THE IMPACT OF MICROBIOTA AND DIET ON ASTHMA

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Review

Introduction:

Asthma prevalence has dramatically increased over the last decade in Westernized societies.

Purpose of the review:

- State-of-the-art of the asthma
- Find new lines of investigation
- Reach to write a Review following the guidelines from editorial BioMed Central

Methodology:

- Database: PubMed, WOS and Google Scholar
- Key words: asthma, diet in asthma, and microbiome
- Publication data: 3-5 years

Background Asthma affects 300 million individuals worldwide, approximately. Usually, asthma manifests in early childhood with a hyper-activation of the T-helper 2 arm of the adaptive immune response, and an over-expression of the proinflammatory cytokines IL-4, IL-5 and IL-13, as well eosinophilia and mast cell infiltration of the airways. Furthermore, asthma development is closely related to the altered microbiota in childhood. The microbiota changes over time in response to environmental factors.

Microbiota

Gut microbiota RESIDENT

Bacteroidetes Firmicutes Proteobacteria Actinobacteria Verrucomicrobia Fusobacteria (lower)

Lung microbiota

EPHEMERAL Bacteroidetes

Firmicutes

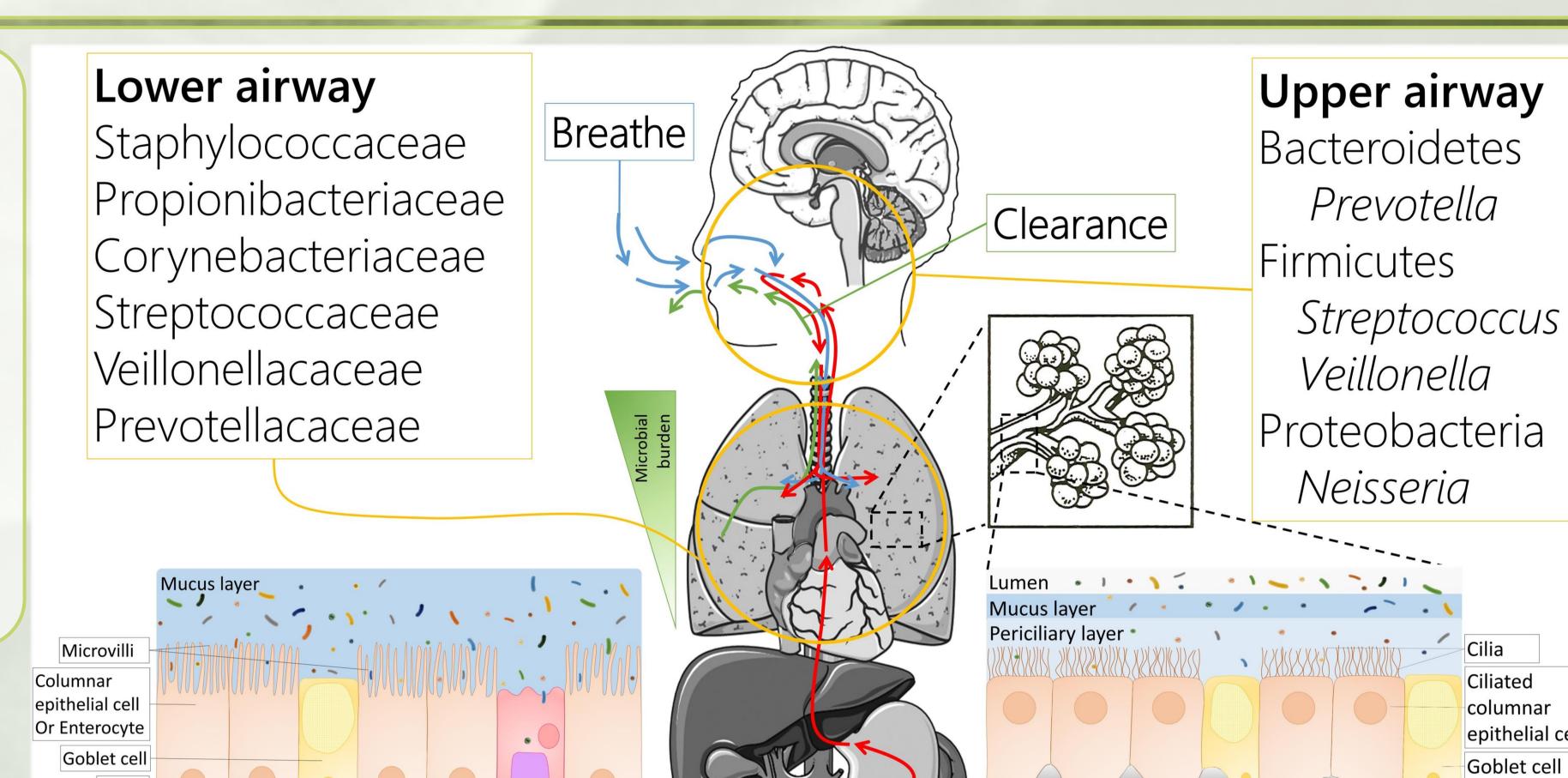
Lactobacillus (1) Proteobacteria (1)

Haemophilus

Pseudomonas Actinobacteria (1)

*Green-marked microbes only modified in asthmatic patients

resemble They microaspiration (gut to lung) and breathing (environment to lung), but at the same time differ due to clearance (coughing, mucocilliary clearance, immune response)



Colon

Lamina propria

Microaspiration

Bacteroidaceae Prevotellaceae Rikenellaceae Lachnospiraceae Ruminococcaceae

Image exclusively elaborated for this poster

Ciliated

columnar

epithelial cel

Goblet cell

Stem cell

The following microbiota modifications during early life are related to increased risk of asthma:

Nasopharyngeal microbiota: dominance of Moraxella, Streptococcus, Corynebacterium, Staphylococcus, Haemophilus, or Alloicoccus

Faecal microbiota: (1) Bacillus fragilis and (1) Faecalibacterium, Rothia, Veillonella, Lachnospira species

Early life critical window

Dysbiosis

Formula feeding

Diet can only modify microbiota in a long-term form applied during pregnancy; other situations (childhood and adult life) does in a transient form.

Small intestine

Lactobacillaceae

Enterobacteriaceae

Diet recommendations to reduce exacerbations:

- Balance of polyunsaturated fatty acids- omega-3 promotes T helper-1 phenotype and stimulate regulatory T cells (Tregs)
- Vitamin D intake- it has a receptor in many immune cells and promote specific Tregs population
- High-fibre consumption- production short-chain fatty acids, energy source for colonocytes, reduce pH and other immune benefits

Mediterranean diet (Intangible Cultural Heritage of Humanity by UNESCO, 2010):

- high intake of unrefined plant-based foods (fruits and fresh vegetables, whole grains, legumes, olive oil and nuts)
- low to moderate consumption of foods of animal origin
- low intake of sugar, refined flour and fast food

Conclusions

- Asthmatics present an increased burden, and higher diversity (number of different species not balanced population) during bronchial hyperreactivity
- Needed to define microbiota in order to diagnose and avoid dysbiosis state

15, 55–63 (2017).

• First efforts focused on gut microbiota -> GUT-LUNG AXIS

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