



## EDITORIAL

# Cystic fibrosis: to ion transport and beyond

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The measurement of sweat electrolytes successfully diagnoses all but a tiny minority of patients with cystic fibrosis (CF). The discovery of a gene for CF, encoding the cystic fibrosis transmembrane conductance regulator (CFTR), has led to the understanding that CFTR is a chloride channel (hence the utility of the measurement of sweat electrolytes) and also regulates other ion channels, in particular the sodium transporter, ENaC. Measurement of transepithelial potential differences suggested that ENaC overactivity is pivotal in CF lung disease [1]. The CF mouse was found to have a poor lower airway phenotype for human CF lung disease, but the ENaC  $\beta$ -subunit overexpressing mouse was a much better model for the devastating human CF lower airway disease [2]. Indeed, CFTR dysfunction, perhaps induced by tobacco smoke exposure, has been implicated in chronic obstructive pulmonary disease (COPD) [3]. After some controversy, a mechanism linking ion transport to CF lung disease was proposed [4]; the airway surface liquid (ASL) height was found to be equal in CF and normal airway cell monolayers if they were subjected to phasic contraction and relaxation, but in the presence of infection with respiratory syncytial virus, CF monolayers had reduced ASL height, and thus impaired mucociliary clearance [5], presumably accounting for the susceptibility to airway infection, inflammation and ultimately death by respiratory failure. So, CF and spin-off CFTR diseases, such as COPD are diseases of ion transport.

If not wrong, this view is perhaps over-simplistic. CFTR is now known to be much more than just an ion channel, a hole in the cell membrane with a bath plug-like structure to block it. CFTR has numerous other functions and a multiplicity of interactions with other proteins [6–8]. It is probably naïve to believe that absence or dysfunction of CFTR only affects ion transport. The model that CFTR dysfunction leads to ion transport problems which lead to infection, inflammation and eventual lung destruction, has already been challenged. Prospective, longitudinal physiological studies have established that infants with CF show evidence of airflow obstruction, independent of any clinical evidence of respiratory infection [9, 10], and this persists into school age [11–13]. Bronchoscopic studies have shown that some aspects of structural airway disease, in particular reticular basement membrane thickening, are independent of infection and inflammation [14]. It has been suggested that CFTR dysfunction may have a direct effect on airway smooth muscle [15]. Studies in CF mice have shown

that there is thickening of the nasal respiratory epithelium and aggregations of lymphoid tissue in the CF mouse nose, even in the absence of any evidence of infection or excessive inflammation [16]. Taken together, these studies suggest there is a component of CF airway disease that is independent of downstream infection and inflammation, and which is very difficult to tie in to deficits in ion transport.

In this issue of the *European Respiratory Journal*, DIF *et al.* [17] report data that confirms this view. Their starting point was that CF mice had increased mucus production and MUC5AC expression (although it should be noted that this point is controversial) [18]. They performed an elegant series of experiments which convincingly show that cytosolic phospholipase A2 $\alpha$  (cPLA2 $\alpha$ ) interacts with CFTR to modulate mucus production, which, critically, is independent of chloride channel function. Thus CFTR<sup>-/-</sup> mice had increased mucus production and MUC5AC expression, exacerbated by *Pseudomonas aeruginosa* infection, but present even in uninfected mice. They showed that mucus production was induced by instillation of cPLA2 $\alpha$ , and abolished in the absence of cPLA2 $\alpha$  activity, whether due to pharmacological inhibition or by cPLA2 $\alpha$  null mutation. Reduction of CFTR expression led to increased cPLA2 $\alpha$  activity and, as a downstream effect, a secondary increase in MUC5AC expression. Crucially, cPLA2 $\alpha$  appears to have no relevance whatsoever to the excessive neutrophilic inflammation characteristic of the CF airway. Furthermore, the CFTR chloride transport inhibitor CFTR<sub>inh</sub>-172 did not lead to an increase in MUC5AC expression, underscoring the dissociation of chloride transport activity from airway mucus production.

Further evidence suggesting there is more to airway disease than just ion transport has come from the recent study of the peroxisome proliferator-activated receptor- $\gamma$  (PPAR- $\gamma$ ) pathway [19]. These investigators report data that also clearly dissociate abnormal chloride and sodium ion transport from the increased mucus production so characteristic of CF (originally called “mucoviscidosis”). The binding of the lipid 15-keto-prostaglandin-2 (15-keto-PGE<sub>2</sub>) to PPAR- $\gamma$  regulates mucus production, and lipidomic studies suggest that dysregulation of this pathway due to reduced 15-keto-PGE<sub>2</sub> (amongst other lipids) may underlie the excessive mucus production in the airways and colons of CFTR<sup>-/-</sup> mice. The synthetic PPAR- $\gamma$  antagonist rosiglitazone reversed around a quarter of the genes up- and down-regulated in CFTR<sup>-/-</sup> mice compared with wild type. Cells treated with CFTR<sub>inh</sub>-172, or derived from CFTR<sup>-/-</sup> mice, showed the expected defect in chloride transport, but this was not reversed by rosiglitazone. What rosiglitazone did do was increase bicarbonate secretion *via* the genes encoding carbonic anhydrases 4 and 6. Unlike the

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findings in the cPLA2 $\alpha$  pathway, PPAR- $\gamma$  is implicated in inflammation; PPAR- $\gamma$  agonists reduce the release of pro-inflammatory cytokines from airway epithelial cells infected with *P. aeruginosa* [20]. Taken together, these two manuscripts clearly establish that CFTR dysfunction exerts adverse effects on the airway independent of sodium and chloride transport. It is probably unsurprising that a multifunctional molecule as complex as CFTR, and an airway disease so multifaceted as that seen in CF, do not bear a simple relationship such as "function X causes disease Y".

Does this mean that efforts to correct ion transport are doomed to be fruitless, and that the use of ion transport as an end-point in CF clinical trials is a waste of time? This would be a radical, and almost certainly incorrect, conclusion. The evidence for the involvement of defective ion transport, in particular ENaC overactivity, in CF lung disease is compelling. Correction of ion transport defects is likely to bring substantial benefits. But before we assume that death from CF lung disease will itself be dead and buried when ion transport is normalised, we should remember that the other functions of CFTR may also need attention; correcting one CFTR function does not mean that everything in the garden is lovely. Indeed, this has been seen in gene therapy trials, in which some correction of chloride transport has been achieved, but correcting sodium hyper-absorption remains uniformly elusive. In the future, correction of aspects of CFTR dysfunction, by gene or other specific molecular therapies, may well lead to CF airway disease ceasing to be a killer of young adults. However, residual CFTR dysfunction may become a factor predisposing to early COPD in the nonsmoker, so there is no excuse for complacency.

#### STATEMENT OF INTEREST

None declared.

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