

ARTICLE CFTR is a negative regulator of $\gamma\delta$ T cell IFN- γ production and antitumor immunity

Yuanyuan Duan^{1,2}, Guangqiang Li^{1,2}, Miaomiao Xu^{1,2}, Xiaofei Qi³, Mingxia Deng^{1,2}, Xuejia Lin^{1,2}, Zhiwei Lei², Yi Hu^{1,2}, Zhenghu Jia^{1,2,4}, Quanli Yang^{1,2,4}, Guangchao Cao^{1,2}, Zonghua Liu^{1,2}, Qiong Wen^{1,2}, Zhenhua Li^{1,2}, Jie Tang³, Wei Kevin Zhang⁵, Pingbo Huang⁶, Limin Zheng⁷, Richard A. Flavell ⁰⁸, Jianlei Hao ^{1,2} and Zhinan Yin^{1,2}

CFTR, a chloride channel and ion channel regulator studied mostly in epithelial cells, has been reported to participate in immune regulation and likely affect the risk of cancer development. However, little is known about the effects of CFTR on the differentiation and function of $\gamma\delta$ T cells. In this study, we observed that CFTR was functionally expressed on the cell surface of $\gamma\delta$ T cells. Genetic deletion and pharmacological inhibition of CFTR both increased IFN- γ release by peripheral $\gamma\delta$ T cells and potentiated the cytolytic activity of these cells against tumor cells both in vitro and in vivo. Interestingly, the molecular mechanisms underlying the regulation of $\gamma\delta$ T cell IFN- γ production by CFTR were either TCR dependent or related to Ca²⁺ influx. CFTR was recruited to TCR immunological synapses and attenuated Lck-P38 MAPK-c-Jun signaling. In addition, CFTR was found to modulate TCR-induced Ca²⁺ influx and membrane potential (V_m)-induced Ca²⁺ influx and subsequently regulate the calcineurin-NFATc1 signaling pathway in $\gamma\delta$ T cells. Thus, CFTR serves as a negative regulator of IFN- γ production in $\gamma\delta$ T cells and the function of these cells in antitumor immunity. Our investigation suggests that modification of the CFTR activity of $\gamma\delta$ T cells may be a potential immunotherapeutic strategy for cancer.

Keywords: CFTR; γδ T cells; TCR; Membrane potential; Antitumor immunity

INTRODUCTION

Ion channels and transporters expressed in the plasma membrane of immune cells participate in the influx (Ca²⁺, Mg^{2+} , Zn^{2+} , and Na^+) and efflux (K⁺ and Cl⁻) of ions to regulate the membrane potential and physiological cell functions, such as gene expression, apoptosis, proliferation, development, and migration. $^{1-7}$ Mutations in ${\rm Ca}^{2+}$ release-activated channel and the Mg²⁺ channel MAGT1 lead to severe combined immunode-ficiency disorders in humans.^{8–10} The membrane potential, which depends on electrical gradients across the cell membrane, is maintained and regulated by K⁺, Na⁺, and Cl⁻ channels.^{11,12} Multiple investigations of immune cells involving membrane potential regulation have focused on K⁺ and TRPM4 channels; these channels transport K^+ and Na^+ and provide the driving force for Ca²⁺ influx following receptor activation that regulates the functions of many enzymes and transcription factors.¹³ However, the effects of CI^- and CI^- channels on membrane potential regulation and the subsequent functional impact on immune cells are still poorly elucidated.

Cystic fibrosis transmembrane conductance regulator (CFTR) is associated with the lethal genetic disease cystic fibrosis, which is mainly caused by bronchial inflammation, viscous secretions, and impaired mucociliary clearance, and is responsible for electrolyte and water transport in the apical membrane of the epithelial cells lining various tissues.^{18–21} CFTR is a unique Cl⁻ channel due to its dual functions as a Cl⁻ channel and a regulator of other ion channels and transporters.²² The opening of Cl⁻ channels in immune cells triggers the efflux of Cl⁻ ions caused by a relatively high [Cl⁻]_i (~38 mM) and an equilibrium membrane potential (approximately -33 mV). Consequently, Cl⁻ efflux depolarizes the membrane potential and inhibits Ca²⁺ influx by regulating voltage-gated Ca²⁺ channels.²³ However, how CFTR regulates the membrane potential and subsequently affects Ca²⁺ influx in immune cells remains unclear.

As a regulator, CFTR modulates several membrane ion transporters in epithelial cells via protein–protein interactions, such as the epithelial Na⁺ channel,²⁴ Ca²⁺-activated chloride channels,²⁵ the outwardly rectifying chloride channel,²⁶ and apical

Correspondence: Jianlei Hao (haojianlei@jnu.edu.cn) or Zhinan Yin (zhinan.yin@yale.edu)

These authors contributed equally: Yuanyuan Duan, Guangqiang Li, Miaomiao Xu

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¹Zhuhai Precision Medical Center, Zhuhai People's Hospital (Zhuhai Hospital Affiliated with Jinan University), Jinan University, Zhuhai 519000 Guangdong, China; ²The Biomedical Translational Research Institute, Faculty of Medical Science, Jinan University, Guangzhou 510632 Guangdong, China; ³Department of Physiology, School of Basic Medical Sciences, Southern Medical University, Guangzhou 510515 Guangdong, China; ⁴The First Affiliated Hospital, Jinan University, Guangzhou 510632, China; ⁵School of Pharmaceutical Sciences, South-Central University for Nationalities, Wuhan 430074 Hubei, China; ⁶Division of Life Science, Hong Kong University of Science and Technology (HKUST), Hong Kong, China; ⁷State Key Laboratory of Oncology in Southern China, Collaborative Innovation Center for Cancer Medicine, Sun Yat-sen University Cancer Center, Guangzhou 510600, China and ⁸Department of Immunobiology, School of Medicine, Yale University, New Haven, CT 06520, USA

K⁺ channels in renal epithelial cells, including renal outer medullary potassium channel (ROMK) 1 and ROMK2.²⁷ Recently, CFTR was reported to participate in immune regulation²⁸⁻³⁰ and potentially be a risk factor for cancer development.³¹ However, the functions and underlying mechanisms of CFTR in T cells have not yet been fully elucidated.

vδ T cells represent an unconventional innate-like T cell subset. which are different from $\alpha\beta$ T cells by their unique characteristics, including their tissue-specific distribution, lower receptor variability, NK receptor expression, and non-MHC-restricted cytotoxicity against target cells.^{32,33} γδ T cells actively perform protective functions against tumors, viral infections, parasites, and bacterial infections.³⁴ v δ T cells also participate in the development of autoimmune and inflammatory diseases via rapid production of proinflammatory cytokines.³⁵ Thus, it is crucial to explore the molecular mechanisms that regulate $\gamma\delta$ T cell effector functions. Effector yo T cells are classified as either IFN-y-producing cells or IL-17-producing cells and regulate immunity in divergent microenvironments. IFN- γ -producing $\gamma\delta$ T cells control processes during herpes, influenza, and other viral infections and during bacterial infections.^{36,37} In our previous publications, we found that $\gamma\delta$ T cells provided an early critical source of IFN-y and played a protective role in antitumor immunity.^{38,39} Furthermore, we reported that the promoter region of the *lfng* gene in $\gamma\delta$ T cells was highly demethylated, which resulted in a rapid IFN-y response upon activation by the transcription factors T-bet and Eomes.⁴ We recently demonstrated mTORC1-mediated IFN-y production by $\gamma\delta$ T cells. Due to the distinctive features of $\gamma\delta$ T cells, several $\gamma\delta$ T cell-based, small-sized clinical trials of tumor immunotherapy using either in vivo expansion or adoptive transfer have been conducted.⁴¹ Clinical trials led by our group using allogenic $\gamma\delta$ T cells showed promising effects on solid tumors.⁴² Although TCR signal transduction from stimuli on the extracellular side to transcription factors in the nucleus of $\gamma\delta$ T cells, including ligand recognition and intracellular signaling cassettes, has been reported,^{43,44} the detailed molecular pathways still need to be explored. The impacts of ion channels, especially Ca²⁺ channels. on $\alpha\beta$ T cell functions have been well documented.

In this regard, we wanted to explore the critical functions of CFTR as a Cl⁻ channel and a regulator in $\gamma\delta$ T cells. In our study, we discovered that CFTR was functionally expressed on the cell surface of $\gamma\delta$ T cells. CFTR negatively regulated IFN- γ production and antitumor immunity in $\gamma\delta$ T cells. Regarding the underlying molecular mechanism, we found that CFTR in $\gamma\delta$ T cells modulated V_m-induced and TCR-stimulated Ca²⁺ influx and subsequently regulated the calcineurin-NFATc1 signaling pathway. In addition, CFTR was recruited to TCR immunological synapses (ISs) and participated in TCR signal transduction by attenuating Lck-P38 MAPK-c-Jun signaling. Our research systematically defined the role of CFTR in the activation of $\gamma\delta$ T cells and the potential contribute to the understanding of $\gamma\delta$ T cell biology and increase awareness of the potential of CFTR in $\gamma\delta$ T cell-mediated cancer immunotherapy.

RESULTS

CFTR is expressed in $\gamma\delta$ and CD4 $^+$ T cells and is polarized toward the IS via the cytoskeleton

To evaluate the relative mRNA expression levels of various Cl⁻ channels, we first performed Q-PCR analysis of primary mouse $\gamma\delta$ T cells and CD4⁺ T cells isolated from the spleen of C57BL/6 mice. We found that the transcripts of CFTR and chloride intracellular channels (CLICs) displayed relatively higher mRNA levels than those of other Cl⁻ channels (Fig. S1A) and treatment with IAA94, which is a potent blocker of CLICs, did not alter IFN- γ production in either $\gamma\delta$ T cells or CD4⁺ T cells (Fig. S1B). Due to the unique functions of CFTR among Cl⁻ channels, we focused on CFTR in

further investigations. We then confirmed the expression of CFTR (band B: core-glycosylated form; band C: fully glycosylated form) in primary mouse $v\delta$ T cells and CD4⁺ T cells by immunoblot analysis (Fig. S1C). Confocal microscopy showed that CFTR was predominantly and evenly distributed in the plasma membrane of resting $\gamma\delta$ T cells and CD4⁺ T cells (Fig. S1D). CFTR^{-/-} mice were used as a negative control for antibody specificity in the immunoblot and confocal analyses. Surprisingly, we found that endogenous CFTR and F-actin relocalized and polarized toward the IS (the arrowhead indicated in Fig. S1E) when $\gamma\delta$ T cells were activated with anti-CD3/CD28-coated Dynabeads, which was different than the distribution seen in resting $v\delta$ T cells (Fig. S1D). It has been reported that in polarized epithelial cells, the cterminal PDZ binding motif of CFTR binds to PDZ domaincontaining proteins, which interact with ezrin to tether this multiprotein complex to the apical actin cytoskeleton.⁴⁶ In addition, actin and ezrin are involved in the formation of the IS.⁴⁷ Given this information, we tested the localization of CFTR and ezrin in stimulated $v\delta$ T cells and found that ezrin was also polarized toward the IS region and colocalized with CFTR (Fig. S1F), suggesting that CFTR is recruited to the IS region through binding to the cytoskeletal complex when yδ T cells are activated. T cell activation drives morphological and functional changes in T cells, including large-scale accumulation of actin and signaling molecules, which form distinct supramolecular clusters termed the IS to stabilize receptor signal transduction.48 Thus, these results indicated the potential role of CFTR in $v\delta$ T cell activation.

CFTR modulates $\gamma\delta$ T cell activation

To define the function of CFTR in $\gamma\delta$ T cell activation, we analyzed $v\delta$ T cell properties in wild-type and CFTR^{-/-} mice. By surface staining and FACS analysis, we observed that there was no difference in the quantity of $\gamma\delta$ T cells between wild-type and $CFTR^{-/-}$ mice (Fig. 1a upper panel), suggesting that CFTR may not be significantly involved in yo T cell development and maintenance. However, CFTR deficiency robustly elevated the size of the population of naturally activated (CD44^{high}CD62L^{low}) γδ T cells (Fig. 1a lower panel), suggesting a negative role for CFTR in $\gamma\delta$ T cell activation in vivo. Notably, although CFTR was also expressed in CD4⁺ T cells, there was no significant phenotype related to $CD4^+$ T cell activation in $CFTR^{-/-}$ mice (Fig. 1b). To further confirm the contribution of CFTR to IS formation when $y\delta$ T cells are activated, we assessed the F-actin distribution by confocal microscopy after stimulating wild-type and CFTR^{-/-} $\gamma\delta$ T cells with anti-CD3/CD28-coated Dynabeads[®]. We observed that significantly more F-actin accumulated in the IS area upon stimulation in the $CFTR^{-/-}$ $\gamma\delta$ T cells than in the wild-type $\gamma\delta$ T cells (Fig. 1c). Given the cytoskeletal control of cell morphology and cell mechanical stiffness, we characterized the cell membrane by directly measuring Young's modulus and the adhesion force with atomic force microscopy and membrane force analysis. As shown in Fig. 1d, the mean Young's modulus of $CFTR^{-/-} \gamma \delta T$ cells was significantly higher than that of wild-type $v\delta T$ cells, which was consistent with our confocal results; in contrast, the mean adhesion force was decreased. Thus, the $CFTR^{-/-} \gamma \delta T$ cells were stiffer than the wild-type $\gamma\delta$ T cells due to the increased F-actin accumulation in the IS region. These results collectively showed a stronger activation status in CFTR^{-/-} y δ T cells.

CFTR^{-/-} $\gamma\delta$ T cells exhibit enhanced IFN- γ production and cytotoxicity against tumors in vitro and in vivo

We then hypothesized that the enhanced T cell activation in CFTR^{-/-} mice might affect subsequent cytokine production. Since IFN- γ and IL-17 released by $\gamma\delta$ T cells have been extensively studied and found to play distinct roles in many disease models, we assessed the alterations in the expression of these two cytokines between CFTR^{-/-} $\gamma\delta$ T cells and wild-type $\gamma\delta$ T cells. By intracellular cytokine staining of splenocytes and peripheral lymph

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Fig. 1 CFTR deficiency resulted in increased activation and more actin accumulation in the immunological synapse (IS) of $\gamma\delta$ T cells. Representative flow cytometry and statistical analyses of CD44 and CD62L expression in WT or CFTR^{-/-} splenic $\gamma\delta$ T cells (**a**) and CD4⁺ T cells (**b**) (n = 3, **p < 0.01). **c** Left: confocal microscopy of WT or CFTR^{-/-} $\gamma\delta$ T cells stimulated with Dynabeads[®] Mouse T-Activator CD3/CD28 and stained with a rabbit anti-Lck antibody followed by an Alexa Fluor 488-conjugated anti-rabbit IgG or with Alexa Fluor 594-conjugated phalloidin (lower panel, arrowheads indicate the immunosynapse area) (scale bar: 5 µm, BF: bright field); Right: quantification of actin accumulation at the IS contact area analyzed with ImageJ (n = 52 per group, ****p < 0.0001). **d** Atomic force microscopy of in vitro-expanded WT or CFTR^{-/-} $\gamma\delta$ T cells with analysis of Young's modulus and the adhesion force (WT: n = 172, CFTR^{-/-}: n = 184, ***p < 0.001)

node cells ex vivo, we found that CFTR deficiency triggered a significant increase in IFN- γ production in $\gamma\delta$ T cells (Fig. 2a) but not in CD4⁺ T cells (Fig. 2b), which was consistent with the impact of CFTR on yδ T cell activation (Fig. 1). In addition, peripheral IL-17 expression was not altered in $CFTR^{-/-} \gamma \delta T$ cells or CD4⁺ T cells (Fig. 2a, b), suggesting a unique mechanism of CFTR-mediated suppression of IFN- γ production in $\gamma\delta$ T cells. In addition, we analyzed the phenotype of thymic $\gamma\delta$ T cells in CFTR^{-/-} mice and found no differences in CD44/CD69 expression or cytokine production between wild-type and CFTR^{-/-} $\gamma\delta$ T cells (Fig. S2), which suggested that CFTR played a regulatory role in the periphery rather than during thymic programming. To dissect the stage of $v\delta$ T cell differentiation or maintenance impacted by CFTR, we sorted naive and naturally active vo T cells separately, polarized them toward the $\gamma\delta$ T1 direction, and analyzed the terminally differentiated cells. The results showed that the $\gamma\delta$ T1 cells derived from sorted wild-type or CFTR^{-/-} naive cells did not exhibit differences (Fig. 2c, upper panel), indicating a redundant role for CFTR in naive $\gamma\delta$ T cell differentiation, and the underlying mechanisms remain to be further studied. In contrast, CFTR^{-/-} $\gamma\delta$ T1 cells had a higher frequency than wildtype $\gamma\delta$ T1 cells when derived from sorted activated cells (Fig. 2c, lower panel). These results indicated that CFTR played an important role in activated $\gamma\delta$ T cell function. Collectively, these findings revealed that CFTR was a negative regulator of $\gamma\delta$ T cell activation and IFN- γ production.

In our previous studies, we found that $v\delta$ T cells, especially those in the Vy4 subset, played a protective role in early tumor immune surveillance via IFN-γ and perforin production.³⁹ Because $\gamma\delta$ T cell activation and IFN- γ production were found to be enhanced in CFTR-deficient mice (Fig. 2a), we hypothesized that CFTR^{-/-} $^{-}$ y δ T cells may exert relatively strong antitumor immunity. To test this hypothesis, we first performed an in vitro killing assay by mixing $\gamma\delta$ T cells and B16 melanoma tumor cells and then monitoring killing in real time. The B16 cells were labeled with calcein-AM and cocultured with the $v\delta$ T cells at a ratio of 1:5, and B16 viability was evaluated and quantified by assessing the fluorescence and density of calcein during a 6-h period. As shown in Fig. S3A, CFTR^{-/-} $\gamma\delta$ T cells displayed more rapid killing activity than wild-type $\gamma\delta$ T cells. To test whether CFTR^{-/-} $\gamma\delta$ T cells have suppressive effects on tumors in vivo, we mixed B16 tumor cells with $\gamma\delta$ T cells, which were expanded and sorted from wild-type or CFTR^{-/-} mice, and separately inoculated the mixtures subcutaneously (s.c.) into the two flanks of B6 TCR $\delta^{-/-}$ mice as described in our previous studies.³⁹ Consistent with the in vitro results, the CFTR^{-/-} $\gamma\delta$ T cells showed significantly more powerful antitumor CFTR is a negative regulator of $\gamma\delta$ T cell IFN- γ production and antitumor... Y Duan et al.

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function than the wild-type $\gamma\delta$ T cells (Fig. 2d). These results collectively demonstrated that CFTR might serve as a negative regulator in $\gamma\delta$ T cell-mediated antitumor immunity, which was consistent with the enhanced IFN- γ production by CFTR^{-/-} $\gamma\delta$ T cells observed (Fig. 2a, c). To test whether the enhanced cytotoxicity of CFTR^{-/-} $\gamma\delta$ T cells against B16 tumor cells was due to elevated IFN- γ production, we blocked IFN- γ with an anti-IFN- γ antibody in the in vitro killing assay, and the results showed that blocking IFN- γ reduced the cytotoxic effect of CFTR^{-/-} $\gamma\delta$ T cells to wild-type levels (Fig. 2e), suggesting that the stronger antitumor function of CFTR^{-/-} $\gamma\delta$ T cells was dependent on elevated IFN- γ production. In addition, we performed an in vitro

Fig. 2 CFTR deficiency led to increased IFN-y release and cytolytic ability against aggressive B16 melanoma. Representative flow cytometry analysis of IFN-γ and IL-17 production by WT or CFTR⁻ peripheral $\gamma\delta$ T cells (**a**) and CD4⁺ T cells (**b**) treated with PMA (50 ng/ml) and Naive (CD44^{low}) $\gamma\delta$ T cells and active (CD44^{low}) $\gamma\delta$ T cells sorted from WT or CFTR^{-/-} mouse splenocytes were selected at the splenocytes were selected. ionomycin (1 µg/ml) stimulation for 6 h is shown (n = 4, *p < 0.05). c T1 subset for 4 days, and flow cytometry analysis of IFN- γ and IL-17 production by with WT or CFTR^{-/-} $\gamma\delta$ T cells treated with PMA (50 ng/ml) and ionomycin (1 µg/ml) stimulation for 6 h was performed (n = 3, **p < 0.01). **d** B16 cells (2×10^5 cells/mouse) were mixed with in vitro-expanded WT or CFTR^{-/-} $\gamma \delta$ T cells (0.5×10^5 cells/mouse) and s.c. injected into B6 TCR $\delta^{-/-}$ mice (as shown in the schematic diagram in the left panel; n = 5 per group). On day 16 post tumor injection, tumors were isolated (left panel, scale bar: 1 cm), and tumor sizes were measured (right panel, paired *t*-test; **p < 0.01). **e** In vitro-expanded WT or CFTR^{-/-} $\gamma\delta$ T cells were cocultured with CFSE-labeled B16 cells at the indicated ratio, and an anti-IFN-y antibody (10 µg/ml) was added to neutralize cytotoxicity, or an isotype control antibody was added. Six hours later, dead B16 cells were assessed by PI staining (***p < 0.001; **p < 0.01; *p < 0.05; ns not significant; n = 3)

killing assay with EL4 lymphoma cells. As shown in Fig. S3B, CFTR^{-/-} $\gamma\delta$ T cells displayed enhanced cell cytotoxicity, which was mediated by IFN- γ , indicating that the killing ability of CFTR^{-/-} $\gamma\delta$ T cells was not B16 melanoma specific. Collectively, these results suggested that CFTR^{-/-} $\gamma\delta$ T cells had enhanced antitumor function mediated by upregulating IFN- γ expression.

CFTR inhibits $\gamma\delta$ T cell IFN- γ production by regulating the TCR signaling cascade and Ca^{2+} influx

We demonstrated that CFTR deficiency resulted in increased activation and IFN-y release in peripheral yo T cells (Figs. 1a and 2a). We further investigated the molecular mechanism underlying CFTR-mediated yo T cell activation. It is well known that the stimulation of T cells mediated by TCR signaling, which is critical for effective host responses to pathogens or tumors, requires the sequential activation and interaction of dozens of kinases and adapter proteins.⁴⁹ Following the engagement of TCR, PLCy-1/ ZAP70/Lck form a TCR transduction complex to induce a cascade of events that spread through the membrane to the cytosol and the nucleus.⁵⁰ Since the TCR signal transduction network requires protein-protein interactions, we first examined whether CFTR binds with TCR signaling molecules. Interestingly, coimmunoprecipitation analysis of wild-type $\gamma\delta$ T cells showed that endogenous CFTR physically interacted with PLCy-1, which bound to Lck and ZAP70, but this interaction did not occur in CFTR^{-/-} $\gamma\delta$ T cells (Fig. 3a). Furthermore, as shown in Fig. 3b, endogenous CFTR colocalized with the TCR transduction complex PLCy-1/ZAP70/Lck in the IS area, but the PDZ binding motif in the c terminus of CFTR was not responsible for PLCy-1 binding (Fig. S1G). Collectively, these results indicated that CFTR, when recruited to the IS region, could be a crucial signaling molecule in the proximal TCR transduction cascade.

We next aimed to evaluate the effect of CFTR deficiency on $\gamma\delta$ T cell TCR signaling. We found no significant difference in the phosphorylation of PLC γ -1 between CFTR^{-/-} $\gamma\delta$ T cells and wild-type $\gamma\delta$ T cells. However, CFTR deficiency in $\gamma\delta$ T cells triggered robust increases in the phosphorylation levels of ZAP70, Lck, P38 MAPK, and c-Jun upon TCR stimulation (Fig. 3c). We then tested whether the augmented phosphorylation of TCR signaling molecules was the reason for the elevated IFN- γ production in CFTR^{-/-} $\gamma\delta$ T cells. Consistent with the results of the FACS analysis of CFTR^{-/-} $\gamma\delta$ T cells, we found that CFTR^{-/-} $\gamma\delta$ T cells produced higher levels of IFN- γ than did wild-type $\gamma\delta$ T cells by ELISA. When the P38 MAPK antagonist SB203580 or the Lck inhibitor PP2 was

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Fig. 3 CFTR participated in TCR signaling via interaction with the PLC γ -1–LCK–ZAP70 signaling complex and contributed to TCR-induced Ca²⁺ influx and membrane hyperpolarization in $\gamma\delta$ T cells. **a** In vitro-expanded WT or CFTR^{-/-} $\gamma\delta$ T cells were immunoprecipitated with an anti-PLC γ -1 antibody, and the immunoprecipitates were immunoblotted with an antibody specific for PLC γ -1, CFTR, Lck, or ZAP70. **b** Confocal microscopy analysis of $\gamma\delta$ T cells stimulated with Dynabeads Mouse T-Activator CD3/CD28 and stained with a mouse anti-CFTR antibody followed by an Alexa Fluor 488-conjugated anti-mouse IgG or with an antibody to PLC γ -1, Lck, or ZAP70 followed by an Alexa Fluor 594-conjugated anti-mouse IgG or with an antibody to PLC γ -1, Lck, or ZAP70 followed by an Alexa Fluor 594-conjugated anti-mouse IgG or with an antibody to PLC γ -1, Lck, or ZAP70 followed by an Alexa Fluor 594-conjugated anti-mouse IgG or with an antibody to PLC γ -1, Lck, or ZAP70 followed by an Alexa Fluor 594-conjugated anti-mouse IgG or with an antibody to PLC γ -1, Lck, or ZAP70 followed by an Alexa Fluor 594-conjugated anti-rabbit IgG was performed (scale bar: 5 µm, BF: bright field). **c** A representative immunoblot analysis (from three independent experiments) of the indicated proteins in the whole-cell lysates of in vitro-expanded WT or CFTR^{-/-} $\gamma\delta$ T cells treated with soluble anti-CD28 (1 µg/ml) antibody stimulation for 2, 5, 10, 15, or 30 min is shown. Band densities were quantified with ImageJ and ratios of indicated intensities were shown. act: actin. **d** ELISA analysis of IFN- γ production by in vitro-expanded and purified WT or CFTR^{-/-} $\gamma\delta$ T cells treated with SB203580 (10 µM) or PP2 (10 µM) for 12 h was performed (n = 3 for each group, ***p < 0.001). **e** The cytoplasmic V_m of WT or CFTR^{-/-} $\gamma\delta$ T (1 × 10⁶ cells for each group) treated with soluble anti-CD3 (10 µg/ml) and anti-CD28 (1 µg/ml) antibody stimulation was measured, and the relative V_m in WT or CFTR^{-/-} $\gamma\delta$ T cells was quantified (n = 3

added to cultures, no significant difference in IFN- γ production was found between WT and CFTR^{-/-} $\gamma\delta$ T cells, suggesting that Lck-P38 MAPK TCR signaling was involved in CFTR-regulated IFN- γ production (Fig. 3d). Together, these results suggested that CFTR negatively regulated TCR signal transduction and TCR-induced IFN- γ production.

Since CFTR modulates the membrane potential and TCR signaling, which subsequently regulate Ca²⁺ influx, we next investigated the contribution of CFTR to V_m-induced and TCR-stimulated Ca²⁺ influx with the V_m indicator DiBAC₄(3) and Ca²⁺ indicator Fluo-4. As expected, CFTR^{-/-} $\gamma\delta$ T cells displayed a lower membrane potential than wild-type $\gamma\delta$ T cells upon TCR stimulation (Fig. 3e). As shown in Fig. 3f, the phosphorylation level of NFATc1 was decreased in CFTR^{-/-} $\gamma\delta$ T cells, which indicated that NFATc1 was activated. In agreement with the data indicating that CFTR negatively regulated TCR transduction and NFATc1 activation, CFTR^{-/-} $\gamma\delta$ T cells exhibited a higher TCR-stimulated Ca²⁺ influx than wild-type $\gamma\delta$ T cells (Fig. 3g).

CFTR inhibits IFN- γ production by regulating the membrane potential and Ca^{2+} influx in $\gamma\delta$ T cells

To further confirm the contribution of CFTR channel activity to $\gamma\delta$ T cell function, we treated wild-type $\gamma\delta$ T cells with a CFTRspecific inhibitor. First, we evaluated whether CFTR was functional in sorted $\gamma\delta$ T cells via the whole-cell patch clamp technique. We treated cells with the CFTR agonist forskolin and the CFTR-specific inhibitor GlyH101 and recorded cAMP-stimulated Cl⁻ conductance. Indeed, surface expression of CFTR in $\gamma\delta$ T cells led to a cAMP-stimulated Cl⁻ current, which was blocked by GlyH101 (Fig. S1H). Next, we confirmed the modulation of CFTR channel activity-mediated IFN-y production by treating wild-type $\gamma\delta$ T cells derived from the mouse spleen with the CFTR-specific inhibitor CFTR_{inh172}. In agreement with our in vivo results (Fig. 2), blocking CFTR activity and Cl⁻ efflux enhanced IFN-y expression in yo T cells (Fig. 4a). In addition, CFTR_{inb172}-treated $v\delta$ T cells exhibited markedly increased cytotoxicity against B16 melanoma cells in vitro (Fig. 4b) and in vivo (Fig. S3D). In addition, an in vitro killing assay with EL4 lymphoma cells showed that CFTR_{inh172}-treated $\gamma\delta$ T cells displayed enhanced cell cytotoxicity, but IAA94-treated yδ T cells did not, indicating the stronger killing ability of the $CFTR_{inh172}\text{-}treated~\gamma\delta$ T cells and inability of CLICs to influence the cytotoxicity of yo T cells (Fig. S3C). We also assessed TCR signaling when CFTR channel activity was blocked. We found that pharmacological inhibition of CFTR did not affect the activity of TCR signaling molecules (Fig. 4c), which suggested that CFTR_{inh172} did not interrupt the interaction between CFTR and TCR molecules and that the CFTR_{inh172}-mediated increase in IFN-γ production was not due to an alteration in ZAP70-P38-c-Jun signaling.

We also observed that CFTR_{inh172} decreased the V_m and hyperpolarized $\gamma\delta$ T cells (Fig. 4d) and that this cell hyperpolarization could be abolished by gramicidin, which was used as a positive control; furthermore, Ca²⁺ influx triggered by an elevated extracellular calcium concentration ([Ca²⁺]_e) was significantly increased in $CFTR_{inh172}\text{-}treated\ \gamma\delta\ T$ cells treated with the sarcoplasmic endoplasmic reticulum Ca²⁺-ATPase pump inhibitor thapsigargin (Fig. 4e). These data suggested that CFTR might affect the V_m and Ca^{2+} influx by regulating an unknown Ca^{2+} channel(s) on the cell surface. Calcineurin is a calcium-dependent serine-threonine phosphatase that activates NFATc1, and we found that FK506, a calcineurin inhibitor, blocked CFTR_{inh172}-induced IFN-y production (Fig. 4f). In addition, the phosphorylation level of NFATc1 was decreased in CFTR_{inh172}treated $\gamma\delta$ T cells (Fig. 4g), suggesting that the CFTR inhibitor, which triggered Ca²⁺ influx, led to calcineurin-NFATc1 activation and IFN-y expression. These results collectively suggested that CFTR inhibited IFN- γ production by $\gamma\delta$ T cells through its ion channel function.

CFTR affects IFN- γ production in human V δ 2 T cells and V δ 2 T cell-mediated cytotoxicity to human K562 leukemia cells

Vy9V δ 2 y δ T cells, which are the dominant subset in human peripheral blood, are known for their high IFN-y production and efficient antitumor activity.⁵¹ To explore the effect of CFTR on Vδ2 γδ T cells, we first evaluated the CFTR-mediated pharmacological inhibition of IFN- γ production in expanded V δ 2 $\gamma\delta$ T cells from the peripheral blood of healthy donors. Consistent with our murine system, CFTR inhibition via CFTR_{inh172} significantly enhanced IFN-γ release (Fig. 5a); in addition, CFTR_{inh172}-treated Vδ2 cells showed relatively high cytotoxicity to human K562 leukemia cells (Fig. 5b). Next, we used a lentiviral system to knockdown CFTR expression in V\delta2 $\gamma\delta$ T cells and observed cytokine production. Before applying shRNA knockdown to VS2 yS T cells, we measured the knockdown efficiencies of 4 different shRNA lentiviral particles. As shown in Fig. 5c, in GFP-positive V δ 2 $\gamma\delta$ T cells, CFTR expression was decreased by approximately 50%, and similar to CFTR pharmacological inhibition, CFTR knockdown increased IFN-v production upon anti-CD3/CD28 antibody stimulation. Collectively, these data indicated that CFTR negatively regulated human Vδ2 γδ T cell function.

DISCUSSION

 $\gamma\delta$ T cells play important roles in antitumor immunity.⁵² Our early studies demonstrated that $\gamma\delta$ T cells rapidly produce IFN- γ at a much higher level than CD4 T cells, even when primed toward the Th1 program in vitro.⁵³ The intrinsic mechanisms involve the expression of T-bet and Eomes and the epigenetically open status of the ifng gene.^{40,54} We further determined that $\gamma\delta$ T cellproduced IFN-y plays critical roles in antitumor immunity and the proinflammatory response in spinal cord injury.^{38,39,55} Subsequent studies by our group have shown that mTORC1-mediated metabolism and downstream signaling are required for IFN-y production by $\gamma\delta$ T cells.⁵⁶ However, the regulation of IFN- γ remains unclear, especially the differences between $\gamma\delta$ T cells and αβ T cells. Ion channels play critical roles in transmembrane ion homeostasis and have been found to play important functions in several types of immune cells.⁶ This is the first study to identify the ion channel CFTR as a negative regulator of IFN-γ production in γδ T cells and antitumor immunity.

We used genetically engineered KO mice/inhibitors and demonstrated that the chloride channel CFTR protein was a negative regulator of IFN-y production in both mouse and human $y\delta$ T cells (Figs. 1, 2, and 5). This indicated that $y\delta$ T cells, although found to have innate features and be more effective than aß T cells, could still be further strengthened by enhancing their activation through deactivation of negative molecules, including CFTR. Thus, our study identified a new member of the negative regulators of T cells, adding to PD-1 and CTLA-4, and provided potentially new targets for cancer immunotherapy based on specifically inhibiting CFTR activities. It is important to consider that several reports have shown contradictory effects of CFTR on different cancers, especially potentially protective effects on In this study, we demonstrated that CFTR leukemia.5/ negatively regulated the cytotoxicity of $v\delta$ T cells to B16 melanoma cells and EL4 lymphoma cells. While recognizing the limitations of this research, especially studying CFTR function in $\gamma\delta$ T cells via analysis of control of tumor cell lines, we believe that more mechanistic studies are necessary to define the precise role of CFTR in specific cell types in a particular cancer.

Another important finding of this study is that CFTR is a bifunctional molecule in $\gamma\delta$ T cells (Fig. S4), with the nonchannel function being especially important (Fig. 3). Upon TCR activation, CFTR was recruited to the immunosynapse, bound to PLC γ -1 and

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Pharmacological inhibition of CFTR significantly increased IFN- γ release and cytotoxic activity in $\gamma\delta$ T cells and triggered V_m-induced Fia. 4 influx. **a** Flow cytometry analysis of splenic $\gamma\delta$ T cells treated with PMA (50 ng/ml) and ionomycin (1 µg/ml) stimulation for 6 h in the Ca^{2} presence of the CFTR inhibitor CFTR_{inh172} (5 μ M) or vehicle DMSO was performed to detect IFN- γ and IL-17 production (n = 4, *p < 0.05). **b** In vitro-expanded $\gamma\delta$ T cells were pretreated with the CFTR inhibitor CFTR_{inh172} (5 μ M) or vehicle DMSO for 2 h and then cocultured with CFSElabeled B16 cells at the indicated ratio. Six hours later, dead B16 cells were identified by PI staining (n = 3, *p < 0.05). **c** A representative immunoblot analysis (from three independent experiments) of the indicated proteins in the whole-cell lysate of in vitro-expanded $\gamma\delta$ T cells stimulated in the presence of the CFTR inhibitor CFTR_{inh172} (5 µM) or vehicle DMSO with anti-CD3 (10 µg/ml) and anti-CD28 (1 µg/ml) antibodies for 2, 5, 10, 15, or 30 min is shown; band densities were quantified with ImageJ and ratios of indicated intensities were shown. act: actin. **d** The cytoplasmic V_m of $\gamma\delta$ T cells treated with the CFTR inhibitor CFTR_{inh172} ($10\,\mu$ M) or vehicle DMSO (1 × 10⁶ cells for each group) followed by gramicidin (10 μ M) was measured, and the relative V_m in WT or CFTR_{inh172}-treated $\gamma\delta$ T cells was quantified (n = 3, **p < 0.01). **e** In vitro-expanded $\gamma\delta$ T cells were pretreated with CFTR_{inh172} (10 μ M) or the vehicle DMSO (1 × 10⁶ cells for each group) for 2 h in Ca²⁺-free medium. Changes in the $[Ca^{2+}]_i$ were monitored by flow cytometry, and the overlay of the Ca^{2+} influx profiles after the addition of $CaCl_2$ (5 mM) to the extracellular medium in the presence of thapsigargin (1 μ M) is shown. The relative peak [Ca²⁺]_i in WT or CFTR_{inh172}-treated $\gamma\delta$ T cells was quantified (n = 3, *p < 0.05). **f** In vitro-expanded $\gamma\delta$ T cells were treated with PMA (50 ng/ml) and ionomycin (1 μ g/ml) for 6 h in the presence of the CFTR inhibitor CFTR_{inh172} (5 µM) or vehicle DMSO with the addition of FK506 and then evaluated to detect IFN- γ and IL-17 production (*p < 0.05, ns not significant, n = 3). **g** Flow cytometry analysis of the phosphorylated NFATc1 level in in vitro-expanded WT or CFTR_{inh172}-treated $\gamma\delta$ T cells stimulated with soluble anti-CD3 (10 µg/ml) and anti-CD28 (1 µg/ml) antibodies for 30 min was performed (n = 3, **p < 0.01). Dotted line: isotype control

directly attenuated Lck-P38 MAPK-c-Jun signaling. This further reminds us that ion channels may transduce previously unknown signals and perform additional functions in immune cells.

On the other hand, we found that CFTR modulated TCRstimulated Ca²⁺ influx and V_m-induced Ca²⁺ influx and subsequently regulated the calcineurin-NFATc1 signaling pathway in $\gamma\delta$ T cells (Fig. 4, Fig. S3). These findings further reinforced the relationships between CFTR and other ion channels. However, the potential candidate activated voltage-gated Ca²⁺ channels in this process have not yet been determined. Hence, CFTR, as an anion channel, needs to be researched further in regard to the comprehensive mechanism of immune cell function regulation.

As many previous studies have indicated, vo and $\alpha\beta$ T cells have distinct developmental and functional features.⁵² For instance, $\gamma\delta$ TCRs recognize a distinct profile of ligands without MHC restriction rather than being activated by self MHC-presented pAgs like $\alpha\beta$ T cells.⁵² In our previous studies, we also provided evidence that $\gamma\delta$ T cells intrinsically produced more IFN- γ than $\alpha\beta$ T cells in several conditions.^{53,54} In this study, we also found that CFTR^{-/-} $y\delta$ T cells but not $\alpha\beta$ T cells showed elevated IFN-y production. This indicated that CFTR effectively suppressed IFN-y production only in $v\delta$ T cells. The molecular basis of this discrepancy between $\gamma\delta$ and $\alpha\beta$ T cells is currently unknown. The possible explanations may involve the inability of CFTR to be recruited to the immunosynapse in $\alpha\beta$ T cells, the compensatory effects of other ion channels, or the distinct machinery of signaling pathways in αβ T cells. Hence, it would be interesting to further investigate the distinct functions of CFTR in specific cell types, which could precisely determine the activity of CFTR in a particular environment.

In summary, this study defined CFTR as a negative regulator of IFN- γ production in $\gamma\delta$ T cells and antitumor immunity. Our investigation suggested that modification of the CFTR activity of $\gamma\delta$ T cells might provide a potential immunotherapeutic strategy for cancer. It is important to further study the detailed molecular mechanisms of CFTR in $\gamma\delta$ T cells, particularly in relation to cell activation and quiescence and the transition between these states. Moreover, the regulation of CFTR in munu for the expression of CFTR and visualization of the spatial localization of CFTR in munu cells are still important areas that remain to be investigated.

MATERIALS AND METHODS

Mice and cell lines

C57BL/6J mice were obtained from Beijing HFK Bioscience Co., Ltd. (Beijing, China). B6.129P2-Cftr^{tm1Unc}/J (CFTR^{-/-}) mice, which have exon 10 replacement, were kindly provided by Dr. Pingbo Huang from the Hong Kong University of Science and Technology (Hong Kong). CFTR⁺⁷⁻ breeders were set up, and CFTR^{+/+} and CFTR^{-/-} littermates were used for experiments. B6.129P2-Tcrd^{tm1Mom}/J (TCR $\delta^{-/-}$) mice were purchased from The Jackson Laboratory (Bar Harbor, ME). All animals were maintained under specific pathogen-free conditions and used at 6-8 weeks of age, and mouse protocols were approved by the Animal Experiment Committee of Jinan University. B16-F0 melanoma cells were purchased from ATCC (Manassas, VA) and were maintained in DMEM supplemented with 10% FBS. EL4 lymphoma cells and K562 leukemia cells were purchased from ATCC (Manassas, VA) and were maintained in RPMI 1640 medium supplemented with 10% FBS.

Antibodies and reagents

The following commercial antibodies were used in this study: a mouse monoclonal anti-CFTR C terminus antibody (clone 24-1, for immunoblotting) from R&D Systems (Minneapolis, MN); an anti-CFTR antibody (clone CF3, for immunofluorescence and FACS) from Novusbio (Littleton, CO); a monoclonal anti-GAPDH antibody (clone 71.1) from Sigma (St. Louis, MO); a β -Actin

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mouse monoclonal antibody (6G3) from Sungene Biotech (Tianjin, China); an anti-ZAP70 (99F2) rabbit mAb, an anti-Phospho-ZAP70 (Tyr319)/Syk (Tyr352) (65E4) rabbit mAb, an anti-PLCy1 antibody, an anti-Phospho-PLCy1 (Tyr783) (D6M9S) rabbit mAb, an anti-Lck antibody, an anti-p38 MAPK antibody, an anti-Phospho-p38 MAP kinase (Thr180/Tyr182) antibody, an anti-phospho-c-Jun (Ser73) (D47G9) XP[®] rabbit mAb, and an anti-Ezrin antibody from Cell Signaling Technology (Boston, MA); an anti-Lck (phospho Tyr394) rabbit polyclonal antibody from GeneTex (Irvine, CA); an anti-NFATc1 (phospho S237) antibody from Abcam (Cambridge, MA); an anti-mouse CD3 mAb (clone 145-2C11), an anti-mouse CD28 mAb (clone PV1), an anti-mouse TCR vδ mAb (clone UC7), an anti-mouse IFN-v antibodv (clone XMG1.2), a FITC-conjugated anti-mouse CD8 antibody, a PerCP-Cy5.5-conjugated anti-mouse CD4 antibody, a PE-conjugated anti-mouse TCR yo mAb, a PE-conjugated anti-mouse CD44 mAb, an APC-conjugated anti-mouse CD62L mAb, a PE-Cy7conjugated anti-mouse CD3 mAb, an APC-conjugated antimouse IL-17 mAb, and an APC-conjugated anti-mouse TNF-a mAb from Sungene Biotech (Tianjin, China); an Alexa Fluor 488conjugated anti-mouse TCR vo antibody and a Brilliant Violet 421[™] anti-mouse IFN-y antibody (clone XMG1.2) from BioLegend (San Diego, CA); and HRP-conjugated AffiniPure donkey antimouse and anti-rabbit secondary antibodies and Alexa Fluor 488- or Alexa Fluor 594-conjugated donkey anti-mouse and antirabbit IgGs from The Jackson ImmunoResearch Laboratory (West Grove, PA). All antibodies were used at the concentration recommended by the manufacturer. Protein A-agarose (sc-2001) was obtained from Santa Cruz (Dallas, TX). Anti-FLAG M2 magnetic beads (M8823) were obtained from Sigma (St. Louis, MO). The CFTR inhibitors CFTR_{inh172} and GlyH101, P38 MAPK inhibitor SB203580, Lck inhibitor PP2, calcineurin inhibitor FK506, and CLIC blocker IAA94 were purchased from Selleck (Houston, TX). The membrane potential probe DiBAC₄(3) (Bis-(1,3-Dibutylbarbituric Acid) Trimethine Oxonol), intracellular calcium indicator Fluo-4, cell viability indicator Calcein-AM, and Dynabeads Mouse T-activator CD3/CD28 for mouse T cell activation were obtained from Thermo Fisher (Carlsbad, CA). A fixation/permeabilization solution kit was purchased from BD Biosciences (San Jose, CA).

Electrophysiology

In vitro-expanded and purified $\gamma\delta$ T cells were subjected to wholecell voltage clamp studies. The pipette solution contained (in mM): 140 CsCl, 2 MgCl₂, 5 HEPES, 1 EGTA, 0.5 Li-ATP, 2 MgATP, and 10 glucose (pH 7.35). The bath solution contained (in mM): 170 Tris-Cl, 10 HEPES, 2.5 CaCl₂, 1 MgCl₂, and 15 glucose (pH 7.4). Reduction of [Cl⁻] in the external solution was performed by replacement of Tris-Cl with Tris-aspartate. the Patch pipettes had resistances of ~2–5 MΩ with these solutions. The procedure and data analysis were similar to those described in a previous protocol.²²

Cell preparation and activation and intracellular cytokine staining Mouse $\gamma\delta$ T cells were expanded and sorted as previously described.³⁹ For expansion of $\gamma\delta$ T cells, day-6 $\gamma\delta$ T cells were used, and the purity was checked (>99%). For analysis of TCR activation, $\gamma\delta$ T cells from wild-type or CFTR^{-/-} mice were activated with soluble anti-CD3 (10 µg/ml) and anti-CD28 (1 µg/ml) antibodies for the indicated times, followed by immunoblot analysis to assess the phosphorylation of PLC γ 1, ZAP70 (Tyr319), Lck (Tyr394), p38 MAP kinase (Thr180/Tyr182), and c-Jun (Ser73). Intracellular cytokine staining was carried out following the protocols in our previous publication.⁶¹ Stained cells were analyzed with a BD FACSVerseTM flow cytometer (San Jose, CA), and the data were analyzed with FlowJo 10.0 software.

The use of blood from healthy adult blood donors was approved by the Institutional Review Boards of Jinan University, Guangzhou.



Fig. 5 Pharmacological and genetic inhibition of CFTR both increased IFN- γ production in human V δ 2 T cells and V82mediated cytotoxicity to human K562 leukemia cells. a Flow cytometry analysis of in vitro-expanded human Vo2 T cells treated with anti-CD3 (10 µg/ml) and anti-CD28 (1 µg/ml) antibody stimulation for 6 h in the presence of the CFTR inhibitor CFTR $_{inh172}$ (5 μM) or vehicle DMSO was performed to detect IFN- γ production (n = 6, paired *t*-test, **p < 0.01). **b** In vitro-expanded human V δ 2 T cells were pretreated with the CFTR inhibitor $\mathsf{CFTR}_{\mathsf{inh172}}$ (5 $\mu\mathsf{M})$ or vehicle DMSO for 2 h and then cocultured with DIR-labeled K562 cells at the indicated ratio. Six hours later, dead K562 cells were identified by PI staining (n = 3, **p < 0.01, ****p < 0.001). **c** Flow cytometry analysis of in vitro-expanded human V δ 2 T cells infected with scramble shRNA or CFTR-specific shRNA lentiviral particles was performed. After 72 h of infection, the cells were stimulated with 5 µg/ml plate-bound anti-human CD3 antibodies and 1 µg/ml soluble anti-human CD28 antibodies before IFN- γ production was detected (upper panels, n = 3, **p < 0.01, ***p < 0.001). The cells were stained with anti-CFTR and PE-anti-mouse IgM (polyclonal) antibodies and analyzed by FACS. Histograms of CFTR expression (gating on GFP⁺CD3⁺ cells) and the MFI of CFTR are shown (lower panels, black line/number: scramble shRNA; red line/ number: CFTR-specific shRNA; dotted line: isotype control. n = 3, ***p < 0.001)

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Human Vδ2 T cells were expanded from PBMCs as previously described.⁵¹ For the expansion of Vδ2 T cells, day-12 Vδ2 T cells were used, and the purity was checked (>95%). For the detection of intracellular cytokines, Vδ2 T cells were stimulated for 6 h with 5 µg/ml plate-bound anti-human CD3 (clone OKT3, eBioscience) and 1 µg/ml soluble anti-human CD28 (clone CD28.2, eBioscience) antibodies in the presence of GolgiPlug[™] (BD Biosciences, San Jose, CA). For intracellular staining, cells were stained with different cocktails of fluorochrome-conjugated monoclonal antibodies specific for CD3 (clones SK7 and UCHT1) or TNF-α (clone MAb11) from BD Biosciences (Heidelberg, Germany), or IFN-γ (clone B27) or TCR Vδ2 (clone B6) from BioLegend (London, UK).

In vitro $\gamma\delta$ T1 cell priming

 $\gamma \delta^+ CD44^{low}$ (naive) and $\gamma \delta^+ CD44^{high}$ (active) T cells were isolated from the spleen of wild-type or CFTR^{-/-} mice. These cells were cultured (at a density of 1 × 10⁶ cells per well) in flat-bottomed 24well plates coated with anti-CD3 antibodies (10 µg/ml) in complete RPMI medium in the presence of anti-CD28 antibodies (1 µg/ml), recombinant mouse IL-12 (10 ng/ml), neutralizing anti-IL-4 antibodies (10 µg/ml), and recombinant mouse IL-2 (2 ng/ml). After 4 days, $\gamma \delta$ T cells were stimulated for 6 h with PMA (50 ng/ ml) and ionomycin (1 µg/ml) in the presence of GolgiPlug[™]. Intracellular cytokines were evaluated according to the manufacturer's instructions (eBioscience).

In vitro killing assay

B16-F0 cells or EL4 cells were labeled with 2 μ M CFSE. Then, the B16 cells or EL4 cells (8000 cells/well) were mixed with expanded $\gamma\delta$ T cells at different ratios for 6 h and analyzed by flow cytometry. The killing capability was evaluated by measuring the percentage of dead B16 or EL4 cells (% of PI⁺CFSE⁺ cells). In addition, an anti-IFN- γ neutralizing antibody (10 μ g/ml) was used to block IFN- γ mediated cytotoxicity, and an isotype control antibody was added to exclude the side effects of the anti-IFN- γ antibody. K562 cells were stained with DiR (Invitrogen) and analyzed by flow cytometry after 6 h of coculture with autologous V δ 2 T cells at different ratios. The killing capability was evaluated by measuring the percentage of dead K562 cells (% of PI⁺DIR⁺ cells).

Tumor models

B16-F0 melanoma cells were mixed with expanded and purified $\gamma\delta$ T cells and subcutaneously injected into TCR $\delta^{-/-}$ mice. Tumor size was monitored as described previously.³⁹

ELISA

In vitro-expanded and purified WT or CFTR^{-/-} $\gamma\delta$ T cells were treated with SB203580 or PP2 for 12 h, and the supernatants were collected for ELISA. The ELISA procedure followed the instructions of the IFN gamma Mouse Uncoated ELISA Kit (Thermo Fisher).

Real-time PCR for gene transcription

Total mRNA was isolated from cells using TRIzol[®] Reagent, and cDNAs were generated by reverse transcription-PCR using M-MLV reverse transcriptase with oligo(dT)₁₅ and TaKaRa Taq[®]. Real-time PCR was performed following the instructions of TB Green Premix Ex Taq II from TaKaRa. The primers for mouse genes were: CFTR-F: GGCACTCCGGTTAAGTAACTC, CFTR-R: TGTCACTTGGTCGAATTTG TTCA; CLCA-1-F: CTAAACATCCGGTCTGCTAGACT, CLCA-1-R: ACC CGTGCGTACACAATCATC; CLCA-4-F: GAAAGAACACTGTCGAAG-CAAC, CLCA-4-R: CAGTCTGGGAATTTGTTTGGGATA; GABA-1-F: GGTC ACAGTGAAAAACCGAATG, GABA-1-R: CGATGTCATCCGTGGTA-TAGCC; CLIC-1-F: CCCAGAGACTGTTCATGGTGC, CLIC-1-R: TCCCGA GGTGTTGGAACTCA; CLIC-3-F: CCTGCTGTACGATGGGGATG, CLIC-3-R: GAAGGCAGAGAACTTGTGGAAG; CLIC-4-F: TTGTGCCCACCCAAG-TACCTA, CLIC-4-R: CATTAGCCTCTGGTCTTGAGTTC; CLIC-5-F: GTG

AAGGCTGGGATCGACG, CLIC-5-R: CCAGGTTGTGTAGATCGGCT; CLIC-6-F: CCCCAGGATGAGGCGATTG, CLIC-6-R: GTCCCTTCAACTCG GGTTCT; LRRC8a-F: GATTGCTGTCTTTGGAGGGAC, and LRRC8a-R: GGCAGGACTGTGGAGTTGG. The primers for the internal control gene β -Actin were Actb-F: AACAGTCCGCCTAGAAGCAC and Actb-R: CGTTGACATCCGTAAAGACC. Real-time PCR was performed on a BioRad CFX Connect cycler. The expression of genes encoding several Cl⁻ channels in mouse $\gamma\delta$ T cells/CD4⁺ T cells was compared after verification that the amplification efficiencies of the different target genes were similar.

shRNA knockdown of CFTR expression in human V δ 2 T cells

shRNA constructs (encoding the GFP gene) targeting CFTR were purchased from Era Biotech (Shanghai, China). The sequences were as follows: shRNA1: GCTCTATCGCGATTTATCTAG, shRNA2: CCTATGTGAGATACTTCAATA, shRNA3: GAACACATACCTTCGATATAT, and shRNA4: GCCAAATGACTGTCAAAGATC. The constructs were used to generate lentiviral particles in HEK293T cells. Day-5 V δ 2 T cells were infected with an shRNA lentivirus, and at 72 h post infection, infected V δ 2 T cells were used for experiments.

Plasmid constructs and transfection

CFTR and CFTR- Δ PDZ were amplified by PCR and inserted into the pcDNA3.1-Myc plasmid (Invitrogen) using the KpnI and Notl restriction enzyme sites. PLCY-1 was amplified and inserted into the p3xFLAG-CMV-7.1 plasmid (Addgene) using the EcoRI and BamHI restriction enzyme sites. After verification by sequencing, the plasmids were cotransfected into HEK293T cells with Lipofectamine 3000 reagent (Invitrogen).

Coimmunoprecipitation

In vitro-expanded and purified $\gamma\delta$ T cells (~1 × 10⁷) were lysed in coimmunoprecipitation buffer (20 mM HEPES, pH 7.4; 175 mM NaCl; 1 mM EDTA; 1 mM DTT; 1 mM PMSF; 0.25% Nonidet P-40; 10% glycerol; and a protease inhibitor mixture (Roche Applied Science)), and the cell lysates were immunoprecipitated with 1 µg of anti-PLC γ 1 antibody and protein A-agarose beads. The immunoprecipitates were washed with co-IP buffer three times, eluted with 1 × SDS loading buffer, and analyzed by western blotting. For transfected HEK293T cells, lysates were immunoprecipitated with anti-FLAG M2 magnetic beads overnight at 4 °C. After three washes, the samples were eluted with 1 × SDS loading buffer and resolved using SDS-PAGE gels.

Immunofluorescence

Cells were fixed by incubation in 4% paraformaldehyde for 10 min and permeabilized in 0.2 M NH₄Cl/PBS plus 0.2% Triton X-100. After being blocked with 3% BSA/PBS, the cells were stained with an appropriate combination of primary antibodies, followed by staining with corresponding fluorophore-conjugated secondary antibodies. The cells were mounted on coverslips and examined under a confocal microscope (Leica). For analysis of the formation of the IS in $\gamma\delta$ T cells, purified $\gamma\delta$ T cells were treated with Dynabeads Mouse T-activator CD3/CD28 for 30 min, followed by immunostaining.

Measurement of the membrane potential and ${\rm Ca}^{2+}$ flux by flow cytometry

For determination of the membrane potential, cells were loaded with 2 μ M DiBAC₄(3) in phenol red-free RPMI 1640 medium at 37 °C and 5% CO₂ for 60 min. The cells were then washed, stained with an APC-conjugated anti-TCR γ/δ antibody, and resuspended at a density of 1×10^6 cells per ml in complete RPMI 1640 medium. After baseline acquisition on an Accuri C6 flow cytometer, the cells were treated with a CFTR antagonist (CFTR_{inh172}) or vehicle (0.1% DMSO), followed by treatment with 10 μ M gramicidin. Data are presented as the DiBAC₄(3) fluorescence (490 nm/516 nm) and were calculated for $\gamma\delta$ T cells gated with FlowJo software. For measurement of Ca²⁺ flux, cells were loaded with 2 μ M Fluo-4 in

PBS at 37 °C and 5% CO₂ for 30 min. The cells were then washed, stained with an APC-conjugated anti-TCR γ/δ antibody, and incubated in phenol red-free complete RPMI 1640 medium at 37 °C and 5% CO₂ for 60 min. After baseline acquisition on an Accuri C6 flow cytometer, the cells were treated with a CFTR antagonist (CFTR_{inh172}) or vehicle (0.1% DMSO), followed by the addition of 10 µg/ml anti-CD3 antibodies and 1 µg/ml anti-CD28 antibodies. Data are presented as the Fluo-4 fluorescence (494 nm/506 nm) and were calculated for $\gamma\delta$ T cells gated with FlowJo software.

Atomic force microscopy

Wild-type and CFTR^{-/-} $\gamma\delta$ T cells were seeded in poly-I-lysine-coated dishes at a density of 1×10^6 cells per 2 ml of medium. The in situ approach was used to measure adhesion force and Young's modulus by performing AFM and force analysis. The petri dishes with cells were directly transferred onto the AFM scanner stage for evaluation without any pretreatment, and whole measurements were conducted according to the protocols of previous studies.⁶²

Statistics

Data are expressed as the mean \pm SEM. Group sizes and the number of replications are stated in the figure legends. Statistical significance was evaluated using GraphPad Prism 6 for Windows (GraphPad). Tumor growth experiments and human V δ 2 T cell stimulation experiments were analyzed with paired *t*-tests. Flow cytometry, patch clamp, Q-PCR, actin quantification, AFM, V_m and Ca²⁺ influx recording, and stimulation experiments were analyzed using unpaired *t*-tests. Killing assay and ELISA results were analyzed using two-way ANOVA with a post hoc Bonferroni test. *p* values less than 0.05 were considered significant. The following annotations are used to show statistical significance: **p* < 0.05, ***p* < 0.01, *****p* < 0.001.

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AUTHOR CONTRIBUTIONS

Y.D., G.L., and M.X. designed and performed experiments and data analysis. X.Q. and J.T. performed the patch clamp experiment and data analysis. Z.J., Q.Y., M.D., and Z.Lei helped with in vitro cell expansion. Y.H. performed the AFM experiment. Z.Li helped with data analysis and drawing of the proposed model. Z.Liu and Q.W. helped with animal breeding. X.L., G.C., W.K.Z., P.H., L.Z., and R.A.F. contributed to data analysis. Z.Y. and J.H. designed the research and wrote the manuscript.

ADDITIONAL INFORMATION

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REFERENCES

- Cahalan, M. D. & Chandy, K. G. The functional network of ion channels in T lymphocytes. *Immunol. Rev.* 231, 59–87 (2009).
- Cai, X., Wang, X., Patel, S. & Clapham, D. E. Insights into the early evolution of animal calcium signaling machinery: a unicellular point of view. *Cell Calcium* 57, 166–173 (2015).
- 3. Clapham, D. E. Calcium signaling. Cell 131, 1047-1058 (2007).
- 4. Feske, S., Concepcion, A. R. & Coetzee, W. A. Eye on ion channels in immune cells. *Sci. Signal.* **12**, 572 (2019).
- 5. Feske, S. Calcium signalling in lymphocyte activation and disease. *Nat. Rev. Immunol.* **7**, 690–702 (2007).

- 6. Feske, S., Skolnik, E. Y. & Prakriya, M. Ion channels and transporters in lymphocyte function and immunity. *Nat. Rev. Immunol.* **12**, 532–547 (2012).
- Shaw, P. J., Qu, B., Hoth, M. & Feske, S. Molecular regulation of CRAC channels and their role in lymphocyte function. *Cell. Mol. Life Sci.* **70**, 2637–2656 (2013).
- Maul-Pavicic, A. et al. ORAl1-mediated calcium influx is required for human cytotoxic lymphocyte degranulation and target cell lysis. *Proc. Natl Acad. Sci. USA* 108, 3324–3329 (2011).
- Shaw, P. J. & Feske, S. Regulation of lymphocyte function by ORAI and STIM proteins in infection and autoimmunity. J. Physiol. 590, 4157–4167 (2012).
- Li, F. Y. et al. Second messenger role for Mg²⁺ revealed by human T-cell immunodeficiency. *Nature* 475, 471–476 (2011).
- Chandy, K. G., DeCoursey, T. E., Cahalan, M. D., McLaughlin, C. & Gupta, S. Voltagegated potassium channels are required for human T lymphocyte activation. *J. Exp. Med.* **160**, 369–385 (1984).
- DeCoursey, T. E., Chandy, K. G., Gupta, S. & Cahalan, M. D. Voltage-dependent ion channels in T-lymphocytes. J. Neuroimmunol. 10, 71–95 (1985).
- Launay, P. et al. TRPM4 regulates calcium oscillations after T cell activation. Science 306, 1374–1377 (2004).
- Launay, P. et al. TRPM4 is a Ca2+-activated nonselective cation channel mediating cell membrane depolarization. *Cell* 109, 397–407 (2002).
- Chimote, A. A. et al. A defect in KCa3.1 channel activity limits the ability of CD8(+) T cells from cancer patients to infiltrate an adenosine-rich microenvironment. *Sci. Signal.* **11**, 527 (2018).
- Crottes, D. et al. Immature human dendritic cells enhance their migration through KCa3.1 channel activation. *Cell Calcium* 59, 198–207 (2016).
- Kuras, Z., Yun, Y. H., Chimote, A. A., Neumeier, L. & Conforti, L. KCa3.1 and TRPM7 channels at the uropod regulate migration of activated human T cells. *PloS ONE* 7, e43859 (2012).
- Riordan, J. R. et al. Identification of the cystic fibrosis gene: cloning and characterization of complementary DNA. *Science* 245, 1066–1073 (1989).
- Chen, J. H., Schulman, H. & Gardner, P. A cAMP-regulated chloride channel in lymphocytes that is affected in cystic fibrosis. *Science* 243, 657–660 (1989).
- Shah, V. S. et al. Airway acidification initiates host defense abnormalities in cystic fibrosis mice. *Science* **351**, 503–507 (2016).
- Pankow, S. et al. F508 CFTR interactome remodelling promotes rescue of cystic fibrosis. *Nature* 528, 510–516 (2015).
- Duan, Y. et al. Keratin K18 increases cystic fibrosis transmembrane conductance regulator (CFTR) surface expression by binding to its C-terminal hydrophobic patch. J. Biol. Chem. 287, 40547–40559 (2012).
- Feske, S., Wulff, H. & Skolnik, E. Y. Ion channels in innate and adaptive immunity. Annu. Rev. Immunol. 33, 291–353 (2015).
- Puga Molina, L. C. et al. CFTR/ENaC-dependent regulation of membrane potential during human sperm capacitation is initiated by bicarbonate uptake through NBC. J. Biol. Chem. 293, 9924–9936 (2018).
- Wei, L. et al. The C-terminal part of the R-domain, but not the PDZ binding motif, of CFTR is involved in interaction with Ca(2+)-activated Cl- channels. *Pflug. Arch.* 442, 280–285 (2001).
- Ogura, T. et al. CIC-3B, a novel CIC-3 splicing variant that interacts with EBP50 and facilitates expression of CFTR-regulated ORCC. FASEB J. 16, 863–865 (2002).
- Welling, P. A. & Ho, K. A comprehensive guide to the ROMK potassium channel: form and function in health and disease. *Am. J. Physiol. Ren. Physiol.* 297, F849–F863 (2009).
- Mueller, C. et al. Lack of cystic fibrosis transmembrane conductance regulator in CD3+ lymphocytes leads to aberrant cytokine secretion and hyperinflammatory adaptive immune responses. Am. J. Respir. Cell Mol. Biol. 44, 922–929 (2011).
- Allard, J. B. et al. Aspergillus fumigatus generates an enhanced Th2-biased immune response in mice with defective cystic fibrosis transmembrane conductance regulator. J. Immunol. 177, 5186–5194 (2006).
- Dorsey, J. & Gonska, T. Bacterial overgrowth, dysbiosis, inflammation, and dysmotility in the Cystic Fibrosis intestine. J. Cyst. Fibros. 16(Suppl 2), S14–S23 (2017).
- Riquelme, S. A. et al. Cystic fibrosis transmembrane conductance regulator attaches tumor suppressor PTEN to the membrane and promotes anti pseudomonas aeruginosa immunity. *Immunity* 47, 1169–81 e7 (2017).
- Holderness, J., Hedges, J. F., Ramstead, A. & Jutila, M. A. Comparative biology of gammadelta T cell function in humans, mice, and domestic animals. *Annu. Rev. Anim. Biosci.* 1, 99–124 (2013).
- Chien, Y. H., Meyer, C. & Bonneville, M. gammadelta T cells: first line of defense and beyond. Annu. Rev. Immunol. 32, 121–155 (2014).
- Serre, K. & Silva-Santos, B. Molecular mechanisms of differentiation of murine proinflammatory gammadelta T cell subsets. Front. Immunol. 4, 431 (2013).

- Sutton, C. E. et al. Interleukin-1 and IL-23 induce innate IL-17 production from gammadelta T cells, amplifying Th17 responses and autoimmunity. *Immunity* 31, 331–341 (2009).
- Ferrick, D. A. et al. Differential production of interferon-gamma and interleukin-4 in response to Th1- and Th2-stimulating pathogens by gamma delta T cells in vivo. *Nature* 373, 255–257 (1995).
- Born, W. K., Yin, Z., Hahn, Y. S., Sun, D. & O'Brien, R. L. Analysis of gamma delta T cell functions in the mouse. J. Immunol. 184, 4055–4061 (2010).
- Gao, Y. et al. Gamma delta T cells provide an early source of interferon gamma in tumor immunity. J. Exp. Med. 198, 433–442 (2003).
- He, W. et al. Naturally activated V gamma 4 gamma delta T cells play a protective role in tumor immunity through expression of eomesodermin. J. Immunol. 185, 126–133 (2010).
- Chen, L. et al. Epigenetic and transcriptional programs lead to default IFN-gamma production by gammadelta T cells. J. Immunol. 178, 2730–2736 (2007).
- Lo Presti, E. et al. Current advances in gammadelta T cell-based tumor immunotherapy. Front. Immunol. 8, 1401 (2017).
- Alnaggar, M. et al. Allogenic Vgamma9Vdelta2 T cell as new potential immunotherapy drug for solid tumor: a case study for cholangiocarcinoma. J. Immunother. Cancer 7, 36 (2019).
- Hayes, S. M., Shores, E. W. & Love, P. E. An architectural perspective on signaling by the pre-, alphabeta and gammadelta T cell receptors. *Immunol. Rev.* 191, 28–37 (2003).
- Rigau, M. et al. Butyrophilin 2A1 is essential for phosphoantigen reactivity by gammadelta T cells. *Science* 367, eaay5516 (2020).
- Bertin, S. et al. The ion channel TRPV1 regulates the activation and proinflammatory properties of CD4(+) T cells. *Nat. Immunol.* 15, 1055–1063 (2014).
- Guggino, W. B. & Stanton, B. A. New insights into cystic fibrosis: molecular switches that regulate CFTR. *Nat. Rev. Mol. Cell Biol.* 7, 426–436 (2006).
- Roumier, A. et al. The membrane-microfilament linker ezrin is involved in the formation of the immunological synapse and in T cell activation. *Immunity* 15, 715–728 (2001).
- Calabia-Linares, C. et al. Endosomal clathrin drives actin accumulation at the immunological synapse. J. Cell Sci. 124(Pt 5), 820–830 (2011).
- 49. Roncagalli, R. et al. Quantitative proteomics analysis of signalosome dynamics in primary T cells identifies the surface receptor CD6 as a Lat adaptor-independent TCR signaling hub. *Nat. Immunol.* **15**, 384–392 (2014).
- Chakraborty, A. K. & Weiss, A. Insights into the initiation of TCR signaling. *Nat. Immunol.* 15, 798–807 (2014).
- Kouakanou, L. et al. Vitamin C promotes the proliferation and effector functions of human gammadelta T cells. *Cell Mol. Immunol.* 17, 462–473 (2019).
- Carding, S. R. & Egan, P. J. Gammadelta T cells: functional plasticity and heterogeneity. *Nat. Rev. Immunol.* 2, 336–345 (2002).
- Yin, Z. et al. Dominance of IL-12 over IL-4 in gamma delta T cell differentiation leads to default production of IFN-gamma: failure to down-regulate IL-12 receptor beta 2-chain expression. J. Immunol. 164, 3056–3064 (2000).
- Yin, Z. et al. T-Bet expression and failure of GATA-3 cross-regulation lead to default production of IFN-gamma by gammadelta T cells. J. Immunol. 168, 1566–1571 (2002).
- 55. Sun, G. et al. gammadelta T cells provide the early source of IFN-gamma to aggravate lesions in spinal cord injury. *J. Exp. Med.* **215**, 521–535 (2018).
- 56. Yang, Q. et al. Roles of mTORC1 and mTORC2 in controlling gammadelta T1 and gammadelta T17 differentiation and function. *Cell Death Differ*. **27**, 2248–2262 (2020).
- 57. Ponzetto, A., Holton, J. & Lucia, U. Cancer risk in patients with cystic fibrosis. Gastroenterology **154**, 2282–2283 (2018).
- Liu, M. et al. Treatment of human T-cell acute lymphoblastic leukemia cells with CFTR inhibitor CFTRinh-172. *Leuk. Res.* 86, 106225 (2019).
- 59. Yamada, A. et al. Risk of gastrointestinal cancers in patients with cystic fibrosis: a systematic review and meta-analysis. *Lancet Oncol.* **19**, 758–767 (2018).
- Abraham, J. M. & Taylor, C. J. Cystic fibrosis & disorders of the large intestine: DIOS, constipation, and colorectal cancer. J. Cyst. Fibros. 16(Suppl 2), S40–S49 (2017).
- Cao, G. et al. mTOR inhibition potentiates cytotoxicity of Vgamma4 gammadelta T cells via up-regulating NKG2D and TNF-alpha. *J. Leukoc. Biol.* 100, 1181–1189 (2016).
- McEwen, G. D. et al. Subcellular spectroscopic markers, topography and nanomechanics of human lung cancer and breast cancer cells examined by combined confocal Raman microspectroscopy and atomic force microscopy. *Analyst* **138**, 787–797 (2013).