 interleukins, which transform β -1 and β -2 growth factors, cysteine leukotrienes and prostaglandin E2. Periostin forms congestion in the basement bronchial membrane in the chronic phase of inflammation, providing its thickening and remodeling. Periostin gene expression is under the control of numerous cytokines and hormones. Periostin expression regulators include bone morphogenetic proteins (type 2 and 4) platelet-derived growth factor, vascular endothelial growth factor, connective tissue growth factor-2, angiotensin II, IL-3, IL-4, IL-6 and IL-13. Literature data of recent years point out that patients with a higher concentration of periostin in the blood serum are characterized by a frequent persistence of bronchial inflammation, also higher probability of connective tissue restructuring of epithelium submucosal layer (remodeling) [8, 9].

In recent years, a special attention is paid to tissue growth factors in the study of processes of remodeling the respiratory tract in asthma. TGF- β 1 is one of cytokines, involved in the process. Produced by epithelial cells of the bronchial mucosa, inflammatory infiltrate cells, TGF- β 1 has a pronounced immunoregulatory, antiproliferative and regenerating effect. This cytokine correlates with increased activity of Th17, aggravating inflammation in the respiratory tract. Moreover, there are data, indicating the inhibitory effect of TGF- β 1 on relaxation of smooth bronchial muscles by induction of shortening smooth muscle fibers and increasing bronchial hyperreactivity [10].

The role of vitamin D (VD) in the development and control of BA remains an active area of research. VD is a pleiotropic hormone, which, along with the regulation of calcium and phosphorus metabolism, has a strong immunomodulatory effect [11]. VD is able to

inhibit the function of T-helpers type 2 (Th2-cells), and also the proliferation and differentiation of B-cells into plasma cells that causes a decrease in the secretion of immunoglobulins E (IgE) [12]. It is known that Th2-response plays a crucial role in all allergic diseases. Therefore, a growing interest in the impact of VD on the pathogenesis of chronic inflammation in BA is justified [13]. In BA VD, by affecting VD receptors (VDR), reduces hypertrophy of bronchial smooth muscles, hyperplasia of goblet cells, subepithelial collagen deposition and fibroblast activity that leads to a lower rate of remodeling process [14, 15].

Thus, study of the relationship between VD concentration and inflammation biomarkers in BA is a relevant objective.

OBJECTIVE OF THE STUDY

Study the relationship of vitamin D level with the serum periostin and TGF- β 1 concentration in children with BA.

MATERIALS AND METHODS

One-stage single-site randomized study involved 80 children aged 6-17 (average age — $12 \pm 3,2$), including girls — 29 (36,0 %), boys — 51 (64,0 %), permanently residing in Ryazan. The children were divided into 2 groups: group 1 (main group) included 40 children with BA, group 2 contained 40 children (control group) (Table 1). The main group is divided into 2 subgroups: 1a — children with moderate asthma ($n = 23$, 57,5 %), 1b — children with a mild disease ($n = 17$, 42,5 %).

The study plan was approved by the local Ethics Committee, FSBEI HE RyazSMU of the Ministry of Health of Russia (Protocol of 09.03.2021). The parents of all the children, taking part in the study,

Table 1. **Characteristics of the study participants**
Таблица 1. **Характеристика участников исследования**

Indicator	Group 1 ($n = 40$)	Group 2 ($n = 40$)	p
Age, years	$12,0 \pm 2,8$	$11,9 \pm 3,3$	0,96
Girls, n (%)	15 (37,5%)	14 (35,0%)	0,12
Boys, n (%)	25 (62,5%)	26 (65,0%)	0,20

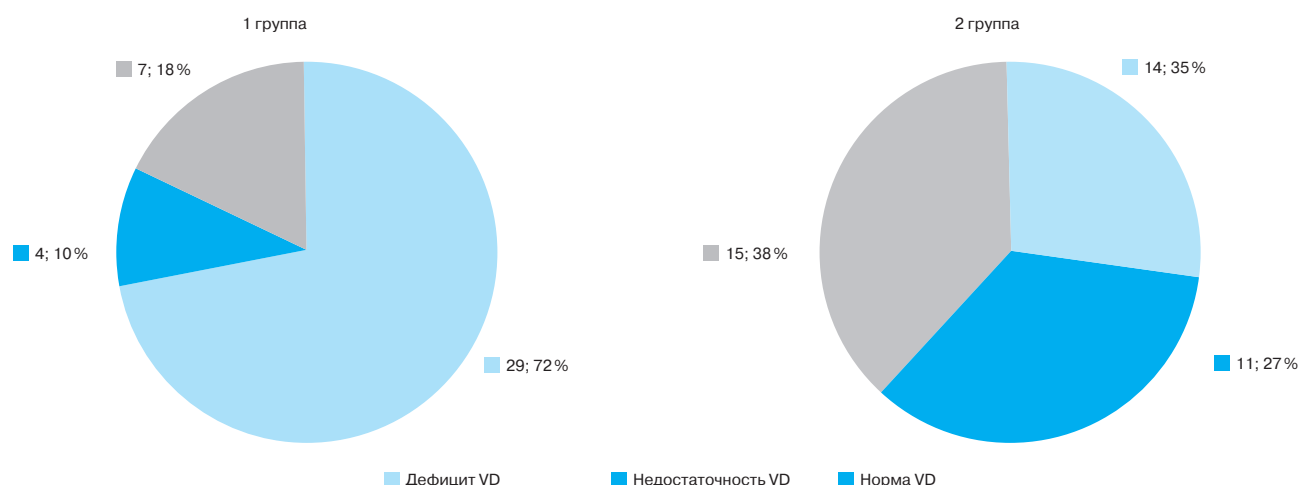


Fig. 1. Provision of the examined children with VD
Рис. 1. Обеспеченность обследованных детей VD

were familiarized with the regulation of the study and signed an informed consent.

SBI «City children's clinic № 3» (chief medical officer – A. O. Burdukova), Central research laboratory of FSBEI HE RyazSMU of the Ministry of Health of Russia (the head of the laboratory – candidate of medical science, associate professor A. A. Nikiforov) were bases to conduct the study.

Criteria for inclusion in the study: the established diagnosis of “bronchial asthma” for at least 1 year, verified according to GINA2022 [16] and Federal clinical guidelines [1]; patient age of 5-17 years; obtaining informed consent from parents and patients for the study.

Exclusion criteria: the presence of malignant neoplasm in the surveyed, an acute disease or exacerbation of other chronic diseases, endocrine or genetic pathology, surgery for the past 4 weeks, intake of anti-spasmodic drugs, disorders of calcium-phosphorus metabolism.

Material sampling was in March-April, 2021. Serum concentration 25(OH) D, periostin, TGF- β 1 were determined by the ELISA method, using “25OH Vitamin D Total ELISA Kit” (DIAsourceImmunoAssaysSA, Belgium),

«ELISA Kit for Periostin», (Cloud-Clone Corp., USA), “ELISA Kit for Transforming Growth Factor Beta 1” (Cloud-Clone Corp., USA) in the Central Research laboratory, FSBEI HE RyazSMU of the

Ministry of Health of Russia with further calculation of the median and interquartile range (Me; 25–75%). The obtained results were evaluated according to the National Program “Vitamin D insufficiency in children and adolescents in the Russian Federation: modern approaches to correction” (2018). Concentration 25(OH)D > 30 ng/mL was considered as the normal level, moderate deficiency – 21–30 ng/mL, severe deficiency – < 20 ng/mL [17]. Periostin value in the serum samples/plasma in 500-fold dilution: 132,4–859,6 ng/mL. TGF- β 1 value in 3-fold dilution in the serum/plasma: 82,4–702,4 pg/mL [18].

Statistical data processing was conducted using MS Excel 2016 and Statistica 6.0 standard software packages. Shapiro-Wilk tests were used to analyze the normality of parameter distribution. Continuous variable was presented as the median (Me) with interquartile range (25–75 percentile). Categorical variables were determined in as a percentage (%). The assessment of differences between groups was conducted using non-parametric Mann-Whitney (U-test) and Pearson's (χ^2) tests with corrections for small samples. Differences were considered significant at $p < 0,05$.

STUDY RESULTS AND THEIR DISCUSSION

The examination revealed that most children had low level of 25(OH)D in the blood serum. At the same time, VD deficiency was recorded twice as often as in the control group ($p = 0,002$) (fig. 1).

The median of 25(OH)D in the group of children with BA was 1,5 times as low as in the control group and corresponded to the deficient condition – 16,7 ng/ml [7,1; 22,8] versus 25,7 ng/ml [17,4; 34,2] in group 2, respectively ($p = 0,017$).

The severity of the disease in children did not depend on the concentration of 25(OH)D in the blood serum in the main study group. Me in children with mild BA was 16,2 ng/ml [13,5; 22,8]), versus 16,8 ng/ml [13,9; 21,6] in the moderate severity ($p = 0,041$). 11,7 % ($n = 2$) of children with mild and 21,7 % ($n = 5$) with moderate asthma had optimal VD vitamin sufficiency. 37,5 % ($n = 15$) of children of the control group had normal VD-status.

Me 25(OH)D did not reach optimal values in any age category of children with BA, however, the rate in group 1 with teenage children was statistically significantly lower than in the control group ($p < 0,05$) (Table 2).

Periostin median in the group of children with BA was statistically much higher than the rate of the control group – 730,0 ng/ml [390,8; 1109,7] versus 536,7 ng/ml [452,0; 666,2] in group 2, respectively ($p = 0,044$) (fig. 2).

In the group of children with BA 30,0 % ($n = 12$) had the increased level of periostin. Periostin Me was 593,0 ng/ml [318,0; 846,3] with mild BA versus 751,0 ng/ml [505,0; 1140,0] with moderate BA ($p = 0,027$). 23,5 % ($n = 4$) of children with mild and 34,7 % ($n = 8$) with moderate asthma experienced the increased level of periostin ($p > 0,05$). 15,0 % ($n = 6$) of children in the control group had the same increased level. This is probably due to the activity of bone metabolism during active growth.

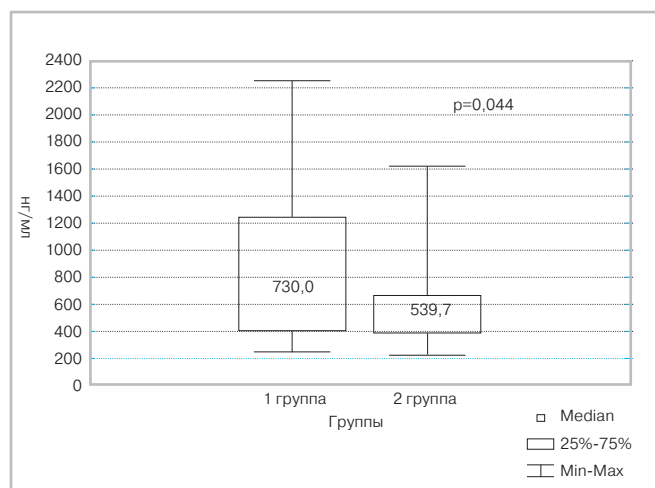


Fig. 2. Median periostin in the blood serum of the examined children (ng/ml)

Рис. 2. Медиана периостина в сыворотке крови у обследованных детей (нг/мл)

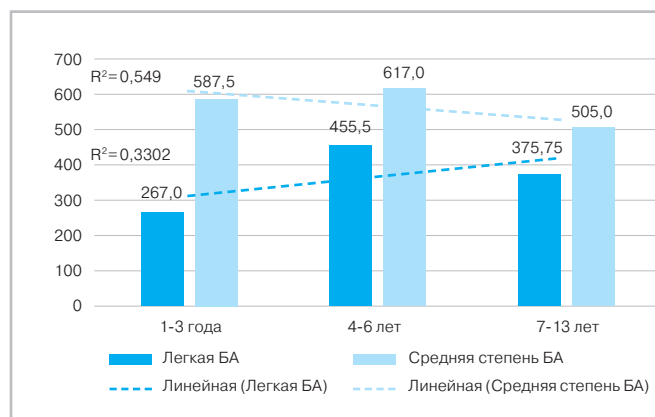


Fig. 3. Median periostin in blood serum in children with asthma depending on the duration of the disease (ng/ml)

Рис. 3. Медиана периостина в сыворотке крови у детей с БА в зависимости от длительности заболевания (нг/мл)

Table 2. Median 25(OH)D in the blood serum of the examined children (ng/ml)

Таблица 2. Медиана 25(OH)D в сыворотке крови у обследованных детей (нг/мл)

Age, years (n)	Group 1 Me [25%; 75 %]	Group 2 Me [25%; 75 %]	p
5–6 years (n=3)	16,8 [16,8; 16,8]	32,7 [25,0; 40,4]	>0,05
7–9 years (n=19)	19,4 [15,7; 30,5]	31,1 [21,1; 40,1]	>0,05
10–14 years (n=36)	16,7 [13,7; 21,9]	25,3 [17,7; 33,5]	>0,05
15–17 years (n=22)	13,5 [11,7; 16,9]	24,4 [16,6; 26,8]	<0,05

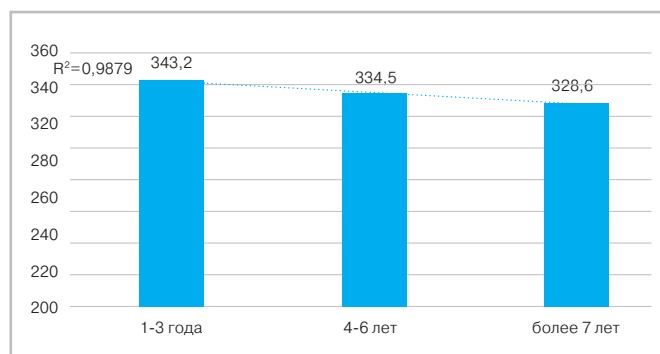


Fig. 4. Median TGF-β1 in blood serum in children depending on the duration of AD (pg/ml) ($p > 0,05$)

Рис. 4. Медиана TGF-β1 в сыворотке крови у детей в зависимости от длительности БА (пг/мл) ($p > 0,05$)

Me of periostin did not depend on the duration of the disease among children with BA, but this rate was higher ($p > 0,05$) with moderate BA (fig. 3).

No clear correlation was found in the analysis of the relationship between the level of periostin and VD (Table 3).

The median of TGF-β1 in both groups of the examined children corresponded to normal values: 309,0 pg/ml [210,9; 408,6] and 355,0 pg/ml [257,4; 426,8], respectively ($p > 0,05$). However, Me TGF-β1 was statistically significantly lower among children with mild BA than in moderate asthma (300,9 [154,5; 342] pg/ml), versus 369,6 [296,1; 455,7] mg/ml, respectively, and lower than in children of the control group (355,0 pg/ml [257,4; 426,8]) ($p < 0,05$).

Me TGF-β1 in the blood serum was within the normal values in children with BA, and the lowest rate was noted with the duration of BA more than

7 years, which may indicate the properly selected anti-inflammatory therapy and high patient compliance ($p > 0,05$) (fig. 4).

Published meta-analysis of recent literature review indicate the presence of low VD sufficiency among pediatric patients with BA. The works by Wang Q et al. (2021) note that the level of 25(OH)D in the blood serum was much lower in children with BA (5 711 participants) than in children without asthma (21 561 people) [19]. The same results were obtained by Russian researchers (S. S. Masalskiy et al., 2018) [20]. The findings coincide with the conclusions of these authors — low VD sufficiency was detected in more than 70% of the examined children with BA.

The relationship between the level of periostin in the blood serum and the presence of BA is also being actively discussed. Inoue T. et al. (2016) noted that the concentration of periostin in the blood serum was higher in children with BA, compared to children with no allergic diseases. The authors also pointed to the need for determining periostin content in the blood serum to diagnose and monitor BA in children [21]. The works by Song J. S. et al. (2015) found that a high level of periostin in the blood serum in children with BA was associated with hyperreactivity of the respiratory tract [22, 23] S. S. Masalskiy and others. (2018) revealed that the level of serum periostin was significantly higher in children with BA, compared to healthy children, and directly correlated with the severity of BA [24]. In our study the level of periostin in the blood serum in children with BA was also statistically much higher than the rate of children in the control group, though, these figures were within the normal values and the concentration of periostin

Table 3. Distribution of children by level 25(OH)D and periostin in blood serum in children with asthma
Таблица 3. Распределение детей по уровню 25(OH)D и периостина в сыворотке крови у детей с БА

Rate	Increase in the concentration of periostin in the blood serum, n (%)	Normal concentration of periostin in the blood serum, n (%)	p
Severe VD deficiency, (n=29)	10 (34,5%)	19 (65,5%)	$> 0,05$
Moderate VD deficiency, (n=4)	1 (25,0%)	3 (75,0%)	$> 0,05$
Norm of VD, (n=7)	1 (14,3%)	6 (85,7%)	$> 0,05$

in children with moderate BA was statistically significantly higher than in a mild degree of the disease, though both the rates were within the normal values.

It is considered that TGF- β 1 plays a central role in the pathogenesis of remodeling the respiratory tract in BA, which can occur at any age, regardless of the severity degree and is caused by complex pathogenetic interactions between different biologically active molecules and external triggers. The work by N.L. Potapova, I.N. Gaymolenko (2019) revealed a significant difference in the content of serum TGF- β 1 in children with BA, compared to the group of healthy children [10]. Our study has not found incidents of the increased content of serum TGF- β 1 in either patients with BA or the controlled group. In this case, patients with mild BA had the level of TGF- β 1 sta-

tistically much lower than in the moderate severity of the disease.

CONCLUSIONS:

1. The incidence of vitamin D deficiency in children with BA from Ryazan was recorded twice as often as in healthy children, and it did not depend on the severity of asthma.
2. Periostin rate in the blood serum increased in proportion to the severity of BA.
3. Vitamin D deficiency may be one of risk factors for developing BA, causing imbalance in the system of periostin and TGF- β .
4. It is advisable to monitor the level of 25(OH) D in the blood serum with subsequent correction in children with BA.

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THE AUTHORS' CONTRIBUTION TO THE WORK

Natalia A. Belykh – development of the concept and design of the study, editing the text of the article.

Inna V. Pisyur – review of publications on the topic of the article, collection of material, statistical data processing, text preparation, writing and editing of the text of the article.

Aleksandr A. Nikiforov, Larisa V. Nikiforova – conducting research, analyzing the data obtained.

ВКЛАД АВТОРОВ В РАБОТУ

Белых Н. А. — разработка концепции и дизайна исследования, редактирование текста статьи.

Пизнур И. В. — обзор публикаций по теме статьи, сбор материала, статистическая обработка данных, подготовка текста, написание и редактирование текста статьи.

Никифоров А. А., Никифорова Л. В. — проведение исследования, анализ полученных данных.

A clinical case of atopic dermatitis with a rapid positive effect from the use of a genetically engineered biological drug in a teenager

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**Ella V. Churyukina^{1,2}, Ekaterina A. Portnyaga³**¹ Rostov State Medical University, Rostov on Don, 344022, Rostov on Don, Nakhichevansky lane, 29, Russia² Kuban State Medical University, 350063, Krasnodar Territory, Krasnodar, Mitrofan Sedin str., 4, Russia³ SM-Clinic, 123022, Moscow, 13 2nd Zvenigorodskaya str., p. 40, Russia

Ella Vitalievna Churyukina — Cand. Sci., Associate professor, Head of Division for Allergic and Autoimmune diseases. Rostov State Medical University. Associate Professor of the Department of Clinical Immunology, Allergology and Laboratory Diagnostics of the Faculty of Advanced Training and Professional Retraining of Specialists of FSBEI HE «Kuban State Medical University» MOH Russia, Krasnodar, Russia, ORCID ID: 0000-0001-6407-6117, e-mail: echuryukina@mail.ru.

Ekaterina Alekseevna Portnyaga — allergist-immunologist, pediatrician SM-Clinic, Moscow, ORCID ID: 0009-0003-5030-917X, e-mail: p_zetta@mail.ru.

Annotation

Introduction. The instructions for the drug dupilumab indicate that the administration of a single loading dose on the first day leads to the rapid achievement of clinically effective concentrations within 2 weeks, which we saw in the example of our patient and her real clinical response to the first injection.

Presentation of a clinical case. Under our medical supervision was a 17-year-old patient with severe atopic dermatitis, resistant to traditional therapy. Heredity for allergic pathology is burdened: the girl's mother suffers from pollen allergy. Initially before the start of therapy: SCORAD — 88 points, EASI — 48.8 points, IGA — 4, blood eosinophils — 11% (1188 cells/ml), total IgE — 1102.0 IU/ml; the content of nitric oxide in exhaled air (FeNO) is 30 ppb. On April 28, 2021, the patient was administered dupilumab at a dose of 600 mg. Assessment of atopic dermatitis control in points upon admission to the hospital 2 weeks after the first administration of dupilumab: on the SCORAD scale — 44.5 points; EASI — 13.8 points; IGA — 2 points, eosinophils — 9% (1070 cells/ml); total IgE — 840 IU/ml; FeNO — 5 ppb.

Conclusion. This clinical observation clearly illustrates the fact that patients with severe atopic dermatitis who don't respond to first-line therapy can achieve positive clinical results after the first use of a recombinant human monoclonal antibody (IgG4).

Keywords: atopic dermatitis, biotherapy, dupilumab.

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For correspondence:

Ekaterina Alekseevna Portnyaga — allergist-immunologist, pediatrician SM-Clinic.

Address: 12 Tikhvinskaya str., sq. 119, 127055, Moscow.

E-mail: p_zetta@mail.ru.

Для корреспонденции:

Портняга Екатерина Алексеевна, врач аллерголог-иммунолог, педиатр МЦ «СМ-Клиника»

Адрес: 127055, г. Москва, ул. Тихвинская 12, кв. 119.

E-mail: p_zetta@mail.ru.

Клинический случай атопического дерматита с быстрым положительным эффектом от применения генно-инженерного биологического препарата у подростка

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Чурюкина Э. В.^{1,2}, Портняга Е. А.³

¹ ФГБОУ ВО «Ростовский государственный медицинский университет» Минздрава России, 344022, г. Ростов-на-Дону, пер. Нахичеванский, д. 29, Россия

² ФГБОУ ВО «Кубанский государственный медицинский университет» Минздрава России, 350063, Краснодарский край, г. Краснодар, ул. Митрофана Седина, д. 4, Россия

³ МЦ «СМ-Клиника», 123022, г. Москва ул. 2-я Звенигородская, д. 13, с. 40, Россия

Чурюкина Элла Витальевна — к. м. н., доцент, начальник отдела аллергических и аутоиммунных заболеваний НИИАП ФГБОУ ВО «Ростовский государственный медицинский университет» Минздрава России, Ростов-на-Дону, Россия; доцент кафедры клинической иммунологии, аллергологии и лабораторной диагностики ФПК и ППС ФГБОУ ВО «Кубанский государственный медицинский университет» Минздрава России, Краснодар, Россия; ORCID ID: 0000-0001-6407-6117, e-mail: echuryukina@mail.ru.

Портняга Екатерина Алексеевна — врач аллерголог-иммунолог, педиатр МЦ «СМ-Клиника», г. Москва, Россия. ORCID ID: 0009-0003-5030-917X, e-mail: p_zetta@mail.ru.

Аннотация

Введение. В инструкции к препарату дупилумаб указано, что введение однократной нагрузочной дозы в первый день приводит к быстрому достижению клинически эффективных концентраций в течение 2 недель, что мы и увидели на примере нашей пациентки и ее реального клинического ответа на введение уже первой инъекции.

Изложение клинического случая. Под нашим медицинским наблюдением находилась пациентка 17 лет с тяжелым течением атопического дерматита, резистентного к традиционной терапии. Наследственность по аллергической патологии отягощена: мать девочки страдает пыльцевой аллергией. Исходно до начала терапии: SCORAD — 88 баллов, EASI — 48,8 балла, IGA — 4, эозинофилы крови — 11% (1188 кл/мл), общий IgE — 1102,0 МЕ/мл; содержание оксида азота в выдыхаемом воздухе (FeNO) — 30 ppb. 28.04.2021 больной введен дупилумаб в дозе 600 мг. Оценка контроля атопического дерматита в баллах при поступлении в стационар через 2 недели после первого введения дупилумаба: по шкале SCORAD — 44,5 балла; EASI — 13,8 балла; IGA — 2 балла, эозинофилы — 9% (1070 клеток/мл); общий IgE — 840 МЕ/мл; FeNO — 5 ppb.

Заключение. Данное клиническое наблюдение наглядно иллюстрирует тот факт, что пациенты с тяжелой степенью атопического дерматита, не отвечающие положительным эффектом на первые линии терапии, могут достичь положительных клинических результатов уже после первого применения рекомбинантного человеческого моноклонального антитела (IgG4).

Ключевые слова: атопический дерматит, биотерапия, дупилумаб.

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INTRODUCTION

Atopic dermatitis is a multifactorial genetically determined inflammatory skin disease, characterized by itching, chronic recurrent course, age characteristics of localization and morphology of lesions [1]. To date, there is a constant growth of atopic diseases in the world, which earliest from of manifestation is exactly atopic dermatitis that develops against the backdrop of genetic predisposition to atopy [2].

Modern understanding of AtD pathogenesis is based on immunological peculiarities of lymphocytic infiltrate with an admixture of dendritic cells, an increased level of inflammatory mediators in the affect-

ed tissues and defects in the skin barrier [3]. It was revealed that the basis of allergic disease is the change in the correlation of subpopulations of T-helper cells in favor of TH2 phenotype [2]. It is T2-inflammation that the basis of a range of diseases, such as bronchial asthma, allergic rhinitis, chronic polypous rhinosinusitis, atopic dermatitis, eosinophilic esophagitis, etc. [4]. IL-4, IL-13 and IL-5 are key cytokines, involved in a cascade of inflammatory responses in T2-inflammation. It is worth noting that a combination of allergic diseases is common in clinical practice: bronchial asthma (BA), allergic rhinitis and AtD [4, 5]. Considering the contribution of the same cytokines to the

development of inflammation in various conditions, it might seem perspective to use one drug for treating various T2-associated diseases in one patient [4, 6].

The appeared information on the pathogenesis of AtD development prompted scientists to search for drugs, inhibiting cytokines, involved in the process of inflammation. Modern immunology is actively studying the role of particular cells and molecules in the pathogenesis of allergic diseases for diagnosing, treating and monitoring the efficiency of therapy [7]. Dupilumab has become the first targeted biological drug to treat AtD. It is a recombinant monoclonal antibody to IL-4 and IL-13 receptor subunit, which suppresses T2-inflammatory response through inhibition of two key cytokines and might be the preferred drug for treating moderate and severe continuously recurrent AtD [8]. Dupilumab is responsible for inhibiting immune disorders, characterized by Th2 phenotype [9, 10]. Clinical trial, conducted for more than 10 years, has confirmed the efficiency and safety of dupilumab for treating AtD [11–13]. A large number of studies demonstrate the efficacy of dupilumab to treat all cases, characterized by Th2-mediated inflammation [4, 14]. In Russia dupilumab is approved for administration in the treatment of patients aged 6 and over for indications “moderate and severe AtD”, under the age 12 years – “bronchial asthma”, under the age of 18 years – “polypous rhinosinusitis” [15]. The drug is available in the prefilled syringe, containing 200 or 300 mg of dupilumab. Dupilumab is administered and can be stored at room temperature (до 25 °C) within 14 days if necessary; injection can be done by either the patient (in the hip or abdomen area) or the caregiver (in the upper arm) [4, 16].

DESCRIPTION OF THE CLINICAL CASE

A girl (17 years), burdened by heredity for allergic pathology (the girl's mother suffers from pollen allergy), has been afflicted with atopic dermatitis from an early age. Skin syndrome debuted from 6 months in the form of hyperemia in the upper extremities. The pediatrician diagnosed “sweating fever”. At the age of 10 months hyperemia occurred in the upper and lower extremities, the allergist diagnosed “atopic der-

matitis, food allergy”. Treatment was prescribed with methylprednisolone aceponate with a positive effect.

Skin syndrome relapsed up to 5 years in spring and autumn with the improvement in summer while being at sea. She was hospitalized due to developed streptoderma at the age of 3.

In 2013 she contacted a dermatovenerologist at the place of residence with a severe exacerbation. Local therapy was prescribed. A temporary positive effect was observed for 1–2 months.

The disease had wave-like nature, topical glucocorticosteroid (GCS) drugs were used with a short-term positive effect in courses for 5 years.

Since January, 2020 there has been a pronounced deterioration after stress from the words of the patient. In the spring 2020 she was hospitalized to the regional hospital at the place of residence with complaints of flushing of the skin, itching, scratching, lichenization in the upper and lower extremities, back and face; she got intravenous administration of dexamethasone (4 mg), 0,9 % solution of sodium chloride, antihistamines, external therapy with GCS with no positive effect.

Due to refractoriness to the therapy, she has repeatedly received inpatient treatment since March, 2020. There was constant intake of non-sedating antihistamines and topical glucocorticosteroids; systemic antihistamines in the hospital.

Allergy testing (21.04.2020) (S-IgE): pollen (meadow grass – 2 class); household (house dust – 3 class); epidermal (cat hair – 3 class, dog hair – 2 class); food (chicken eggs – 3 class).

There are no cats and dogs in the patient's house, no chicken eggs consumed and regular wet cleaning.

In December, 2020, due to persistent rash, the patient went to the hospital at the new place of residence, where she got therapy with no positive effect. She received standard care in FSBI “State Scientific Centre of Dermatovenereology and Cosmetology”, the Ministry of Health of Russia in conditions of 24-hour hospital in December, 2020. In March, 2021 due to the appearance of new rashes, she returned to the FSBI “SSCDC”, the Ministry of Health, where with regard to severity and prevalence of the skin patho-

Table 1. **Comparative data of clinical manifestations**
Таблица 1. **Сравнительные данные клинических проявлений**

Date	SCORAD	EASI	IGA
26.04.21	88 points	48,8 points	4 points
11.05.21	44,5 points	13,8 points	2 points

Table 2. **Comparison of laboratory data**
Таблица 2. **Сравнение данных лабораторных показателей**

Date	IgE total	Eosinophils	FeNO
27.04.21	1102,0 IU/ml	11 % (1188 cells/ml)	30 ppb
11.05.21	840 IU/ml	9 % (1070 cells/ml)	5 ppb

logical process Repeated therapy is recommended under the conditions of 24-hour hospital.

Since 24.03.21 she has got immunosuppressive therapy after another hospitalization (100 mg of Cyclosporin 1 capsule twice a day). Given torpidity and severity of the skin pathological process, it is suggested to consider prescribing the patient genetically engineered biological therapy in the case of failure of immunosuppressive therapy.

One month later, 26.04.2021, the patient was hospitalized in Rostov-on-Don for examining and starting genetically engineered biological therapy with dupilumab, regarding ineffective therapy.

The assessment of atopic dermatitis control in points upon admission to the hospital before dupilumab administration: on the SCORAD scale — 88 points; EASI — 48,8 points; IGA — 4 points.

The patient complained about skin rashes in the scalp area, face, neck, torso, upper and lower extremities, accompanied by severe itching, a feeling of tightness and burning of the skin.

Laboratory indicators: blood eosinophils (CBC) for 27.04.2021 — 11 % (1188 cells/microlitre). Total IgE — 1102,0 IU/ml.

27.04.2021: there was the measurement of exhaled nitric oxide (FeNO) — 30 ppb, with the norm from 0 to 19 ppb. And yet, the patient never reported signs of suffocation or coughing fits.

28.04.2021 dupilumab was administered at a dose of 600 mg (initial dose).

In 2 weeks, the patient was hospitalized to administer the second dose of the drug.

The assessment of atopic dermatitis control in points upon admission to the hospital 2 weeks after the administration of the first dose of dupilumab (Table 1): on the SCORAD scale — 44,5 points; EASI — 13,8 points; IGA — 2 points.

Laboratory indicators in 2 weeks (Table 2): eosinophils — 9 % (1070 cells/ml); total IgE — 840 ME/мл; FeNO — 5 ppb.

The patient gave consent to receive biological therapy and use the obtained results of the examination for scientific purposes.

Currently, the patient continues biological therapy. Atopic dermatitis is under control.

DISCUSSION

Literature data show that dupilumab, used in adolescents, causes significant improvements of symptoms in the 16th week (randomized placebo-controlled clinical trial (LIBERTY AD ADOL), in which 251 adolescents with severe and moderate atopic dermatitis received 300 mg of dupilumab every 4 weeks) [17]. In our clinical case significant improvement of clinical and laboratory parameters were observed within a week after starting biological therapy, using dupilumab. What allows to verify the drug efficacy and need for further research to produce new effective and convenient treatments of atopic dermatitis. Some studies have data on the early positive effect of dupilumab

application not only in atopic dermatitis, but also in severe bronchial asthma; the effect was noted in the first 2 weeks of the treatment [18].

It has previously been shown that dupilumab both decreases the incidence of new allergy and improves pre-existing allergic conditions [19]. It has also been proven that blocking signaling pathway of IL-4/IL-13 reduces the concentration of many of type 2 inflammatory markers, including IgE, periostin and multiple pro-inflammatory cytokines and chemokines (e.g., eotaxin, TARC) as well as decreases the level of fractional exhaled nitric oxide (FeNO) – the marker of inflammation in the lungs [20]. Due to a decrease in FeNO rate in the clinical case, we can assume that in the future FeNO rates should be evaluated in patients with atopic dermatitis to predict therapy efficiency in the prevention of bronchial asthma.

Biological therapy in its breakthrough action is comparable to antibiotic therapy and vaccination, innovative at the time. Undoubtedly, prospects of biological therapy in allergology are huge and not limited to symptom control. Dupilumab is a successful representative of its class of drugs with a highly favourable

profile of efficacy and safety, enabling it to occupy an important niche in therapy of allergic diseases [21].

CONCLUSION

Dupilumab therapy has enabled the patient to achieve a significant improvement during atopic dermatitis:

1. There was clinical improvement on SCORAD, EASI and IGA evaluation index in two or more times;
2. A moderate decline in the level of eosinophils was noted during therapy: initially – 1188 cells/microlitre, in 2 weeks – 1070 cells/ml;
3. There was a decrease in the level of total IgE: initially – 1102,0 IU/ml, in 2 weeks – 840 IU/ml;
4. Taking the drug contributed to a reduction in inflammation activity (reduction of FeNO from 30 to 5 ppb);
5. There was improvement in quality of life.

This clinical case makes us think about the prevention of bronchial asthma within the concept of “atopic march” by acting on key targets in immunopathogenesis of T2-associated diseases.

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THE AUTHORS' CONTRIBUTION TO THE WORK

Ella V. Churyukina – curation, treatment of patients, review of literature, collection and analysis of literary sources, writing and editing the article.

Ekaterina A. Portnyaga – curation of patients, diagnostic monitoring of patients, review of literature, collection and analysis of literary sources, preparation and writing of the article.

All authors made a substantial contribution to the conception of the work, acquisition, analysis, interpretation of data for the work, drafting and revising the work, final approval of the version to be published and agree to be accountable for all aspects of the work.

ВКЛАД АВТОРОВ В РАБОТУ

Чурюкина Э. В. — курация, лечение пациента, обзор литературы, сбор и анализ литературных источников, написание текста и редактирование статьи.

Портняга Е. А. — курация пациента, диагностический мониторинг пациента, обзор литературы, сбор и анализ литературных источников, подготовка и написание текста статьи.

Все авторы подтверждают соответствие своего авторства международным критериям ICMJE, все авторы внесли существенный вклад в разработку концепции, подготовку статьи, прочли и одобрили финальную версию перед публикацией.

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Jacobsen's syndrome: case report

RAR — научная статья

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Liudmila Yu. Barycheva^{1,2}, Leila I. Bachieva¹, Natal'ja A. Koz'mova¹
¹ Stavropol State Medical University, 310, Mira St., Stavropol, 355017, Russia.² Regional Children's Clinical Hospital, 3, Semashko St., Stavropol, 355029, Russia.

Liudmila Yurievna Barycheva — Dr. Sci. (Med.), Professor, Head of the Department of Immunology with a course of continuing professional education, Stavropol State Medical University; Allergist-Immunologist, Regional Children's Clinical Hospital; ORCID ID: 0000-0002-4069-0566, e-mail: for_ludmila@inbox.ru.

Leila Ibragimovna Bachieva — 2nd year resident at the Department of Faculty Pediatrics, Stavropol State Medical University, ORCID ID: 0009-0008-7785-4676, e-mail: bachleila@mail.ru.

Natal'ja Aleksandrovna Koz'mova — assistant of the Department of Immunology with a course of continuing professional education, Stavropol State Medical University; Allergist-Immunologist, Regional Children's Clinical Hospital; ORCID ID: 0000-0003-0971-5347, e-mail: n-koz'mova@mail.ru.

Annotation

Introduction. Jacobsen syndrome (JS) is a rare genetic disease associated with the deletion of chromosome 11q, characterized by multiple malformations, hematological and immune disorders. The development of immunodeficiency in JS is often underestimated, which leads to recurrent infectious complications.

Presentation of a clinical case. The article presents a clinical case of a patient with a deletion of chromosome 11q and combined immunodeficiency. Our patient had recurrent infections, cytopenic syndrome, combined immunodeficiency, as well as other clinical manifestations of Jacobsen syndrome.

In addition to a decrease in serum immunoglobulins, a deep deficiency of the T-cell link of immunity with a low content of T-lymphocytes, recent emigrants from the thymus, has been established.

Conclusions. The peculiarity of the presented clinical case is that with a relatively small amount of deletion 11q, the child realized a complete clinical phenotype of the disease and a deep combined immunodeficiency. The article was written to improve doctors' knowledge about this rare form of congenital immunodeficiency.

Keywords: Jacobsen syndrome, del 11q, combined immunodeficiency.

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Барычева Л. Ю.^{1,2}, Бачиева Л. И.¹, Козьмова Н. А.¹
¹ ФГБОУ ВО «Ставропольский государственный медицинский университет», 355017, г. Ставрополь, ул. Мира, д. 310, Россия² ГБУЗ СК «Краевая детская клиническая больница», 355029, г. Ставрополь, ул. Семашко, д. 3, Россия

Барычева Людмила Юрьевна — д. м. н., профессор, заведующий кафедрой иммунологии с курсом дополнительного профессионального образования, Ставропольский государственный медицинский университет, врач аллерголог-иммунолог, ORCID ID: 0000-0002-4069-0566, e-mail: for_ludmila@inbox.ru.

For correspondence:

Lyudmila Yurievna Barycheva, Dr. Sci. (Med.), Professor, Head of the Department of Immunology with a course of continuing professional education, Stavropol State Medical University; Allergist-Immunologist, Regional Children's Clinical Hospital.

Address: 310, Mira St., Stavropol, 355017, Russia.

E-mail: for_ludmila@inbox.ru.

Для корреспонденции:

Барычева Людмила Юрьевна, д. м. н., профессор, заведующий кафедрой иммунологии с курсом дополнительного профессионального образования, Ставропольский государственный медицинский университет, врач аллерголог-иммунолог.

Адрес: 355017, г. Ставрополь, ул. Мира, д. 310, Россия.

E-mail: for_ludmila@inbox.ru.

Бачиева Лейла Ибрагимовна — ординатор 2-го года кафедры факультетской педиатрии, Ставропольский государственный медицинский университет, ORCID ID: 0009-0008-7785-4676, e-mail: bachleila@mail.ru.

Козьмова Наталья Александровна — ассистент кафедры иммунологии с курсом дополнительного профессионального образования, ORCID ID: 0000-0003-0971-5347, e-mail: n-kozlova@mail.ru.

Аннотация

Введение. Синдром Якобсена (СЯ) — редкое генетическое заболевание, связанное с делецией хромосомы 11q, характеризующееся множественными пороками развития, гематологическими и иммунными расстройствами. Развитие иммунодефицита при СЯ часто является недооцененным, что приводит к рецидивирующим инфекционным осложнениям.

Изложение клинического случая. В статье приведено клиническое наблюдение пациента с делецией хромосомы 11q и комбинированным иммунодефицитом.

У нашего пациента отмечались рецидивирующие инфекции, цитопенический синдром, комбинированный иммунодефицит, а также другие клинические проявления синдрома Якобсена. Кроме снижения сывороточных иммуноглобулинов, установлен глубокий дефицит Т-клеточного звена иммунитета с низким содержанием Т-лимфоцитов — недавних эмигрантов из тимуса.

Заключение. Особенностью представленного клинического случая является то, что при сравнительно небольшом объеме делеции 11q у ребенка реализовался полный клинический фенотип заболевания и глубокий комбинированный иммунодефицит. Статья написана для улучшения знаний врачей об этой редкой форме врожденного иммунодефицита.

Ключевые слова: синдром Якобсена, синдром делеции 11q, комбинированный иммунодефицит.

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INTRODUCTION. Jacobsen syndrome (JBS; MIM 147791), also known as the syndrome of terminal deletion 11q, is a rare genetic disease, caused by the loss of continuous set of genes in the long arm of chromosome 11 [1, 2, 3]. The disease occurs in 1 per 100 000, with a ratio of women to men 2:1 [4, 5]. 85 % of recorded cases arise by mutation de novo [4].

Since the first description of Jacobsen syndrome in 1973 there have been more than 200 recorded cases, characterized by deletions and affecting telomeric regions of chromosome 11. The size of the deletion is 7–20 Mb and the proximal breaking point is inside or closer to the telomeric end of the sub-band 11q23.3 [6, 7, 8]. This terminal haploinsufficiency might affect the function of more than 100 different genes. The diagnosis of complete syndrome is established when BSX, NRG1, ETS-1, FLI-1 and RICS (ARHGAP32) genes are involved in the deletion. Patients with smaller deletions have a partial phenotype [2, 6]. Variability of phenotype-genotype might be associated with incomplete penetrance as well as with other genes of interest, located on 11q, such as TIRAP, FLI-1, NFRKB, THYN1 and SNX19 [9].

The disease covers a wide range of clinical manifestations. Studies show that 97 % of patients have from mild to severe mental retardation. The degree of neurocognitive disorder is closely associated with the size of the deletion [5, 10].

Platelet abnormalities occur in 88,5–94% of cases. There is noted neonatal thrombocytopenia that may resolve with time, and platelet dysfunction of long-lasting nature [11]. There is an increased number of small megakaryocytes (micromegakaryocytes) and their delayed maturation in the bone marrow [12].

Congenital heart defects, most often, defects of the interventricular septum and left-side obstructive lesions, are found in 56 % of patients, and they are the most common cause of death [5, 7]. Hypoplastic left heart syndrome, one of the most severe congenital heart diseases, is described in 5–10 % of patients with Jacobsen syndrome (compared to 0,02 % in the general population) [13]. Studies in humans and mice showed that ETS-1 gene, located in a “cardiac critical area” on the terminal part of chromosome 11, is the cause of congenital heart disease [2, 13].

Craniofacial dysmorphism (> 40 %) is often manifested in the form of trigonocephaly, ocular hypertelorism, strabismus, eyelid ptosis, coloboma of the iris and wide nose bridge. There is skin syndactyly on the hands, abnormal palmar crease and hypoplastic thenar space. Feet are short, flat with syndactyly of the 2nd and 3rd toes [5, 7].

In 2004 P. D. Grossfeld and colleagues conducted the prospective analysis of 110 patients with the syndrome of terminal deletion 11q. There were no clear signs of immunodeficiency, no recorded life-threaten-

List of abbreviations/список сокращений:

BSX:	brain specific homeobox
ETS-1:	proto-oncogene 1, transcription factor
FLI-1:	proto-oncogene, ETS transcription factor
JAM3:	junctional adhesion molecule 3
KREC:	kappa-deleting recombination excision circle
NFRKB:	nuclear factor related to kappaB binding protein
NRGN:	neurogranin
RICS:	Rho GTPase activating protein 32
SNX19:	sorting nexin 19
THYN1:	thymocyte nuclear protein 1
TIRAP:	TIR domain containing adaptor protein
TREC:	T-cell receptor excision circle
ADP:	adenosine diphosphate
IVIG:	intravenous immunoglobulin
CHD:	congenital heart disease
DIVS:	defect of the interventricular septum
DISS:	defect of the interstitial septum
IUGR:	intrauterine growth retardation
DC:	disturbed circulation
AS:	Apgar score
SCIG:	subcutaneous immunoglobulin
JS:	Jacobsen syndrome
FC:	functional class
CKD:	chronic kidney disease

ing and (or) opportunistic infections in the studied cohort. However, recurrent episodes of otitis media and (or) sinusitis were frequent and observed in 42 out of 78 patients (54 %) [5].

The first immune defect, recorded in this syndrome, was antibody deficiency [14]. A number of studies have noted a decrease in all classes of immunoglobulins (IgA, IgM, IgG) and disorder of specific antibody formation in response to vaccination with pneumococcal polysaccharide vaccine, which is typical for patients with common variable immunodeficiency [15]. The mechanism by which the terminal deletion of chromosome 11 contributes to the development of immunodeficiency is not fully understood. It is assumed that immunodeficiency occurs mainly

due to ETS (ETS-1) or FLI-1 gene loss. ETS-1 is highly expressed in NK-cells, B- and T-lymphocytes and involved in the development of NK-cells, differentiation of T- and B-lymphocytes [16, 17, 18].

Only in 2020 the disease was considered as congenital immune system defect and included in the classification of primary immunodeficiencies, namely in the group of combined primary immunodeficiency states with syndromic manifestations [19].

DESCRIPTION OF THE CLINICAL CASE

Here we present a clinical case of a patient aged 7 with diagnosed Jacobsen syndrome. The patient's parents gave consent to the use of information, including the child's photo, in research and publication.

The boy K. Was born from the second pregnancy, second birth, in the 37th week of gestation. Heart disease was recorded parenterally at 18 weeks. Birth weight is 2030 g, length – 46 cm, AS (Apgar score) – 7/8 points. Tetralogy of Fallot is diagnosed in the neonatal period (aortic dextroposition, membranous ventricular septal defect up to 6,9 mm, secondary defect of interatrial septum up to 5,0 mm, interventricular septum hypertrophy). Circulatory failure – 1B. FC 2 (NYHA). There is intrauterine growth retardation of grade III, abnormalities of the facial skeleton. Radical correction of double-outlet right ventricle was carried out at the age of 10 months.

Due to congenital anomaly, the child has been examined by a geneticist, inherited metabolic diseases are excluded, the normal male karyotype is established (46 XY), dysplastic phenotype has been ascertained: facial dysmorphism, trigonocephaly, hypertelorism, antimongoloid slant, ptosis, highly arched palate, anomaly of the external auditory canal on the left, sparrow chest (fig. 1).

Psychomotor retardation, restless behaviour, sleep disruption and autoaggression were observed from the first month of life. The patient was observed by the neurologist with the diagnosis “Symptomatic epilepsy”. Mental retardation. Behavioral disorders. Alalia. At the age of 7 the patient has sign language, says single words, does not fix attention; there is motor awkwardness and a significantly limited scope of actions.

Polycystic left kidney, 1-2 degree of chronic kidney disease, is diagnosed at the age of 5. At an early



Рис. 1. Признаки лицевого дисморфизма: тригоноцефалия, широкая переносица, гипертелоризм глаз, птоз, низко посаженные ушные раковины, тонкая верхняя губа

Fig. 1. Signs of facial dysmorphic disorder: trigonocephaly, wide bridge of the nose, hypertelorism of the eyes, ptosis, low-set auricles, thin upper lip

age the by experienced frequent respiratory infections — ARVI (up to 9 times per year), pleural and purulent otitis media, bronchitis, poor weight gain. Till the age of 3 there were 4 reported episodes of purulent otitis; he was hospitalized twice with bronchopneumonia, got repeated antibiotic therapy (at least 4 per year). From 1 year 4 months complete blood count has revealed thrombocytopenia in the range of $78-88 \times 10^9/l$, leukopenia — $3,3-1,6 \times 10^9/l$. At the age of 3 the boy came to the attention of the allergist-immunologist of the Regional Children's Clinical Hos-

pital in Stavropol, there was a significant reduction in TREC (T-cell receptor excision circle) rates, all populations of T-lymphocytes, hypogammaglobulinemia (Table 1). Given a distinctive phenotype, primary immunodeficiency, DiGeorge syndrome, was suspected. However, parathyroid hormone and ionized calcium were within reference values. Substitution therapy with intravenous immunoglobulins (IVIG) was initiated, followed by subcutaneous immunoglobulin (SCIG); preventive antimicrobial therapy (cotrimoxazole, fluconazole, azithromycin) was prescribed.

To clarify the diagnosis in FSBI Rogachev National Medical Research Center of Pediatric Hematology, Oncology and Immunology, fluorescence in situ hybridization FISH on chromosome 22 was conducted, however, 22q11.2 deletion was not found. The child's blood was sent for NGS-sequencing (next generation sequencing), "immunological panel". There was a two-fold reduction in reads of all FLI-1 gene exons, localized on chromosome 11 that pointed to deletion of one of the two gene copies. When specifying the size of the defect by molecular genetic microarray, there was identified terminal deletion on the region of chromosome 11 11q24.2q25 at a size of 10084933 base pairs with coverage of 44 genes in the imbalance region as well as microduplication of chromosome 16 16p13.11 at a size of 14435773 base pairs (31 genes) and chromosome 22 22q13.31q13.33 with a length of 6730554 base pairs (52 genes) that indicated an unbalanced translocation between 11q and 22q with high probability. The child got the confirmed rare syndromal immunodeficiency: Jacobsen syndrome. Paris-Trousseau thrombocytopenia. The number of genes in the imbalanced region included the genes, responsible for the development of immune responses (TIRAP, FLI-1, JAM3).

In order to estimate our patient's immune dysfunction, we have analyzed levels of immunophenotyping and serum immunoglobulins during the 4-year follow-up (Table 1). There is a critical decrease in TREC and KREC ("BT test" ("Generium", Russia)), persistent T-cell immunodeficiency with a decrease in the level of total T-lymphocytes, T-helper cells, T-cytotoxic lymphocytes. A reduction in naive T-helper

Table 1. Investigation of subpopulations T-, B-lymphocytes, immunoglobulins and platelets in a patient with Jacobsen syndrome
Таблица 1. Исследование субпопуляций Т- и В-лимфоцитов, иммуноглобулинов и тромбоцитов у пациента с синдромом Якобсена

Indicators	3 years	5 years	6 years	7 years	Norm
Leukocytes*10 ⁹ /l	3,8	2,23		3,5	6,1–9,9
Lymphocytes/μl	1216	820	590	1,2	1,5–7,0
T-cell CD3 ⁺ /μl	510	500	650	420	900–4500
Helper T-cell CD3 ⁺ CD4 ⁺ /μl	310	300	460	290	500–2400
Naive helper T-cell CD3 ⁺ CD4 ⁺ CD45 ⁺ CD197 ⁻ /μl		46,5	53		200–2500
Memory helper T-cell TEMRA CD3 ⁺ CD4 ⁺ CD45 ⁺ CD197 ⁻ /μl		15,3	66,6		0,025–25
Cytotoxic T-cell CD3 ⁺ CD8 ⁺ /μl	150	140	160	110	300–1600
Naive cytotoxic T-cell CD3 ⁺ CD8 ⁺ CD45 ⁺ CD197 ⁺ /μl		34,1	40,3		42–1300
Memory cytotoxic T-cell TEMRA CD3 ⁺ CD8 ⁺ CD45 ⁺ CD197 ⁻ /μl		58,7	37,9		57–340
B-cell CD19 ⁺ /μl	280	100	120	120	200–2100
Naive B-cell IgD ⁺ IgM ⁺ CD27 ⁻ /μl		75	91		147–431
Switched memory B-cell IgD ⁻ IgM ⁻ CD27 ⁺ /μl		5	5,2		31–94
NK-cell CD16 ⁺ CD56 ⁺ /μl	360	250		160	100–1000
TREC/μl	1,9	2,9	0		30–327
KREC/μl	48,1	36	20,9		75–541
IgA (g/l)	0,31	0,27	0,28	0,41	0,9–1,9
IgM (g/l)	0,34	0,17	0,12	0,2	0,8–1,9
IgG (g/l)	2,9	5,52	8,79	9,0	8,7–11,7
Platelets*10 ⁹ /l	60	80	78	95	204–356

cells and T-cytotoxic T-lymphocytes as well as naive B-lymphocytes and switched memory B-lymphocytes (Switched memory B-cell) was shown. The reduction of these indicators was combined with hypoinmunoglobulinemia.

Levels of B-lymphocytes, serum IgA and IgM remained low throughout the follow-up period. IgG levels were normalized due to regular replacement IVIG/SCIG therapy.

Given the presence of FLI-1 gene deletion, causing the development of thrombocytopenia with dense platelet granule defect, there was functional platelet study — a decrease in platelet aggregation with ristocetin up to 29 % and adrenaline to 38% was noted, aggregation with ADP (adenosine diphosphate) was within normal limits — 36 %.

Serious systemic infectious diseases were not revealed against the background of preventive therapy.

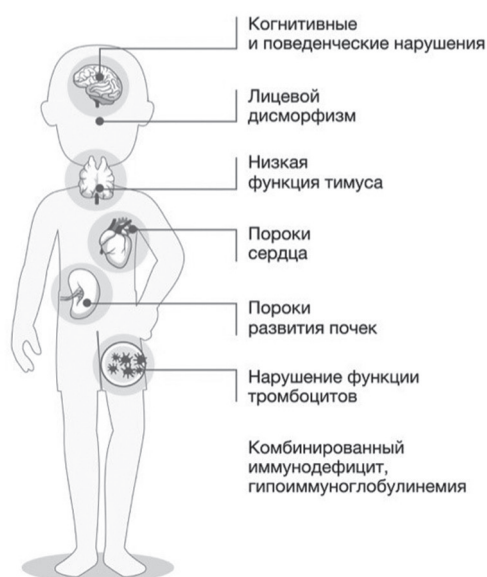


Fig. 2. Clinical features of Jacobsen syndrome
Рис. 2. Клинические особенности синдрома Якобсена

DISCUSSION

Jacobsen syndrome is a rare form of a genetic disease that has recently been classified as primary combined immunodeficiency [15, 19].

The presented case corresponded to the complete clinical phenotype of the disease, despite the genetic traits of partial Jacobsen syndrome. Our patient experienced typical dysmorphic features such as low growth, microcephaly, facial skeleton anomaly, congenital heart disease, chronic kidney disease, mental retardation, cytopenic syndrome (fig. 2).

In recent years, researchers have focused on genotype-phenotype correlations as well as candidate gene reading, responsible not only for cognitive impairment and multiple malformations in patients with Jacobsen syndrome, but also for immune defects [20, 33]. The deletion level in Jacobsen syndrome may vary (fig. 2). Its level in our clinical case is much lower, in comparison with the deletions, described in other sources (fig. 3).

Three genes in the deleted region 11q24.2q25 were associated with the defect in immune regulation (FLI-1, TIRAP, JAM3).

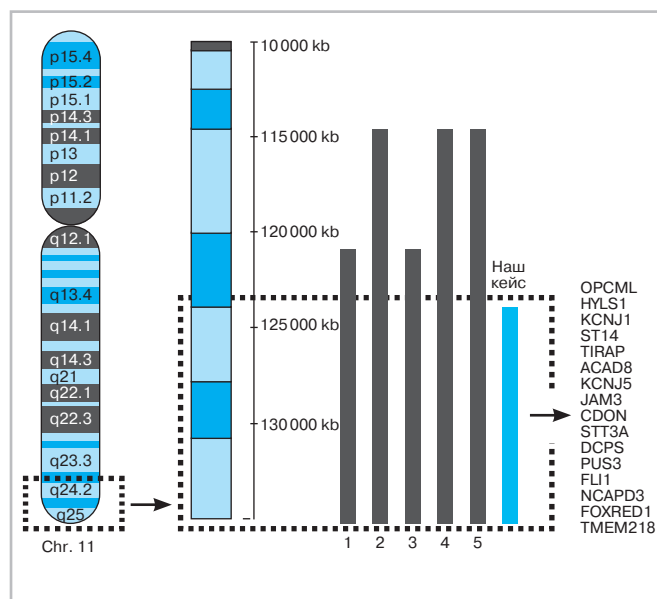


Fig. 3. Deletion 11q24.2q25 in a patient with partial Jacobsen syndrome. On the left is chromosome 11 with a critical region of deleted genes in comparison with deletions described in available sources [5, 20, 21, 22, 24]. On the right are the deleted genes

Рис. 3. Делеция 11q24.2q25 у пациента с частичным синдромом Якобсена. Слева — хромосома 11 с критической областью удаленных генов в сравнении с делециями, описанными в доступных источниках [5, 20, 21, 22, 24]. Справа — удаленные у пациента гены

It is known that FLI-1 encodes a transcription factor, specific to erythroblast transformation. However, FLI-1 heterozygous deletion might cause dysmegakaryopoiesis, functional disorders of T-lymphocytes, deficiency of T-helper cells and a low level of serum IgM [23, 24, 25] that is confirmed in experimental models [26]. It was found that FLI-1 gene modulates a marginal zone follicular B-cell development in mice [27]. Previously it was shown that patients with Jacobsen syndrome are quite often the holders of FLI-1 gene deletion [2, 28, 29, 30]. FLI-1 gene haploinsufficiency has been suggested as a genetic change, responsible for immune system defects in Jacobsen syndrome.

The number of gene in the imbalance area also included TIRAP genes (607948, 611162, 614382),

encoding signaling protein of Toll-like receptors — TLR2 и TLR4. For the past two decades, their key role in reactions of the innate immune response on bacterial lipopolysaccharides has been determined and their relevance in antitumor immunity has been confirmed [31].

JAM3 gene is part of the family of connective-tissue adhesion molecules with high expression on the cell surface of T-cytotoxic lymphocytes and activated cells. Besides, there is a large number of JAM3 molecule, presented on megakaryocytes and platelets that implies its part in the inflammatory process, mediated by monocytes [2].

Immunological data on patients with Jacobsen syndrome in the world is significantly limited, however, there are accumulated reports on serious disorders in the process of maturation and differentiation of all compartments of T- и B-lymphocytes in the syndrome of terminal deletion 11q [32].

Our patient has experienced persistent impairment of the T-cell component of immunity and antibody formation for 4 years of observation.

Previous studies have showed that T-lymphocyte reduction and [28, 32] their functional defects [15, 20] are observed in the majority of patients with 11q disorder that is often accompanied by hypersensitivity to persistent herpes viruses CMV, HSV1, VZV, human papilloma virus [34].

In the work of Baronio M. and colleagues (2022) 66,7 % of patients with JS had a defined reduction in CD3⁺-cells, 58,3 % — T-helpers. Naive T-helpers were low in 45,4% of patients, TREC rates — in 88,9 % [32].

Data on a decrease in the total number of B-lymphocytes as well as immunoglobulins in patients with JS were first published back in 1998 [33]. Several studies have shown hypogammaglobulinemia with reduced IgG, IgA, IgM and disorder of specific antibody formation in response to pneumococcal polysaccharide vaccine that is compatible with the phenotype of common variable immunodeficiency [15]. The literature describes clinical cases of adult patients with humoral immunodeficiency manifestations, which exacerbated over time [23, 34]. For example, a girl with JS has suffered from recurrent

sinopulmonary infections since the age of 18, she experienced a decreased IgG level, a low number of switched memory B-cells as well as disorder of specific antibody formation [34]. A low number of B-lymphocytes with IgD⁺IgM⁺CD27⁺ phenotype is also noted in other studies [15] which has coincided with the obtained data.

Depending on eliminated genes, various immunological phenotypes can be observed, however, their correlation with deletions of particular genes, situated in 11q region, are not well understood. Trachsel T. And colleagues (2022 г.) described a patient with JS and severe primary immunodeficiency, who had decreased antibody titers against *Haemophilus influenza*, content of B-lymphocytes and switched memory B-cells. Another patient with heterozygous deletion of TIRAP, FLI-1, NFRKB, THYN1 and SNX19 suffered a severe bacterial infection, had predominantly a low number of switched memory B-cells [30]. There is an opinion that clinical manifestations of immunodeficiency in patients with JS can have varying degrees of severity, however, they increase with age in the absence of treatment [23, 28, 34, 35].

CONCLUSION

Patients with partial 11q deletion have a high risk of inborn immunity errors due to loss the genes, responsible for immune responses, which function is only being studied. The presented clinical case in conjunction with literature data demonstrates the importance of immunological observation of patients with JS. We have shown that even quite small 11q deletion might cause the formation of severe combined immunodeficiency, that's why patients with JS require a regular immunological screening by lymphocyte immunophenotyping as well as defining serum immunoglobulins. It should be noted that immunological disorders might develop over time and require to be re-tested. In case of quantitative and qualitative defect of T- and B-lymphocytes as well as serious infectious complications, it is necessary to consider preventive administration of antibacterial agents and replacement therapy with immunoglobulin preparations.

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Liudmila Yu. Barycheva — concept of the article, text development, literature review, editing.

Leila I. Bachieva — literature review, text development, translation into English.

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Барычева Л. Ю. — концепция публикации, написание текста, обзор литературы, редактирование.

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FOLLOWING THE X ALL-RUSSIAN CONGRESS OF THE ASSOCIATION OF PEDIATRIC ALLERGOLOGISTS AND IMMUNOLOGISTS OF RUSSIA (APAIR)

По следам X Всероссийского Конгресса Ассоциации Детских Аллергологов и Иммунологов России (АДАИР)

Diagnostic significance of the determination of antinuclear antibodies in children with autoimmune hepatitis

SCO — краткое сообщение

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A. A. Zhuzhula¹, O. V. Kurbatova¹, S. V. Petrichuk¹, D. V. Parakhina¹, M. A. Snovskaya¹, G. B. Movsisyan¹, E. L. Semikina^{1,2}, A. S. Potapov^{1,2}, A. P. Fisenko¹

¹ Federal State Autonomous Institution "National Medical Research Center for Children's Health" of the Ministry of Health of the Russian Federation, 119991, Moscow, Lomonosovsky prospect, 2, building 1, Russia

² The State Education Institution of Higher Professional Training The First Sechenov Moscow State Medical University under Ministry of Health of the Russian Federation, 119991, Moscow, st. Trubetskaya, d. 8, building 2, Russia

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Диагностическая значимость определения антинуклеарных антител у детей с аутоиммунным гепатитом

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Жужула А. А.¹, Курбатова О. В.¹, Петричук С. В.¹, Парахина Д. В.¹, Сновская М. А.¹, Мовсисян Г. Б.¹, Семикина Е. Л.^{1,2}, Потапов А. С.^{1,2}, Фисенко А. П.¹

¹ Федеральное государственное автономное учреждение «Национальный медицинский исследовательский центр здоровья детей» Министерства здравоохранения Российской Федерации, 119991, г. Москва, Ломоносовский пр., д. 2, стр. 1, Россия

² Федеральное государственное автономное образовательное учреждение высшего образования «Первый Московский государственный медицинский университет имени И.М. Сеченова» Министерства здравоохранения Российской Федерации (Сеченовский Университет), 119991, г. Москва, ул. Трубецкая, д. 8, стр. 2, Россия

Ключевые слова: дети, аутоиммунный гепатит, фиброз печени, антинуклеарные антитела, HEp-2, аутоиммунитет, иммунофлуоресценция.

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INTRODUCTION

Autoimmune hepatitis (AIH) is an autoimmune inflammatory liver disease, causing the formation of autoaggressive antibodies to own hepatocytes [1]. AIH is considered a rare disease, its prevalence is 3–17 cases per 100000 population in Europe and the USA [2, 3, 4]. In Russia the percentage of AIH is 1,5–7 % [2, 4, 5] in the structure of chronic hepatitis in children. The disease often occurs in children aged 6–10, and it predominantly affects females [4, 5].

AIH etiology is not fully understood [4]. The violation of immunoreactivity in genetically predisposed individuals is considered a major cause of AIH development [2]. AIH triggers might be Epstein-Barr, hepatitis (A, B, C, D, G) and herpes simplex viruses, bacteria, drugs (interferon drugs, non-steroid anti-inflammatory drugs, nitrofurans, etc.), environmental factors [2, 4]. In the presence of a genetic predisposition there is impaired immune regulation, manifested by the defect in the function of T-cells [4]. The syn-

thesis of B-cells of the IgG class antibodies intensifies with AIH, which might cause the membrane destruction of normal hepatocytes [4].

One of the criteria for AIH diagnosis is the presence of antinuclear antibodies (ANA) in the blood serum, that are detected in 65–78 % of patients with AIH [5, 6]. ANA is not detected in about 10 % of patients with AIH, but their absence does not exclude the diagnosis [6]. The “gold standard” and main screening method of laboratory diagnosis for detecting ANA in the blood serum is the reaction of indirect immunofluorescence (RNIF) on the cell line HEp-2 [3, 7]. The test result is information on the final titer of ANA and the type of nucleus fluorescence and cytoplasm of HEp-2 cells [7].

Detecting ANA in adults with AIH is accompanied by homogeneous (34–58 %), granule (21–34 %) and cytoplasmic (27 %) types of fluorescence or a combination of several types [6]. It is shown that the highest ANA titers in adults with AIH are noted in concomitant syndrome with primary biliary cirrhosis [6]. ANA studies in AIH against the backdrop of pathogenetic treatment in childhood have not been conducted.

Thus, the objective of our study was to define the diagnostic significance of a titer and types of ANA fluorescence in children with AIH during treatment.

MATERIALS AND METHODS

The study involved 77 children with AIH (42 girls and 35 boys), who were screened and treated in FSAI “NMRC for Children’s Health” of the Ministry of Health of the Russian Federation. 65 of them were diagnosed with AIH type 1, 8 children – AIH type 2, 4 – seronegative AIH. Patients with AIH type 1 had comorbid diseases: AIH + primary sclerosing cholangitis (PSC) in 20 % of cases, AIH + autoimmune cholangitis (AIC) – in 12 %, AIH + multiple autoimmune pathologies – in 15 %. In the treatment dynamics 16 children were screened in 6 months – 1 year of standard therapy. The patient age changed from 1,91 to 17,97 years. All the children under treatment were examined on the standard protocol, including complete blood count (hematological automatic analyzer Sysmex XN 550, Japan), biochemical blood test (AU680, USA), ANA on the cell line HEp-2 by RNIF (Immco Diagnostics, Inc., USA).

RNIF method is based on incubation of patients’ blood serum in serial dilutions with epithelial cells of adenocarcinoma in the person’s larynx (HEp-2) in

wells of a glass slide. After removing unbound components with the wash buffer, FITC conjugate was added to wells of a glass slide, which coloured bound antibodies. The result of the reaction was evaluated using fluorescence microscope Nikon Eclipse Ni (Japan) at a magnification of $\times 40$. A maximum titer of detecting ANA and type of cell fluorescence were analyzed. ANA titers $< 1/160$ were considered as normal values, with the titer $1/160$ the response was low-positive, $1/320$ – $1/640$ – moderate-positive, $1/1280$ and higher – high-positive. We considered any detected type of ANA fluorescence as a positive result. To diagnose the liver fibrosis stage, the method of liver transient elastography was used on FibroScan F502 device (EchoSence, France). METAVIR scale was used to evaluate the severity of the degree of liver fibrosis: F0 stage – density in the range 1,5–5,8 kPa; F1 – 5,9–7,2 kPa; F2 – 7,3–9,5 kPa; F3 – 9,6–12,5 kPa; F4 stage (liver cirrhosis) – density 12,6 kPa and more [8, 9]. The distribution by the stages of liver fibrosis was: F0 – 11 patients (14 %), F1 – 14 (18 %), F2 – 13 (17 %), F3 – 9 (12 %), F4 – 30 детей (39 %).

Statistical processing of the results was carried out using Statistica 10.0 program (StatSoft, USA), Excel (Microsoft, USA), IBM SPSS Statistics 25 (USA). Descriptive statistics of quantitative traits is presented in the format: median [lower and upper quartiles] – Me [$Q_{0,25}$ – $Q_{0,75}$].

RESULTS

Conducted analysis of ANA study results in children with AIH revealed that 8 children (10 %) had the titer $1/160$, 14 (18 %) – $1/320$, 9 (12 %) – $1/640$, 19 (25 %) – $1/1280$, 15 (20 %) – $> 1/2560$, and 12 patients out of 77 (15 %) did not have antibodies.

Analysis revealed that children with AIH often had the following types of ANA fluorescence: cytoplasmic (74 %), granular (63 %) and homogeneous (57 %). Combined, dots in the nucleus and nucleolar type of fluorescence were 32 %, 9 % and 8 % of incidence, respectively. There were more often cytoplasmic (63 %), homogeneous (57 %) and granular (54 %) types of fluorescence in the high-positive ANA titer ($> 1/1280$) than in the moderate-positive one ($1/320$ – $1/640$). Nucleolar type of fluorescence (5 children – 8 %) was detected only in the moderate-positive titer. Dots in the nucleus were less common, but in any type of fluorescence (6 children – 9 %). High titers of ANA fluorescence in children were identified in 69 % with an isolated variant of

AIH type 1, while with a combination of AIH and PSC, AIC, multiple autoimmune pathologies high titers were detected in 34 % of children.

Analysis of titers and types of fluorescence with the severity by the degree of liver fibrosis has revealed phase dependence: from fibrosis stage F0 to F2 there is a decrease in the proportion of children with the high-positive ANA titer ($p < 0,001$), and from fibrosis stage F2 to F4 the proportion of patients with the high-positive ANA titer ($p < 0,05$) increases. We have not revealed the dependence of the fluorescence type on the liver fibrosis stage in this sample of patients.

Analysis of ANA content in 16 children during treatment showed that the titer of ANA fluorescence decreased in 10 patients: in 5 of them – became negative ($< 1/160$), in 4 – the fluorescence titer decreased to the minimum, in 1 – was negative at the onset of treatment and remained the same; in 6 children ANA titer did not change or became higher.

A decrease in the antibody titer in children (10 people – 63 %) was reliably ($p < 0,05$) associated with the reduction in measures of disease activity: ALT – from 43 [20; 156,7] to 19 [12; 50,3]; AST – from 35,7 [27,1; 75,8] to 25,7 [20; 33] and in the relative (from 0,45 [0,2; 0,6] to 0,1 [0,09; 0,11]) and absolute number of immature granulocytes (from 0,03 [0,01; 0,03] to 0,005 [0; 0,01]). However, there was tendency to decreasing CRP, GGT, ALP, WBC, neutrophils, ESR as well as a trend towards increasing albumin, MCHC, RDW-CV.

6 (37 %) children did not experience considerable dynamics in the titer of ANA fluorescence on the background of therapy; there was also a significant reduction during treatment: ALT – from 129,5 [92,3; 333,4] to 30,5 [16,5; 48]; AST – from 148 [74,4; 192,9] to 41,5 [24,4; 55,0], reduced absolute number of immature granulocytes, increased albumin – from 38,1 [36,8; 40,3] to 41 [40,8; 46]. There was tendency to

decreased GGT, ALP, WBC, neutrophils, % of immature granulocytes, ESR, increased MCHC, RDW-CV. However, biochemical parameters in this group were significantly higher than in the children with decreased ANA titer.

All the children, who did not experience a decrease in the titer during treatment, had determined liver cirrhosis (F4) on METAVIR scale, the course of the disease was accompanied by the presence of other gastrointestinal diseases: erosive bulbit, gastroduodenitis, erosive gastritis, reflux esophagitis, ulcerative colitis. With remaining ANA titer, the type of fluorescence remained too, new types did not appear.

DISCUSSION

Thus, analysis of the study results shows that ANA have been identified in 85 % children with AIH that exceeds ANA detection in adults (78 %) with AIH [10]. Unlike adult with identified higher ANA titers in case of combined AIH and PBC, no such correlation was found in children with crossed syndrome [6].

It was found that children with AIH often had the following types of ANA fluorescence: cytoplasmic, granular and homogeneous. The comparison of the data obtained on prevailing types of fluorescence in children with AIH is consistent with the data on detected pathognomonic types of ANA fluorescence with AIH in adults obtained by K. L. Raykhelson [6]. The most frequently identified types of fluorescence with AIH are associated, according to the website <https://www.anapatterns.org/>, with the appearance of antibodies to the cytoplasm, to nucleoprotein and chromatin [11].

It is interesting to note that the proportion of children with the high-positive ANA titer ($p < 0,001$) decreases from fibrosis stage F0 to F2 and increases from fibrosis stage F2 to F4 ($p < 0,05$). The onset of the disease is apparently associated with a more inten-

sive formation of ANA that is confirmed by our data on detecting higher ANA titers in children with AIH in the absence of liver fibrosis (F0). In liver cirrhosis (F4) the proportion of children with high-positive ANA titers is also higher than in F1 – F3. This may be caused by the presence of a genetic predisposition to the disease and requires further study [12].

With a dynamic follow-up of patients during therapy, most children experienced a decrease in ANA titer one year after starting treatment that was associated with clinical efficacy of therapy. This fact is

consistent with data, obtained by L.P. Ananyeva, on a decrease in ANA titers against the background of pathogenetic treatment in adults with autoimmune diseases [13].

CONCLUSION

Determination of ANA in children with AIH is of great clinical and diagnostic importance. ANA titer correlates with the severity of the disease. Determination of ANA titer can be used as additional criteria to evaluate efficacy of AIH therapy in children.

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Serotype diversity of *Streptococcus pneumoniae* in children with chronic bronchopulmonary pathology in the pre-vaccination and post-vaccination periods

SCO — краткое сообщение

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Серотиповой состав *Streptococcus pneumoniae* у детей с хронической бронхолегочной патологией в довакционный и поствакционный периоды

<https://doi.org/10.53529/2500-1175-2024-1-41-43>**Комягина Т. М., Тряпочкина А. С., Алябьева Н. М., Лазарева А. В., Фисенко А. П.***Федеральное государственное автономное учреждение «Национальный медицинский исследовательский центр здоровья детей» Министерства здравоохранения Российской Федерации, 119991, г. Москва, Ломоносовский проспект, д. 2, с. 1, Россия***Ключевые слова:** хроническая бронхолегочная патология, дети, *Streptococcus pneumoniae*, серотип, ПКВ13.**Для цитирования:** Комягина ТМ, Тряпочкина АС, Алябьева НМ, Лазарева АВ, Фисенко АП. Серотиповой состав *Streptococcus pneumoniae* у детей с хронической бронхолегочной патологией в довакционный и поствакционный периоды. *Аллергология и иммунология в педиатрии*. 2024; 1: 41-43. <https://doi.org/10.53529/2500-1175-2024-1-41-43>

INTRODUCTION. *Streptococcus pneumoniae* plays an important role in the development of respiratory bacterial infections among children in their early years, people with chronic diseases and elderly people. *Streptococcus pneumoniae* (*S. pneumoniae*, *pneumococcus*) is the main etiological factor of severe invasive infections (bacteremia, meningitis) and the most common causative agent of acute otitis media, sinusitis as well as community-acquired pneumonia [1]. Nasopharynx epithelium is considered the initial colonization place of *S. pneumoniae* in young children, where it may be detected as a part of the commensal flora [2]. Despite the fact that nasopharyngeal colonization by pneumococcus does not often precede the development of the infectious process, carriage creates source of infection and may be the initial stage of the disease [3].

The main method of fighting pneumococcal infection is vaccinal prevention, which started in the Russian Federation in 2014 using 13-valent pneumococcal conjugate vaccine (PCV13).

It includes serotypes, important for pediatric practice (1, 3, 4, 5, 6A, 6B, 7F, 9V, 14, 18, 19A, 19F, 23F); it is characterized by high immunogenicity and also provides long-lasting immunity and immunological memory [4]. Vaccination causes elimination of vaccine serotypes and an increase in prevalence of non-vaccine, previously rare serotypes among pathogens of invasive infections as well as with carriage of *S. pneumoniae* [5,6].

The problem of pneumococcal infection in children with cystic fibrosis (CF) and congenital bronchial and pulmonary malfunctions (CM) has not been studied enough as *S. pneumoniae* is secreted together with other bacterial pathogens in more than 80 % of cases. However, bacterial infections are generally the leading cause of death just in patients with cystic fibrosis, and bacterial agents might be associated with morbidity and mortality in children with CM [7]. So, understanding the serotype structure of pneumococci and its changes as well as the vaccination effect on serotype strain composition of *S. pneumoniae* with these pathologies is essential to determine treatment tactics of these children.

OBJECTIVE of our study: to determine the serotype composition of *Streptococcus pneumoniae* isolates in children with chronic bronchopulmonary pathology prior to mass vaccination against pneumococcal infection and in post-vaccination periods.

MATERIALS AND METHODS.

Over the period of 2011-2021 there was the study of 140 isolates of *S. pneumoniae*, obtained from respiratory specimens (aspirates, bronchial lavage water, sputum) of 86 children (61,4 %) with congenital bronchial and pulmonary malfunctions and 54 children (38,6 %) with cystic fibrosis, observed in FSAI "NMRC for Children's Health" of the Ministry of Health of the Russian Federation. The age of children varied from 0,3 to 17,8 years (the median of 6,5 years). The informed consent of the parents and legal representative was obtained. Demographic and clinical characteristics were examined according to the medical records of patients. Despite the lack of data on the vaccination status of the children screened, official sources for 2015–2018 reports on PCV13 vaccine coverage from 20,9 to 55,3 % of children, born in Moscow [8, 9].

All the isolates were identified using traditional bacteriological methods (colony morphology, α -hemolysis on blood agar, optochin test (Bio-Rad, France) and latex agglutination technique (express test Remel Europe Ltd, UK)).

Pure culture of pneumococci was serotyped in latex agglutination or quelling reactions by Neufeld using specific pool, group and factor serums (Statens Serum Institut, Danmark), and also through molecular typing by multiplex polymerase chain reaction method [10]. Pneumococci were considered untypable if they gave no reaction with any pool serum. *S. Pneumoniae* serotypes, which polysaccharides were included in PCV13, were regarded as "vaccine", all other serotypes were seen as "non-vaccine" (non-PCV13). Statistical data processing was performed using IBM SPSS Statistics 25 software package.

RESULTS AND DISCUSSION. During isolate typing we identified 29 different *S. Pneumoniae* sero-

types, two isolates (1,4 %) were assigned to untypable. Most pneumococci (65, %; 91/140) were related to 11 various vaccine serotypes (1, 3, 4, 6A, 6B, 9V, 14, 18C, 19A, 19F, 23F) in the ten-year study period. The proportion of non-vaccine strains was 35% (49/100) and it was presented by 18 serotypes (6C, 6D, 8, 9N, 10A, 11A, 15B/C, 16F, 20, 22F, 23A, 23B, 28A, 28F, 31, 34, 35C, 35F) as well as two untypable isolates. 19F (22,1 %; 31/140); 3 (11,4%; 16/140); 14 (7,9 %; 11/140) and 23F (7,9 %; 11/140) were leading among vaccine serotypes. 11A serotype (6,4 %; 9/140) prevailed over non-vaccine strains. When considering pre-vaccination (pre-PCV13) and post-vaccination (post-PCV13) periods separately, we revealed the change in the incidence of vaccine and non-vaccine serotypes in time. PCV13 serotypes peaked in 2012, entering pre-vaccination period (2011–2014). This year 27,5 % of specimens have had vaccine serotypes (25/91). It is noteworthy that most of these isolates were obtained from children with CM (68 %; 17/25). 14,19F and 23F (46,9 %; 31/66) were the predominant vaccine serotypes in pre-PCV13-period. These findings are consistent with the results, previously observed in our center for children with the nasopharyngeal carriage of *S. pneumoniae* [6]. Among non-vaccine serotypes 11A (9,1 %; 6/66) occurred more often in children with chronic bronchopulmonary pathology in pre-PCV13-period. However, non-vaccine serotype 15 B/C was most commonly detected in children with the nasopharyngeal carriage [6].

We observed a growing trend in non-vaccine serotypes among children with chronic bronchopulmonary pathology in post-PCV13-period. Their number increased by 4,8 %, and most of the specimens (36,7 %; 18/49) with non-PCV13-serotypes fell on 2016–2017, i.e. right after implementing the vaccine in a wide clinical practice that is consistent with the results of previous studies [6]. It was observed, mainly, in the group of children with CM, who had an increase in the number of strains with non-vaccine serotypes from 32,5 % to 45,7 % after 2014. In addition to increasing number of non-vaccine strains, there was a growth in their serotype variety in the postPCV13-period.

Apart from the identified in the pre-vaccination period (11A, 16F, 6C, 15B/C, 10A, 28A, 35C), 23B, 34, 31, 22F, 28F, 20, 6D, 35F serotypes appeared. Two specimens, related to untypable, were also detected in the post-vaccination period.

CONCLUSIONS. The obtained data prove the impact of PCV on the circulation of vaccine and non-vaccine *S. pneumoniae* strains. An increase in the

number and serotype variety of non-vaccine strains after implementing PCV13 in the Russian national immunization programme requires continuous monitoring of *Streptococcus pneumoniae* population. The emergence of strains with new non-vaccine serotypes in children with chronic bronchopulmonary pathology shows the need for further study of pneumococcal population, including their serotype variety and sensitivity to antimicrobial drugs.

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Characteristics of subpopulation composition of peripheral blood lymphocytes in children with different forms of congenital ichthyosis

SCO — краткое сообщение

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D. G. Kuptsova¹, N. N. Murashkin^{1,2,3}, S. G. Makarova^{1,4}, R. A. Ivanov^{1,2}, K. O. Avetisyan¹, T. V. Radigina¹, O. V. Kurbatova¹, S. V. Petrichuk¹

¹ National Medical Research Center for Children's Health, Moscow, Russian Federation

² Central State Medical Academy of Department of Presidential Affairs, Moscow, Russian Federation

³ Sechenov First Moscow State Medical University, Moscow, Russian Federation

⁴ Lomonosov Moscow State University, Moscow, Russian Federation

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Характеристика субпопуляционного состава лимфоцитов периферической крови у детей с различными формами врожденного ихтиоза

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Купцова Д. Г.¹, Мурашкин Н. Н.^{1,2,3}, Макарова С. Г.^{1,4}, Иванов Р. А.^{1,2}, Аветисян К. О.¹, Радыгина Т. В.¹, Курбатова О. В.¹, Петричук С. В.¹

¹ ФГАУ «Национальный медицинский исследовательский центр здоровья детей» Министерства здравоохранения РФ, 119991, г. Москва, Ломоносовский проспект, д. 2, с. 1, Россия

² ФГБУ ДПО «Центральная государственная медицинская академия» Управления делами Президента РФ, Москва, Россия

³ ФGAOYBO «Первый Московский государственный медицинский университет имени И. М. Сеченова» Министерства здравоохранения РФ (Сеченовский университет), Москва, Россия

⁴ ФГБОУ ВО «Московский государственный университет имени М. В. Ломоносова», Москва, Россия

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Congenital ichthyosis (CI) is a large heterogeneous group of rare genetic skin diseases with immune disorders and malformation of keratinization of the skin [1]. Defective differentiation of keratinocytes and abnormal formation of the epidermal barrier are observed in these diseases [1, 2]. The resulting dysfunction of the skin barrier causes an increase in transepidermal water loss (TEWL-value of the skin barrier function) and inflammation [3, 4]. Impaired keratinization of the skin regardless of CI form is clinically characterized by dryness, peeling, hyperkerato-

sis and erythema [1, 2]. Studies in recent years have shown that clinical manifestations of CI and progression of the disease is also caused by uncontrolled activation of immune system cells and inflammatory mediators in the skin [3, 5, 6]. Regarding patients with CI, it was found that a strong activation of cytokines in the IL-17 and TNF α families is closely associated with the severity of the disease and change in TEWL value [7, 8].

CI covers the spectrum of syndromic and non-syndromic dermatosis with a different genetic basis [1,

6]. Non-syndromic ichthyoses include the most described and common phenotype — ichthyosis vulgaris (IV), caused by *FLG* gene mutation (encoding filaggrin). The most prevalent rare non-syndromic CI forms are lamellar ichthyosis (LI, *TGM1* gene mutation, encoding transglutaminase 1 and others) and congenital ichthyosiform erythroderma (CIE; multiple genes), known collectively as autosomal recessive congenital ichthyosis as well as keratinopathic ichthyosis (KPI) with mutations in *KRT1*, *KRT2* and *KRT10* genes (genes, encoding keratin proteins) [1, 9, 10]. Syndromic forms of CI are characterized by the monogenic type of inheritance, low prevalence and include Netherton syndrome (NS), which is characterized by a large number (> 80) of *SPINK5* gene mutations and specific clinical manifestations [1, 7].

The relevance of the study of the immune system role in developing chronic inflammation with ichthyosis is driven by a search for efficient targeted methods of treating children with various forms of CI. Nowadays, there is still a huge need for safer and more efficient treatment methods of ichthyosis in children [1, 5, 11]. Complex blood immunophenotyping in more patients with ichthyosis will help characterize the immune profile of different disease forms and choose targeted therapy in children with CI.

STUDY OBJECTIVE: to determine value characteristics of major and small lymphocyte populations in children with various forms of congenital ichthyosis.

MATERIALS AND METHODS. 96 children were screened with non-syndromic (n = 65) and syndromic (n = 31) forms of congenital ichthyosis, including Netherton syndrome (NS — group 1, n = 20), X-linked ichthyosis (group 2, n = 11), lamellar ichthyosis (LI — group 3, n = 17), ichthyosiform erythroderma (CIE — group 4, n = 16), keratinopathic ichthyosis (KPI — group 5, n = 15) and ichthyosis vulgaris (IV — group 6, n = 14). The age of the screened children was from 2 months to 18 years. The examination was carried out on the basis of the laboratory of experimental immunology and virology, screening and treatment of patients — in the Department of Dermatology and Allergology of FSAI NMRC for Children's Health" of the Ministry of Health of the Russian Federation.

Diagnosis and subtype of CI were made based on the results of molecular genetic studies with NGS.

The study is approved by the local ethics committee (Protocol № 6 of 17.06.2021). Written informed consent was obtained from the children's parents during examination.

All the patients had immunophenotyping of peripheral blood lymphocytes by flow cytometry on «Novocyte» cytofluorimeter (ACEA Biosciences, USA), using monoclonal antibodies (Beckman Coulter, USA). The method of step-by-step gating in CD45⁺ region determined the composition of: T-lymphocytes (CD3⁺), T-helpers (CD3⁺CD4⁺), cytotoxic T-lymphocytes (CD3⁺CD8⁺), B-lymphocytes (CD3⁺CD19⁺), NK-cells (CD3⁺CD16⁺CD56⁺), regulatory T-cells (CD4⁺CD25^{high}CD127^{low} — Treg), activated T-helpers (CD4⁺CD25⁺CD127^{high} — Thact), Th17-lymphocytes (CD3⁺CD4⁺CD161⁺ — Th17) and Th2-lymphocytes (CD3⁺CD4⁺CD294⁺ — Th2).

Since the study involves children of different age, to assess changes in major and small lymphocyte populations, deviations of individual indicators from the level of the age norm were calculated by the formula:

$$X_n = (X_{\min} - X) / 0,01 \times (X_{\max} - X_{\min}), \text{ где}$$

X_n — the value of the individual indicator, standardized on the age norm; X — the value of the studied indicator; X_{\max} — the upper limit of the age norm; X_{\min} — the lower limit of the age norm. The range of the age norm was accepted as 100 %.

Statistical processing of the data obtained was performed using Statistica 10.0 program. Descriptive statistics of quantitative trait is presented in the format: median (lower and upper quartiles) — Me ($Q_{0,25}$ — $Q_{0,75}$). The non-parametric Mann-Whitney test was used to evaluate significance of differences between groups. Differences were considered statistically significant at $p < 0,05$.

RESULTS AND DISCUSSION

The first stage of the study included analysis of the percentage of major and small lymphocyte populations in children with syndromic and non-syndromic forms of congenital ichthyosis. Regardless of CI form, the children had an increase in the concentration of activated T-helpers relative to the values of the age norm. Analysis revealed a significant increase in the

relative composition of Th17-lymphocytes in the group of children with syndromic forms of the disease as to rates in the group with non-syndromic ones: for the relative composition of Th17 (% from LF) deviation from the norm was 84,3 (30–203) % versus 29,0 (–2,3–77) %, $p = 0,005$; for the relative composition of Th17 (% from CD4) – 111,8 (22,4–221) % versus 18,2 (–12–80) %, $p = 0,001$.

Composition analysis of major lymphocyte populations in children with Netherton syndrome, X-linked ichthyosis, lamellar ichthyosis, ichthyosiform erythroderma, keratinopathic ichthyosis and ichthyosis vulgaris showed that the relative and absolute composition of T-lymphocytes, T-helpers, NK-cells, B-lymphocytes was mainly within the age reference values, however, there were statistically significant differences and a large range between the forms of the disease. Patients with Netherton syndrome experienced a reduction in the percentage of cytotoxic T-lymphocytes by 12,5 % (–21–42,6) below the level of the age norm. A similar decrease in the relative composition of CD8⁺ T-cells was detected in children with lamellar ichthyosis: –10,8 (–27,1–20,7) %. The group of children with X-linked ichthyosis showed a decline in the relative and absolute composition of B-lymphocytes ($p < 0,05$), and the group with KPI – a decrease in the composition of B-lymphocytes below the age norm. Children with vulgar ichthyosis are characterized by a significant reduction in the composition of NK-cells relative to the age norm and indicators of children with lamellar ichthyosis ($p = 0,025$) and keratinopathic ichthyosis ($p = 0,002$).

The greatest changes in the lymphocyte composition were identified in the analysis of small populations of CD4⁺ T-cells in the peripheral blood in children with different forms of CI. composition content of activated T-helpers by 1,8–3,9 times in regard to the values of the age norm. As for regulatory T-cells, the largest increase in the composition of this population was shown in children with Netherton syndrome and was 126 (46–200) %—for the relative value (% CD4) and 155 (–5–349) % — for the absolute

one (cells/ μ L). The relative composition of Treg was within normal range in the other groups of children with CI, while the absolute composition of the population was increased in the groups with LI, CIE, KPI и IV. It is worth noting that there was a large spread of Treg content in all forms of congenital ichthyosis.

The content analysis of Th17-lymphocytes in children with different forms of CI showed that the increased population was observed in children with Netherton syndrome: for the relative composition (% CD4) by 112 (65–209) %; for the absolute one (cells/ μ L) by 168 (43–342) %. An increase in the absolute number of Th17 was also found in the group of children with ichthyosiform erythroderma and was 190 (76–311) % in regard to the age norm. A rise in the composition of Th2- lymphocytes was revealed with NS, CIE and KPI. The highest growth in the composition of Th2-lymphocytes relative to the values of the age norm was detected in the group of patients with keratinopathic ichthyosis at 200 (131–393) % for the relative composition and at 399 (172–571) % for the absolute one.

Thus, the study has allowed to determine characteristic deviations of the values of major and small lymphocyte populations for different forms of congenital ichthyosis in children. A significant increase in the composition of activated T-helpers and Th17-lymphocytes was shown for pediatric patients with Netherton syndrome, ichthyosiform erythroderma and lamellar ichthyosis, and a rise in the composition of regulatory T-cells — for children with Netherton syndrome, keratinopathic ichthyosis, ichthyosiform erythroderma and ichthyosis vulgaris.

CONCLUSION

The obtained data on the state of cellular immunity in children with different forms of congenital ichthyosis expands the understanding of the disease immunopathogenesis and may serve as a basis for choosing targeted biological therapy and thus allows to improve patients' condition and to predict the course of the disease.

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Antinuclear antibodies in children with Wilson's disease

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O. V. Kurbatova¹, A. A. Zhuzhula¹, S. V. Lapin², M. A. Snovskaya¹, D. I. Kozlova^{3,4}, S. V. Petrichuk¹, D. G. Kuptsova¹, D. A. Kuznetsova², G. B. Movsisyan¹, A. D. Komarova¹, T. V. Radygina¹, A. B. Guslev^{2,3}, I. V. Kholopova², E. L. Semikina^{1,5}, S. G. Makarova^{1,6}, A. S. Potapov^{1,5}, A. P. Fisenko¹

¹ National Medical Research Center for Children's Health, Moscow, Russian Federation, 119991, Lomonosovsky Prospekt, 2, p. 1, Moscow, Russia

² Federal State Budgetary Educational Institution of Higher Education PSPbSMU named after acad. I. P. Pavlova Ministry of Health of Russia, St. Petersburg, Russia

³ Federal State Budgetary Healthcare Institution St. Petersburg Clinical Hospital 1 of the Russian Academy of Sciences, St. Petersburg, Russia

⁴ Federal State Budgetary Institution of Science Institute of Evolutionary Physiology and Biochemistry named after I. M. Sechenov Russian Academy of Sciences, St. Petersburg, Russia

⁵ Sechenov First Moscow State Medical University, Moscow, Russian Federation

⁶ Lomonosov Moscow State University, Moscow, Russian Federation

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Антинуклеарные антитела у детей с болезнью Вильсона

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Курбатова О. В.¹, Жужула А. А.¹, Лапин С. В.², Сновская М. А.¹, Козлова Д. И.^{3,4}, Петричук С. В.¹, Купцова Д. Г.¹, Кузнецова Д. А.², Мовсисян Г. Б.¹, Комарова А. Д.¹, Радыгина Т. В.¹, Гуслев А. Б.^{2,3}, Холопова И. В.², Семикина Е. Л.^{1,5}, Макарова С. Г.^{1,6}, Потапов А. С.^{1,5}, Фисенко А. П.¹

¹ ФГАУ «Национальный медицинский исследовательский центр здоровья детей» Министерства здравоохранения РФ, 119991, г. Москва, Ломоносовский проспект, д. 2, с. 1, Россия

² ФГБОУ ВО ПСПбГМУ им. акад. И.П. Павлова Минздрава России, г. Санкт-Петербург, Россия

³ Федеральное государственное бюджетное учреждение здравоохранения «Санкт-Петербургская клиническая больница №1» Российской академии наук, г. Санкт-Петербург, Россия

⁴ Федеральное государственное бюджетное учреждение науки «Институт эволюционной физиологии и биохимии им. И.М. Сеченова» Российской академии наук, г. Санкт-Петербург, Россия

⁵ ФГАОУ ВО «Первый Московский государственный медицинский университет имени И.М. Сеченова» Министерства здравоохранения РФ (Сеченовский университет), Москва, Россия

⁶ ФГБОУ ВО «Московский государственный университет имени М.В. Ломоносова», Москва, Россия

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ВВЕДЕНИЕ

Wilson-Konovalov's disease (WD) is a hereditary condition, transmitted as an autosomal recessive trait and associated with impaired excretion of copper from the body. Excessive accumulation of copper causes combined lesion of parenchymal organs (pri-

marily liver) and brain [1, 2]. There is a description of over 700 *ATP7B* protein mutations (Cu^{++} transporting beta polypeptide; beta-polypeptide cooper-transporting ATPase), which may cause cooper metabolism disorder [1, 2, 3]. Missense variant is the most prevalent pathogenic variant, leading to replacement

of histidine by glutamine in 1069 position of the encoded protein *c.3207C>A (p.H1069Q)* [4]. It is also found that this H1069Q variant is associated with late neurological manifestations [5].

The clinical picture of Wilson's disease is distinguished by the polymorphism: in 40–45 % of cases the disease debuts with liver damage, developing at the age 5–18; rarer (in 30 %) — with the development of neurological and psychiatric disorders [6]. The symptoms of the disease include: jaundice, edema of shin, increased abdominal volume, esophageal varicose veins, propensity to form bruising and extended hemorrhage. Some patients with WD may have only minor abnormalities in biochemical parameters of liver function with no other symptoms over the years. Other patients with WD experience a fast progression of liver inflammation in the form of chronic hepatitis with high activity, pronounced jaundice, necrosis of hepatocytes and rapid transformation into cirrhosis [6]. Considering diversity of clinical symptoms, there are difficulties in differential diagnosis of Wilson's disease with other liver diseases [7]. In the early stages of hepatocellular damage, the involvement of endoplasmic reticulum, mitochondria, peroxisome and nucleoli in the process, combined with decreased activity of mitochondrial enzymes causes lipid peroxidation, the accumulation of triglycerides, and further — necrosis of hepatocytes. Malondialdehyde, produced by lipid peroxidation, stimulates collagen synthesis, contributing to fibrogenesis [8].

There is the description of disturbances in the immune system in patients with WD, progressing with increasing liver fibrosis stage and age of patients [9]. Subpopulation composition of lymphocytes in children with WD changes and is characterized by involvement of T-cellular immunity in the process of liver fibrosis, and changes are similar to those in other liver diseases [10]. For instance, different stages of fibrosis in non-alcoholic fat liver are accompanied by the accumulation of T-cell and NK-cell subpopulations with different functions and phenotypes in the liver tissues that usually causes pro-inflammatory effects [11].

The process of liver fibrosis is accompanied by changes in the profile of circulating cytokines in chronic liver disease, by the concentration of which it is possible to carry out differential diagnostics between fibrosis stages [12, 13].

The clinical picture of chronic inflammation of the liver in WD is less distinguishable from liver dam-

age of another genesis and causes the development of liver cirrhosis without pathogenetic therapy. This points to the need for detecting WD in all patients with chronic damage of hepatic parenchyma [14].

Characteristic of chronic hepatitis in WD is a moderate increase in biochemical markers of cytotoxicity, cholestasis and bilirubin exchange with a high level of structural changes in hepatic parenchyma on the results of morphological examination of liver biopsy slides [15].

Clinically WD may occur like autoimmune hepatitis with a detected high level of serum immunoglobulins and nonspecific autoantibodies. Hence, it is also necessary to exclude Wilson's disease in patients with autoimmune hepatitis with inefficiency of corticosteroid therapy [15]. However, the detection rate of autoantibodies in WD has not been studied, only isolated cases of their detection are described that often causes a false positive diagnosis of "autoimmune hepatitis" [16].

STUDY OBJECTIVE: to evaluate the presence of antinuclear antibodies, parameters of cellular immunity and the composition of circulating cytokines in children with WD.

MATERIALS AND METHODS. As part of diagnostic measures for screening children with liver diseases, treated in FSAI "NMRC for Children's Health" of the Ministry of Health of the Russian Federation, there was examination of 46 children with clinical signs of autoimmune hepatitis. All the children got evaluation of the liver fibrosis stage by the method of transient liver elastography on FibroScan F502 (EchoSence, France). METAVIR scale was used to diagnose LF stage [17]. Complete blood test (automated hematology Sysmex XN 550, Japan), biochemical blood test (AU680, USA), immunophenotyping of peripheral blood lymphocytes (CYTOMICS FC500, Beckman Coulter, USA), the determination of antinuclear antibody (ANA) on the cell line HEp-2 using the reaction of indirect immunofluorescence (RNIF, Immco Diagnostics, Inc, USA). Determining the level of circulating cytokines was performed using enzyme-linked immunosorbent assay MILLIPLEX Human Cytokine, based on Luminex technology (Merck Millipore, Germany). To confirm Wilson's disease, molecular genetic study was carried out by the Sanger method of sequencing on 3500xL Genetic Analyzer (Applied Biosystems, USA). All changes in the reference AT-

P7B gene sequence were described according to the HGVS nomenclature in accordance with the accepted recommendations [18]. Statistical calculations were carried out using Statistica 10.0 program (USA).

RESULTS AND DISCUSSION

Out of 46 examined children, 11 (24 %) were diagnosed with “Wilson’s disease”, based on molecular genetic testing. The following pathogenic variants of ATP7B gene were identified, causing the development of WD: *c.2304dup (p.M769HfsTer26)*, *c.2998G>A (p.G1000R)*, *c.3002T>G (p.V1001G)*, *c.3036dup (p.K1013QfsTer15)*, *c.3472_3482del (p.G1158FfsTer2)* и *c.3207C>A (p.H1069Q)*. The variant *c.3207>A (H1069Q)* was more common as a part of compound-heterozygous mutation in the presented sample of patients, and as a part of homozygous mutation only in two children. 35 children (76%) got verified diagnosis of “autoimmune hepatitis”.

ANA on the cell line HEp-2 were found in 4 out of 11 children with WD diagnosis, using RNIF, moreover, 3 children had a low titer of ANA (1/160), one patient had a high one (1/2560). Only the facts of low ANA titers in patients with WD were previously reported [15]. The type of fluorescence in all children with WD was similar— nuclear granular (AC-2,4). It is interesting to note that only in one child with revealed ANA, the pathogenic variant *c.3207>A (H1069Q)* was in homozygous state. *H1069Q* mutation in the homozygous state causes the development of severe hepatic insufficiency, depression, dysarthria and tremor at an earlier age than a mutation in the compound-heterozygous state [5, 19].

According to transient liver elastography, the child with a high ANA titer (1/2560) did not have liver fibrosis of F0 stage. Perhaps such a high ANA titer of the child is associated with sensitization to cow’s milk protein (class 3), egg (class 2), wheat (class 3), total gE = 3043 IU/mL. In children with low ANA titer liver fibrosis ranged from F0 to F2. Hence, the presence of ANA in children with WD

did not depend on liver fibrosis stage in this sample of patients.

The comparison of laboratory values of children with WD and the presence of ANA with no antibodies revealed that the former had a significantly lower concentration of albumin in serum than the latter: Me = 65,3 [63,55; 66,85] g/l versus Me = 68 [67; 69,3] g/l ($p = 0,042$).

Children with WD are characterized by increased concentration of T-cells through the population of T-helpers with a decrease in cytotoxic T-lymphocytes, B-lymphocytes and NK-cells against the backdrop of a rise in Thact, Th17-lymphocytes and Treg [20]. The level of Treg, Th17-lymphocytes, activated T-helpers in the group of children with and without ANA did not have statistical difference and was characterized by a large data range, especially in the group of children without ANA. Nevertheless, we can note tend to decrease the relative amount of Treg and to increase the relative amount of Th17-lymphocytes in the group of children with ANA. Perhaps a large spread of parameters of cellular immunity is due to the fact that the group of children without ANA had children with different fibrosis stages, including F3-4 stage, in which the level of Th17-lymphocytes is significantly higher than in the early stages of liver fibrosis in children with WB [9].

The evaluation of the level of circulating cytokines revealed that the level of IL27 was much higher in the group of children with ANA than in the one without ANA: Me = 12097 [11028; 13299] pg/ml versus

Me = 8338 [8,8; 10559] pg/ml ($p = 0,024$). Yet, IL22 and $TNF\alpha$ were significantly lower than in the group without ANA and were: $TNF\alpha$ — Me = 60 [53,55; 64,86] pg/ml versus Me = 130 [66,6; 164,5] pg/ml ($p = 0,024$); IL22 — Me = 20,3 [16,4; 22,4] pg/ml versus Me = 41,9 [34,8; 592,9] pg/ml ($p = 0,012$). In addition, there was tend to a higher concentration of IL4 and IL9 in children with ANA. It is worth noting a large spread in values of circulating cytokine levels that might also be associated with the fact that

the children with WD in this sample experienced different liver fibrosis stages, and the change in the cytokine profile of patients is related to liver fibrosis stages [12, 13].

The detection of antinuclear antibodies in patients with accumulation diseases is justified in terms of pathophysiology: damaged tissues of the body, including those associated with reactions to accumulating toxic components, are capable of inducing immune response to tissue damage. Liver damage as well as in Wilson's disease, can be a prime example of such reactions due to the fact that there is a considerable number of lymphocytes and macrophages in the liver tissue, which are the main producers of cytokines. Hence, assessment and monitoring of immune re-

sponses in WD is a promising marker of the severity of the condition and therapy efficacy. A promising area of research is broadening methods of drug therapy, with the inclusion of immunomodulators, metabolites and antioxidants, optimizing immune responses and the cytokine profile of patients [21].

CONCLUSION

The presence of ANA in children with Wilson's disease may indicate the attachment of an autoimmune component to the congenital genetic disease. A larger study is required regarding the frequency of ANA disclosure in children with WD, the association of ANA with liver fibrosis and the presence of specific pathogenic variants in *ATP7B* gene.

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