Supplemental Data

Table S1

Table S1. Genotypes of Human CF Donors		
	Donor ID	CFTR Genotype
RNA-seq		
	CFB0219	$\Delta F508/3659 delC$
	CFB0319	$\Delta F508/N1303K$
	CFB0419	$\Delta F508/\Delta F508$
	CFB2518	$\Delta F508/1336K$
	CFB2818	$\Delta F508/R560T$
	CFB3218	$\Delta F508/\Delta F508$
Other studies		
	CFB0219	$\Delta F508/3659 delC$
	CFB1419	$\Delta F508/3659 delC$
	CFB1519	$\Delta F508/2184$ -del-A
	CFB1719	$\Delta F508/N1303K$
	CFB2019	$\Delta F508/3849 + 10kbC > T$
	CFB2219	$\Delta F508/\Delta F508$
	CFB2319	$\Delta F508/R75X$
	CFB2419	$\Delta F508/E60X$
	CFB2619	$\Delta F508/unknown$
	CFB2819	$\Delta F508/\Delta F508$
	CFB2919	$\Delta F508/\Delta F508$
	CFB3019	$\Delta F508/G542X$
	CFB3119	$\Delta F508/\Delta F508$
	CFB0420	$\Delta F508/\Delta F508$
	CFB0520	$\Delta F508/\Delta F508$
	CFB0720	$\Delta F508/\Delta F508$



Fig. S1. Effect of TNF α and IL-17 on amiloride-sensitive and DIDS-sensitive I_{SC} in CF airway epithelia. Primary differentiated human CF airway epithelia were treated with TNF α , IL-17, or both for 24 hours. All epithelia were also exposed to a triple combination of CFTR modulators comprising elexacaftor, tezacaftor, and ivacaftor. Epithelia were assayed in Ussing chambers and short-circuit current (I_{SC}) was recorded. A) Changes in I_{SC} after addition of apical amiloride. B) Changes in I_{SC} after addition of apical DIDS. Each data point represents an epithelium from a different donor. N = 5 different donors. Data are shown as the mean ± SEM. Statistical significance was tested using repeated-measures ANOVA and post-test Tukey's. **P* < 0.05. Amil = amiloride; DIDS = 4,4'-diisothiocyano-2,2'-stilbenedisulfonic acid.



Fig. S2. Effect of genotype on CFTR activity in TNFα+IL-17-treated epithelia. Primary differentiated human airway epithelia were treated with TNFα+IL-17 for 24 hours. CF epithelia were also exposed to a triple combination of CFTR modulators comprising elexacaftor, tezacaftor, and ivacaftor. Epithelia were assayed in Ussing chambers and short-circuit current (I_{SC}) was recorded. **A**) Comparison of CFTR-mediated I_{SC} in non-CF versus CF epithelia exposed to CFTR modulators. N = 7 non-CF, and 5 CF donors. **B**) CFTR-mediated I_{SC} in CF donors either homozygous ($\Delta F/\Delta F$) or heterozygous ($\Delta F/Other$) for $\Delta F508$ allele and treated with CFTR modulators. The dotted line marked by green arrow indicates ΔI_{SC} in CF epithelia in the absence of CFTR modulators. N = 16 homozygous, and 5 heterozygous donors. Each data point represents an epithelium from a different donor. Data are shown as the mean ± SEM. Statistical significance was tested using mixed-effects ANOVA and post-test Tukey's. **P* < 0.05; ***P* < 0.01; *****P* < 0.0001.



Fig. S3. TNF α +IL-17 increase expression of selected mucin (*MUC*) genes. Δ F508-CF epithelia were exposed to TNF α +IL-17 for 48 hours and RNA-seq was performed. N = 6 different donors. Statistical significance was tested using paired Student's *t* test. **P* < 0.05, ***P* < 0.01. TPM = transcripts per million.