

Protein kinase-independent activation of CFTR by phosphatidylinositol phosphates

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The cystic fibrosis transmembrane conductance regulator (CFTR) is a chloride channel that is expressed in many epithelia and in the heart. Phosphorylation of CFTR by protein kinases is thought to be an absolute prerequisite for the opening of CFTR channels. In addition, nucleoside triphosphates were shown to regulate the opening of phosphorylated CFTR. Here, we report that phosphatidylinositol 4,5-bisphosphate (PIP2) activates human CFTR, resulting in ATP responsiveness of PIP₂-treated CFTR. PIP₂ alone is not sufficient to open CFTR, but ATP opens nonphosphorylated CFTR after application of PIP₂. The effect of PIP₂ is independent of protein kinases, as PIP₂ activates CFTR in the complete absence of Mg. Phosphatidylinositol and phosphatidylinositol monophosphate activate CFTR less efficiently than PIP2. PIP2 application to phosphorylated CFTR may inhibit the CFTR chloride current. We suggest that regulation of CFTR by PIP₂ is a previously unrecognized, alternative mechanism to control chloride conductance.

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INTRODUCTION

The cystic fibrosis transmembrane conductance regulator (CFTR), the gene defective in the autosomal recessive disease cystic fibrosis (CF), was first cloned and sequenced in 1989 (Riordan *et al*, 1989). CFTR is expressed in the heart (Nagel *et al*, 1992) and in different exocrine glands, for example, the pancreas, airways, gastrointestinal tract and sweat glands (Riordan, 1993; Quinton, 1999; Sheppard & Welsh, 1999). Therefore, abnormalities in CFTR interfere with vital exocrine and absorptive functions in CF patients (Quinton, 1999; Sheppard & Welsh, 1999). Although its amino-acid sequence placed it into the family of ABC transporters (Riordan *et al*, 1989; Hyde *et al*, 1990), most of which mediate active transport across membranes, CFTR functions as an anion channel enabling the passage of chloride or other anions along their electrochemical gradient. CFTR is subject to complex regulation that requires interaction with nucleoside triphosphates

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and phosphorylation by protein kinases (Anderson *et al*, 1991; Nagel *et al*, 1992; Riordan, 1993; Hwang *et al*, 1994; Gadsby *et al*, 1995; Gadsby & Nairn, 1999; Nagel, 1999; Sheppard & Welsh, 1999).

CFTR contains multiple sites of phosphorylation by cyclic AMP (cAMP)-dependent protein kinase (PKA) and protein kinase C (PKC) (Riordan *et al*, 1989; Riordan, 1993; Gadsby & Nairn, 1999). It was also shown that the membrane-associated cGMP-dependent protein kinase isoform II (French *et al*, 1995; Szellas & Nagel, 2003) and tyrosine kinases (reviewed by Gadsby & Nairn, 1999) are able to phosphorylate CFTR.

Recently it was reported that in human sweat gland ducts, endogenous CFTR is activated by heterotrimeric G proteins via a cAMP-independent pathway (Reddy & Quinton, 2001). This Gprotein-mediated activation of CFTR is dependent on hydrolysable nucleoside triphosphates and Mg, indicating that phosphorylation is involved in the activating process although it is independent of the cAMP cascade.

The question arose whether alternative pathways exist to activate CFTR, apart from protein phosphorylation. In this study, we tested the effect of phosphatidylinositol 4,5-bisphosphate (PIP₂) on excised inside-out plasma membrane patches of *Xenopus laevis* oocytes, heterologously expressing human CFTR. Previously it was shown that PIP₂ plays an important role in the regulation of ion channels and transporters (Hilgemann *et al*, 2001; Hardie, 2003). For example, Na⁺/Ca⁺ exchangers (Hilgemann & Ball, 1996), all inwardly rectifying potassium channels (Huang *et al*, 1998) or the epithelial sodium channel (ENaC) (Yue *et al*, 2002), are reported to be activated by direct interaction with PIP₂. Unpublished experiments indicated that CFTR chloride channels were insensitive to PIP₂ (Hilgemann *et al*, 2001). Here we report a robust activation of nonphosphorylated CFTR by PIP₂, whereas phosphorylated CFTR may be inhibited by PIP₂.

RESULTS

Basal cAMP concentration, and therefore PKA activity in *Xenopus* oocytes, is normally low so that CFTR response to MgATP is marginal when excised from an oocyte (Weinreich *et al*, 1997). Figure 1A shows a very small effect of MgATP on membrane current in an excised inside-out giant membrane patch from an oocyte expressing human CFTR. This small CFTR activity might be due to the effects of basal [cAMP] and the resulting low kinase

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Fig 1 | PIP₂ activates chloride current by unphosphorylated CFTR. (A) Giant patch experiment. Several PIP₂ additions activate CFTR, followed by the decrease of the signal in consequence of PIP₂ washout. Bars indicate changes of the bath solution. PIP₂ = 1 μ M (hatched bar) or 10 μ M (black bar), MgATP = 500 μ M, membrane potential (V_m) = 0 mV. The arrow identifies zero current. Inset: Magnification of the beginning of the experiment showing increase of chloride current by prolonged MgATP application. Scale bars: 5 pA/10 s. 20 Hz filtered. (B) Single-channel experiment demonstrating activation of 9–11 channels by PIP₂ addition, followed by decay after PIP₂ washout and complete closing after ATP removal. PIP₂ = 1 μ M, MgATP = 500 μ M, V_m = 50 mV.

activity of the native intact oocyte before excising the membrane patch (Chan *et al*, 2000), but see alternative explanation below. Subsequent application of 1 μ M PIP₂, however, activates a large chloride current (Fig 1A). On the basis of voltage pulses, it was proved that the reversal potential ($-85 \pm 4 \text{ mV}$, n = 4) of the PIP₂-induced current is that of a chloride current as it is comparable to the value calculated for the employed [Cl⁻] gradient. This PIP₂-induced chloride current is not observed in control oocytes that are not expressing CFTR (n=4; data not shown). A further argument for the CFTR specificity of the PIP₂ effect is the CFTR-typical conductance ($5.3 \pm 0.7 \text{ pS}$ with [Cl⁻] gradient at 20–80 mV, n=8, for PIP₂-activated channels, compared to

 5.2 ± 0.3 pS, n = 9, for PKA-activated channels) and the ATP dependence of PIP₂-activated single channels (Fig 1B). Therefore, the PIP₂-induced chloride conductance is attributed to activation of CFTR, but not of other Cl- channels, for example, the oocyteendogenous Ca2+-activated chloride channel (Takahashi et al, 1987). Our repeated observation that prolonged ATP application to nonphosphorylated and not PIP₂-treated CFTR may lead to an increase of the small CFTR current, as shown in the inset of Fig 1A, may be explained by an ATP-induced increase of PIP₂. Such a generation of PIP2 from endogenous precursors via membranebound lipid kinases was previously described for excised membrane patches from heart cells (Hilgemann & Ball, 1996). The CFTR current observed before PIP₂ application (ATP-induced current increase in Fig 1A), which may be ascribed to basal oocyte-endogenous PKA activity, may therefore alternatively be caused, at least partially, by an endogenous small concentration of PIP_2 in the oocyte membrane.

The observed increase of the chloride current on addition of PIP₂ to CFTR-containing membrane patches depends on PIP₂ concentration. At lower concentrations (1 μ M, n = 7) in the bath solution, PIP₂ addition exclusively led to a current increase during application (Fig 1); however, at 10 μ M PIP₂ (n = 15) an exclusively increasing chloride current was observed in 73% of patches, and in the residual 27% a chloride current increase was followed by a decrease during PIP₂ application. This antagonistic effect of PIP₂ on CFTR is noticeably different from the purely stimulating effect of PKA on CFTR or from the effect of PIP₂ on K_{ATP} channels (Baukrowitz *et al*, 1998), where it was shown that PIP₂ application slowly antagonized the inhibition of K_{ATP} channels by ATP.

In accordance with the membrane solubility of PIP₂, its effect on CFTR was not completely reversible (Fig 1). The effect of removing PIP₂ from the bath solution was variable and was dependent on PIP₂ concentration. When removing 10 μ M PIP₂ we observed in 6 of 15 experiments a still continuing increase of chloride channel activity as long as ATP was present. Alternatively, removal of 10 μ M PIP₂ from the bath solution could also lead to a decrease (40%) or no change in the chloride current (20%).

In contrast, on removing $1 \mu M PIP_2$ from the bath, a decay of the chloride current was observed in 100% of patches (Fig 1). This 'rundown' after PIP₂ application appeared similar to the rundown after PKA withdrawal, but is probably caused by decreasing membrane concentration of PIP₂, for example, by dilution or because of enzymatic digestion.

As expected from the known ATP dependence (Weinreich *et al*, 1999), MgATP withdrawal from phosphorylated CFTR chloride channels yielded a current decay with time constants of about $\tau_1 = 2.9 \pm 0.2 \text{ s}$ and $\tau_2 = 24 \pm 2.5 \text{ s}$ (n = 6). The time constants of ATP withdrawal-induced channel closing were faster ($\tau_1 = 0.8 \pm 0.3 \text{ s}$, $\tau_2 = 9 \pm 1.7 \text{ s}$, n = 2) for CFTR activated by 1 μ M PIP₂ than for CFTR activated by 10 μ M PIP₂ in the bath solution ($\tau_1 = 2.3 \pm 0.4 \text{ s}$, $\tau_2 = 20 \pm 1.9 \text{ s}$, n = 7), similar to the previously described dependence of the closing rates on CFTR phosphorylation and rundown (Weinreich *et al*, 1999).

Next we investigated the effect of $10\,\mu$ M PIP₂ on phosphorylated CFTR. Compared to unphosphorylated CFTR, where application of $10\,\mu$ M PIP₂ led to activation of the chloride current in 100% of the patches, the addition of $10\,\mu$ M PIP₂ to PKAactivated CFTR resulted in variable effects. A further activation of



Fig 2 | Inhibitory effect of PIP₂ on phosphorylated CFTR. Bars indicate the bath solution. PIP₂ = 10 μ M, MgATP = 500 μ M. PKA (100 U/ml) addition activates a large CFTR chloride current. This current is first increased by application of PIP₂, directly followed by a strong decrease.



Fig 3 | Comparison of the stimulating effects of PI, PIP and PIP₂ on unphosphorylated CFTR. Bars indicate bath solution. MgATP = $500 \,\mu$ M (hatched bar) or $5 \,m$ M (black bar).

the chloride current was observed in only 18% of the experiments (n = 17). The stimulating effect was relatively small (maximally +23%), due to already strong activation by PKA. In 47% of the patches an inhibition of PKA-activated current (Fig 2) and in 35% no apparent change of the PKA-activated chloride current (data not shown) was observed.

To test the influence of the number of free phosphate groups at the inositol ring, we added to the patch either phosphatidylinositol (PI), which has no phosphate residue, or phosphatidylinositol monophosphate (PIP) with one phosphate group at position 4, or PIP₂ with two phosphate groups at positions 4 and 5 of the inositol ring. Fig 3 shows that already PI had a clear stimulating effect on unphosphorylated CFTR, PIP was similar to PI, and PIP₂ yielded the strongest stimulation of CFTR chloride current.

An obvious question is whether PIP_2 alone can open unphosphorylated CFTR, that is, in the absence of nucleoside triphos-



Fig 4|ATP is needed to open PIP₂-treated CFTR channels. Bars indicate changes of the bath solution. PIP₂=1 μ M, MgATP = 500 μ M.

phates. Application of 1 or 10 µM PIP₂ alone, however, resulted in no change of electrical current (Fig 4). If ATP was applied after the application of $1 \mu M PIP_{2}$, CFTR opened very fast (Fig 4). Channel activity, which increased 9.2 (± 2) -fold, was then followed by a decrease of about $75 \pm 4\%$ (*n*=6) (Fig 4), as described above for PIP₂ removal from the bath. Similar to CFTR activated by protein kinases, CFTR activated by PIP₂ also needs ATP to open. An initial application of PIP₂ in the absence of ATP, however, is sufficient to allow ATP to open CFTR chloride channels rapidly (Fig 4). This result already suggests that no protein phosphorylation is involved in PIP₂-mediated activation of CFTR, but it does not yet unambiguously rule out this possibility. We addressed this question by applying PIP₂ in the complete absence of Mg^{2+} or other divalent cations, that is, conditions that are known to abolish any kinase activity. Alternatively, we tested the effect of PIP₂ on CFTR in the absence of ATP by replacing it with the nonhydrolysable ATP analogue adenylyl imido diphosphate (AMP-PNP).

Although Mg-free conditions sometimes cause problems with the baseline stability of the patch current, we could unambiguously demonstrate that PIP₂ application in the presence of ATP (0 Mg) always leads to an increase of CFTR chloride current (n = 11), with both 1 and $10 \,\mu\text{M}$ PIP₂ (Fig 5A). In most cases, the increase of CFTR-mediated chloride current is directly followed by a decrease. This antagonistic effect of PIP₂ on CFTR is demonstrated in Fig 5A by a second treatment of the patch with PIP₂. The activation of CFTR by PIP₂ in the absence of Mg strongly argues against a PIP₂-mediated protein phosphorylation of CFTR. Although we previously concluded, from experiments with 500 µM AMP-PNP, that phosphorylated CFTR may not be opened by application of the nonhydrolysable ATP analogue AMP-PNP (Nagel et al, 1992; Weinreich et al, 1999), it was recently demonstrated that AMP-PNP may open phosphorylated CFTR when used at high concentrations (5 mM) (Aleksandrov et al, 2000). We recently confirmed this observation for PKA-activated human CFTR in giant patches of oocyte membrane (data not shown). Fig 5B shows that CFTR in PIP₂-treated membrane patches does not open on application of 500 µM AMP-PNP but rapidly opens on application of 5 mM AMP-PNP, in the complete absence of ATP, whereas 5 mM AMP-PNP has no effect on CFTR before PIP₂ application. This result is further proof that no protein phosphorylation takes place during PIP₂ activation of CFTR.



Fig 5 | PIP₂ activates CFTR without protein phosphorylation. (A) CFTR activation in the absence of Mg. Bars indicate bath solution. TrisATP = 5 mM. (B) PIP₂ activates CFTR in the absence of ATP. Bars indicate bath solution. PIP₂ = 1 μ M, MgATP = 500 μ M, AMP-PNP = 500 μ M (hatched bar) or 5 mM (black bar). Before PIP₂ application, neither ATP nor AMP-PNP is able to open CFTR. AMP-PNP at 5 mM, but not 500 μ M, opens PIP₂-activated unphosphorylated CFTR.

DISCUSSION

Interaction between PIP₂ and ion channels or transporters was described previously in several reports and reviews (Hilgemann et al, 2001). With this report, we add another ion channel to the growing list of membrane proteins regulated by phosphoinositides. For several ion-translocating membrane proteins, a direct interaction with PIP₂ was demonstrated. All inward rectifying K⁺ channels, the Na+/Ca2+ exchanger and the epithelial sodium channel directly interact with PIP2. In Kir6.2 channels, a conserved carboxy-terminal region was identified as interacting with cell membranes. From systematic mutagenesis, it was proposed that a lipid interaction domain could be identified that shows structurally significant features of phospholipid interacting pleckstrin homology (PH) domains (Cukras et al, 2002b). Positively charged amino acids in the amino terminus (Nterminus) of Kir6.2 were also shown to affect regulation by ATP and PIP₂ (Cukras et al, 2002a). A conserved region with some homology to PH domains is also present within the N-terminus of CFTR. Interestingly, Kirk and co-workers found that the N-terminal tail and especially four conserved acidic amino acids play an important role in the positive regulation of CFTR activity by directly binding to the R-domain (Naren *et al*, 1999) or with NBD2 (Fu *et al*, 2001).

Before trying to pinpoint the region on CFTR that is interacting with PIP_{2} , it is necessary to exclude the possibility that PIP_{2} mediates its effect indirectly via a protein kinase. Two lines of evidence support a direct CFTR-PIP₂ interaction: treatment of membrane patches with $\ensuremath{\mathsf{PIP}}_2$ in the complete absence of ATP did not open CFTR but resulted in an activation of CFTR as subsequently demonstrated by fast ATP-induced opening of CFTR. As the time constants were at least as fast as those measured for ATP-induced opening of phosphorylated CFTR (Weinreich et al, 1999), it seems unlikely that ATP addition first triggered a PIP₂mediated phosphorylation and then the opening of CFTR. Even more compelling are the results with PIP₂ and ATP in the complete absence of divalent cations. Because hydrolytic enzymes require Mg^{2+} or other divalent cations, phosphorylation by kinases is abolished in their absence (Schultz et al, 1996). For CFTR, we show stimulating effects of PIP₂ even under Mg²⁺-free conditions, indicating that activation of CFTR by PIP₂ cannot be attributed to protein phosphorylation, for example, by a membrane-bound PIP₂-dependent kinase. The fact that PIP₂ alone has no effect on the patch current, but needs application of ATP to induce a chloride current, is additional evidence for the CFTR specificity of the PIP₂ effect. This argument is strengthened by the CFTR-typical gating and single-channel conductance observed in PIP₂-treated CFTR-expressing membrane patches.

Until now, CFTR channel activity under physiological conditions was thought to have an absolute requirement for protein phosphorylation and nucleoside triphosphates. However, a recent report showing dATP-induced activation of carefully controlled, nonphosphorylated CFTR already indicated that protein phosphorylation may not be an absolute requirement to open CFTR chloride channels (Aleksandrov *et al*, 2002). Even more surprising is the recent demonstration that human CFTR, endogenously expressed in the sweat duct, may conduct chloride ions simply by the addition of cytoplasmic glutamate, in a phosphorylation- and ATP-independent manner (Reddy & Quinton, 2003).

Our studies indicate that also under virtually physiological conditions CFTR may be activated by a pathway that does not involve protein phosphorylation. The addition of PIP₂ to the ATPcontaining bath solution activated a strong CFTR chloride conductance, smaller but still similar to activation by PKA (18% of PKA-activated conductance with 10 μ M PIP₂, n = 6, and 7% of PKA-activated conductance with $1 \mu M PIP_2$, n=8; data not shown). In contrast to activation of CFTR by glutamate (Reddy & Quinton, 2003), activation of CFTR by PIP₂ still needs ATP to open the CFTR chloride channel. As we found no effect of glutamate on CFTR in excised membrane patches, with or without ATP (n = 6; data not shown), we conclude that glutamate does not activate CFTR directly. We suggest a possible involvement of phosphatidylinositols in transmitting the signal from glutamate receptors to CFTR in the sweat duct, even if this seems in contrast to the reported ATP independence of activation by glutamate (Reddy & Quinton, 2003).

Already the reported cAMP-independent activation of CFTR by G proteins in the sweat duct (Reddy & Quinton, 2001) provided evidence for the involvement of alternative kinase phosphorylation

in CFTR activation. Although this elegant study could not pinpoint which kinase led to activation of CFTR, it seemed most probable to presume a protein kinase. From our results we suggest the possibility that activation of G proteins in the sweat duct stimulates one or several lipid kinase(s). A kinase-induced increase of PIP₂ in the membrane should then activate CFTR. Alternatively, our results would also predict activation of CFTR if PIP₂ content of the membrane increases by inhibition of PIP₂ degradation.

We suggest that there is a previously unrecognized alternative way to regulate CFTR by membrane components, such as phosphatidylinositides. We speculate that the recently reported cAMP-independent activation of CFTR in sweat gland ducts (Reddy & Quinton, 2001) is due to G-protein-mediated production of PIP₂ via lipid kinase(s) and binding of PIP₂ to unphosphorylated CFTR.

METHODS

Materials. Egtazic acid (EGTA), ethylenediaminetetraacetic acid (EDTA), MgCl₂, D,L-aspartic acid, MgATP, TrisATP, L-glutamic acid, PI, PIP, Tricain and collagenase IA were obtained from Sigma (Deisenhofen, Germany). *N*-methyl-D-glucamine, HCl and CaCl₂ were obtained from Merck (Darmstadt, Germany). HEPES was purchased from AppliChem (Darmstadt, Germany). AMP-PNP and PIP₂ were obtained from Sigma (Deisenhofen, Germany) or Roche (Mannheim, Germany). PKA catalytic subunit was obtained from Promega (Wisconsin, USA). Oocytes and human CFTR cRNA were prepared as described before (Weinreich *et al*, 1997). The oocytes were incubated for 3–4 days at 16–18 °C in modified Ringer's solution containing 100 µg/ml gentamycin.

Solutions. A holding potential of 0 mV (50 mV for single-channel experiments) and a [Cl-] gradient of 4/156 mM (bath/pipette) was applied as previously reported by Nagel *et al* (1992). The Mg^{2+} containing bath solution was composed of 5 mM EGTA, 10 mM HEPES, 2 mM MgCl₂ and 160 mM NMG, and was titrated to pH 7.4 with D,L-aspartic acid. The Mg²⁺-free bath solution was composed of 2 mM EDTA, 5 mM EGTA, 4 mM HCl, 10 mM HEPES and 160 mM NMG, and was titrated to pH 7.4 with D,L-aspartic acid. The pipette solution was composed of 10 mM HEPES, 150 mM NMG and 2 mM MgCl₂, and was adjusted to pH 7.4 with HCl. PI was obtained as a solution (10 mg/ml in chloroform); PIP or PIP₂ was dissolved in a mixture containing chloroform, methanol, water and 1N HCl in a ratio of 20:10:1:1 or 20:9:1:0.1, respectively, to a final concentration of 10 mM. The phosphoinositides were dried under argon or nitrogen, and then sonicated or vortexed in bath solution directly before use.

Patch clamp experiments. Giant patch experiments were performed as described (Hilgemann & Lu, 1998; Weinreich *et al*, 1999). All the experiments were performed at room temperature $(20-22 \degree C)$ with a standard holding potential of 0 mV for giant patch and 50 mV for single-channel experiments. Short test pulses in the range of -90 to +30 mV, or +20 to +80 mV for single-channel experiments, were performed to obtain the *I–V* relationship. The electrical measurements were performed with an Axopatch 200B (Axon instruments) or a Patch Clamp L/M EPC7 amplifier (List-Medical, Darmstadt, Germany). Clampfit 8.0 and Microcal Origin 6.1 were used to perform data analysis. To reduce data points for the analysis of giant patch experiments, only every tenth point was used for the figures.

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REFERENCES

- Aleksandrov AA, Chang X, Aleksandrov L, Riordan JR (2000) The nonhydrolytic pathway of cystic fibrosis transmembrane conductance regulator ion channel gating. *J Physiol* **528**: 259–265
- Aleksandrov AA, Aleksandrov L, Riordan JR (2002) Nucleoside triphosphate pentose ring impact on CFTR gating and hydrolysis. *FEBS Lett* **518**: 183–188
- Anderson MP, Berger HA, Rich DP, Gregory RJ, Smith AE, Welsh MJ (1991) Nucleoside triphosphates are required to open the CFTR chloride channel. *Cell* **67:** 775–784
- Baukrowitz T, Schulte U, Oliver D, Herlitze S, Krauter T, Tucker SJ, Ruppersberg JP, Fakler B (1998) PIP2 and PIP as determinants for ATP inhibition of KATP channels. *Science* **282:** 1141–1144
- Chan KW, Csanady L, Seto-Young D, Nairn AC, Gadsby DC (2000) Severed molecules functionally define the boundaries of the cystic fibrosis transmembrane conductance regulator's NH(2)-terminal nucleotide binding domain. J Gen Physiol **116:** 163–180
- Cukras CA, Jeliazkova I, Nichols CG (2002a) The role of NH2-terminal positive charges in the activity of inward rectifier KATP channels. *J Gen Physiol* **120:** 437–446
- Cukras CA, Jeliazkova I, Nichols CG (2002b) Structural and functional determinants of conserved lipid interaction domains of inward rectifying Kir6.2 channels. *J Gen Physiol* **119**: 581–591
- French PJ, Bijman J, Edixhoven M, Vaandrager AB, Scholte BJ, Lohmann SM, Nairn AC, de Jonge HR (1995) lsotype-specific activation of cystic fibrosis transmembrane conductance regulator-chloride channels by cGMPdependent protein kinase II. *J Biol Chem* **270**: 26626–26631
- Fu J, Ji HL, Naren AP, Kirk KL (2001) A cluster of negative charges at the amino terminal tail of CFTR regulates ATP-dependent channel gating. J Physiol 536: 459–470
- Gadsby DC, Nairn AC (1999) Control of CFTR channel gating by phosphorylation and nucleotide hydrolysis. *Physiol Rev* **79:** S77–S107
- Gadsby DC, Nagel G, Hwang TC (1995) The CFTR chloride channel of mammalian heart. *Annu Rev Physiol* **57:** 387–416
- Hardie RC (2003) Regulation of trp channels via lipid second messengers. Annu Rev Physiol 65: 735–759
- Hilgemann DW, Ball R (1996) Regulation of cardiac Na⁺, Ca²⁺ exchange and KATP potassium channels by PIP2. *Science* **273:** 956–959
- Hilgemann DW, Lu CC (1998) Giant membrane patches: improvements and applications. *Methods Enzymol* **293**: 267–280
- Hilgemann DW, Feng S, Nasuhoglu C (2001) The complex and intriguing lives of PIP2 with ion channels and transporters. *Sci STKE* **2001:** RE19
- Huang CL, Feng S, Hilgemann DW (1998) Direct activation of inward rectifier potassium channels by PIP2 and its stabilization by Gβγ. *Nature* **391**: 803–806
- Hwang TC, Nagel G, Nairn AC, Gadsby DC (1994) Regulation of the gating of cystic fibrosis transmembrane conductance regulator C1 channels by phosphorylation and ATP hydrolysis. *Proc Natl Acad Sci USA* **91**: 4698–4702
- Hyde SC, Emsley P, Hartshorn MJ, Mimmack MM, Gileadi U, Pearce SR, Gallagher MP, Gill DR, Hubbard RE, Higgins CF (1990) Structural model of ATP-binding proteins associated with cystic fibrosis, multidrug resistance and bacterial transport. *Nature* **346**: 362–365
- Nagel G (1999) Differential function of the two nucleotide binding domains on cystic fibrosis transmembrane conductance regulator. *Biochim Biophys Acta* **1461:** 263–274
- Nagel G, Hwang TC, Nastiuk KL, Nairn AC, Gadsby DC (1992) The protein kinase A-regulated cardiac Cl⁻ channel resembles the cystic fibrosis transmembrane conductance regulator. *Nature* **360**: 81–84
- Naren AP, Cormet-Boyaka E, Fu J, Villain M, Blalock JE, Quick MW, Kirk KL (1999) CFTR chloride channel regulation by an interdomain interaction. *Science* **286:** 544–548
- Quinton PM (1999) Physiological basis of cystic fibrosis: a historical perspective. *Physiol Rev* **79:** S3–S22

- Reddy MM, Quinton PM (2001) cAMP-independent phosphorylation activation of CFTR by G proteins in native human sweat duct. *Am J Physiol Cell Physiol* **280:** C604–C613
- Reddy MM, Quinton PM (2003) Control of dynamic CFTR selectivity by glutamate and ATP in epithelial cells. *Nature* **423**: 756–760
- Riordan JR (1993) The cystic fibrosis transmembrane conductance regulator. Annu Rev Physiol 55: 609–630
- Riordan JR *et al* (1989) Identification of the cystic fibrosis gene: cloning and characterization of complementary DNA. *Science* **245**: 1066–1073
- Schultz BD, Bridges RJ, Frizzell RA (1996) Lack of conventional ATPase properties in CFTR chloride channel gating. J Membr Biol **151:** 63–75
- Sheppard DN, Welsh MJ (1999) Structure and function of the CFTR chloride channel. *Physiol Rev* **79:** S23–S45

- Szellas T, Nagel G (2003) Apparent affinity of CFTR for ATP is increased by continuous kinase activity. *FEBS Lett* **535**: 141–146
- Takahashi T, Neher E, Sakmann B (1987) Rat brain serotonin receptors in *Xenopus* oocytes are coupled by intracellular calcium to endogenous channels. *Proc Natl Acad Sci USA* **84:** 5063–5067
- Weinreich F, Wood PG, Riordan JR, Nagel G (1997) Direct action of genistein on CFTR. *Pflugers Arch* **434**: 484–491
- Weinreich F, Riordan JR, Nagel G (1999) Dual effects of ADP and adenylylimidodiphosphate on CFTR channel kinetics show binding to two different nucleotide binding sites. J Gen Physiol **114:** 55–70
- Yue G, Malik B, Eaton DC (2002) Phosphatidylinositol 4,5-bisphosphate (PIP2) stimulates epithelial sodium channel activity in A6 cells. *J Biol Chem* **277:** 11965–11969