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⊗ Cystic Fibrosis Lung Disease and Immunometabolism Targeting the NLRP3 Inflammasome

The CFTR (cystic fibrosis transmembrane conductance regulator) protein regulates airway mucus viscosity and surface fluid pH (1). Defective CFTR leads to severe chronic airway infection and inflammation that destroys the structural support of bronchi. Correction of the basic defect by highly effective modulators of CFTR function is fueling the hope of a cure or effective control for people with cystic fibrosis (CF) (2). However, structural damage to CF airways is likely to persist even in those individuals whose CFTR function is restored. Furthermore, inflammatory mediators such as oxidants and proteases can suppress CFTR function (3, 4). If all individuals are to fully benefit from CFTR modulator therapies, we will need a better understanding of chronic inflammatory pathways associated with CF lung disease.

Chronic lung disease remains the leading cause of morbidity and mortality in CF (5). The cardinal feature of CF lung disease is the presence of bronchiectasis: floppy cystic airways bearing abundant purulent secretions. The appearance and severity of bronchiectasis is strongly associated with the abundance of airway secretion neutrophil elastase (6). One of the key cytokines driving neutrophilic inflammation in the CF lung is IL-1 β (7).

The regulation of IL-1 β synthesis and its activation involves a complex interplay of cellular danger sensors, metabolic reprogramming, and post-transcriptional protein processing (8). The study published in this issue of the *Journal* by McElvaney and colleagues (pp. 1381–1391) documents these interactions in CF lung and peripheral blood neutrophils and provides key information pointing to potential new antiinflammatory strategies (9). These immunometabolic changes in neutrophils are independent of CFTR function.

Neutrophil immunometabolism is altered by the CF airway environmental signals known as damage-associated molecular

patterns and pathogen-associated molecular patterns, particularly bacterial LPS (Figure 1). Once engaged by LPS, the neutrophil increases transcription of the key metabolic protein, PKM2 (pyruvate kinase M2 isoform). PKM2 undergoes post-translational modification, forming a transcriptional complex with PHD3 (prolyl hydroxylase-3), P300, and HIF1 (hypoxia inducible factor-1) to induce several glycolytic proteins (10). Neutrophil metabolism of glucose is thus reprogrammed toward glycolysis (known as the Warburg effect), rather than oxidative phosphorylation through the Krebs cycle (tricarboxylic acid cycle and citric acid cycle). The HIF1 complex binds to the HRE (HIF1-responsive element) and increases neutrophil PKM2, HIF1, and pro-IL-1 β synthesis. Increased glycolysis was confirmed in the CF neutrophil cytosol by high PKM2, lactate, and succinate as well as low pH.

The current study reveals that the amount of circulating LPS in CF blood is sufficient to increase neutrophil PKM2 and pro-IL-1 β but not to activate pro-IL-1 β . In contrast, the amount of LPS in the CF lung is sufficient to assemble the NLRP3 inflammasome and initiate caspase-1-dependent cleavage and activation of pro-IL-1 β . Interestingly, the levels of IL-1 β in CF airway secretions strongly correlate with neutrophil burden and patient outcomes. The addition of an NLRP3 inflammasome diarylsulfonylurea-containing inhibitor (MCC950), blocked the LPS-dependent IL-1 β production both *in vitro* and in animal models of LPS exposure and *Pseudomonas aeruginosa* airway infection.

The NLRP3 inflammasome is a protein complex comprising an intracellular sensor (Nod-1-like receptor), procaspase-1, and the inflammasome adaptor protein ASC, an activating adaptor for procaspase-1 (11). Oligomerization of the NLRP3 inflammasome into a functional complex activating IL-1 β occurs in the presence of mitochondria-derived reactive oxygen species, phagolysosomal destabilization, or bacterial cytotoxins that induce potassium efflux (12). Although the exact mechanism(s) by which the NLRP3 inflammasome is assembled remains unknown, the importance of this complex and its effects on IL-1 β activation in CF lung disease is strongly supported by the results of this study as well as the work of previous investigators (13).

Inhibition of the IL-1 β and/or the NLRP3 inflammasome represents an attractive goal in the quest to find new efficient strategies to control CF inflammation and possibly improve pathogen clearance. Several molecules capable of preventing the

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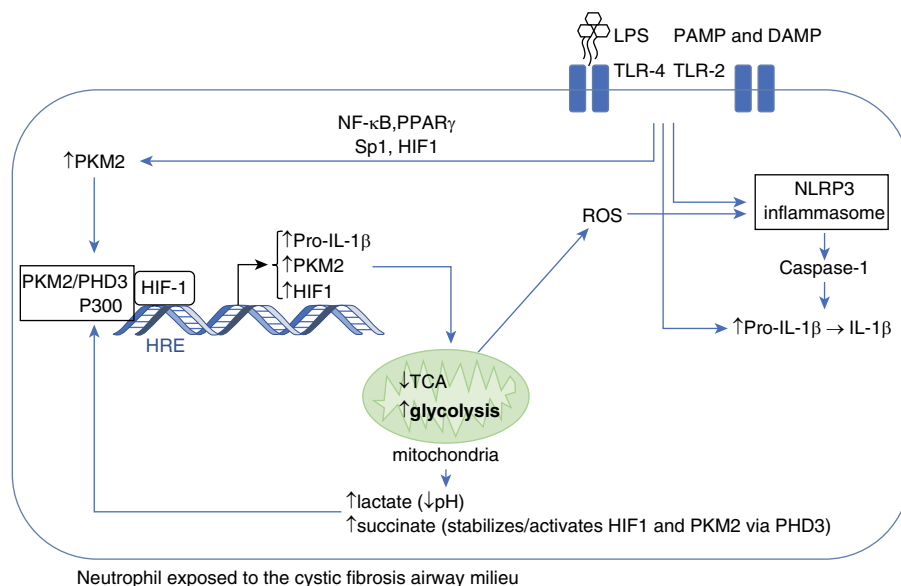


Figure 1. Neutrophil immunometabolism in cystic fibrosis. On exposure to LPS in the cystic fibrosis airway, the neutrophil increases its cytoplasmic levels of the M2 isoform of PKM2 (pyruvate kinase) and pro-IL-1 β . Simultaneously, the NLRP3 inflammasome is activated through several mechanisms. PKM2 in association with PHD3 (prolyl hydroxylase-3) is stabilized and in conjunction with the coactivator P300 facilitates HIF1 (hypoxia inducible factor-1)-dependent transcription that further increases the synthesis of pro-IL-1 β , PKM2, and HIF1, while reprogramming mitochondria to deviate glucose from the tricarboxylic acid cycle (TCA) to the glycolytic pathway. Increased glycolysis results in production of lactate, acidification of the cytoplasm, and accumulation of succinate, which helps stabilize the HIF1 complex. Mitochondria release reactive oxygen species (ROS) that further activate the NLRP3 inflammasome, which converts pro-IL-1 β to active IL-1 β through caspase-1-mediated cleavage. DAMP = damage-associated molecular patterns; NF- κ B = nuclear factor- κ B; PAMP = pathogen-associated molecular patterns; PPAR γ = peroxisome proliferator-activated receptor- γ ; TLR = toll-like receptors.

metabolic reprogramming toward glycolysis exist, some of which could be quickly amenable to clinical trials (14). The current study reveals that blocking glycolysis with 2-deoxyglucose decreases neutrophil PKM2, a key factor that may be linked to increased NLRP3 inflammasome activity. Whether inhibition of glycolysis alone will be sufficient to control CF neutrophilic lung inflammation remains unknown. Alternatively, anakinra, an inhibitor of the IL-1 β receptor in use for many years in the treatment of rheumatoid arthritis, has a favorable safety profile and has shown benefit in murine models of CF lung disease (13). The antiinflammatory effects of MCC950 in this study further suggest that the NLRP3 inflammasome and its resultant release of IL-1 β represent key targets worthy of further investigation.

The novelty of linking inflammation with metabolic reprogramming in the neutrophil certainly provides an exciting advancement in our knowledge of CF airway inflammation. However, further work is needed before translation of these findings into clinical studies. The current study includes a short-term intervention of NLRP3 suppression in murine models of LPS exposure and lung infection. The degree to which the bacterial pathogen lung burden was suppressed was modest and was measured at an early time point. Lung disease in CF is a chronic lifelong condition. There is reason to believe that inhibition of glycolytic pathways or of IL-1 β over several years could be associated with significant adverse events. The chronic use of IL-1 β in rheumatoid arthritis can be associated with a mildly increased risk of serious infection (15), a fact that may give pause to clinicians treating patients who already are at risk of life-threatening infectious complications. Effective inhibition of glycolytic pathways

could also have unwanted effects on tissues known for their high glucose uptake and metabolism, such as the heart.

In summary, McElvaney and colleagues have clearly demonstrated that the neutrophil undergoes marked changes in immunometabolism during its encounter with the CF lung environment and that these changes have profound effects on the cell's capacity to enhance IL-1 β -dependent inflammation, a cytokine strongly correlated with parameters of poor patient outcome (9). The study highlights the importance of neutrophil metabolic reprogramming characterized by the Warburg effect and NLRP3 inflammasome assembly, both of which represent potential areas of novel therapeutic strategies for CF lung disease. ■

Author disclosures are available with the text of this article at www.atsjournals.org.

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⊗ Promotion of a Protease–Antiprotease Imbalance in the Airways through Chronic Vaping

There is ongoing controversy in regard to the safety of electronic cigarettes (e-cigarettes) and vaping. Although originally promoted to help facilitate smoking cessation, a number of significant concerns have been highlighted, not least the uptake of these devices by previously nonsmoking youths and their high transfer to traditional smoking as a result of nicotine addiction (1).

Electronic nicotine delivery systems are aerosol-generating devices that heat, not burn, a solution containing a complex mixture of solvents and flavoring (a number of which have known toxicity) in addition to nicotine, the final composition in the aerosol of which is determined by temperature (2). Of note, the fine particles delivered by e-cigarettes are similar in size and concentration to tobacco smoke, and although the composition differs, the pattern of particle deposition in the lungs is similar (3). A number of studies now report e-cigarette exposure to be associated with airway irritation and inflammation, as well as mucus hypersecretion, and have been linked to an exacerbation of symptoms in those with chronic airways diseases such as cystic fibrosis, asthma, and chronic obstructive pulmonary disease (4).

Proteases and their inhibitors play pivotal regulatory roles in most physiological processes required for life. They act as nature's molecular scissors, processing other biological molecules leading to the synthesis, activation, or degradation of functionally important peptides and proteins, and play a vital role in tissue remodeling and wound healing. However, when the normally exquisite control of their action is lost, proteases can be key triggers or amplifiers of

many important human diseases such as cancer, cardiovascular disease, rheumatoid arthritis, sepsis, and neurological disorders including Alzheimer disease and multiple sclerosis, with many proteases well-recognized as potential biomarkers of disease and/or therapeutic targets (5–7). In chronic airways diseases such as cystic fibrosis, chronic obstructive pulmonary disease, and bronchiectasis, a protease–antiprotease imbalance has long been associated with tissue injury and disease progression. Aberrant proteolytic activity resulting from high levels of neutrophil elastase (NE), in particular, is widely associated with episodes of acute exacerbation and pulmonary decline (8–10).

E-cigarette vapor extract has been shown to stimulate the release of MMP-9 (matrix metalloprotease-9) and interleukin 8 (CXCL8) from isolated neutrophils, as well as increase in NE and MMP-9 activity (11). MMP-9 and CXCL8 release caused by e-cigarette vapor extract prepared from different e-cigarette brands were found to be similar to, or in excess of, a cigarette smoke extract response. In addition, MMP-9 and CXCL8 was increased after exposure to e-cigarette vapor extract with and without nicotine, suggesting the involvement of other proinflammatory constituents.

In this issue of the *Journal*, Ghosh and colleagues (pp. 1392–1401) investigate the effect of chronic e-cigarette use on the protease–antiprotease balance in the airways of vapers (12). The study recruited never-smokers, current tobacco smokers, and e-cigarette users (vapers), with the latter group including both never-smokers and former tobacco smokers. Protease levels were quantified in BAL samples, as well as from immune cells stimulated with e-liquid components.

Protease levels were measured using Western blotting and activity by hydrolysis of peptide-based substrates (\pm protease class inhibitors), with gelatin zymography also used to assess the activity of MMP-2 and MMP-9. Importantly, serum nicotine, cotinine, and

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