Abnormalities of airway epithelial function and the implications of the discovery of the cystic fibrosis gene

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Of all the functional abnormalities present in cystic fibrosis, those affecting the lung are the most serious. In 1989 research workers in Canada and the United States reported the structure of the gene responsible for 70% of cystic fibrosis among white people. I shall explore what is known about abnormal lung function in cystic fibrosis at the cellular and molecular level and discuss how the recently discovered genetic information may be related to the cellular defects. Although it is too early to speak of a treatment for cystic fibrosis, the increased understanding of underlying mechanisms suggests ways in which treatment may be approached in the future.

By means of a painstaking combination of chromosome walking and jumping and complementary DNA (cDNA) hybridisation over half a million base pairs on the long arm of chromosome 7 band q 31 have been sequenced. The putative protein coded for by the cDNA for the cystic fibrosis locus consists of 1480 amino acids. The tri-nucleotide codon for the 508th amino acid from the N terminus (phenylalanine) was found to be deleted in 70% of patients with cystic fibrosis. On the basis of analogy and homology a tentative structure for the protein coded for by the gene has been deduced (figure 1). The protein has been named the cystic fibrosis transmembrane regulator.

This protein has two related motifs, each consisting of six membrane spanning regions joined by a long polypeptide chain. It is proposed that very little (5%) of the protein is exposed externally and that some 80% is in the cytosol. Two features of the cystolic domain are important. The first is a highly charged R domain with 16 potential sites for phosphorylation and the second two nucleotide binding folds, which may bind or even hydrolyse ATP. The phenylalanine deletion (ΔF508) occurs in the first nucleotide binding fold.

Other proteins with similar structures and a fair degree of homology are already known, principally the P glycoproteins or multiple drug resistance (MDR) proteins. These proteins are responsible for the energy dependent efflux of cytotoxic drugs from cells. Increased expression of multiple drug resistance proteins after exposure to cytotoxic drugs may be a major cause of drug resistance in cancer chemotherapy.

Figure 1. Hypothetical structure of the cystic fibrosis transmembrane regulator. Two sets of six membrane spanning domains are joined by a long polypeptide chain. Two nucleotide binding folds (NBF) bind ATP. In cystic fibrosis the missing phenylalanine is in the nucleotide binding fold nearest to the N terminus. The R domain contains 16 potential phosphorylation sites. Note that very little of the protein is exposed to the external (upper) surface of the cell.

Transepithelial ion transport in the respiratory system

What goes awry in the respiratory system in cystic fibrosis and how is this related to the presence of the cystic fibrosis transmembrane regulator protein with phenylalanine deleted at position 508? It is obvious, but worth restating, that biological systems have no way of pumping water. Water can be made to move across biological barriers only by osmosis—provided, of course, that the barriers are water permeable. Movement of fluid therefore depends on solute transport, usually of inorganic ions. In the lungs surface liquid elaborated in the alveoli and small airways is moved by ciliary action to more proximal regions. As fluid does not accumulate in the proximal regions even though there is a 4000 fold reduction in total volume and a comparable reduction in surface area, fluid must be reabsorbed as a consequence of ion transport. These same processes are vitally important at birth for converting fluid filled lungs to respiratory organs.

How do cells lining the respiratory system,
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Abnormalities of epithelial ion transport in cystic fibrosis

The first insightful observation in relation to the transport function in cystic fibrosis airway epithelium was that the potential difference across the nasal mucosa and bronchi of patients with cystic fibrosis was around twice that of subjects not suffering from cystic fibrosis. On its own this result is difficult to interpret because either increased sodium absorption or reduced chloride conductance at the apical surface would give this result.

When amiloride is superfused into the apical surface of nasal epithelia in vivo the potential...
fibrosis cells than in non-cystic fibrosis cells. By application of various agents that activate adenylate cyclase (beta agonists, forskolin) to monolayer cultures of airway epithelial cells chloride secretion was stimulated in normal but not cystic fibrosis cells. In cystic fibrosis monolayers beta agonists further enhanced sodium absorptive processes. These differences could not be explained by differences in the generation of cAMP or in the amount of cAMP dependent protein kinases in the tissue.

Bradykinin and A23187, both of which increase intracellular calcium concentration (Ca	extsuperscript{2+}), are able to elicit transient chloride secretory responses in cystic fibrosis epithelia. It is, however, possible that raised Ca	extsuperscript{2+} activates basolateral K	extsuperscript{+} channels, leading to hyperpolarisation of the apical membrane, thus increasing the electrochemical potential for efflux, even though the membrane has a low chloride conductance. Figure 3 provides a summary of the comparisons in transporting properties between cystic fibrosis and normal respiratory epithelium, plus the effects of different types of secretagogues.

Possible consequences of transporting abnormalities in cystic fibrosis
Mucociliary clearance depends on the functioning of cilia within the sol phase lining the respiratory passages. Not only is fluid moved toward the proximal airways but mucus and entrapped foreign particles are moved to a position from which they can be eliminated from the lung. Excessive solute transport out of the airways, with consequent fluid absorption, may reduced the sol phase, depositing thick, viscid mucus on the cilia. Mucus deposits are difficult to dislodge and provide a breeding ground for Staphylococcus aureus and Pseudomonas aeruginosa. Although this is a reasonable hypothesis there is little definitive evidence to support it. The reason is a technical one—it is difficult to sample fluid from the airways without evaporative losses, making accurate determination of composition very difficult.

The chemical composition of secreted mucus is known to be altered in cystic fibrosis, with increased sulphation of high molecular weight glycoconjugates. As cystic fibrosis is a single gene defect it becomes increasingly difficult to explain how properties as diverse as mucus composition and chloride conductance are related to a single abnormal form of the cystic fibrosis transmembrane regulator, though abnormal chloride transport may increase the concentration of intracellular sulphate and increase substrate driven sulphation of glycoproteins.

Molecular correlates of abnormal transporting properties
Patch clamping is a relatively new technique that allows the functional activity of single ion channels to be examined. Theoretically the concept is straightforward. First devise a way

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**Figure 3** Summary of the differences between airway epithelium in normal and cystic fibrosis tissues. In cystic fibrosis sodium absorption (Na) is exaggerated and chloride secretion (Cl) is blocked, even in the presence of secretagogues that increase cyclic AMP (cAMP). In normal epithelium secretagogues increase chloride secretion whereas in cystic fibrosis sodium absorption is enhanced. Excessive dehydration may remove the sol phase, depositing mucus flakes with adherent bacteria on the cilia.
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of isolating a single ion channel so that its activity can be monitored, without interference from the thousands of other ion channels present in a cell membrane. Secondly, find a way of detecting whether the channel is open or closed. This is done by monitoring the flow of ions through the channel with sensitive current amplifiers. Ion channels allow millions of ions to pass each second, giving rise to currents in the picoampere (10^{-12} A) range. By measuring the relation between the current and the voltage across the channel and taking account of the ion concentration either side of the membrane you can assess channel selectivity. An all important factor is the reversal potential (the potential at which the current through the channel reverses its direction); application of the Goldman-Hodgkin-Katz equation and comparison of actual and theoretical values allow the channel to be designated as selective for chloride, sodium, potassium, or calcium. To isolate a single ion channel a fire polished glass electrode with a tip diameter of a few microns is gently pressed against the surface of a cell. A self-sealing process occurs such that the resistance between the conducting fluid in the electrode and the bathing fluid outside is very large (gigaohm seal). The patch so isolated will contain a few ion channels, or preferably a single one. Recordings are made of the currents flowing across the patch either in the cell attached mode or by pulling the patch off the cell.

Figure 4 shows a patch clamp record from a chloride ion channel from a human cystic fibrosis sweat gland cell. Diagrammatic representation of major findings in cystic fibrosis and airway epithelial cells is given in figure 5, which shows idealised diagrams of patch clamp recordings. Figures 5a and 5b shows the usual features seen with patch clamping. The current flowing through a particular type of ion channel is constant at a given voltage—that is, the channel conductance is characteristic of the channel, and channels are either open or closed. If two channels are present in a patch and are open simultaneously then the current flowing is twice that of a single channel. By contrast, the duration of the open and closed periods varies even for a single channel. In the simplest systems analysis of patch clamp records shows that open and closed time histograms are fitted by single exponentials, suggesting that a single process is responsible for transitions from open to closed.

Two groups have reported their findings on chloride channels in airway epithelial cells using the patch clamp technique. In the cell attached configuration chloride channels were rarely seen unless the cell was activated by agents that increased cAMP. After stimulation there was an increased probability of chloride channel opening in normal cells (fig 5c) but not in cystic fibrosis cells (fig 5d). Again with the cell attached configuration, the calcium ionophore A23187 was able to activate chloride channels in both normal and cystic fibrosis cells. Isolated patches from cells activated by cAMP generating secretagogues showed chloride channel activity (fig 5e.i), whereas patches from similarly treated cystic fibrosis cells were electrically silent (fig 5e.ii). Chloride channels could be activated in cystic fibrosis patches either by strong depolarisation or by exposure of the cytosolic face to calcium ions (fig 5e.iii). The effects of these last two manoeuvres are not
immediate or necessarily reversible and it is not clear how activation is achieved. Kunzelmann et al. have shown that if isolated patches are formed at 37°C rather than at room temperature there is immediate activation of chloride channels in cystic fibrosis cells without the need to apply strong depolarisation. He has argued that regulation of the chloride channel is defective rather than the channel itself. Possibly the cytosol in cystic fibrosis contains a tonic inhibitor of the chloride channel that is removed once an inside out patch is formed. Finally, as shown in figure 5f, when apical membrane patches are held at room temperature and at a voltage inhibiting channel activation, addition of ATP and the catalytic subunit of protein kinase A causes channel activation in normal but not cystic fibrosis membranes. Thus there appears to be a phosphorylation defect at a stage later than either cAMP generation or the activation of protein kinase A. There are two other findings with a bearing on this problem. Firstly, the patch clamp findings have now been confirmed in an immortalised, virally transformed epithelial line developed from cystic fibrosis airway cells. This establishes that the distinctive cystic fibrosis phenotype is preserved in cells after many generations of culture in vitro and therefore cannot be related to the infected conditions in cystic fibrosis airways. Secondly, apical membrane chloride channels incorporated into lipid bilayers show properties similar to those in normal membranes, but undergo a rapid rundown. Activity can be restored by addition of the phosphorylating “cocktail” including the catalytic subunit of protein kinase A, indicating that a phosphorylation event is concerned in the functioning of airway chloride channels. Protein kinase C can also activate chloride channels in airway epithelial cells, but again regulation by this Kinase is defective in cystic fibrosis. Whatever method has been used to activate cystic fibrosis chloride channels in vitro their properties have proved to be identical to those of normal channels—that is, conductance of around 50 pS at 0 mV, an outwardly rectifying current-voltage relationship, and a Cl⁻ to Na⁺ selectivity ratio of 10:1.

Far less is known about sodium channel mechanisms in airway cells. The mean channel lifetime for sodium channels in cystic fibrosis cells is twice that of normal channels, providing a correlate for the increased sodium reabsorption in cystic fibrosis. It has also been shown that the macroscopic Kd for amiloride is increased and there is an altered relation between sodium concentration and sodium transport in cystic fibrosis epithelium. These latter studies were made with sweat duct epithelium, however, not airway epithelium. Nevertheless, a picture is emerging of altered molecular characteristics affecting both chloride and sodium channels in cystic fibrosis. From the patch clamp data it is difficult to argue that the sodium channel effects are a result of aberrant chloride transport, yet both alterations in channel properties are a consequence of a single gene defect.

**Future Therapeutic Strategies**

An experiment that many must now be attempting is to transfect various cells with the cDNA for cystic fibrosis transmembrane regulator. This approach may show whether or not the cystic fibrosis transmembrane regulator is the chloride channel. This seems a little unlikely as altered sodium channel characteristics are retained in isolated patches. More probably it will be shown that transfection of cystic fibrosis cells with the cystic fibrosis transmembrane regulator cDNA will correct the ion channel abnormalities. In the long term transfection of airway epithelium in patients with cDNA in a retrovirus vector may be possible. Insufflation of the appropriate material into the respiratory tract may be easily arranged and considerable improvement might be envisaged even from transfection of only a fraction of the affected cells. Major obstacles need to be overcome before ethical permission is likely to be given, not the least of which is the possibility of switching on cellular oncogenes.

An alternative approach is to correct the transport irregularities with agents active from the apical surface. Nebulisation provides a route of administration that is likely to confine the pharmacological effect to the airways. Amiloride, an epithelial sodium channel blocker, is an obvious candidate and has been tried. Fourteen volunteer adults with cystic fibrosis took part in a study in which amiloride solution or control vehicle was administered by nebuliser in a double blind crossover study with two 25 week treatment periods. The mean (SEM) loss in forced vital capacity was reduced significantly from 3.4 (1.1) ml/day with vehicle to 1.4 (0.7) ml/day with amiloride. Furthermore, the rheological properties of the secretions returned to virtually normal values during amiloride administration, though bacterial contamination was unmodified. This trial raises the possibility that prophylactic treatment for young patients with cystic fibrosis before acute respiratory symptoms have developed may be worthwhile. Analogues with greater potency or a longer duration—for example, benzamil—might be more useful.

As we have seen, evidence suggests that, at least in airway epithelia, chloride secretion fails in response to secretagogues working via cAMP whereas agents that increase Ca²⁺ are still effective. Simultaneous inhibition of sodium absorption and stimulation of chloride secretion may prove even more effective than the former alone. There is an added incentive to develop this strategy because when amiloride blocks the apical sodium channels in airway epithelia a profound hyperpolarisation of the apical membrane results, exactly what is required to promote chloride exit from the cell provided that apical chloride channels can open. Thus a small increase in chloride conductance may have an amplified effect on chloride secretion in the presence of amiloride. Several ways of increasing chloride conductance may be envisaged, via novel agents that raise Ca²⁺ or development of chloride openers working independently of normal regulatory mechanisms. Agents of the former
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type are known (for example, thapsigargin) but are only experimental tools. Bradykinin promotes chloride secretion in many types of epithelia and does not require cAMP for its action. Bradykinin receptors are usually located basolaterally but in airways they are located apically. Bradykinin, however, also causes bronchoconstriction, clearly undesirable in cystic fibrosis, but is rapidly metabolized by tissue peptidases. Some thought should be given to developing analogues with appropriate pharmacodynamic and pharmacokinetic properties that may promote chloride secretion in airway epithelium.

In view of the similarity between the cystic fibrosis transmembrane regulator and multiple drug resistance proteins mentioned earlier we may ask whether the cystic fibrosis transmembrane regulator performs a similar function, expelling unwanted endogenous substances from cells that otherwise alter the functioning of other systems, such as the behaviour of sodium and chloride channels. The number of endogenous materials which, in excess, might have this effect is obviously large but the possibility that a tonic inhibitor of the chloride channel exists is supported by the immediate activation of these channels when the patches are removed from cells at 37°C. Moreover, the volume of distribution of many drugs is increased in cystic fibrosis, indicating that they can penetrate spaces from which they are normally excluded. Is this too a manifestation of an altered function of the cystic fibrosis transmembrane regulator? New information about the cystic fibrosis transmembrane regulator may promote questions where there is already a body of clinically relevant information containing important clues. For example, the immunosuppressive drug cyclosporin has been shown to label multiple drug resistance proteins and possibly the cystic fibrosis transmembrane regulator is also labelled; if it is, is it its function altered? This could be studied in patients receiving transplants and cyclosporin to see whether they show any membrane characteristics of cystic fibrosis and whether these disappear as cyclosporin dosage is reduced. Multiple drug resistance proteins are increased in cells, particularly epithelial cells, when they are exposed to natural cytotoxic drugs, such as vinblastine and vincristine. If the cystic fibrosis transmembrane regulator can also be induced, a search for non-toxic inducers may lead to a useful treatment as cystic fibrosis may be a missense mutation, where activity of the abnormal cystic fibrosis transmembrane regulator is not destroyed but impaired.

The cystic fibrosis transmembrane regulator with phenylalanine deleted at position 508 accounts for only 70% of mutations. Other haplotypes, accounting for the other 30% of mutations, are likely to be discovered soon. This information will provide further clues about which structural features of the cystic fibrosis transmembrane regulator are essential for function.

Summary

Details of ion transporting abnormalities in cystic fibrosis airway epithelium are now known. The central hypothesis, that excessive drying of the airway surfaces is a primary event that leads to all the manifestations of the respiratory insufficiency in cystic fibrosis, is not proved. The hypothesis is, however, consistent with the known transporting abnormalities and is strengthened by the modest clinical improvement produced by a strategy designed to correct the transporting abnormalities. The discovery of the cystic fibrosis gene, together with the presumed structure of the protein product, provides a focal point that must eventually connect the functional abnormalities with the genetic defect. The cellular function of the cystic fibrosis transmembrane regulator must now be the major target in research on cystic fibrosis. Strategies for treatment based on known cellular and molecular abnormalities are beginning to emerge but will be undoubtedly more focused once the responsibility of the cystic fibrosis transmembrane regulator is known.

Addendum

Since this review was completed the prediction made under “Future therapeutic strategies” has been realised. Complementation studies have shown that the introduction of plasmids containing cystic fibrosis transmembrane regulator cDNA into cultured cystic fibrosis cells does indeed restore a chloride conductance response to cAMP. Additionally, it now appears that the cystic fibrosis transmembrane regulator fails to be glycosylated in cystic fibrosis cells, being retained in the golgi and not transferred to the cell membrane. A plasma membrane location of the cystic fibrosis transmembrane regulator is consistent either with an MDR like function or with its being a direct regulator of ion channel activity.

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