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BACTERIAL SUPERINFECTION OF THE RESPIRATORY VIRUSES

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REZUMAT

SUPRAINFECTIA BACTERIANA A INFECTIILOR RESPIRATORII VIRALE

Actualitatea temei. Suprainfecția bacteriană a infecțiilor respiratorii virale este un fapt confirmat, atât prin diagnosticul clinic, cât și cel molecular. *Material și metodă.* Au fost luate în considerare studiile care au recrutat pacienți de vârstă pediatrică în vederea identificării virale, bacteriene, precum și coinfecția viro-bacteriană și suprainfecția bacteriană a infecțiilor respiratorii virale premonitorii. *Rezultate.* Studiile au consemnat creșterea incidenței patogenilor virali odată cu dezvoltarea tehnicilor de detecție moleculară, modificarea etiologiei bacteriene odată cu introducerea vaccinării conjugate antipneumococice. Coinfecția viro-bacteriană sau prezența concomitentă a mai multor tulpini virale cresc severitatea bolii, cu o staționare de lungă durată în unitățile de terapie intensivă. *Concluzii.* Suprainfecția bacteriană a infecțiilor respiratorii virale rămâne un fapt, cu implicare terapeutică și evolutivă. **Cuvinte cheie:** patogenii respiratori virali și bacterieni, infecția respiratorie virală, coinfecția viro-bacteriană, suprainfecția bacteriană.

SUMMARY

BACTERIAL SUPERINFECTION OF THE RESPIRATORY VIRUSES

Theme topicality. Bacterial superinfection of the respiratory viruses is a confirm fact, either by clinical diagnosis, and molecular one. *Material and methods.* It were considered some studies which have been recruited pediatric patients for viral and bacterial identification, also for viral and bacterial coinfection and superinfection of the premonitory viral respiratory infection. *Results.* The studies have been recorded an increase incidence of viral pathogens with the improvement of molecular detection techniques, and the bacterial etiology changes with the introduction of conjugate pneumococcal vaccin. Viral and bacterial coinfection or the concomitant presence of a multiple viral strains increase the severity of disease, with a long length-of-stay in pediatric intensive care units. *Conclusions.* Bacterial superinfection of viral respiratory infection remains a fact, with therapeutic and evolutive implication. **Keywords:** viral and bacterial respiratory pathogens, viral and bacterial coinfection, bacterial superinfection.

Introduction

Viral infection-bacterial infection relationship (coinfection, superinfection). In prevaccinal era (conjugate pneumococcal vaccin), *S. Pneumoniae* was considered bacteriologic leader causing lower airways infection, respectively pneumonia (community-acquired)⁴. *S. Pneumoniae* has followed, as frequency, by *Haemophilus influenzae*, *Streptococcus pyogenes*, *Staphylococcus aureus* and *Moraxella catarrhalis*. Enteric gram-negative bacteria and those intracellular and atypical (*Mycoplasma pneumoniae* și *Chlamydia pneumoniae*)^{13,15,25} were also reported. With introduction of multi-valent and conjugate haemophilus and pneumococcal vaccine, the frequency of community-acquired pneumonia was significant reduced in countries with a complete national program of vaccination. In developing countries, with an incomplete vaccine program, bacteria remains the main etiologic factor of community-acquired pneumonia.

With the improvement of molecular diagnosis, viruses with respiratory tropism are detected with a higher frequency in children diagnosed with pneumonia, especially those younger than 5 years. The most frequent viruses are respiratory syncytial virus, influenza, parainfluenza, rhinovirus and adenovirus¹². The newest molecular techniques have allowed identification of metapneumovirus, human coronavirus, and human bocavirus. In industrialized countries up to 81 % children with community-acquired pneumonia claim viral etiology, especially respiratory syncytial virus (up to 48 %).

It exists more and more evidences regarding the important role of respiratory viruses in facilitating bacterial colonization in child. Most of those refer to lower airways infection, respectively pneumonia (community-acquired pneumonia). The mechanism is not totally known, either the facilitating bacterial colonization is producing directly, or viruses contribute of development of severe forms of

disease. Viruses make a vulnerability of host respiratory epithelium, thus favoring the initiation of respiratory bacterial infection. If the bacterial etiology is significant influenced by conjugate pneumococcal vaccine, *S. pneumoniae* infection being the leader of bacterial infection, the viral etiology being influenced only by improvement of viral techniques detection, the most frequent viruses have been influenza, respiratory syncytial virus, parainfluenza and adenovirus^{30,31}. And the invasive pneumococcal infection has observed mostly in the context of viruses¹⁹. The proof of viral and bacterial coinfection is certain, in or not directly relationship with bacterial superinfection of respiratory viral infection. A trial made in South Africa which included children with conjugate pneumococcal vaccination has proven a decrease of pneumonia with *S. pneumoniae* incidence, also of viral pneumoniae, just in context of respiratory viruses and pneumococcus relationship^{26,27}. Much more than that, the Australian¹² statistical data have shown a decrease of viral pneumonia incidence as effect of pneumococcal vaccination. The respiratory syncytial virus has recorded as favorable factor of bacterial colonization with *H. influenzae* and *S. aureus* in 25 % of hospitalized patients with severe form of A type influenza infection during 2000 pandemia²³. The patients which have presented viral and bacterial coinfection have presented a severe evolution, with ventilator support and long length-of-stay in intensive care unit^{23,24}.

Methodology. The relationship between naso-pharyngeal pathogens and the form of severity of lower airways infection has studied in a couple of clinical studies. The naso-pharyngeal pathogens could be considered as predictive factors for pneumonia severity in child. A study made in USA has demonstrated a link between respiratory syncytial virus infection and long hospitalization, respiratory failure and intensive treatment. In Vietnam, children with radiologic confirmed pneumonia were detected with pneumococcal infection in a much larger proportion than other etiology, and the pneumococcus presence was 15 times more in patients with viral and bacterial coinfection. Other studies, in Kenya and Holland have reported a nasopharyngeal load of respiratory syncytial virus in children with severe forms of disease.

In comparison with many studies with the same target – the determination of community-acquired pneumonia etiology, a few studies have included healthy children or asymptomatic ones, as control group. The naso-pharyngeal carriage was frequent associated with healthy children or asymptomatic ones. Are required studies which must use supplement investigation to certify that asymptomatic carriage is a fact in healthy population, and its involvement in community-acquired pneumonia become a challenge.

Although the conjugate pneumococcal vaccination has reduced a lot the hospitalization rate of community-ac-

quired pneumonia in industrialized countries, pneumococcus remains an important pathogen regarding non-vaccin serotype with severe and complicate pneumonia potential (inclusively, empiema).

The incidence of viral infection, bacterial infection, viral and bacterial coinfection and bacterial superinfection varies according to level of income country, patient age and utilized sample (naso-pharyngeal fluid, sputum, broncho-alveolar fluid)³².

Results. Start with 1993-1995 period, in Finland²², a study which included children younger than 18 years, diagnosed with community-acquired pneumonia has shown the presence in naso-pharyngeal aspiration of viral pathogen in 62 percent, and of a bacterial pathogen in 53 percent. The pathogens frequency was: *S. pneumoniae* 37 %, respiratory syncytial virus 29 %, human rhinovirus 24 %, parainfluenza 10 %, *H. influenzae* 9 %, adenovirus and *M. pneumoniae* each one 7 %, influenza and *Moraxella catarrhalis* each one 4 %.

The other study²⁹, performed in 1999-2000 period, made in USA, which included children 6 weeks-17 years old, diagnosed with pneumonia, has showed the following pathogens in naso-pharyngeal aspirate: *S. pneumoniae* 44 %, influenza 21 %, *M. pneumoniae* 14 %, respiratory syncytial virus and parainfluenza each 13 %, *Chlamydia pneumoniae* 9 %, adenovirus 7 %, rhinovirus 3 %.

In 2001-2002 period were performed two studies:

- in Italy¹⁰, children 3 months-16 years old, diagnosed with pneumonia, used naso-pharyngeal swab and blood culture as a detection methods (*M. pneumoniae* 27 %, *S. pneumoniae* 18 %, human parainfluenza 12 %, influenza 9 %, human metapneumovirus 5 %);
- in Japan³⁶, children 1 month-13 years old, diagnosed with pneumonia, used naso-pharyngeal swab and blood culture as a detection methods (respiratory syncytial virus 48%, *S. pneumoniae* 36 %, *H. influenzae* 26 %, influenza 22 %, *M. pneumoniae* 17 %, parainfluenza 14 %).

In 2003-2005 period, in Switzerland⁵, was initiated a study which recruited children 2 months-5 years old, diagnosed with pneumonia. The results revealed the presence in the naso-pharyngeal aspirate of the following respiratory pathogens: bacteria, 72 percent (*S. pneumoniae*, 46 %; *M. pneumoniae*, 11 %; *C. pneumoniae*, 7 %) and viruses, 67 percent (human rhinovirus, 20 %; influenza, 14 %; respiratory syncytial virus and human metapneumovirus, each 13 %; adenovirus, 7 %).

Other study, performed in 2004-2006 period, in Spain⁸, recruited children younger than 3 years, diagnosed with pneumonia, use for pathogen detection naso-pharyngeal aspirate. The study showed the following results: viral and bacterial coinfection (30 %), viruses 67 percent (respiratory syncytial virus 20 %, human bocavirus and

rhinovirus each 14 %, human metapneumovirus 12 %, parainfluenza 11 %, influenza 7 %).

A study from USA³⁴, performed between 2005-2007 period, with patients younger than 3 years, revealed the following:

- for diagnosis of airways infection, with samples from blood and naso-pharyngeal swab, were identified viral pathogens in 90 percent (human rhinovirus 44 %, adenovirus 30 %, respiratory syncytial virus 23 %, parainfluenza 18 %, human metapneumovirus 15 %) and bacteria in 3 percent (*S. pneumoniae* 2 %);
- for asymptomatics, with samples from naso-pharyngeal swab, were identified viral pathogens in 52 percent (human rhinovirus 33 %, adenovirus 16 %, human metapneumovirus 7 %, respiratory syncytial virus 4 % and parainfluenzae 3 %).

The same differentiation, children with respiratory infection and asymptomatics, was performed in a study from Holland²⁰, in 2007-2009 period, children younger than 6 years, with samples collected from nasal fluid:

- for airways infection, the viral pathogens were present in 72 percent (respiratory syncytial virus 26 %, human rhinovirus 20 %, adenovirus 9 %, metapneumovirus and influenza each 6 %);
- for asymptomatics, the viral pathogens were present in 26 percent (human rhinovirus 16 %, adenovirus 9 %, human coronavirus 5 %, influenza 3 % and respiratory syncytial virus 1 %).

In 2006-2007 period, in Finland¹⁶, was initiated a study which recruited children 6 months-15 years old, diagnosed with pneumonia. The results revealed the sputum presence of the following pathogens: bacteria, 91 percent (*S. pneumoniae*, 50 %; *H. influenza*, 38 %; *S. aureus*, 13 %) and viruses, 72 percent (human rhinovirus, 30 %; bocavirus, 18 %; metapneumovirus, 14 %, parainfluenza, 8 %; respiratory syncytial virus, 7 %).

A study performed in Holland³⁵, in 2008-2011 period, with patients 3 months-16 years old, showed the following:

- for airways infection, with samples from blood and naso-pharyngeal swab, were identified bacterial pathogens (*S. pneumoniae* 28 %, *Moraxella catarrhalis* 23 %, *M. pneumoniae* and *H. influenzae* each 16 %, *S. aureus* 10 %);
- for asymptomatics, with samples from blood and naso-pharyngeal swab, were identified bacterial pathogens (*S. pneumoniae* 28 %, *M. pneumoniae* 21 %, *S. aureus* 21 %, *Moraxella catarrhalis* 18 %, *H. influenzae* 15 %).

Other study performed in USA¹⁸, between 2010-2012, with patients younger than 18 years, evidenced the following:

- for pneumonia diagnosis, with samples from blood and naso-pharyngeal swab, were identified viral pathogens in 66 percent (respiratory syncytial virus 28 %, human rhinovirus 27 %, human metapneumovirus 13 %, adenovirus 11 % and influenza 7 %), bacterial pathogens in 8 percent (*M. pneumoniae* 8 % and *S. pneumoniae* 4 %);
- for asymptomatics, with samples from naso-pharyngeal swab, were identified viral pathogens represented by human rhinovirus in 17 percent (other viruses, less than 3 % percent); viral and bacterial coinfection was identified in 7 percent.

A study performed in Sweden³³, in 2011-2014 period, in children younger than 5 years, has shown the following results:

- in children with pneumonia, from naso-pharyngeal aspirate, were identified viral pathogens in 81 percent (respiratory syncytial virus 32 %, human metapneumovirus and human rhinovirus each 23 % and adenovirus 15 %)
- in asymptomatic, from naso-pharyngeal swab, were identified viral pathogens in 56 percent (human rhinovirus 27 %, human bocavirus 21 %, coronavirus 12 %, adenovirus 7 %, respiratory syncytial virus 6 %).

Discussions. As can be seen in the presented studies the frequency of respiratory pathogens varies from country to country, according with income level, vaccination program, detection methods used, samples used. Over the decades, with the conjugate pneumococcal vaccine initiation, the *S. pneumoniae* frequency significant decreased (from 37 % to 4 %). According as, with the improvement of molecular detection techniques, the frequency of viral pathogens increased. The last years studies propose as target the viral and bacterial coinfection detection.

The distribution of identified pathogens in community-acquired pneumonia is variable from country to country^{1,11,17,21,28}. Thus:

- for viral pathogens
 - respiratory syncytial virus was detected in a high percent in Kenya (34 %), Nigeria (30,4 %) and USA (28 %) and almost insignificant in Gambia (4 %);
 - rhinovirus was detected in a high percent in USA (27 %) and much less in Great Britain (8,5 %)
 - human metapneumovirus, the most detected in USA (13 %) and insignificant in Great Britain (0,7 %)
 - influenza, in the same percent of detection (between 2 and 7,4 %), knowing the fact that both the virus and the vaccine have the strongest impact in community
 - bocavirus was reported in a less percent and only

- in a few countries (3,3 %, in Great Britain and 4 %, in Gambia)
- adenovirus presented the most important detection in USA (11 %)
- parainfluenza, with the big rate of detection in Nigeria (19,5 %)
- for bacterial pathogens
 - *S. pneumoniae*, detected in an extreme percent in Gambia (91 %), and in the opposite percent in USA (4 %)
 - *H. influenzae*, detected in 23 percent in Gambia and 2,3 percent in Great Britain
 - *A Streptococcus*, detected in 10,5 percent in Great Britain and 1 percent in USA
 - *S. aureus*, detected in an important percent only in Nigeria
 - *M. pneumoniae*, (8-9,9 percent)
 - *Moraxella catarrhalis*, in 2,3 percent in Great Britain
 - *Klebsiella pneumoniae*, 15,3 percent in Nigeria and insignificant one in Great Britain (0,8 percent).

The results per se are influenced by sample methods of detection of respiratory pathogens. This stage of diagnosis (molecular diagnosis) become very important, especially in the situation when clinics and radiologic exam are not able to sustain the diagnosis of pneumonia (community-acquired).

The British Thoracic Society and The Infectious Disease Society of America guidelines recommend the PCR viral tests or immunofluorescence, while for bacterial detection the methods varied according to the forms of gravity of infection.

In Great Britain are used the following tests: blood culture, real-time PCR, naso-pharyngeal fluid, pleural culture, pneumococcal antigen test, endotracheal tube aspirate, broncho-alveolar fluid.

In USA are used the following tests: blood culture, naso-pharyngeal fluids, oro-pharyngeal fluids, pleural culture, broncho-alveolar fluid, culture from endotracheal tube aspirate.

In Gambia are used the following tests: culture from broncho-alveolar and pleural fluid for non-molecular serotypes, uniplex and multiplex PCR, multilocular sequence typing, molecular serotyping.

The viral diagnosis has been revolutionized in the last 20 years, with ADN detection. With the routinely multiplex PCR technology use in community-acquired pneumonia diagnosis in child, the presence of multiple viral pathogens become a fact, with a rate of 30-40 percent of 4 different types at a single case⁷.

The significance of this multiple presence remain unclear⁶. For respiratory syncytial virus the coinfection with other viruses creates an increase severity of disease⁹. Recent identified, human bocavirus was associated with a rate of 83 percent of coinfection, when the pathogen represents the etiologic factor per se or an exacerbation factor, or an accidental detection. This context remains unclear. The significance threshold utilization for real-time qPCR¹⁴ for estimative viral load has significance in association with evidence clinics of a viral infection.

The determination of viral etiology represents a challenge in countries where the molecular diagnosis is not routinely used. There are some studies which demonstrated a high specificity and a negative predictive value in detection of parainfluenza and adenovirus from naso-pharyngeal aspirate, but a discordance between broncho-alveolar fluid and naso-pharyngeal fluid in detection of bacterial infection.

Concluzii. It is widely accepted the idea of bacterial infection that follow a viral one. The ideal bacterial identification is that one which utilized a sample directly obtains by lung without any host flora contamination. It is very interesting the fact that the viral identification from broncho-alveolar aspirate has a lower rate comparing with naso-pharyngeal aspirate. The blood culture has a limited place (because a higher rate of false positive results). Stupum cultures have a limit regarding the modality of collection (after hypersaline solution nebulization or thoracic wall percussion, with the possibility of contamination from upper airways). Conversely, the serologic tests are considerate the diagnosis gold-standard for bacterial pathogens (*M. pneumoniae* and *S. pneumoniae*). The last generation is considered the multilocular sequence, and for the future the technology of isothermal amplification.

These molecular detection techniques brought major changes in the etiology of respiratory airways infection. What was believed in the past (bronchiolitis, due to respiratory syncytial virus, does not known the bacterial superinfection) seems to be not longer support today (respiratory syncytial virus not only are the favorizant factor for bacterial superinfection, but the coinfection produced a severe form of disease).

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