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Statins Enhance Clearance of Apoptotic Cells through Modulation of Rho-GTPases

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Clearance of apoptotic cells (efferocytosis) is a key modulator of the inflammatory response, and is dysregulated in several chronic inflammatory lung diseases, including chronic obstructive pulmonary disease and cystic fibrosis. Rho-GTPases are signaling molecules that differentially regulate efferocytosis: RhoA and its downstream effector Rho kinase inhibit efferocytosis, whereas Rac-1/Cdc42 stimulate it. Statins are 3-hydroxyl-3methylglutaryl coenzyme A (HMG-CoA) reductase inhibitors with broad antiinflammatory effects, largely based on their ability to block prenylation and membrane localization of Rho-GTPases, including RhoA. Therefore, we hypothesized that statins would enhance efferocytosis by inhibiting RhoA. In vitro, lovastatin increased efferocytosis by both human and murine macrophages. The positive effect of lovastatin on efferocytosis was prevented by mevalonate, farnesylpyrophosphate, and geranylgeranylpyrophosphate, and was mimicked by specific inhibitors of farnesyltransferase and geranylgeranyltransferase. These results suggested that lovastatin acted by inhibiting HMG-CoA reductase and subsequent downstream prenylation events. In support of this notion, lovastatin decreased membrane localization of RhoA, but had no effect on expression of macrophage efferocytosis receptors. To determine whether lovastatin enhanced efferocytosis in vivo, mice were challenged intratracheally with apoptotic thymocytes, in the presence and absence of lovastatin or the Rho kinase inhibitor (Y-27632), and clearance of these apoptotic thymocytes was assessed. Both lovastatin and Y-27632 enhanced efferocytosis in vivo. These findings suggest that statins enhance efferocytosis in vitro and in vivo, and may ultimately play

an important therapeutic role in diseases where efferocytosis is impaired and inflammation is dysregulated.

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Prostacyclin Synthase in Smokingrelated Lung Disease

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Background: Prostacyclin is a major product of the cyclooxygenase (COX) pathway with both potent vasodilatory and antimitogenic properties. Previous studies have demonstrated an increase in cyclooxygenase 2 (COX-2), prostaglandin E₂, cytosolic phosphoplipase A2 (cPLA₂), and vascular endothelial growth factor (VEGF), but decreased prostacyclin synthase (PGI₂S) expression within lung tumors. Lung-specific overexpression of PGI₂S has protective effects in both chemical- and tobacco smokeinduced murine models of lung tumorigenesis. Lastly, PGI₂S expression within lung tumors may correlate with survival. Here, we hypothesize that an imbalance in COX-2, PGE2, and PGI2S may be important to the pathogenesis of other smoking-related lung diseases such as chronic obstructive pulmonary disease and that tobacco smoke may alter PGI₂S expression. Design: We evaluated PGI₂S expression in both normal and emphysematous human lung tissue samples by: (1) immunohistochemistry, (2) quantitative PCR, (3) Western analysis, and (4) both 6-keto PGF1α (the stable metabolite of prostacyclin) and PGE₂ levels by ELISA. In addition, we treated primary human pulmonary microvascular endothelial cells (HPMVEC) with varying concentrations of cigarette smoke extract (CSE). Treated cells were assessed for PGI₂S, COX-2, VEGF, and cPLA₂ gene expression, and 6Keto-PGF1α and PGE₂ levels. Measurements and Results: In this study, we observed that human lung emphysema tissue exhibited lower PGI₂S expression in the endothelium than normal lung tissue. Furthermore, in HPMVEC, CSE suppresses PGI₂S gene expression while potently inducing COX-2, CPLA₂, and VEGF expression. Pretreatment with potent antioxidants did not alter this imbalance with the exception of N-acetylcysteine (NAC), which reversed the induction of COX-2 by CSE. Conclusions: The main finding of this investigation is the demonstration of decreased PGI₂S protein and mRNA in the lungs from patients with emphysema. CSE reduces the expression of the PGI₂S gene and increases the expression of VEGF, COX-2, and cPLA2 in HPMVEC. Decreased PGI₂S expression is likely multifactorial, including oxidant stress, loss of alveolar capillary endothelial cells, nitric oxide-related suppression, and altered transcriptional control. In vitro work by our laboratory has demonstrated that PGI₂S gene expression in lung carcinoma cell lines is partially controlled through methylation silencing. Our demonstration that PGI₂S is diminished in both human emphysema tissue and in vitro by CSE, as well as similar observations in carcinomas of the lung, suggest that this imbalance may be an early tobacco smoke-induced event and relevant to the pathogenesis of both diseases.

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IFN-γ-dependent DNA Injury and/ or Apoptosis Are Critical in Cigarette Smoke-induced Murine Emphysema

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There is mounting evidence that Tc1 immune responses contribute to the pathogenesis of pulmonary emphysema. This includes studies demonstrating increased number of IFN-y-producing inflammatory cells and increased expression of IFN-y target genes in lungs from patients with chronic obstructive pulmonary disease. Although cigarette smoking is a key risk factor for pulmonary emphysema, the roles of IFN-γ and apoptosis in these responses are poorly understood. We hypothesized that IFN-ydependent apoptosis plays an important role in the pathogenesis of cigarette smoke (CS)-induced murine emphysema. To test this hypothesis, we exposed male C57BL/6 wild-type and IFN- γ null (-/-) mice to the CS from 2R4 research cigarettes, 20 puffs/ session, two sessions/d, 5 d/wk for 6 mo. Emphysema was then assessed with the NIH Image Analysis program, which provided morphometric measurements of alveolar chord length. DNA injury and apoptosis-like cell death were evaluated with TUNEL assays. After 6 mo, the mean chord length was increased by 36.8% in CS-exposed mice compared with nonsmoking control