

compelling basic science and animal model observations would provide a solid foundation upon which to implement additional trials of “anti-coagulants” in IPF. In conclusion, we have answered the question of the utility of warfarin in progressive IPF, but many questions remain.

Author disclosures are available with the text of this letter at www.atsjournals.org.

IMRE NOTH, M.D.
University of Chicago
Chicago, Illinois

MITCHELL OLMAN, M.D.
Cleveland Clinic
Cleveland, Ohio

ON BEHALF OF THE IPFNET INVESTIGATORS

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A Young Hispanic with c.1646G>A Mutation Exhibits Severe Cystic Fibrosis Lung Disease: Is Ivacaftor an Option for Therapy?

To the Editor:

Cystic fibrosis (CF) is a lethal autosomal recessive inherited disease caused by the loss or dysfunction of the cystic fibrosis transmembrane conductance regulator (CFTR) chloride (Cl⁻) channel activity resulting from mutations (1, 2). More than 1,600 mutations, which can be broadly grouped into six classes, have been identified in the CFTR gene (3). The incidence of CF and the frequency of specific mutations have been found to vary among ethnic populations (4, 5).

We presented a clinical case of a young Hispanic patient with CF heterozygous for c.1521_1523delCTT (p.Phe508del or F508del)/c.1646G>A (p.Ser549Asn or S549N) mutations (please note that the legacy names for the mutations will be used in this letter). This individual was diagnosed with CF at 5 months of age after multiple hospitalizations due to dehydration, failure to thrive, and emesis. She also had a prolonged and persistent cough thought secondary to a diagnosis of pertussis at 4 months of age. At diagnosis, she was also noted to have a hypochloremic, hypokalemic metabolic alkalosis. She was below the fifth percentile on all aspects of her growth curve. Sweat chloride levels were abnormal with values of 87.1 mmol/L (right arm) and 78.2 mmol/L (left arm). She was empirically given pancreatic replacement enzymes and fat soluble vitamin replacement. By the age of 3 years, she was found to be colonized by two strains of nonmucoid *Pseudomonas aeruginosa* as well as a mucoid *Pseudomonas* that was her predominant isolate by 4 years of age. Over the past 2 years, she has had three additional hospitalizations for pulmonary exacerbations (April 2010, April 2011, and October 2011). Pulmonary function tests (PFTs) have continued to worsen with FEV₁ values of 50 to 60% predicted (the most recent FEV₁ prior to hospitalization in October 2011 was 51% predicted) and FEF_{25–75} values in the low end of 20 to 40% predicted (the most recent value was 20% predicted). She has had increasing pulmonary symptoms and has developed worsening sinusitis. Chest computed tomography scan completed in October 2011 showed areas of extensive bronchiectasis and fibrotic change, predominantly affecting the upper lobes. Areas of disease were observed in the lower lobes as well (Figure 1A and Figure E1 in the online supplement). There are numerous areas of “tree-in-bud” opacities present essentially involving all lobes of both lungs (Figure 1A).

F508del is the most common CFTR mutation, with more than 90% of patients with CF carrying it on at least one allele. F508del-CFTR protein is insufficiently folded and is trapped in the endoplasmic reticulum and targeted for degradation (1). Like c.1652G>A (p.Gly551Asp or G551D), S549N mutation occurs in the signature sequence (LSGGQ) of the first nucleotide-binding domain on the CFTR protein (Figure E2A). S549N mutation was first reported by Curtis and colleagues in a Pakistani individual who was homozygous for S549N and demonstrated severe malnutrition, growth retardation, and advanced pulmonary disease (6). S549N mutation was found to be prevalent among Hispanic, African American, and Asian populations (4).

We characterized S549N-CFTR at the protein level to understand its molecular characteristics and the associated severe disease phenotype and, more importantly, to explore using mutation-specific therapy for medical interventions. S549N-CFTR was expressed as a mature form of CFTR (band C) with levels comparable to wild-type CFTR (Figure 1B). Surface labeling assay and fluorescence microscopy data showed that S549N-CFTR was expressed at the plasma membrane of cells (Figures E3A and E3B). However, S549N-CFTR lacks Cl⁻ channel function (Figures E2B and E2C). These molecular characteristics correlate well with the severe CF phenotype we observed and suggest that, like G551D, S549N-CFTR is probably a regulation mutant.

Author Contributions: The manuscript was written by S.Y. and W.Z. supervised by A.P.N. and D.C.S. The project was designed and supervised by A.P.N. S.Y. conducted site-directed mutagenesis and surface labeling, F.A.I.K. performed iodide efflux assays, and H.P. performed immunoprecipitation and Western blotting experiments. A.R. conducted *I_{sc}* measurements. K.A. performed immunofluorescence microscopy studies. C.A.D., S.S., and D.C.S. did the clinical studies. J.C.K. assisted in site-directed mutagenesis. All authors discussed the results and commented on the manuscript.

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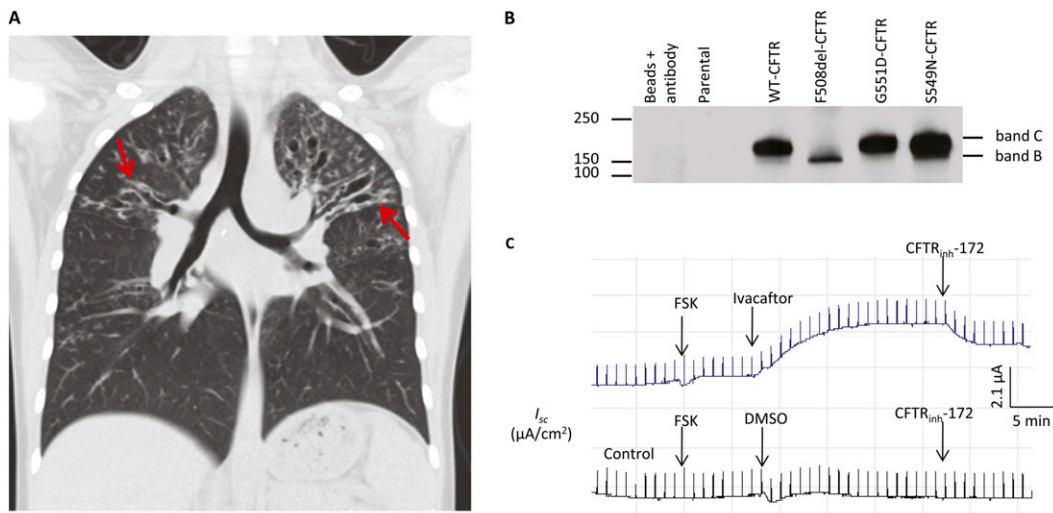


Figure 1. S549-CFTR is a regulation mutant and associates with a severe cystic fibrosis (CF) lung disease in a young Hispanic patient with CF. (A) Chest computed tomography (coronal view) shows extensive bilateral upper lobe bronchiectasis (arrows, left > right) with relatively less involvement of the lower lung fields. (B) S549N-CFTR has normal protein expression levels. HEK-293 cells were transiently transfected with pcDNA3 containing WT, S549N, G551D, or F508del cDNA, lysed 48 hours after transfection, immunoprecipitated with α -24-1 monoclonal antibody and Western

blotted for CFTR by using α -NBD-1-R polyclonal antibody. (C) Ivacaftor (100 μ M) restored Cl^- channel function of CFBEo $^-$ cells. The cells were grown on Costar Transwell permeable supports until they reached resistance of more than 1,400 Ω and then mounted in an Ussing chamber. FSK (10 μ M) was used to activate CFTR channel function and CFTR $_{\text{inh}}-172$ was used to verify the observed signal were CFTR mediated. DMSO (the solvent for Ivacaftor) was used as a negative control. See Reference 12 for technical details. DMSO = dimethyl sulfoxide; WT = wild-type; FSK = forskolin.

Ivacaftor (VX-770) is a CFTR potentiator that can restore the defective gating of G551D and several other regulation mutants (7–9) and has been approved by U.S. Food and Drug Administration for treating patients with CF aged 6 and older with G551D mutation (10). We expressed S549N-CFTR in polarized human cystic fibrosis bronchial epithelial cells (CFBEo $^-$) and tested the effect of ivacaftor on CFTR channel function. Our results showed that ivacaftor restored the defective channel function of S549N-CFTR (Figure 1C), suggesting that ivacaftor has potential clinical benefit for patients with CF with S549N mutation. Our biochemical findings are consistent with data from a recent study in which ivacaftor was found to potentiate the channel open probability and total Cl^- transport of S549N-CFTR expressed in Fischer rat thyroid cells (9). To further confirm S549N is a regulation mutant and can be rescued by CFTR potentiators, we also tested the effect of another known CFTR potentiator, P1, on the channel function of S549N-CFTR expressed in CFBEo $^-$ cells or human embryonic kidney (HEK-293) cells and found that P1 exerted a potentiating effect similar to that of ivacaftor (Figures E4 and E5).

Because this subject in our study is heterozygous for F508del/S549N mutations, she would be an ideal candidate for the ongoing combinational trials using VX-809 (11) and ivacaftor and, most likely, will benefit from these trials.

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SUNITHA YARLAGADDA, B.Sc., M.Sc.

WEIQIANG ZHANG, Ph.D.

HIMABINDU PENMATS, Ph.D.

AIXIA REN, Ph.D.

KAVISHA ARORA, B.S.

ANJAPARAVANDA P. NAREN, Ph.D.

*The University of Tennessee Health Science Center
Memphis, Tennessee*

FATIMA ANMOL I. KHAN, B.S.

*The University of Tennessee Health Science Center
Memphis, Tennessee*

and

*Christian Brothers University
Memphis, Tennessee*

CATHERINE A. DONNELLAN, M.D.

SAUMINI SRINIVASAN, M.D.

DENNIS C. STOKES, M.D., M.P.H.

*The University of Tennessee Cystic Fibrosis Care and
Research Center at LeBonheur Children's Hospital
Memphis, Tennessee*

JOHN C. KAPPES, Ph.D.

*The University of Alabama at Birmingham
Birmingham, Alabama*

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Rod-like Bacteria and Recurrent Venous Thromboembolism

To the Editor:

A 52-year-old male developed abdominal pain and diarrhea (five to six times daily) accompanied by fever (38.5°C) for 3 days. He was admitted with the preliminary diagnosis of acute enteritis. One week after hospitalization, abdominal pain alleviated, but chest tightness and swelling of the right lower limb occurred. Chest computed tomography and ultrasonography showed acute pulmonary embolism and deep vein thrombosis (DVT) of the right lower limb. After anticoagulation therapy with low-molecular-weight heparin, he was discharged after symptoms improved. However, DVT recurred several times. Abdominal ultrasonography showed suspected thrombosis in the left portal vein, and oral warfarin was administered. Six months after venous thromboembolism (VTE), this patient was again admitted to our hospital due to DVT in the left lower limb. In addition to routine examination and immune examination, electron microscopy was performed to observe the morphology of blood cells, revealing that white blood cell count increased ($11.5 \times 10^9/L$; range: $4-10 \times 10^9/L$); CD_3^+T : 50.6% (range: 60–85%). Levels of CD_4 , CD_8 , CD_{16+56} , and CD_{19} were normal. Expressions of IL-1, IL-2, IL-2 receptor, and tumor necrosis factor- α were normal. IL-4 was 108.9 pg/ml (normal: <39.2), IL-6 was 36.85 pg/ml (normal: <31.2), and IL-8 was 101.41 pg/ml (normal: <48.7).

Electron microscopy showed apoptosis and nucleus disappearance of phagocytes. Rod bacteria-like microorganisms were identified in the phagocytes (Figures 1A–1D). CD_3^+T cells were reduced, but the levels of IL-4, IL-6, and IL-8 increased, suggesting Th1/Th2 imbalance.

Infection causes venous thrombosis in multiple organs (1). Patients with acute pulmonary embolism and chronic thromboembolic pulmonary hypertension had compromised function of CD_3^+T cells and CD_8^+T cells, respectively. The increase in CD_4^+T/CD_8^+T ratio suggested that the occurrence of VTE was related to the immune dysfunction (2, 3). The patient had sometimes withdrawn the intake of warfarin from the first onset of VTE. This may also be a reason of VTE recurrence, and the compromised immune function is an internal factor of VTE recurrence. There is internal correlation between immune disorder and coagulation dysfunction. Patients with inflammatory bowel disorders, Behçet's disease, HIV infection, and other infections exhibit an increased risk of VTE (4). Smeeth and colleagues (5) reported that infectious diseases in a community setting are linked to a transient increase in the risk of VTE, so infection may be a triggering factor in VTE recurrence.

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LEMING WANG, M.D., PH.D.
XIAOYU ZHANG, M.D.
QIANGLIN DUAN, M.D.

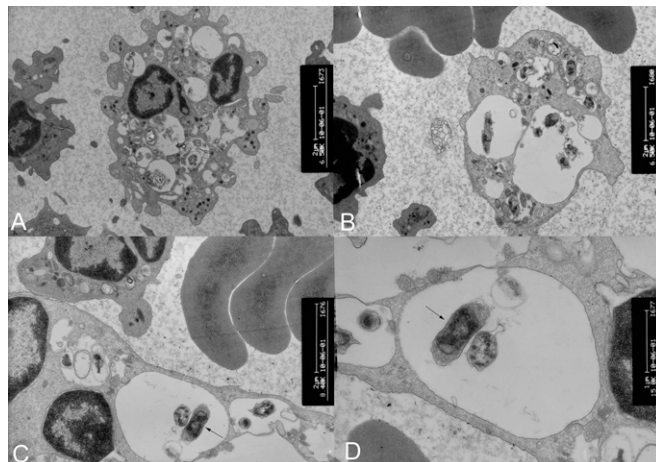


Figure 1. Findings of the neutrophil damage and rod-shaped bacteria-like microorganisms under electron microscope. (A) Apoptotic phagocytes. (B) The nucleus of phagocytes disappeared. (C) Rod-shaped bacteria-like microorganisms were identified in the cytoplasmic cavity of phagocytes (arrow). (D) Rod-shaped bacteria-like microorganisms at a higher magnification (arrow).

WEI LV, M.D.
ZHU GONG, M.D.
YUAN XIE, M.D.
AIBIN LIANG, M.D., PH.D.
YENAN WANG, M.D.
*Tongji Hospital
Shanghai, China
and
Medicine School of Tongji University
Shanghai, China*

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Proposed Alternative Scoring for SAQLI Domain D

To the Editor:

The Sleep Apnea Quality of Life Index (SAQLI) is a disease-specific quality-of-life (QOL) measure that was first developed and evaluated over 10 years ago (1, 2) and has been widely used since that time. One advantage of the SAQLI over other disease-specific QOL measures is that it allows for patients to select up to five “important” symptoms. This provides a tailored