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A novel drug series optimized to address cystic fibrosis and other CFTR deficiency diseases of human airways

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Pharmacologic activation of the cystic fibrosis transmembrane conductance regulator (CFTR) has transformed cystic fibrosis (CF) therapy. Other, more common airway diseases can also be associated with CFTR deficiency. For example, individuals with one dysfunctional CFTR variant (i.e., CF carriers), as well as those with acquired CFTR deficiency, are predisposed to both non-CF bronchiectasis and chronic rhinosinusitis, raising the possibility that CFTR stimulation in these settings could provide clinical improvement. This study describes a new triazolo-thiadiazine-based compound series optimized to augment mutant and wildtype CFTR function when administered topically to airway epithelium. Mechanism of action appears attributable—at least in part—to phosphodiesterase 4 inhibition (PDE4i), with effects on other PDEs also noted. Together with a growing body of previous and emerging evidence, our results suggest a novel therapeutic strategy for treating people with CF (PwCF) who lack access to effective modulator therapy—and addressing common diseases such as chronic bronchiectasis and rhinosinusitis in the non-CF population.

Over the past several years, drug discovery has dramatically benefited people with cystic fibrosis^{1–7}. Highly effective modulator treatment (HEMT) is available in the US to >90% of individuals with the disease, and the FDA label includes approvals for over 300 mutations in the cystic fibrosis transmembrane conductance regulator (CFTR), most of which have been established as disease-causing variants. “Triple combination therapy” (comprised of elexacaftor (VX-445), tezacaftor (VX-661), and ivacaftor (VX-770; ETI) or vancacaftor (VX-121), tezacaftor, and deutivacaftor (VX-561)⁸) confer improvement in CF lung function, rhinosinusitis, quality of life, and other endpoints (including markers of CFTR activity such as sweat chloride) within days to a few weeks^{8–10}. As with any new therapy, a subset of PwCF fail HEMT (e.g., ~5% of individuals encoding the common F508del variant may not benefit from ETI) or exhibit limiting toxicity (for example, hepatic impairment) that precludes use^{6,9–11}. In addition, hundreds of disease-causing CFTR variants are not FDA-approved for modulator treatment, including refractory defects of folding and maturational processing^{1,12}. Based on successful precedent achieved with ETI, several novel compounds with superior efficacy are being advanced by academic and pharmaceutical laboratories^{8,13–15}. A recent survey examined over 650 CFTR variants associated with clinical disease for modulator responsiveness¹⁶ using cell systems considered predictive of pulmonary benefit in vivo^{14,5,17–21}. That study revealed many CFTR mutations will

require new and distinct activators specifically designed to improve lung function in patients with cystic fibrosis.

Individuals with CF experience tissue damage throughout the respiratory tract, including nasal passages, sinuses, lung parenchyma, and submucosal glands^{1,7,22–31}. Involvement of the upper airways is very common (>90% of PwCF have radiographic evidence of chronic sinusitis). Impaired mucociliary clearance, airway obstruction, bioelectric abnormalities, immunologic hyperactivity, and chronic infection (by *P. aeruginosa*, *H. influenza*, *S. aureus*, *B. cepacia* and other pathogens) characterize both nasal and lower CF respiratory tissues^{6,7}. HEMT strongly palliates—but does not resolve—existing CF lung and sinonasal pathology. While it has been suggested that CFTR modulators given at or before birth may prevent airway disease from occurring, longitudinal studies will be required to formally test that possibility. In either case, improved small molecules are actively being sought based on the view that a therapeutic ceiling has not yet been achieved with existing modulators^{14,15}. Moreover, data in primary airway epithelial cells—model systems that correlate with clinical benefit—indicate potential for activation of the common F508del variant well above what can be achieved with HEMT alone^{14,15,32,33}.

Respiratory failure due to bronchiectasis is the primary cause of morbidity and mortality in people with CF³⁴. As above, pulmonary tissues exhibit many of the same clinical findings as the CF sinonasal tract, and

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pathogenic mechanisms are often viewed as comparable^{6,7}. Epithelial monolayers from both nasal and lower airways have been used as tools for understanding and advancing access to HEMT, further supporting the relationship between nasal passages and modulator activity in lungs^{35–38}. In addition, the importance of improved treatment for chronic rhinosinusitis in CF has been emphasized as a means to prevent “seeding” of lower pulmonary tissues and to improve outcomes post-lung transplantation^{39–46}. Chronically infected nasal sinuses serve as a bacterial reservoir that infects the lower respiratory tract, impairs pulmonary graft viability, and causes failure of attempts to eradicate pulmonary pathogens in PwCF. Findings such as these indicate mechanistic similarities and strong interrelationships between CF lower and upper airway disease, with implications for other, more common illnesses.

A pulmonary condition that exhibits pathologic features with close similarity to CF lung disease—non-cystic fibrosis bronchiectasis (NCFB)—afflicts millions worldwide. Comparable clinical findings include abnormal airway dilation, diminished mucociliary clearance (MCC), and infection by similar (and recalcitrant) bacterial pathogens⁴⁷. Endobronchial damage, associated inflammatory changes, and pulmonary remodeling in NCFB can lead to respiratory failure and death^{48–50}. Although NCFB has traditionally been viewed as mechanically unrelated to CFTR dysfunction, individuals who are silent carriers of CF (heterozygous with one pathogenic CFTR variant and one wildtype CFTR allele) are at higher risk of developing NCFB^{51–53}. In addition, persistent inflammation, hypoxia, cigarette smoke exposure, and other toxins have been shown to contribute to an acquired loss of CFTR in conditions such as NCFB^{54,55}. Abnormalities of the CFTR airway bioelectric phenotype⁵⁴ and a variety of mutations in CFTR have been associated with NCFB. Moreover, recent case studies indicate potential usefulness of CFTR modulators for treating a subset of individuals with non-CF bronchiectasis⁵⁶.

In the same context, approximately 1 in 8 Americans suffers from intermittent, non-CF chronic rhinosinusitis (CRS), which (like NCFB) has been increasingly linked to both mutant CFTR and diminished wildtype CFTR activity^{55,57–72}. Non-CF CRS accounts for >10 million coded office visits annually in the US⁷³ and is a cause of significant morbidity, including persistent and/or severe facial pain, headache, rhinorrhea, anosmia, absenteeism from work, and substantial debility. From a clinical perspective, predisposition to non-CF CRS is associated with heterozygosity for CFTR mutations, indicating that relative CFTR deficiency and diminished mucociliary clearance are partly responsible for pathogenesis^{59–62}. Moreover, in semblance to NCFB, chronic airway infection/inflammation impairs CFTR biogenesis and ion transport, further supporting a therapeutic role for CFTR modulators in the treatment of this disease^{55,57,58,63,69–71,74}.

Clinical cystic fibrosis and “forme frustes” of CF such as NCFB and CRS are clearly caused—at least in part—by inherited or acquired CFTR deficiency. We are developing a new class of compounds that activate both WT and mutant CFTR through robust, PKA-dependent R-domain phosphorylation. The drugs are intended for use as monotherapy or in combination with available CFTR modulators for treating PwCF who exhibit suboptimal clinical response, poor modulator tolerance, or those encoding CFTR variants that remain “off label” for emerging modulator therapy. Because the new compounds strongly activate both mutant and wildtype CFTR, contributions to treatment of non-CF airway diseases attributable to CFTR deficiency and inadequate mucociliary clearance (for example, CRS and NCFB) are also anticipated.

Results

Overview

We previously screened 300,000 agents from a library of bioactive compounds as a means to rescue N1303K or W1282X CFTR, two relatively common variants with measurable (albeit low level) function⁷⁵. The screen was conducted using Fischer rat thyroid (FRT) cells stably transduced with CFTR encoding an extracellularly tagged horseradish peroxidase (HRP) for monitoring cell surface localization. The FRT model has been successful in many CFTR pharmaceutical drug discovery programs from both academic

and commercial laboratories^{4,17,21,37,76}, leading to several FDA-registered compounds. Following confirmation and dose-dependence analysis, as well as *in silico* (modified ‘PAINS’) profiling to exclude candidate drugs with inherent liabilities⁷⁷, active scaffolds were identified. Based on a high compound “dropout” rate associated with drug screening in FRT cells, candidate molecules were tested using complementary cell systems. The most promising agents were advanced to validation in primary airway epithelial monolayers grown on permeable supports—a model shown previously to predict *in vivo* clinical benefit and recognized by the FDA for the purpose of drug registration^{4,21}.

HDCF drug analysis and CFTR ion transport

Lead agents resulting from the original compound library screen and subsequent medicinal chemistry optimization⁷⁵ identified a drug series that strongly augments wildtype CFTR activity as judged by changes in short circuit current (Isc) in FRT cells (Fig. 1). Two examples of this series—termed HDCF104 and uHTS159—are reported in detail here. Functional assays in primary human bronchial epithelial monolayers encoding WT CFTR also showed robust enhancement of CFTR transepithelial transport (Fig. 2). Importantly, compounds in the newly identified class acutely stimulate many disease-causing CFTR variants. Unlike CFTR “corrector” agents, which rescue folding but do not augment channel gating, drugs described in this report block phosphodiesterase activity, enhance PKA-dependent phosphorylation of CFTR, and stimulate ion transport (see below). Data showing use of the new drug series against a number of CFTR variants are provided in the following sections.

G551D CFTR activation by uHTS159

An FDA-approved CFTR modulator (VX-770) was specifically developed by the pharmaceutical industry as a “potentiator” to rescue the G551D CFTR channel gating defect^{37,78}. Figure 3 shows uHTS159 activates G551D CFTR independent of VX-770 and can strongly enhance VX-770 potentiation. In terms of efficacy as a single agent, uHTS159 is roughly equivalent to VX-770. When saturating concentrations of the clinically approved potentiator are applied, uHTS159 promotes further short circuit current, significantly surpassing the effects of VX-770 alone.

HDCF stimulation of F508del CFTR following ETI

The FRT model has been viewed by pharmaceutical groups and FDA as predictive of *in vivo* clinical response—particularly when a test compound augments mutant CFTR function to >10% wildtype activity levels (e.g., Fig. 3)^{4,16,17,21,37}. As described above, primary CF airway epithelial cells are also well-established for predicting patient benefit^{4,19–21}. Figure 4 provides an example of significant (10–20 $\mu\text{A}/\text{cm}^2$) CFTR-dependent anion current increase conferred by HDCF104 following pharmacocorrection of the common F508del CFTR protein processing defect (present in ~90% of PwCF), indicating substantial and clinically meaningful enhancement of CFTR well above what can be accomplished with ETI alone. Importantly, HDCF104 elevates Isc to levels that are nearly comparable to wildtype CFTR activity in this model system. In the same context, note that although VX-770 activates F508del CFTR even in the absence of forskolin, total ion transport is significantly lower than what can be achieved by combination treatment of VX-770 with HDCF104. While levels of constitutively active PKA in primary human airway cells (Fig. 4) are therefore sufficient for stimulation of F508del CFTR by VX-770, augmenting PKA-dependent R-domain phosphorylation provides much stronger enhancement of protein function.

Findings regarding N1303K CFTR

N1303K is the second most common class 2 CFTR defect, and a variant for which ETI has recently been FDA-approved based on clinical findings showing favorable—albeit less than maximal—lung function improvement⁷⁹. Figure 5 shows activation of N1303K CFTR by HDCF104 in primary airway epithelial monolayers beyond what can be accomplished with ETI alone. In Fig. 5a–h, nasal airway epithelial cells from an individual

Fig. 1 | Dose-dependent activation of wildtype CFTR in FRT cells by two HDCF analogs (HDCF104 and uHTS159). Ussing chamber analysis of Fischer rat thyroid (FRT) cells shows stimulation of WT CFTR. **a, b** Escalating doses of HDCF104 demonstrate strong WT CFTR activity with $EC_{50} = 101 \mu\text{M}$ (95% CI $91.7 \mu\text{M}$ to $112.3 \mu\text{M}$) as judged by dose-dependence profile. In similar fashion, **(c, d)** uHTS159 displays robust enhancement of WT CFTR with $EC_{50} = 14.4 \mu\text{M}$ (95% CI $12.5 \mu\text{M}$ – $17.2 \mu\text{M}$). Data is representative of 3 biological replicates and expressed as mean \pm SD. Maximal stimulation of wildtype CFTR by saturating concentrations of forskolin in this model is 300 – $500 \mu\text{A}/\text{cm}^2$. Amiloride ($100 \mu\text{M}$) was added to block epithelial sodium channels; forskolin ($5 \mu\text{M}$) augments cellular cAMP and activates CFTR by PKA-dependent phosphorylation of the regulatory domain; inh-172 ($10 \mu\text{M}$) is a CFTR inhibitor. **e** Structures of the compounds. Statistics were by use of GraphPad Prism 10 software with the function of Sigmoidal, 4PL, with X as concentration. “HDCF” denotes the medicinal chemistry laboratory in which new compounds were synthesized (directed by Dr. Huw Davies). Panels **(a, c)** plotted with Microsoft Excel; **(b, d)** plotted (and compound EC_{50} calculated) using GraphPad Prism 10 software; Panel **(e)** created with ChemDraw; panels arranged and presented using Affinity Designer software.

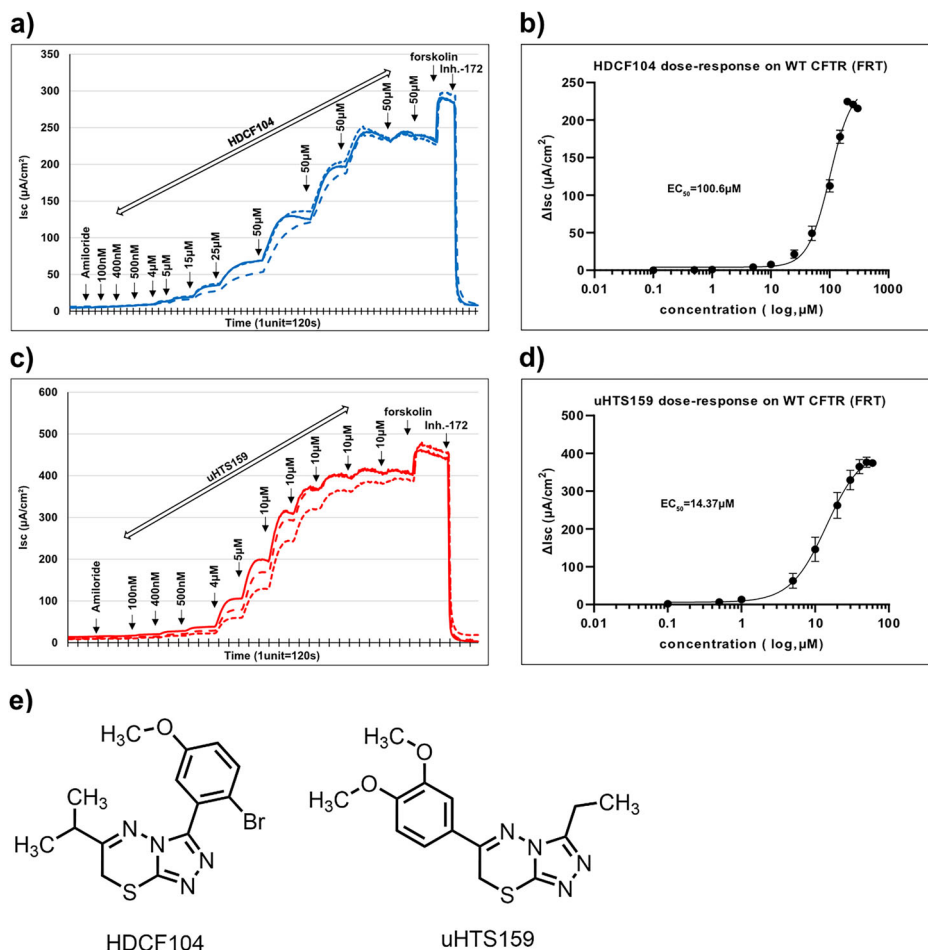


Fig. 2 | Dose-dependent activation of wildtype CFTR in primary human bronchial epithelial cells by two HDCF analogs. **a** Escalating doses of HDCF104 provide robust WT CFTR activation with $EC_{50} = 4.16 \mu\text{M}$ (95% CI $2.748 \mu\text{M}$ to $5.572 \mu\text{M}$). **b** Data summarized by dose-activity profile. **c, d** In similar fashion, escalating doses of uHTS159 demonstrate activation of WT CFTR with $EC_{50} = 0.71 \mu\text{M}$ (95% CI $0.558 \mu\text{M}$ to $0.864 \mu\text{M}$). Data is representative of 4 biological replicates and expressed as mean \pm SD. Addition of amiloride ($100 \mu\text{M}$), forskolin ($10 \mu\text{M}$), inh-172 ($10 \mu\text{M}$), and UTP (stimulates non-CFTR chloride channels as a control for monolayer integrity; $100 \mu\text{M}$) are shown. HBEC = primary human bronchial epithelial cells. Statistics were by use of GraphPad Prism 10 software with the function of Sigmoidal, 4PL, with X as concentration. Panels **(a, c)** plotted with Microsoft Excel; **(b, d)** plotted (and compound EC_{50} calculated) using GraphPad Prism 10 software; panels arranged and presented using Affinity Designer.

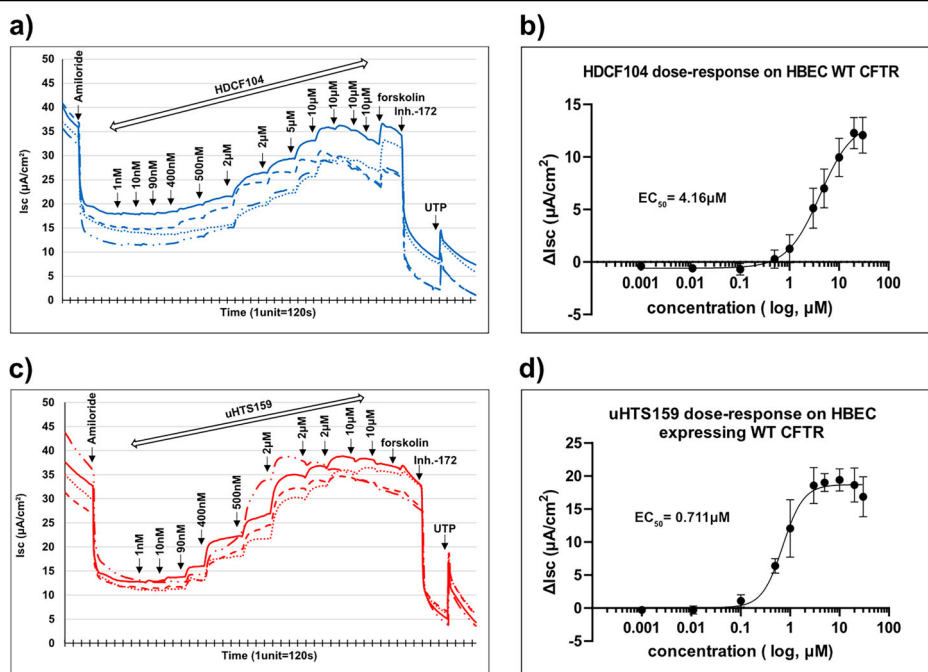


Fig. 3 | G551D rescue by uHTS159 in FRT cells. Activation of G551D CFTR in the FRT model by uHTS159 (10 μ M) is shown, including substantial potentiation of I_{sc} beyond VX-770 alone irrespective of the order in which compounds are added. **a** Representative tracings; **b** summary of 8 biological replicates per condition expressed as mean \pm SEM. $p < 0.01$ for maximal activation of G551D CFTR by forskolin and VX-770 vs addition of uHTS159 with forskolin and VX-770. Statistics were by Excel software, ANOVA, single factor analysis function. Figures plotted with Microsoft Excel; plots arranged and presented using Affinity Designer software.

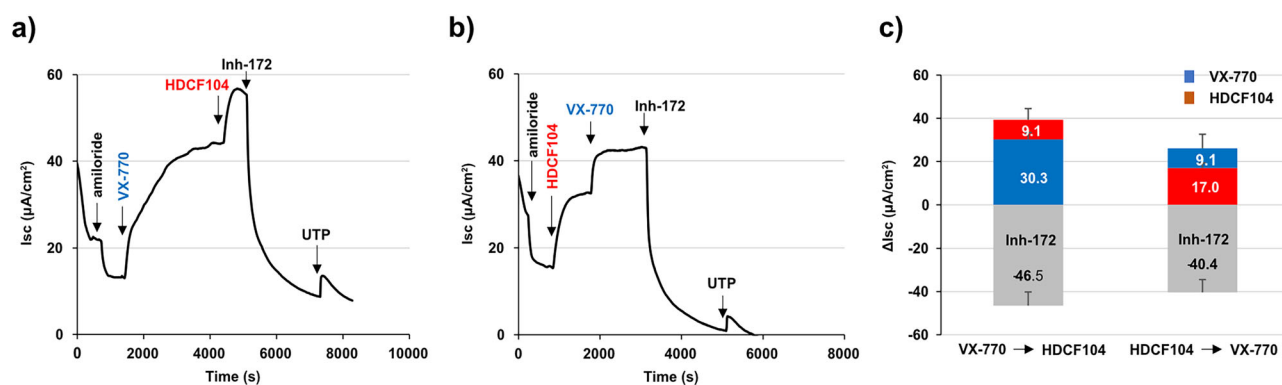
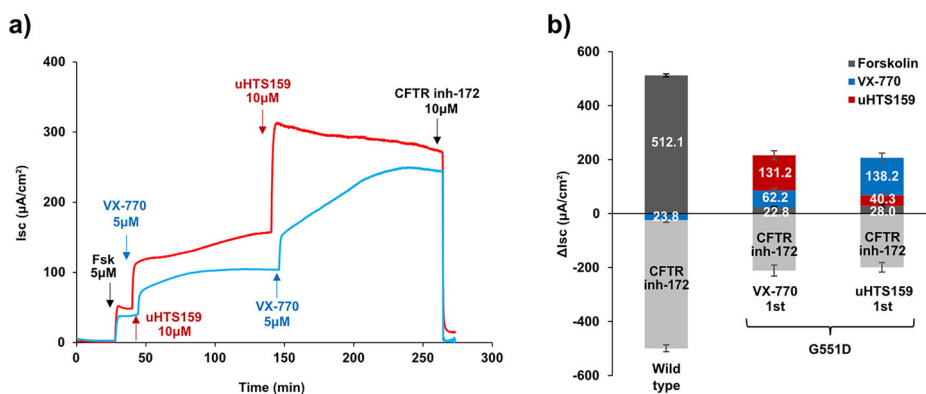


Fig. 4 | Rescue of F508del CFTR using HDCF104 following pharmacocorrection by VX-661 and VX-445. Activity in homozygous F508del human bronchial epithelial monolayers is shown. **a, b** Representative Ussing assay tracings. **c** Summary data using saturating concentrations of either VX-770 or HDCF104 added first. $n = 4$ biological replicates per condition; bars depict standard deviation (SD). Acute

treatments shown are amiloride (100 μ M), VX-770 (5 μ M), HDCF104 (20 μ M), and inh-172 (10 μ M). $p = 6.8 \times 10^{-6}$ for HDCF104 plus VX-770 versus VX-770 alone (HDCF compound added first). Statistics were by Excel Data Analysis ToolPak, T -test tool. Figures plotted with Microsoft Excel; plots arranged and presented using Affinity Designer software.

with genotype N1303K/N1303K exhibited an additional 4–5 μ A/cm² stimulation when corrector treatment was followed by HDCF104. Nasal airway epithelium from a patient with N1303K/G542X genotype (Fig. 5i, j) also responded to the HDCF compound. G542X is expected to be “null,” and findings indicate ~ 4 μ A/cm² stimulated CFTR current in the sample being tested. This level of rescue when combined with ETI amounts to $\sim 30\%$ of wildtype CFTR in primary nasal airway epithelium, a value seen as indicative of strong clinical respiratory improvement^{4,19–21,79}.

HDCF104 and treatment of other CFTR mutations

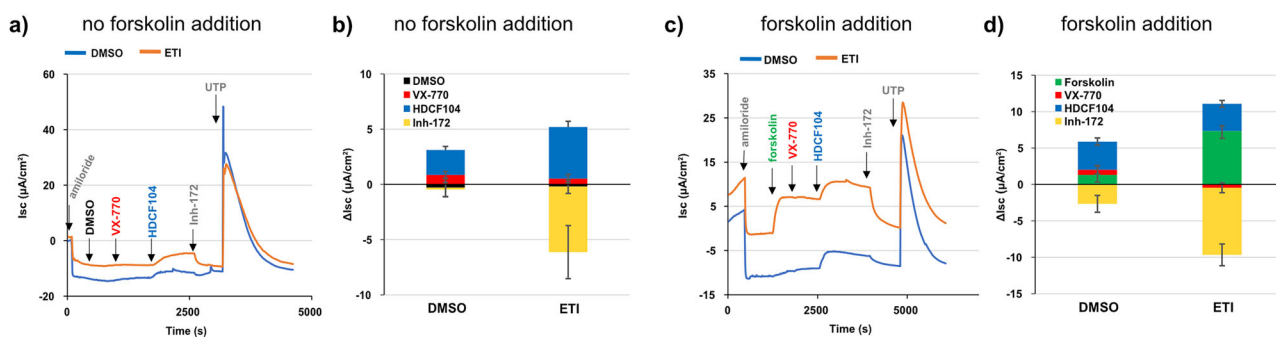
With regard to levels of CFTR activation shown in Figs. 4 and 5, it is reasonable to expect that combination drug treatment may ultimately be required to maximally benefit many refractory CF variants. For example, as introduced above, although ETI has been FDA-approved for N1303K CFTR, a recent clinical trial demonstrated substantive but only partial clinical benefit in vivo among patients with this abnormality⁷⁹. In other words, the magnitude of FEV1 rescue in human subjects with N1303K was less than what has been observed for several CFTR variants following triple combination therapy, and there is no question that further improvements in CFTR activation would be desirable⁷⁹. Drugs in the HDCF series appear well suited to achieve that objective. In the same context, we evaluated a series of common CFTR mutations, many of which have recently been approved for ETI despite comparatively modest in vitro CFTR activation. The magnitude of lung function improvement for these variants in human subjects is not yet known. However, based on preclinical findings from predictive model systems it is likely that—in semblance to N1303K—much greater pulmonary improvement will be possible than with triple combination therapy

alone. Our findings indicate that HDCF104 confers a robust increase in CFTR function when combined with ETI for many of these CFTR abnormalities, and may be useful to improve clinical outcomes (Table 1).

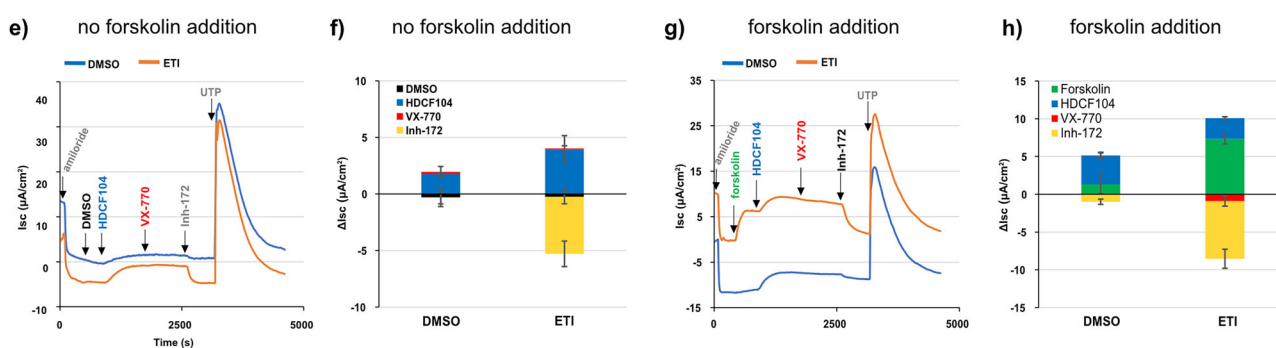
HDCF mechanism of action

Because the HDCF agents were identified by unbiased compound library screening, their mechanism of action was not initially known. Wildtype CFTR and many variants, including less responsive mutations such as N1303K (Fig. 5), as well as CFTR proteins exhibiting severe gating or other defects (Fig. 3 and Table 1) are activated by this drug class. Based on such findings, we initially anticipated a mechanism that involved direct binding of drug to the CFTR protein itself. However, the broad spectrum of activity observed for HDCF agents prompted us to explore a variety of possible contributing pathways. In one series of experiments, a specific PKA inhibitor (H89), which diminishes PKA-induced CFTR phosphorylation and activation, was used to investigate mechanism of action. Electrophysiologic (Ussing) tracings (Fig. 6a, b) demonstrated strong stimulation of WT CFTR by uHTS159 or HDCF104 in a CF bronchial epithelial cell line (CFBE41o⁺). Unexpectedly, a brief (1 h) pretreatment with H89 had minimal impact on VX-770 potentiation, whereas CFTR activation by HDCF analogs was abolished, suggesting a mechanism distinct from that of VX-770 (a drug that directly binds and improves gating of certain CFTR variants but with modest effects on wildtype CFTR in the absence of forskolin/cAMP pre-stimulation) (ref. 7, Fig. 6c), and unpublished observations. Acute addition of H89 partially blocked VX-770-stimulated current, which then recovered spontaneously (Fig. 6c, red tracings). These findings suggest VX-770 opens CFTR by virtue of conformational change that does not confer robust

Addition of acute VX-770 before HDCF104 in the presence or absence of forskolin



Addition of HDCF104 before acute VX-770 in the presence or absence of forskolin



Alternative interventions (1μM VX-661+1 μM Corr4a, 48hr pretreatment)

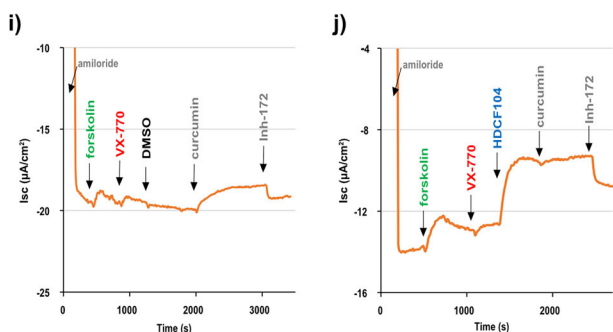


Fig. 5 | Rescue of N1303K CFTR by HDCF104 in primary nasal airway epithelium. a–h HDCF104 activates short circuit current in N1303K/N1303K homozygous and (i, j) N1303K/G542X heterozygous nasal airway epithelial monolayers. Drug additions include amiloride (100 μM), forskolin (5 μM), HDCF104 (10 μM), curcumin (40 μM), Inh-172 (10 μM), and VX-770 (5 μM). UTP, which activates non-CFTR dependent anion transport, was tested at 100 μM. Corrector 4a has been

shown to stabilize CFTR. Curcumin potentiates certain CFTR variants such as W1282X (Wang et al., *PLoS One*, 2016). $n = 3$ biological replicates per condition. ETI (48 h) = elxacaftor (3 μM)—tezacaftor (3 μM)—ivacaftor (5 μM). $p < 0.03$ for HDCF104 plus ETI versus previous addition. Statistics were by Excel Data Analysis ToolPak, T -test too. Figures plotted with Microsoft Excel; plots arranged and presented using Affinity Designer software.

R-domain phosphorylation, and operates in a manner partially independent of the classical CFTR activation mechanism. The results using VX-770 also contrast studies of acute H89 addition followed by forskolin (Fig. 6a–c, red and black tracings), in which forskolin-stimulated CFTR chloride current is abrogated.

To further investigate the role of phosphorylation during CFTR activation by the HDCF class, we conducted biochemical measurements of the CFTR regulatory domain (R-domain). PKA-dependent R-domain phosphorylation represents a key first step during CFTR ion channel gating⁸⁰. In particular, CFBE cells expressing WT CFTR were treated with uHTS159 and surface CFTR biotinylated, followed by immunoprecipitation. CFTR was then probed with four phospho-sensitive anti-CFTR antibodies, as well as an anti-CFTR antibody raised against a nonphospho-epitope. Western

blot and percent R-domain phosphorylation by PKA were determined (Fig. 6d, e), and showed enhanced phosphate addition to serine at R-domain position 813, visualized by diminished binding of UNC217 antibody. The same CFTR position (serine 813) was recently studied using Cryo-EM by Fiedorczuk K et al. and shown to be in close proximity to a PKA-C binding site, suggesting this location may be particularly important for initiating R-domain phosphorylation and CFTR channel activity⁸⁰. H89 treatment led to decreased CFTR phosphorylation as judged by increased antibody binding at the same epitope. A comparable alteration was noted at serine 700 after uHTS159 treatment. Historically, certain phospho-epitopes (serines 737 and 768) have been identified as inhibitory⁸¹ of CFTR gating. These two positions appeared less sensitive to uHTS159-induced phosphate addition. In contrast, saturating concentrations of forskolin strongly enhanced

Table 1 | A series of CFTR variants tested in the FRT model

	10 μ M Forskolin		25 nM Forskolin		
	DMSO	+HDCF104	DMSO		
			VX-661 + VX-445 + VX-770		+HDCF104
Wildtype	100.0%	100.0%			
F508del		2.4%	4.1%	32.0%	105.4%**
I507del	0.0%	0.0%	0.0%	0.0%	0.0%
R334W	7.3%	7.9%	6.3%	5.9%	8.9%**
R560T	0.0%	0.0%	0.4%	1.1%	1.2%
R1066C	0.1%	0.0%	0.0%	0.7%	1.2%*
V520F	0.4%	0.3%	0.4%	10.5%	20.6%*
A559T	0.0%	0.0%	0.0%	0.1%	0.3%
Y569D	0.0%	0.1%	0.3%	0.2%	0.7%
L1065P	0.2%	0.1%	1.0%	34.2%	57.8%*
L467P	0.6%	1.8%	2.0%	1.8%	2.1%
L558S	0.4%	1.8%	4.5%	5.4%	12.0%*
I1234V	2.8%	2.9%	6.6%	7.4%	11.5%*
L927P	12.0%	1.1%	8.8%	5.2%	21.1%**
L227R	0.1%	0.9%	3.6%	0.4%	4.7%*
Q359K/T360K	13.9%	0.8%	4.7%	5.9%	38.1%*
A561E	0.6%	0.6%	1.5%	3.4%	8.4%**
R560K	0.0%	0.0%	0.0%	1.9%	0.0%
W1098R	0.4%	0.5%	0.9%	1.1%	2.5%*
W57G	0.5%	0.5%	1.1%	1.0%	1.2%
H609R	2.6%	0.0%	8.0%	18.4%	53.8%**
P99L	1.3%	5.0%	18.6%	18.6%	45.8%*
D513G	2.2%	2.8%	7.8%	30.8%	64.2%**
Y563D	0.0%	0.0%	0.0%	0.1%	0.0%
S1118F	3.8%	1.7%	7.2%	6.8%	23.1%**
R560S	0.3%	0.0%	0.5%	0.8%	1.7%*
A613T	3.2%	2.2%	6.2%	21.4%	26.5%
G91R	1.1%	1.8%	0.3%	20.2%	37.5%*
L102R	0.1%	0.2%	0.9%	17.5%	18.6%
M1101R	0.0%	0.0%	0.8%	0.5%	1.1%

Percent of wildtype CFTR activity in the presence or absence of ETI (24 h with VX-661 and VX-445, followed by acute addition of VX-770 during Ussing chamber measurements +/- HDCF104 are shown). Lines in bold font indicate activation of mutant CFTR beyond a 10% threshold of wildtype function often used in the FRT model to predict clinical benefit. Note: These experiments utilized a much lower forskolin concentration (25 nM) than the pharmacologic levels (5–10 μ M) commonly used for testing new modulator compounds^{16,17,21,37,89}. Stimulation of CFTR in FRT cells using nominal forskolin represents a stringent model for evaluating drug activation of mutant CFTR. * $p < 0.05$; ** $p < 0.005$ for HDCF104 stimulation following ETI versus vehicle; $n = 3$ –4 biologic replicates per condition; statistics were by t -test.

phosphorylation of the two inhibitory sites (not shown), indicating the mechanistically complex process of CFTR domain rearrangement and channel opening elicited by forskolin as opposed to HDCF104 or uHTS159. Together, the findings point to CFTR stimulation by HDCF compounds as dependent on PKA phosphorylation at specific positions within the regulatory domain.

We next examined whether PDE inhibitors would influence WT CFTR activation by HDCF compounds in primary human bronchial epithelial cells (HBEC). Administration of IBMX (a broad-range PDE inhibitor) prior to HDCF104 (10 μ M) completely eliminated CFTR ion transport elicited by HDCF104 (Fig. 7a, b). Rolipram, a specific PDE4 blocker, also abolished HDCF104 stimulation (Fig. 7c). Similar findings were obtained at higher

concentrations of HDCF104 (100 μ M, Fig. 7d–f). Cilostamide (a specific PDE3 inhibitor) did not activate CFTR either prior to or after HDCF104 administration (Fig. 7g, h). These data indicate that compounds in the HDCF104 class⁷⁵, which have been specifically optimized for CFTR activation, act as PDE inhibitors to elevate cAMP intracellularly and compartmentally^{82–85} to phosphorylate the CFTR regulatory domain.

We also evaluated the HDCF series by a conventional cell-free enzymatic assay using purified PDE protein in which luciferase serves as an ATP “sink” over the linear range of luminescence. Because ATP is a substrate for PKA, inhibition of PDE4 (with a resultant rise in cAMP and PKA activity) elicits a decrease in ATP available to luciferase and decreased assay luminescence (Fig. 8). The inhibitory profile of HDCF104 showed IC_{50} of 7.5 μ M for purified PDE4B—a concentration that agrees well with half-maximal activation levels of HDCF104 needed to stimulate wildtype CFTR in primary airway cells (Fig. 2). We note that drug concentrations of this magnitude are commonly used for mucosal administration to the human respiratory tract (unlike oral agents, where greater potency may be required but with an associated risk of systemic toxicity). For example, levels of approved antibiotics or other compounds administered by aerosol to cystic fibrosis lungs are often at a concentration range of 100–200 μ M in CF airway secretions in vivo^{86,87}.

Finally, for many other PDE inhibitors—including those approved for clinical use—cross-reactivity with more than one PDE isoform is commonly observed. As case in point, in addition to strong blockade of PDE4s, modest inhibition of PDE5 by HDCF104 has also been identified. A plan for nebulized airway delivery of HDCF104 is intended to minimize PDEi-related off-target toxicities. Extensive safety analysis of the compound in two mammalian species has been presented to FDA as part of pre-IND interactions. The findings to date support future use of aerosolized HDCF104, including absence of PDE-related or other safety concerns.

Discussion

This report describes a novel drug scaffold developed specifically for mucosal activation of CFTR and well suited for testing as innovative therapy in airway diseases caused by deficient CFTR. Compounds in this class are intended to promote mucus clearance in people with CF or non-CF lung and sinonasal conditions. For engineered FRT cells and primary airway epithelial monolayers (two models recognized by FDA as relevant to registration of CFTR activators), drugs in the HDCF class markedly augment mutant CFTR activity beyond what can be achieved with HEMT alone (e.g., variants such as F508del, G551D, and many others), while also providing a means to strongly stimulate wildtype CFTR. Importantly, the compounds also increase short circuit current in cell monolayers encoding variants for which CFTR modulators are not considered “highly active” (e.g., N1303K⁷⁹; also Table 1). Agents in the HDCF series exhibit several advantages relevant to treatment of CF and non-CF respiratory conditions including: (1) novel mechanism of action that does not overlap with currently approved CFTR modulators, (2) topical (mucosal) route of administration intended to minimize systemic exposure, (3) the ability to activate mutant CFTR (including “corrected” F508del CFTR) to levels that exceed what can be achieved with modulators alone, and (4) expected amelioration of immune-mediated pulmonary injury (see below).

In our studies, we used H89, a selective PKA inhibitor, to determine whether the new compound class acts through a mechanism involving PKA and CFTR phosphorylation. Administration of H89 following either HDCF104 or uHTS159 abolished CFTR stimulation, supporting a mechanism dependent upon R-domain phosphorylation (Fig. 6a, b). Moreover, we show that the compound series directly inhibits purified PDE4 enzyme (Fig. 8) and biochemically phosphorylates the CFTR R-domain (Fig. 6). In addition, HDCF agents stimulate CFTR activity in a dose-dependent manner using relevant cell-based systems (an effect expected to increase cAMP/PKA and promote R-domain phosphorylation) (Figs. 1 and 2) and exhibit a cross-reactive mechanism of action with other PDE inhibitors, including rolipram, a specific PDE4i (Fig. 7). In unpublished studies, we have shown that HDCF compounds elevate cellular cAMP

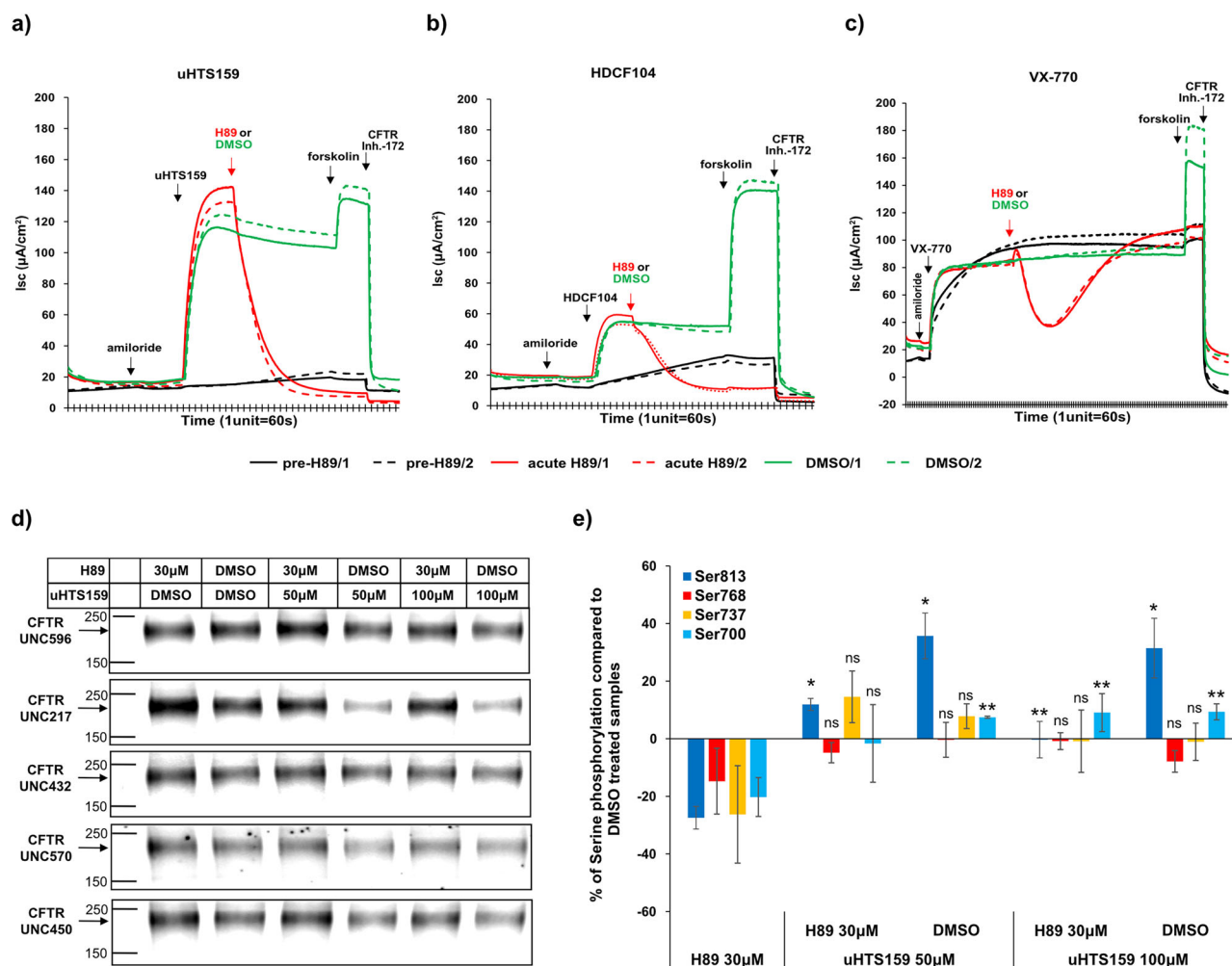


Fig. 6 | Wildtype CFTR activation and R-domain phosphorylation by novel CFTR activators. **a, b** CFTR-dependent chloride currents stimulated by saturating concentrations of uHTS159 or HDCF104 in CFBE41o- cells were inhibited by acute or more chronic addition of H89, a specific PKA blocker. **c** VX-770 displays a distinct activation pattern that was only transiently inhibitable by acute H89. **d** Western blot analysis demonstrates dose-dependent phosphorylation of WT CFTR by uHTS159 utilizing a phospho-sensitive CFTR antibody (UNC217) in comparison with nonphospho-sensitive CFTR detection (UNC596; an antibody recognizing an epitope in NBD2). Phosphorylation diminishes UNC217 affinity at a specific CFTR R-domain epitope (aa 807–819). One additional PKA phosphorylation site assayed by this approach

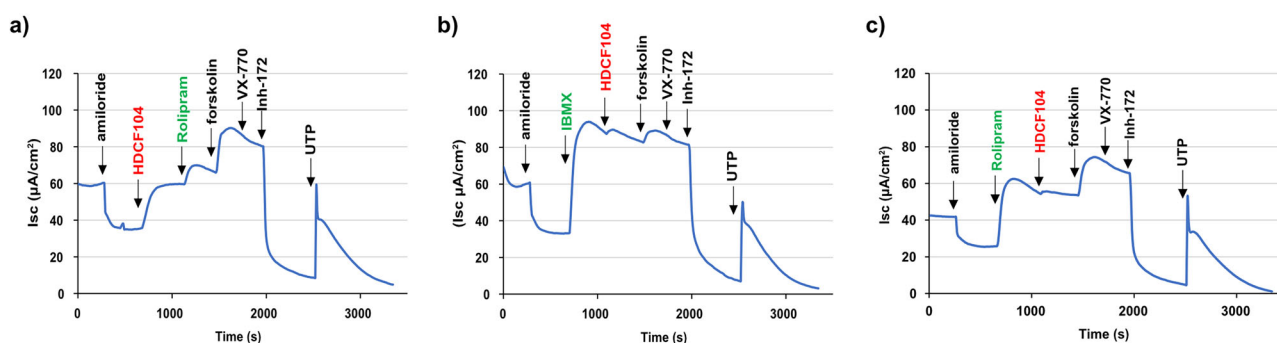
showed evidence of uHTS159 dependent phosphorylation (UNC450, aa 696–705). **e** Density of protein bands was quantified with Image Lab software (Bio-Rad) and plotted as % phosphorylation normalized to UNC596 and vehicle control (DMSO) (Methods). *n* = 3 biological replicates per condition. Significance levels determined by one-way ANOVA. **p* ≤ 0.006; ***p* ≤ 0.04; NS not significant compared with H89 alone. Statistics were by Excel software, ANOVA single factor function. Panels (a–c) plotted using Microsoft Excel; (d) blot images generated with Image Lab 5.2 software, edited with Adobe Photoshop 2025, and labeled using Microsoft PowerPoint; (e) plotted with Microsoft Excel. All plots were arranged and presented using Affinity Design software.

in epithelial cells and fail to activate ΔR CFTR – findings that provide additional support for dependence on R-domain phosphorylation. Future studies using CFTR constructs encoding amino acid substitutions at consensus PKA sites may offer a means to characterize the R-domain conformational changes that underlie CFTR folding and gating after treatment with HDCF compounds.

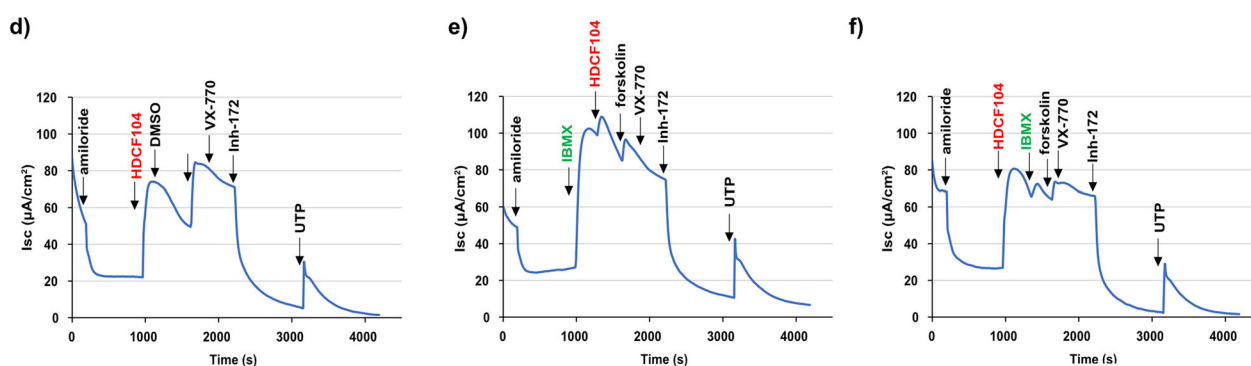
A functional threshold in primary airway epithelial cells of 10–30% wild-type CFTR activity is often viewed in the field as predictive of respiratory benefit among PwCF^{4,20,21,37,88,89}. In vitro studies of sinonasal epithelia show that establishing a benchmark can be somewhat imprecise, particularly due to significant variability of wildtype CFTR within cell monolayers obtained from nasal airways. In our laboratory, 10–15 μA/cm² is frequently observed in epithelial monolayers derived from non-CF nasal tissue samples, although much larger CFTR currents can be obtained in cells from certain individuals^{4,21,36,37}. As described above, model systems such as FRT lines have also been very useful for predicting respiratory improvement, and are recognized by the FDA for this purpose, particularly at CFTR activation levels

above 10% of wildtype^{4,16,17,20,21}. These thresholds suggest that for commercial and academic groups developing CFTR activators for rare variants, the impact of a molecule like HDCF104 could be transformational as a means to maximally stimulate mutant CFTR. Among gating mutations such as G551D, we show that HDCF104 can function either as monotherapy or provide a strong complement to FDA-approved modulators, leading to activity far above 10% of wildtype when combined with ETI (Fig. 3). For other variants, including N1303K (Fig. 5 and Table 1), a combination of drug treatments (similar to what has been required for F508del rescue) will likely be needed to achieve maximal benefit. Data from several laboratories, including recent clinical findings from our group⁷⁹, indicate that N1303K stimulation by ETI occurs at levels that provide incomplete lung function improvement. Our current results demonstrate that HDCF104 confers activation of the N1303K variant to significantly above 30% of wildtype when combined with available modulators (Fig. 5). For PwCF who fail, are ineligible, or cannot gain access to HEMT, compounds in the HDCF104 series are anticipated to provide considerable value.

Sequential addition of HDCF104 at 10µM concentration with PDE inhibitors



Sequential addition of HDCF104 at 100µM concentration with PDE inhibitors



Specific PDE3 inhibitor (Cilostamide) does not influence CFTR activation by HDCF104

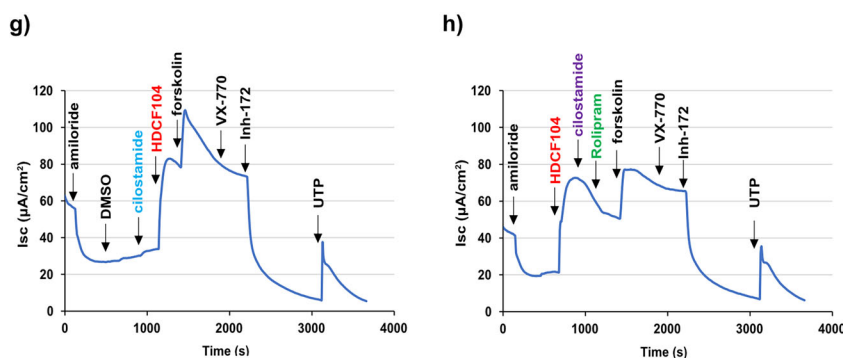


Fig. 7 | HDCF104 is blocked by pretreatment with PDE4 inhibitory compounds.

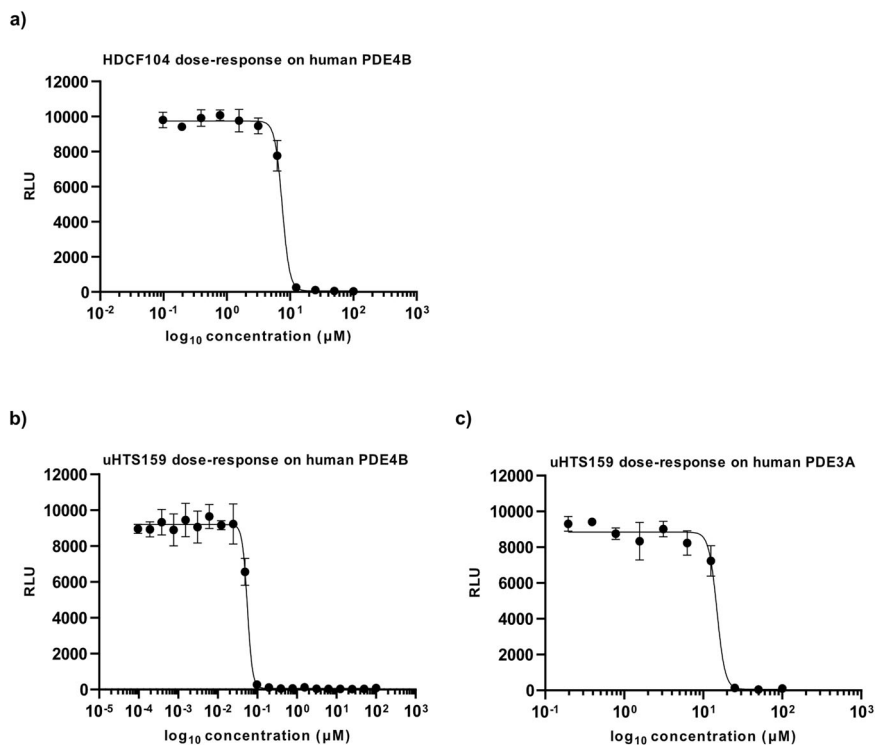
a–h Loss of HDCF104 activity demonstrated following pretreatment with IBMX or rolipram, but not a PDE3 inhibitor (cilostamide). Pretreatment with HDCF104 attenuated the activity of rolipram and IBMX. Compounds added acutely include amiloride (100 µM), forskolin (5 µM), Inh-172 (10 µM), and UTP (100 µM). To assure saturating drug concentrations, IBMX (broad spectrum PDE inhibitor) was

added at 200 µM, whereas rolipram (PDE4 specific inhibitor) and cilostamide (PDE3 inhibitor) were tested at 10 µM and 1 µM, respectively. HDCF104 was added at 10 µM in Panels (a) through (c), and 100 µM in Panels (d) through (h). The studies were conducted in non-CF primary nasal airway epithelial cells and show robust wildtype CFTR activation mediated by HDCF104. All panels were plotted using Microsoft Excel; plots were arranged and presented using Affinity Design software.

Enzymatically and pharmacologically distinct members of the PDE family—including PDE subtypes 1–11—are found in nearly all mammalian cells or tissues, and hydrolyze/inactivate cyclic nucleotides. Blockade of certain PDEs elicits compartmental elevation of cyclic AMP, which (in a manner dependent on PDE category, tissue, cell type, etc.) can strongly enhance protein kinase A activity and stimulate CFTR through phosphorylation of specific positions within the regulatory or other protein domains^{90,91}. PDE4is have been previously recognized as activators of wildtype CFTR in airway epithelium^{90,92–97}. Even within the same PDE4 subclass, however, differences in airway epithelial partitioning, PDE4i

affinity, subcellular compartmentalization, plasma membrane permeability, cellular compound clearance, and other features may influence the magnitude and duration of effects on CFTR gating – including distinctions from use of forskolin to stimulate cAMP/PKA. We believe the application of high throughput assays and discovery of new PDE4is specifically tailored for CFTR activation in primary airway epithelium, as described here, will be essential for optimizing interventions that address CFTR deficiency in a number of human disease states. A large body of clinical, biophysical, biochemical, and genetic evidence supports that contention^{51–65,68–72,82–85}. As indicated above, emerging data establishes that several common muco-

Fig. 8 | Purified PDE enzyme assays to test inhibition by HDCF104 or uHTS159. **a** Dose-response curve of HDCF104 as a blocker of PDE4B (IC_{50} of HDCF104 = $7.5 \pm 0.32 \mu\text{M}$). **b** Shows approximately two orders of magnitude lower IC_{50} of uHTS159 compared to HDCF104. **c** More modest inhibition of PDE3A was observed (in the μM range). (IC_{50} of uHTS159 = $0.06 \pm 0.0021 \mu\text{M}$ as an inhibitor of PDE4B and $14.9 \pm 1.57 \mu\text{M}$ for PDE3A). HDCF104 did not demonstrate inhibition of PDE3A at concentrations up to $100 \mu\text{M}$ (data not shown). IC_{50} of compounds calculated using GraphPad Prism 10 software, which was also employed for plotting all panels. Panels were arranged and presented with Affinity Designer software.



obstructive respiratory conditions, including non-CF bronchiectasis (NCFB) and CRS, share key pathogenic and mechanistic features with CF lung disease^{51–65,68–70,72}. For patients with these illnesses, stimulation of CFTR and improved mucociliary clearance are likely to result from treatment with agents such as those described here.

PDE4 blockade has been established as a clinically effective means to diminish immune hyper-responsiveness within human lungs^{98–107}. In recent preclinical studies, PDE4 inhibitors were shown to blunt inflammation in models of asthma, mediate bronchodilation and suppress inflammatory response associated with COPD, diminish immune activity of eosinophils in human bronchial specimens from subjects with bronchitis, repress inflammation in human airway tissue, and mediate numerous additional anti-inflammatory effects of this sort^{98–107}. From the standpoint of the current findings, anti-inflammatory effects attributable to PDE4i blockade would be in addition to ways in which CFTR activation, itself, may diminish cytokine release from airway cells¹⁰⁸. AstraZeneca markets the oral PDE4 inhibitor roflumilast for treatment of COPD^{101,109–111} as a broad-spectrum compound that blunts several inflammatory pathways. The oral drug has been significantly limited by gastrointestinal and other systemic sequelae, including pronounced diarrhea. Side effects due to oral (systemic) PDE4is have led commercial developers to advance aerosolized PDE3 and PDE4 blockers for treating inflammation in COPD. Such drugs have shown considerable promise—without significant GI or other limiting complications^{98,108,112–119}.

Mucosal administration of CFTR activators represents an emerging area of interest for treatment of chronic respiratory conditions such as CF. Although wildtype CFTR stimulation by PDE4is is well described, activation of disease-causing CFTR variants has been less well studied in the CF modulator era. Recent work from Hanrahan and colleagues demonstrated that RPL554 (a PDE3»PDE4 inhibitor) enhanced G551D and a number of other mutant CFTRs in the FRT model^{120–122}. Based on high-throughput compound analysis, Beekman et al. recently concluded that certain PDE4i molecules rank among the most promising for stimulating mutant CFTR¹²³. Findings presented in the current study highlight the effectiveness of PDE4i compounds for activating both wildtype and variant CFTR at levels expected to provide clinical improvement.

In summary, we describe a novel class of CFTR gating enhancers discovered using unbiased cell-based screening. The compounds exhibit robust activity in both FRT and primary airway cell models with a molecular target (CFTR) already well validated by an extensive body of genetic, biophysical, and clinical literature. Drugs described here are expected to have a therapeutic role not only in cystic fibrosis pulmonary and sinonasal disease, but also much more common illnesses associated with CFTR deficiency such as non-CF bronchiectasis and chronic sinusitis. The mechanism of action for the new drug series has been identified as phosphodiesterase blockade, and lead agents have been optimized specifically for topical use in human airway epithelium to activate CFTR-dependent mucociliary clearance. Importantly, since both mutant and wildtype CFTR are strongly enhanced, both CF and non-CF muco-obstructive illnesses are appropriate targets for the new drug class. Compounds in the same mechanistic category have already been shown to provide clinically beneficial anti-inflammatory properties in diseases such as COPD, but have not been well tested in CF, NCFB, or CRS. Future clinical development is merited to determine the extent to which mucosal CFTR activators can be used to improve treatment of inherited and acquired CFTR deficiency in human airways.

Methods

Cell culture

Fischer rat thyroid (FRT) cells expressing diverse CFTR variants were cultured in Coon's modification of Nutrient Mixture F-12 Ham media (Sigma, F6636) supplemented with 2.68 g/l sodium-bicarbonate in the presence of 5% fetal bovine serum (Gibco). The cells were maintained at 37 °C under humidified, 5% CO₂ – 95% air. For electrophysiology studies, polarized cell cultures were established, with FRT cells seeded onto Transwell permeable supports (Corning, 3470). Cells were cultured for 5 days until a mature monolayer was established (with transepithelial resistance of at least 400 Ω × cm²).

Primary nasal or bronchial epithelial cells carrying diverse CFTR variants (hNEC N1303K/N1303K patient K and hNEC N1303K/G542X patient H, SickKids, Toronto, Canada; hBEC F508del/F508del product code EP57AB, Geneva Switzerland; hBEC healthy donor FC0035, Lifeline Cell Technology, Frederick, MD; hNEC healthy donor patient 21.A2, University

of Alabama, Birmingham) were expanded using PneumaCult-Ex Plus medium (StemCell Technology; 05040) on collagen-coated flasks, followed by seeding onto collagen-treated Transwell inserts. Resulting monolayers were cultured for 2–4 days under submerged conditions, and subsequently transitioned to air-liquid interface (ALI) utilizing PneumaCult ALI medium (StemCell Technology; 05001). Cells were then maintained at 37 °C under humidified, 5% CO₂–95% air until well-differentiated epithelium was established, typically within 3 weeks of seeding.

CFBE41o⁻ cells with stable expression of WT CFTR¹²⁴ were cultured in MEM medium supplemented with L-glutamine and 10% FBS. Cells for surface CFTR biotinylation studies or electrophysiology assays were seeded onto Transwell permeable supports (Corning, 3450 or Corning, 3470) and grown until well polarized under submerged conditions.

FRT clonal cell lines with stable expression of CFTR variants

Fischer Rat Thyroid cells (FRT) were used to establish an FRT Flip-In host cell line utilizing pFRT/lacZeo vector. WT or mutant human CFTR cDNAs were incorporated at a single insertion site according to the manufacturer's protocol (Thermo Fisher Scientific). Cell clones demonstrating comparable levels of CFTR mRNA expression (within 0.5–1.5 fold of wildtype) were selected from each CFTR variant pool to minimize differences in response after modulator treatment (a confounder that may otherwise result from distinct CFTR protein expression levels).

Transient expression of CFTR variants in FRT cells

Rare CFTR missense variants were selected from among those listed in the CFTR2 mutation database (<https://cftr2.org>). CFTR constructs encoding these molecular defects contained nucleotide changes identical to those found in patients (including methionine at position 470) and were cloned into the pDNA5/FRT mammalian expression vector (Thermo), as previously described¹⁷. Expression was under regulatory control of the CMV promoter. Variants were transiently expressed and studied by short-circuit current (Ussing chamber) assays to investigate CFTR function.

Electrophysiology

A total of 1.2×10^5 FRT cells were seeded and cultured on each Transwell permeable support (Corning 3470). One day after seeding, cells on each filter were transfected with 0.5 µg of CFTR plasmid using Lipofectamine 3000 (Thermo). The media was changed 72 h post-transfection, and cells were treated with the indicated CFTR modulators for 24 h. CFTR functional assays were performed on day 5 post-seeding in the Ussing chamber¹⁷. Eight filter supports were analyzed for each CFTR variant by Ussing chamber experiments, and each experiment included wildtype CFTR-expressing cells as positive controls.

Ussing chamber analysis

Monolayers were mounted onto an EasyMount Ussing Chamber System (Physiologic Instruments) and bathed apically in low chloride Ringer's solution containing: 140 mM Na-gluconate, 1.2 mM NaCl, 25 mM NaHCO₃, 3.33 mM KH₂PO₄, 0.83 mM K₂HPO₄, 1.2 mM CaCl₂, 1.2 mM MgCl₂, and 10 mM D-glucose (pH 7.4). Basolateral solutions contained 120 mM NaCl, 25 mM NaHCO₃, 3.33 mM KH₂PO₄, 0.83 mM K₂HPO₄, 1.2 mM CaCl₂, 1.2 mM MgCl₂, and 10 mM D-glucose (pH 7.4) to establish a chloride gradient. Temperature of bathing solutions was maintained at 37 °C and stirred by bubbling through 5% CO₂/95% O₂. Test compounds were dissolved in DMSO at 1000X concentration and diluted as described. For bioelectric measurements, once a baseline had been established, 100 µM amiloride (MilliporeSigma, A7410) was applied to both the apical and basolateral surfaces to inhibit epithelial sodium channel (ENaC) activity. CFTR protein was activated by applying forskolin (MilliporeSigma, F3917), typically 5 µM (FRT cells) or 10 µM (primary cells), followed by addition of test compound. In some experiments, VX-770 (SelleckChem, S1144) was added apically at 5 µM. At the end of each recording, CFTR_{inh}-172 (10 µM; MilliporeSigma C2992) was

applied at the mucosal surface to inhibit CFTR currents. In primary cell experiments, UTP (stimulates non-CFTR chloride channels; VWR, 0145) was added to confirm monolayer integrity. Rolipram (Selleck S1430), Cilostamide (Enzo BML-PD125) and IBMX (MilliporeSigma, I7018) were used to determine importance of phosphodiesterase activity. Across all experiments, short circuit current was recorded under voltage clamp conditions with change in I_{sc} following acute treatment calculated and expressed as mean ± SD.

CFTR phosphorylation

CFBE41o⁻ cells expressing WT CFTR were seeded onto permeable supports to generate polarized monolayers grown under immersed conditions (typically ~6 days until mature). Test compounds (e.g., uHTS159, ChemSpace US Inc., CSSS00160668684) and/or PKA inhibitor (H89 2HCl, Selleckchem, S1582) were applied bilaterally for 15 min. At the end of each incubation, monolayers were transferred to ice-cold PBS and oxidation of glycan chains followed by biotinylation with Hydrazide-LC-Biotin (Thermo Scientific, 21340)¹²⁵. Cells were next lysed in RIPA buffer supplemented with protease and phosphatase inhibitor, and equal amounts of total protein immunoprecipitated using Streptavidin magnetic beads (Thermo Scientific (Pierce), 88817). Immunoprecipitated proteins were resolved on 8–16% Novex, Tris-Glycine SDS-PAGE (Thermo Fischer Scientific, XP08165BOX) and transferred to nitrocellulose membrane (Bio-Rad, 1620112). CFTR proteins were blotted with specific antibodies UNC596, UNC217, UNC432, UNC570, UNC450 (CFTR Antibody Distribution Program, UNC-Chapel Hill, Dr. Martina Gentsch), followed by detection with HRP-goat-anti-mouse antibody (Abcam, ab205719). Specific bands were visualized with SuperSignal ELISA Femto Maximum Sensitivity Substrate (Thermo Scientific, 37074). The ChemiDoc MP System was used for imaging. Band densities were quantified with ImageLab 5.2.1 software, and normalized to corresponding bands from UNC596 (as loading control) followed by normalization to DMSO-treated conditions. Results were plotted and expressed as mean ± SEM. Single-factor ANOVA was used to determine significance.

IC₅₀ determination and in vitro PDE enzyme activity assay

PDE-Glo Phosphodiesterase Assay (Promega, V1361), recombinant human PDE4B (Abcam, ab125582), and recombinant human PDE3A (Abcam, ab125582) were used in a 384-well plate format to determine IC₅₀ of test compounds according to the manufacturer's protocol. Recombinant PDE enzymes were titrated to establish optimal concentrations (determined for PDE4B or PDE3A as 3 ng/reaction) and reaction times (identified as 20 min). Test compound was added to each plate in serial dilution and relative luminescent signal was detected using a BioTek Synergy Neo2 plate reader. For calculating IC₅₀ of test compounds, relative luminescent signal was plotted against concentration with GraphPad Prism 10 software (Sigmoidal, 4PL, X as concentration function).

Statistical methods

Descriptive statistics (mean, median, standard deviation and standard error of mean) were applied to in vitro continuous data (e.g., I_{sc}). All in vitro studies were conducted in paired fashion (same day, same cell passage). A *p*-value (α) of less than or equal to 0.05 was considered statistically significant, based on single-factor ANOVA, paired *t*-tests and/or Mann-Whitney analysis.

Data availability

All data is provided within the manuscript.

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Competing interests

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Additional information

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