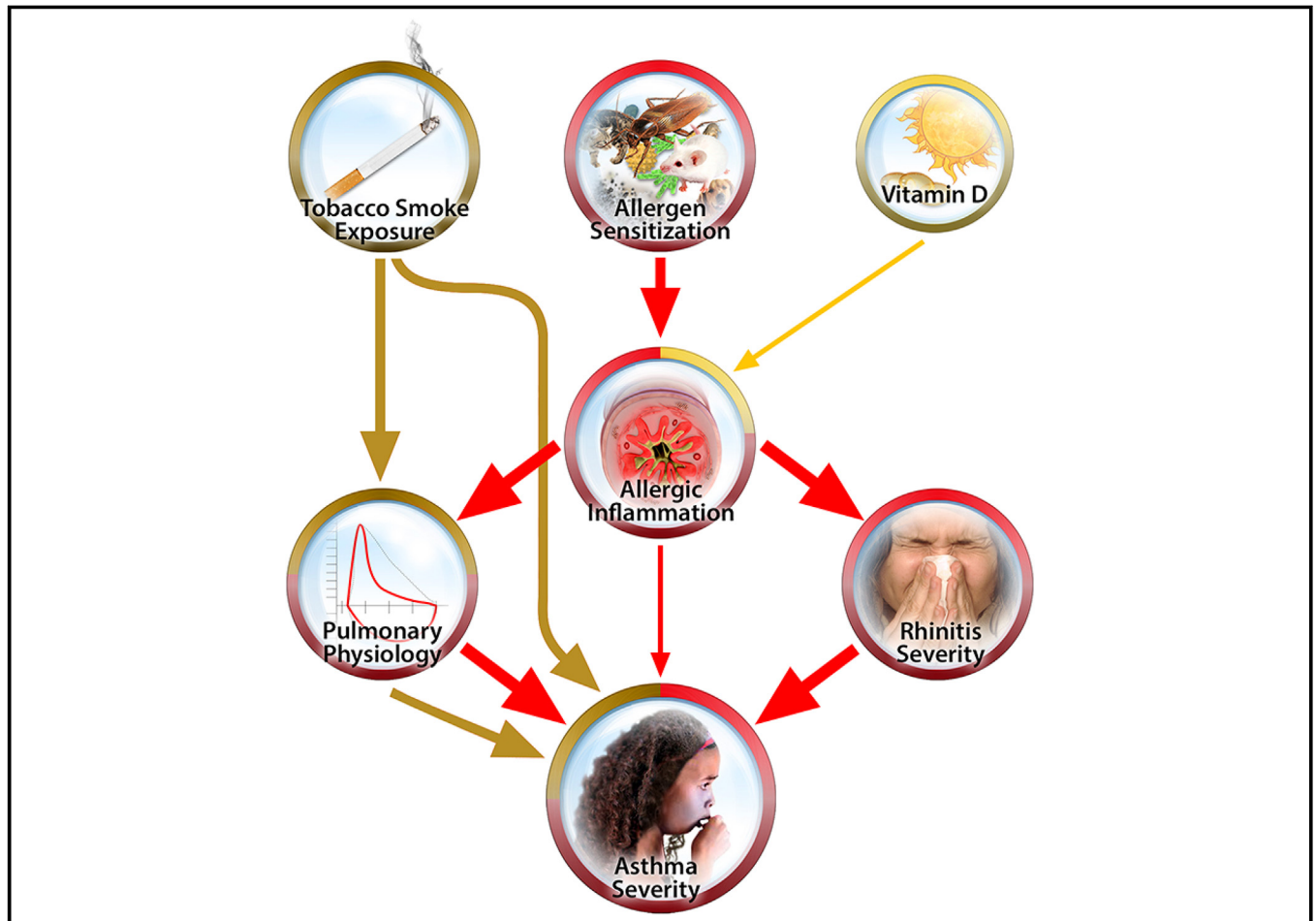


Pathways through which asthma risk factors contribute to asthma severity in inner-city children



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GRAPHICAL ABSTRACT



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of Physicians and Surgeons, Columbia University, New York; ^kChildren's National Health System and the George Washington University School of Medicine and Health Sciences, Washington; ^lthe University of Wisconsin School of Medicine and Public Health, Madison; and ^mthe National Institute of Allergy and Infectious Diseases, Bethesda.

Background: Pathway analyses can be used to determine how host and environmental factors contribute to asthma severity. **Objective:** To investigate pathways explaining asthma severity in inner-city children.

Methods: On the basis of medical evidence in the published literature, we developed a conceptual model to describe how 8 risk-factor domains (allergen sensitization, allergic inflammation, pulmonary physiology, stress, obesity, vitamin D, environmental tobacco smoke [ETS] exposure, and rhinitis severity) are linked to asthma severity. To estimate the relative magnitude and significance of hypothesized relationships among these domains and asthma severity, we applied a causal network analysis to test our model in an Inner-City Asthma Consortium study. Participants comprised 6- to 17-year-old children (n = 561) with asthma and rhinitis from 9 US inner cities who were evaluated every 2 months for 1 year. Asthma severity was measured by a longitudinal composite assessment of day and night symptoms, exacerbations, and controller usage.

Results: Our conceptual model explained 53.4% of the variance in asthma severity. An allergy pathway (linking allergen sensitization, allergic inflammation, pulmonary physiology, and rhinitis severity domains to asthma severity) and the ETS exposure pathway (linking ETS exposure and pulmonary physiology domains to asthma severity) exerted significant effects on asthma severity. Among the domains, pulmonary physiology and rhinitis severity had the largest significant standardized total effects on asthma severity (-0.51 and 0.48 , respectively), followed by ETS exposure (0.30) and allergic inflammation (0.22).

Although vitamin D had modest but significant indirect effects on asthma severity, its total effect was insignificant (0.01).

Conclusions: The standardized effect sizes generated by a causal network analysis quantify the relative contributions of different domains and can be used to prioritize interventions to address asthma severity. (J Allergy Clin Immunol 2016;138:1042-50.)

Key words: Asthma, children, inner-city, allergy, sensitization, inflammation, lung function, pulmonary physiology, rhinitis, environmental tobacco smoke exposure

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Inner-city children experience a high burden of asthma symptoms and morbidity despite guidelines-directed care.^{1,2} The National Institutes of Health/National Institute of Allergy and Infectious Diseases sponsored the Inner-City Asthma Consortium Asthma Phenotypes in the Inner City (APIC) study to investigate how host and environmental factors contribute to asthma severity among children carefully monitored while prospectively receiving optimal, guidelines-based care. Results of the primary objectives of the APIC study are presented in the articles titled “Asthma Phenotypes in the Inner City: Distinguishing Characteristics of Difficult-to-Control Asthma in Children”³ and “Asthma Phenotypes in Inner-City Children,”⁴ which appear in this issue of the journal.

On the basis of clinical and mechanistic evidence in the published literature that we have detailed in this article’s [Online Repository](http://www.jacionline.org) at www.jacionline.org, we developed a conceptual model to describe how 8 risk-factor domains (allergen sensitization, allergic inflammation, pulmonary physiology, stress, obesity, vitamin D, environmental tobacco smoke [ETS] exposure, and rhinitis severity) are linked to asthma severity (Fig 1). This model allows us to conceptualize which domains have a direct influence on asthma severity and/or an indirect influence on asthma severity by acting through another domain(s). We then tested this conceptual model in the APIC cohort and data set using a causal network analysis, a quantitative approach that is commonly used to describe the effects of genomics, metabolomics, or biochemical pathway influences on disease phenotypes but, to our knowledge, has never been applied in the context of

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evaluating the safety of inhaled corticosteroid + long-acting beta-agonist versus inhaled corticosteroid alone in children (ClinicalTrials.gov Identifier: NCT01462344). R. S. Gruchalla is employed by the Center for Biologics Evaluation and Research and has consultant arrangements with the Massachusetts Medical Society. M. Kattan has received a grant from the NIH-NIAID and is on the advisory board for Novartis Pharma. S. J. Teach has received grants from the NIH-NIAID, Novartis, Patient-Centered Outcomes Research Institute, Fight for Children Foundation, EJF Philanthropies, and the NIH-National Heart, Lung, and Blood Institute; has consultant arrangements with Novartis; and has received royalties from Up To Date. J. E. Gern has received grants from the NIH and GSK; has consultant arrangements with GSK, Genentech, Amgen, Novartis, Janssen, and Regeneron; has received payment for the development of educational presentations from Boehringer-Ingelheim; and has stock/stock options in 3V BioSciences. W. W. Busse has received a grant from the NIH-NIAID; has received partial study funding and provision of study drug and placebo from Novartis; is a member of the Data Safety Monitoring Boards for Boston Scientific and Circassia; is a member of the Study Oversight Committee for ICON; and has consultant arrangements with Novartis, GSK, Genentech, Roche, Pfizer, Merck, Boehringer-Ingelheim, Sanofi, AstraZeneca, Teva, Tekeda, Aerocrine, and 3M. The rest of the authors declare that they have no relevant conflicts of interest.

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