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IV 研究成果の刊行に関する一覧 別冊

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Functional STAT3 deficiency compromises the generation of human T follicular helper cells

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T follicular helper (Tfh) cells are critical for providing the necessary signals to induce differentiation of B cells into memory and Ab-secreting cells. Accordingly, it is important to identify the molecular requirements for Tfh cell development and function. We previously found that IL-12 mediates the differentiation of human CD4⁺ T cells to the Tfh lineage, because IL-12 induces naive human CD4⁺ T cells to acquire expression of IL-21, BCL6, ICOS, and CXCR5, which

typify Tfh cells. We have now examined CD4⁺ T cells from patients deficient in IL-12Rβ1, TYK2, STAT1, and STAT3 to further explore the pathways involved in human Tfh cell differentiation. Although STAT1 was dispensable, mutations in *IL12RB1*, *TYK2*, or *STAT3* compromised IL-12–induced expression of IL-21 by human CD4⁺ T cells. Defective expression of IL-21 by STAT3-deficient CD4⁺ T cells resulted in diminished B-cell helper activity in vitro. Importantly, muta-

tions in *STAT3*, but not *IL12RB1* or *TYK2*, also reduced Tfh cell generation in vivo, evidenced by decreased circulating CD4⁺CXCR5⁺ T cells. These results highlight the nonredundant role of STAT3 in human Tfh cell differentiation and suggest that defective Tfh cell development and/or function contributes to the humoral defects observed in STAT3-deficient patients. (*Blood*. 2012;119(17):3997-4008)

Introduction

The generation of robust Ab responses is crucial for the correct functioning of the immune system. The importance of this is apparent in diseases that result from dysregulated humoral immune responses. For example, immunodeficient states and autoimmune disorders can develop as a consequence of impaired or exaggerated Ab responses, respectively. Thus, it is imperative to identify factors that control Ab responses. Early studies found that T cells play an important role in initiating Ab responses (reviewed in Tangye et al¹). This was mediated by instructive signals in the form of cell–cell contacts and secretion of soluble mediators such as cytokines. More recently, a subset of CD4⁺ T cells with specialized B-cell helper capabilities was identified that is now referred to as T follicular helper (Tfh) cells.^{2,3} Tfh cells are identified by several characteristics that also serve functional roles. Thus, Tfh cells express the chemokine receptor CXCR5,^{2,3} which facilitates their

positioning to B-cell follicles in secondary lymphoid tissues, and the transcription factor Bcl-6,⁴ which is required for the commitment of naive CD4⁺ T cells to the Tfh lineage.⁵⁻⁷ Tfh cells also express the costimulatory molecules CD40L, ICOS, OX40, and members of the SLAM family, as well as the cytokine IL-21,^{2-4,8-12} all of which play important roles in the induction of T cell–dependent (TD) B-cell activation and differentiation.

Because of the importance of Tfh cells in regulating Ab responses, much work has been performed to determine the requirements for their differentiation from naive CD4⁺ T cells. It was initially found that IL-21 was required for the development of murine Tfh cells.^{13,14} This was later expanded to include IL-6 and IL-27.¹⁵⁻¹⁷ However, conflicting findings have been made about the relative importance of IL-6 and IL-21 to murine Tfh cell formation^{18,19}; this may reflect redundancy because these cytokines, as

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Table 1. Primary immunodeficient patients

Disease	Patient ID	Mutation/genotype	
MSMD (IL-12Rβ1 deficiency)	IL12RB1#1	1745_1746insCA+1483 + 182-1619-1073del	
	IL12RB1#2	628_644dup	
	IL12RB1#3	R173P	
	IL12RB1#4	1791 + 2T > G	
	IL12RB1#5	C198R	
	IL12RB1#6	1623_1624delinsTT	
MSMD + viral infection (STAT1 deficiency)	STAT1#1	1928insA (homozygous)	
	STAT1#2	P696S (homozygous)	
	STAT1#3	P696S (homozygous)	
MSMD only (STAT1 deficiency)	STAT1#4	Q463H/WT	
	STAT1#5	L706S/WT	
	STAT1#6	L706S/WT	
AD-CMC	STAT1gof#1	A267V	
	STAT1gof#2	A267V	
	STAT1gof#3	A267V	
AD-HIES	STAT3#1	R382Q	
	STAT3#2	V637M	
	STAT3#3	R382Q	
	STAT3#4	H437P	
	STAT3#5	Q644P	
	STAT3#6	S465F	
	STAT3#7	Y657N	
	STAT3#8	R382W	
	STAT3#9	L706M	
	STAT3#10	L706M	
	STAT3#13	V463del	
	STAT3#14	V463del	
	STAT3#15	R593P	
	STAT3#16	V463del	
	AR-HIES (incl infection with mycobacteria, viruses and fungi)	TYK2#1	550_553GCTTdel (homozygous)
	MSMD + viral infection	TYK2#2	2292-2301del

MSMD indicates Mendelian susceptibility to mycobacterial disease; WT, wild type; AD-CMC, autosomal dominant chronic mucocutaneous candidiasis; AD-HIES, autosomal dominant hyper-IgE syndrome; AR-HIES, autosomal recessive hyper-IgE syndrome; and gof, gain-of-function

well as IL-27, can operate through STAT3.^{20,21} We and others previously showed that IL-12 is the key cytokine implicated in the differentiation of human Tfh cells in vitro.^{11,22} IL-6, IL-21, IL-23, and IL-27 also induce human Tfh-like cells in vitro, albeit to a much lesser extent than IL-12.^{11,17,22} We have now extended these observations by investigating the molecular requirements for the differentiation of naive human CD4⁺ T cells into Tfh cells. This was achieved by studying patients with primary immunodeficiencies resulting from mutations in *IL12RB1*, *STAT1*, *STAT3*, and *TYK2*. IL-12-mediated induction of human Tfh-like cells was abolished in the absence of IL-12Rβ1 or TYK2, and significantly reduced in CD4⁺ T cells deficient in STAT3 function. In contrast to the effects of IL-12, induction of Tfh cells by IL-6, IL-21, IL-23, and IL-27 was completely dependent on STAT3. These studies indicate that multiple cytokine pathways are involved in the differentiation of human Tfh cells, and IL-12 most efficiently induces human Tfh cells predominantly in a STAT3-dependent manner. This defect in generating Tfh cells from STAT3 mutant (STAT3_{MUT}) CD4⁺ T cells would contribute to impaired TD humoral immune responses observed in patients with *STAT3* mutations. In contrast, the ability of non-IL-12 cytokines to induce Tfh cell function is sufficient to elicit intact Ab responses in persons with impaired IL-12R signaling.

Methods

Human patient samples

Patients with mutations in *IL12RB1*, *STAT1*, *TYK2*, and *STAT3* have been previously described (Table 1²³⁻²⁸). PBMCs were isolated from these patients and healthy donors (Australian Red Cross). Tonsils were obtained from St Vincent's Hospital, Sydney. All studies were approved by Institutional Human Research Ethics Committees, and all participants gave written informed consent in accordance with the Declaration of Helsinki.

Antibodies

Alexa-647-conjugated anti-IL-21, biotinylated anti-ICOS, PE-anti-CD4, Pacific Blue-anti-CD4, peridinin chlorophyll protein complex (PerCP)/cyanine 5.5-anti-CD45RA, anti-IFNγ, and FITC-anti-CD45RA were purchased from eBiosciences. Alexa-647-anti-CXCR5, allophycocyanin-anti-CD38, FITC-anti-CD20, PE-anti-CD4, anti-CD27, PerCP-anti-CD3 mAb, and streptavidin-PerCP were purchased from Becton Dickinson. Allophycocyanin-anti-CD4 was purchased from Caltag, and FITC-anti-CCR7 was purchased from R&D Systems.

CD4⁺ T-cell isolation

CD4⁺ T cells were isolated from healthy donors or immunodeficient patients with the use of Dynal beads.²³ Peripheral blood (PB) CD4⁺ T cells were labeled with anti-CD4, anti-CD45RA, and anti-CCR7, and naive

Table 2. Primers for qPCR

Gene	Primers	UPL probe	Amplicon size, bp
<i>BCL6</i>	fwd: gagctctgttgattcttagaactgg rev: gcctgcttcacagtccaa	9	110
<i>TBX21</i>	fwd: tgtggccaagttaatacagca rev: tgacaggaatgggaacatcc	9	77
<i>IL21</i>	fwd: aggaaccacctccacaaa rev: gaatcacatgaaggcatgtt	7	68
<i>IFNG</i>	fwd: ggcattttgagaattggaag rev: ttggatgctctggtcactt	21	112
<i>GAPDH</i>	fwd: cctgctcctcctgttcgac rev: acgaccaaatccgtgactc	60	112

CD45RA⁺CCR7⁺ CD4⁺ T cells were isolated (> 98% purity) with the use of a FACSAria (BD Biosciences).

Cell cultures

Naive PB CD4⁺ T cells were labeled with CFSE (Molecular Probes) and cultured with T-cell activation and expansion beads (anti-CD2/CD3/CD28; Miltenyi Biotec) alone (nil culture) or under Th1 (IL-12 [20 ng/ml; R&D systems]), Th2 (IL-4 [100 U/ml]), or Th17 (IL-1 β [20 ng/ml; Peprotech]), IL-6 (50 ng/ml; PeproTech), IL-21 (50 ng/ml; PeproTech), IL-23 (20 ng/ml; eBioscience), anti-IL-4 (5 μ g/ml), and anti-IFN γ (5 μ g/ml; eBioscience)^{23,29} polarizing conditions, or with IL-6, IL-21, IL-23, or IL-27 (50 ng/ml; eBioscience) alone. After 4 or 5 days, expression of intracellular cytokines, transcription factors, and surface phenotype of cells determined.

T- and B-cell coculture assays

Naive CD4⁺ T cells were activated for 5 days (see previous section), treated with mitomycin C (100 μ g/ml; Sigma-Aldrich) and then cocultured at a 1:1 ratio (50 \times 10³/200 μ L/well) with sort-purified allogeneic naive (CD20⁺CD27⁻CD38^{inter}) tonsillar B cells.^{11,29} After 7 days Ig secretion was determined by ELISA.²⁹

Cytokine and transcription factor expressions

Activated CD4⁺ T cells were restimulated with phorbol 12-myristate 13-acetate (100 ng/ml) and ionomycin (750 ng/ml) for 6 hours, with Brefeldin A (10 μ g/ml) added after 2 hours. Cells were then fixed with formaldehyde, and expression of cytokines was detected by intracellular staining.^{23,29} RNA was extracted with the use of RNeasy kit (QIAGEN) and transcribed into cDNA with the use of random hexamers and Superscript III (Invitrogen). All quantitative PCR (qPCR) primers (Integrated DNA Technologies) were designed with Roche UPL Primer Design Program. Primer sequences, Roche UPL probes, and size of each amplicon are listed in Table 2. qPCR was performed with Roche LightCycler 480 Probe Master Mix and Roche Lightcycler 480 System with the following conditions: denaturation at 95°C for 10 minutes; amplification for 45 cycles at 95°C for 10 seconds, 65°C for 30 seconds, and 72°C for 5 seconds; and cooling at 40°C for 30 seconds. All reactions were standardized to *GAPDH*.

Results

Patients deficient for IL-12R β 1 have altered differentiation of CD4⁺ T cells in vivo

IL-12 can induce human naive CD4⁺ T cells to differentiate into IL-21-expressing cells that resemble T_H cells in vitro.^{11,22} To investigate this function of IL-12 further, we examined patients with homozygous or compound heterozygous null mutations in *IL12RB1*.²⁶ We first determined the frequency of total CD4⁺ T cells and CD4⁺ T cells with a naive (CD45RA⁺CCR7⁺), memory (CD45RA⁻CCR7^{+/-}), or T_H (CXCR5⁺) phenotype in healthy

donors (age range, 16-64 years) and patients deficient for IL-12R β 1 (Figure 1A-D,F). Patients deficient for IL-12R β 1 had a normal frequency of CD4⁺ T cells (Figure 1; Table 3). In contrast, they had a significant increase in the frequency of naive and a corresponding significant decrease in memory CD4⁺ T cells (Figure 1A-E; Table 3). When the phenotype of CXCR5⁺ T cells was analyzed, ~90% were found within the CD45RA⁻ (ie, memory) subset (Table 3^{2,3,11}). Therefore, we analyzed the frequency of both CXCR5⁺CD45RA⁻ and CD45RA⁺ T cells in healthy donors and in patients deficient for IL-12R β 1. No significant difference was observed for the frequency of circulating CD4⁺ CXCR5⁺CD45RA⁻ or CD45RA⁺ T_H-like cells in healthy donors and patients deficient for IL-12R β 1 (Figure 1A-B,F; Table 3).

Naive CD4⁺ T cells from patients deficient for IL-12R β 1 are unable to differentiate into IL-21-expressing cells in response to IL-12

To assess the potential of CD4⁺ T cells deficient for IL-12R β 1 to differentiate into T_H-like cells, we examined the ability of naive cells to express IL-21 in vitro. Naive CD4⁺ T cells were cultured with T-cell activation and expansion beads alone (nil) or with IL-12 (Th1). After 5 days, cells were restimulated with phorbol 12-myristate 13-acetate/ionomycin, and the expression of IL-21 and IFN γ was then determined. Naive CD4⁺ T cells from either healthy donors or patients deficient for IL-12R β 1 expressed little IL-21 or IFN γ when cultured under neutral (nil) conditions (Figure 1G-J). However, when normal naive CD4⁺ T cells were cultured under Th1-polarizing conditions (ie, with IL-12), IL-21- and IFN γ -expressing cells were readily detectable (Figure 1G,I,J). In contrast, IL-12 failed to induce IL-21 or IFN γ in naive CD4⁺ T cells deficient for IL-12R β 1 (Figure 1H-J). Next, we questioned whether naive CD4⁺ T cells deficient for IL-12R β 1 could differentiate into IL-21-expressing cells in response to other cytokines and signaling pathways. Accordingly, naive CD4⁺ T cells from healthy donors and patients deficient for IL-12R β 1 were subjected to Th2 (IL-4) and Th17 (IL-1 β , IL-6, IL-21, IL-23) polarizing conditions or were cultured in the presence of IL-6, IL-21, IL-23, or IL-27. A small frequency of IL-21-expressing cells could be generated from both normal and IL-12R β 1-deficient naive CD4⁺ T cells activated with IL-21 or IL-27 (Figure 1I). Similarly, although IL-12 could not induce IFN γ in naive CD4⁺ T cells deficient for IL-12R β 1, the ability of IL-27 to induce IFN γ was unaffected by *IL12RB1* mutations (Figure 1J). Taken together these results indicate that, although IL-12-induced IL-21 expression is abrogated by *IL12RB1* mutations, other cytokines and their associated signaling pathways that induce IL-21 (eg, IL-21 and IL-27, albeit to a lesser extent than IL-12) are intact, which is consistent with normal Ab responses to infection and vaccinations in these patients.^{30,31}

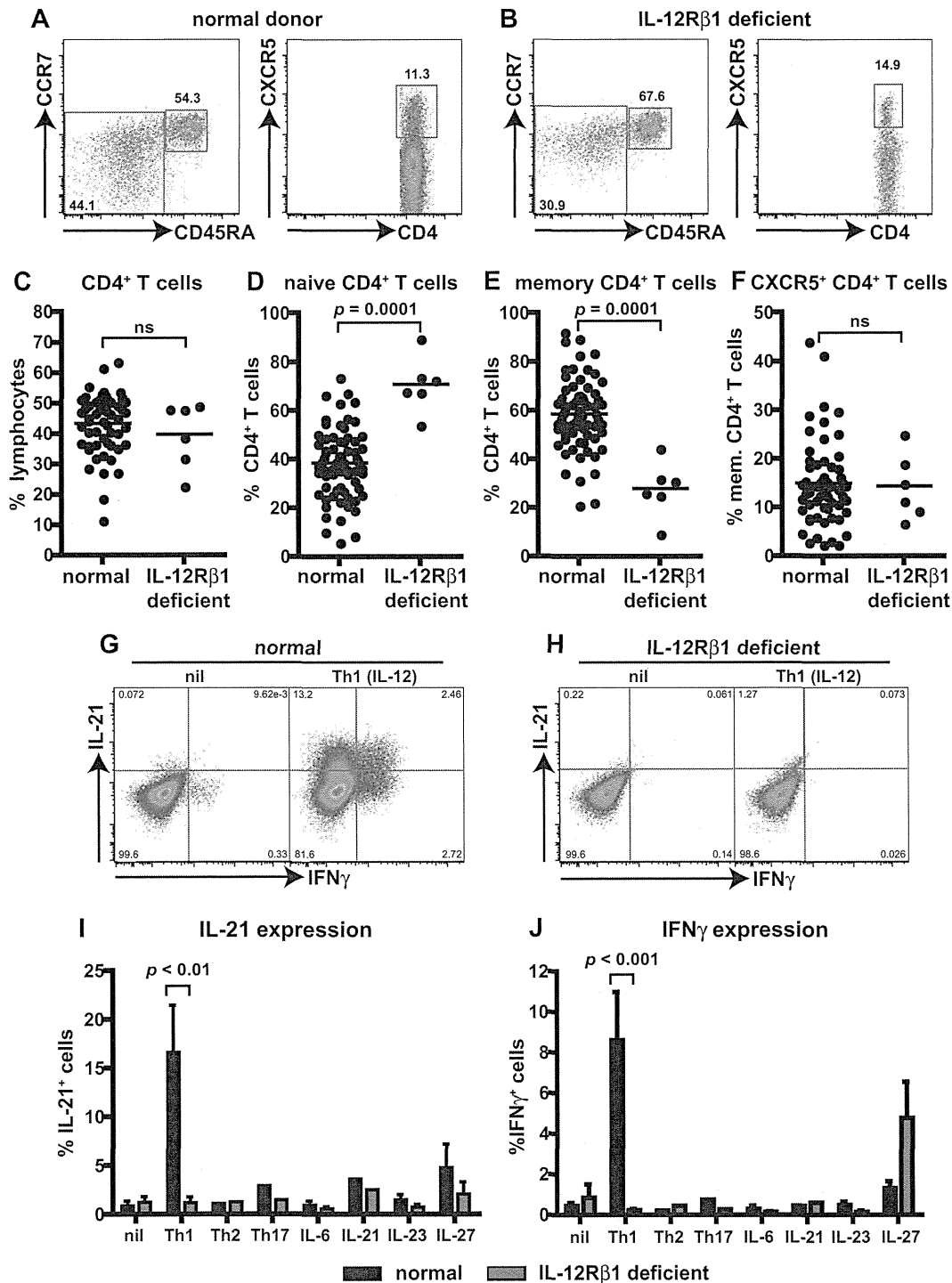


Figure 1. Naive CD4⁺ T cells deficient for IL-12Rβ1 fail to differentiate into IL-21–expressing cells in response to IL-12. (A–F) The frequency of total CD4⁺ T cells, and naive (CD45RA⁺CCR7⁺), memory (CD45RA⁺CCR7^{-/-}), and CXCR5⁺CD45RA⁻ CD4⁺ T cells, in PBMCs was determined for healthy donors and patients deficient for IL-12Rβ1. (A–B) Representative dot plots from 1 donor and 1 patient. (C–F) The frequency of (C) total, (D) naive (CD45RA⁺CCR7⁺), (E) memory (CD45RA⁺CCR7^{-/-}), and (F) CXCR5⁺CD45RA⁻ CD4⁺ T cells from all healthy donors (total CD4⁺ T cells, n = 54; naive CD4⁺ T cells, n = 70; memory CD4⁺ T cells, n = 70; CXCR5⁺CD45RA⁻ CD4⁺ T cells, n = 61) and patients deficient for IL-12Rβ1 examined (n = 6). (G–J) Naive CD4⁺ T cells isolated from healthy donors (n = 5) and patients deficient for IL-12Rβ1 (n = 5) were cultured for 5 days under neutral (nil); polarizing Th1, Th2, or Th17 conditions; or in the presence of IL-6, IL-21, IL-23, or IL-27, and intracellular expression of IL-21 and IFN γ were then determined. (G–H) Representative dot plots of IL-21 and IFN γ expression by activated naive CD4⁺ T cells from 1 donor and 1 patient deficient for IL-12Rβ1. (I–J) Percentage of activated normal and IL-12Rβ1–deficient naive CD4⁺ T cells induced to express (I) IL-21 or (J) IFN γ in response to the indicated culture. The values represent the mean \pm SEM.

Analysis of cytokine responsiveness in STAT-deficient human CD4⁺ T cells

The cytokines that induce IL-21 in human naive CD4⁺ T cells (IL-12, IL-6, IL-21, IL-23, IL-27)^{11,22} function by activating

JAK/STAT signaling pathways. These cytokines phosphorylate STAT1 (IL-6, IL-12, IL-21, IL-23, IL-27), STAT3 (IL-6, IL-12, IL-21, IL-23, IL-27), STAT4 (IL-12, IL-23), and STAT5 (IL-12).^{20,21,24,32-35} We confirmed these studies by showing that these

Table 3. CD4⁺ T-cell subsets in primary immunodeficiencies

PID	CD4 ⁺ T cells	Naive CD4 ⁺ T cells	Memory CD4 ⁺ T cells	CXCR5 ⁺ CD45RA ⁺ CD4 ⁺ T cells	CXCR5 ⁺ CD45RA ⁻ CD4 ⁺ T cells
IL12RB1 (n = 6)	40 ± 4.4	71 ± 4.7	28 ± 4.7	1.6 ± 0.46	14 ± 2.8
STAT1 (n = 6)	40*	41 ± 10	56 ± 10	3.9 ± 0.68	11 ± 2.3
STAT3 (n = 14)	41 ± 3.1	65 ± 3.8	33 ± 3.7	1.5 ± 0.29 (n = 11)	8.7 ± 1.6 (n = 11)
TYK2 (n = 2)	41*	53 ± 24	43 ± 27	2.8 ± 1.2	15 ± 5.4
Controls	43 ± 1.4 (n = 54)	39 ± 1.7 (n = 70)	59 ± 1.7 (n = 70)	2.8 ± 0.24 (n = 61)	15 ± 1.1 (n = 61)

Values are mean ± SEM percentage.

PID indicates primary immunodeficiency.

*Analysis was only performed on 1 patient.

IL-21–inducing cytokines predominantly activate STAT1, STAT3, and STAT4 in human naive CD4⁺ T cells (data not shown), suggesting that these transcription factors are important in inducing T_H cells in humans. To investigate this further, we took advantage of additional primary immunodeficiencies that result from mutations in components of several cytokine-signaling pathways, namely *STAT1*, *TYK2*, and *STAT3*.

STAT1 is dispensable for IL-12–induced expression of IL-21 in human CD4⁺ T cells

STAT1-deficient patients and healthy donors had comparable frequencies of naive and memory CD4⁺ T cells (Figure 2A-D; Table 3). The proportions of CXCR5⁺CD45RA⁻ or CD45RA⁺ T_H cells in STAT1-deficient patients were also comparable with healthy donors (Figure 2A-B,E; Table 3). The ability of STAT1-deficient CD4⁺ T cells to differentiate into IL-21⁺ cells in response to Th1-polarizing conditions was then tested. Because of the limited numbers of cells available from these rare patients, total CD4⁺ T cells were examined. When CD4⁺ T cells from healthy donors or STAT1-deficient patients were activated under neutral conditions, IL-21 and IFN γ production could be detected; however, the frequencies of cytokine-expressing cells was lower in STAT1-deficient patients (Figure 2F,H; IL-21: normal, 9.1% ± 6.2%; STAT1, 2.7% ± 1.8%; IFN γ : normal, 11.3% ± 3.8%; STAT1, 1.6% ± 0.9%). Despite these differences in the nonpolarizing cultures, expression of IL-21 and IFN γ in both normal and STAT1-deficient CD4⁺ T cells increased after culture with IL-12 (Figure 2F,H; IL-21: normal, 30.5% ± 7.9%; STAT1, 13.2% ± 6.7%; IFN γ : normal, 31.3% ± 7.9%; STAT1, 12.3% ± 2.5%). Although there appeared to be a reduced ability of IL-12 to enhance expression of IL-21 and IFN γ in the absence of STAT1, compared with normal CD4⁺ T cells (Figure 2F,H), when the effect of IL-12 was expressed as fold-change relative to nonpolarizing cultures, STAT1-deficient CD4⁺ T cells responded comparably to normal cells, that is, ~4- to 6-fold increase in IL-21 (Figure 2G) and ~3- to 6-fold induction in IFN γ expression (Figure 2I). Together these results indicate that IL-12–induced IL-21 and IFN γ in CD4⁺ T cells is independent of STAT1 signaling.

Mutations in *STAT3* compromise the generation of CD4⁺ memory and CXCR5⁺ T_H cells

To investigate further the signaling pathways involved in the differentiation of IL-21–expressing cells we used patients with autosomal-dominant hyper IgE syndrome resulting from heterozygous mutations in *STAT3*.^{36,37} Patients heterozygous for these mutations display impaired, but not abolished, STAT3 function with ~25% residual signaling.^{36,37} As detailed above, STAT3 is activated by many cytokines, including those that induce IL-21

expression in human CD4⁺ T cells (ie, IL-6, IL-12, IL-21, IL-23, IL-27).^{20,21,24,32-35} The frequency of total CD4⁺ T cells was comparable between healthy donors and STAT3_{MUT} patients (Figure 3C). Compared with healthy donors, STAT3_{MUT} patients have a significant increase in the frequency of naive and a significant decrease in the frequency of memory CD4⁺ T cells (Figure 3A-F; Table 3). Analysis of CXCR5⁺ T cells within the CD45RA⁺ and CD45RA⁻ fractions revealed significant decreases in both of these compartments in STAT3_{MUT} patients compared with healthy donors (Figure 3A-B,F; Table 3). Thus, mutations in *STAT3* cause a ~50% reduction in the frequency of circulating CXCR5⁺CD45RA⁻ and CD45RA⁺ T_H cells (Figure 3F; Table 3).

STAT3 mutations partially impairs IL-12–induced expression of IL-21 in naive CD4⁺ T cells

The observations that (1) IL-12 is the main driver of human T_H cell differentiation in vitro,^{11,22} (2) IL-12 is capable of activating STAT3,^{21,32-35} and (3) STAT3_{MUT} patients have a contracted population of circulating CD4⁺ CXCR5⁺CD45RA⁻ and CD45RA⁺ T_H-like cells (Figure 3E) led us to investigate the consequences of *STAT3* mutations on the ability of IL-12 to induce IL-21 in human naive CD4⁺ T cells. No differences were observed in the low frequencies of IL-21⁻ and IFN γ –expressing cells detected in cultures of naive CD4⁺ T cells from healthy donors or STAT3_{MUT} patients after stimulation under neutral conditions (Figure 4A-D). However, induction of IL-21 expression by IL-12 was significantly reduced in STAT3_{MUT} naive CD4⁺ T cells compared with normal naive CD4⁺ T cells (Figure 4A-C; normal, 20.9% ± 4.5%; STAT3_{MUT}, 5.2% ± 1.3%). In fact this defect was most pronounced when the frequency of IL-21⁺IFN γ ⁻ cells were determined (Figure 4A-B). This analysis found a significant ($P < .01$) decrease in IL-21⁺IFN γ ⁻ cells but not IL-21⁺IFN γ ⁺ ($P > .05$) cells in STAT3_{MUT} patients compared with healthy donors after Th1 polarization (IL-21⁺IFN γ ⁻: normal, 16.7% ± 3.2%; STAT3_{MUT}, 2.3% ± 0.5%; IL-21⁺IFN γ ⁺: normal, 4.8% ± 1.5%; STAT3_{MUT}, 3.3% ± 1%). In contrast to IL-21, expression of IFN γ in CD4⁺ T cells in response to IL-12 was unaffected by *STAT3* mutations; in fact it was higher than normal CD4⁺ T cells (Figure 4D). This increase in IFN γ production by STAT3_{MUT} CD4⁺ T cells, however, did not contribute to decreased IL-21 production because neutralizing IFN γ in these cultures had no effect on IL-21 expression, and there was a positive correlation between IL-21 and IFN γ expression in both normal and STAT3_{MUT} naive CD4⁺ T cells after Th1 polarization (data not shown). It was possible that induction of IL-21 by IL-12 resulted from not only a direct effect of IL-12 but also the effects of other cytokines induced by IL-12 that act in an autocrine manner to further promote IL-21 expression. To address this, we examined IL-21 and IFN γ expression in normal and STAT3_{MUT} CD4⁺ T cells by qPCR after 24 and 48 hours of

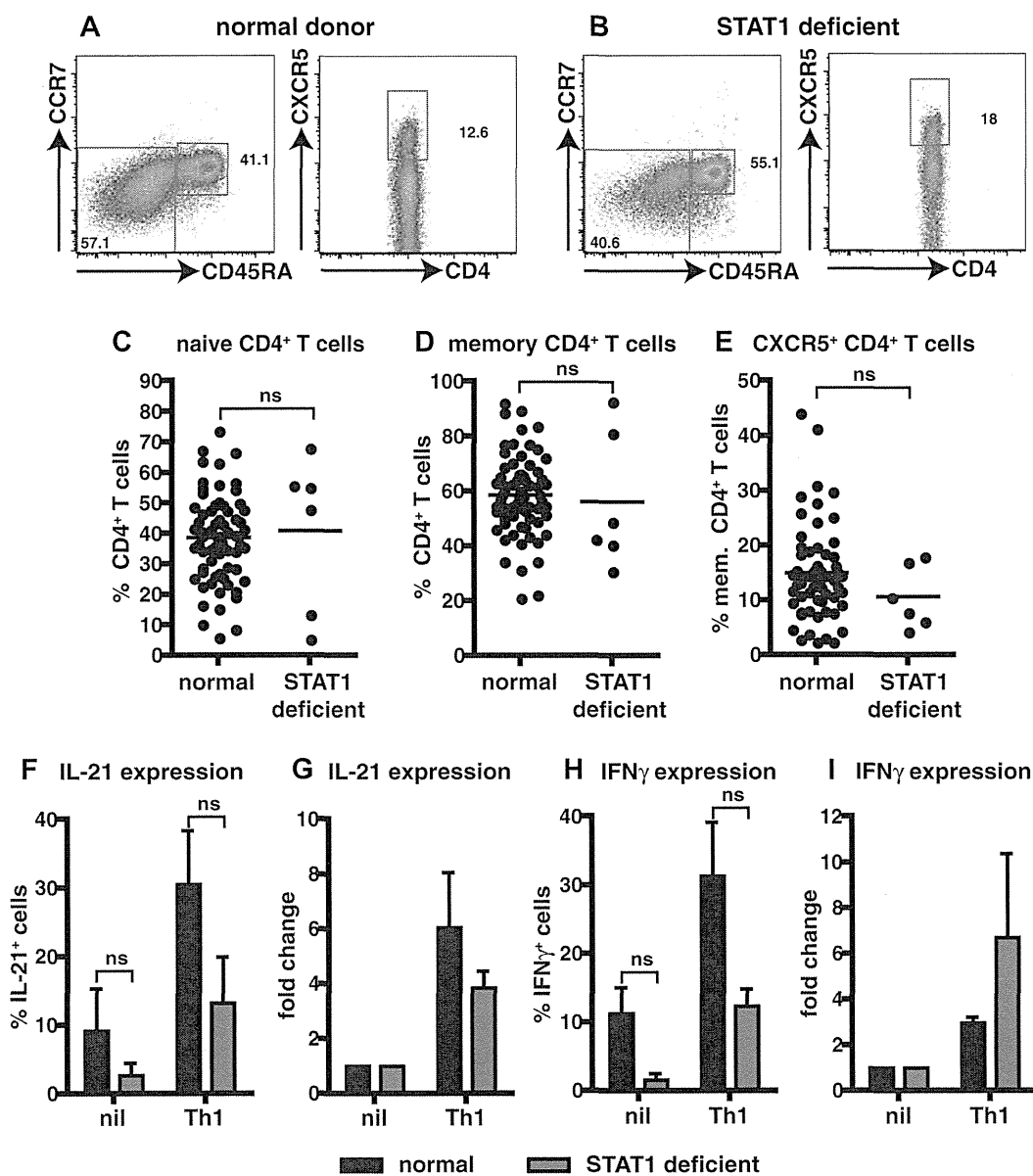


Figure 2. STAT1 is dispensable for IL-12-induced expression of IL-21 in human CD4⁺ T cells. (A-E) The frequency of naive (CD45RA⁺CCR7⁺), memory (CD45RA⁺CCR7^{-/-}), and CXCR5⁺CD45RA⁻ CD4⁺ T cells in PBMCs was determined for healthy donors and STAT1-deficient patients. (A-B) Representative dot plots from 1 donor and 1 STAT1-deficient patient. (C-E) The frequency of (C) naive (CD45RA⁺CCR7⁺), (D) memory (CD45RA⁺CCR7^{-/-}), and (E) CXCR5⁺CD45RA⁻ CD4⁺ T cells from all healthy donors (total CD4⁺ T cells, n = 54; naive CD4⁺ T cells, n = 70; memory CD4⁺ T cells, n = 70; CXCR5⁺CD45RA⁻ CD4⁺ T cells, n = 61) and STAT1-deficient patients (n = 6) was examined. (F-I) Total CD4⁺ T cells isolated from healthy donors and STAT1-deficient patients (n = 3) were cultured for 5 days under neutral (nil) or Th1-polarizing (ie, IL-12) conditions, and expression of intracellular IL-21 (F-G) and IFN γ (H-I) was then determined. The graphs in panels F and H show the frequency of cytokine-positive cells; those in panels G and I depict cytokine expression after Th1 polarization as fold-increase relative to the nil culture in each experiment. The values represent the mean \pm SEM (n = 3).

stimulation with or without IL-12. Expression of *IL21* was substantially increased by IL-12 in normal CD4⁺ T cells within 24 hours of stimulation, compared with cells cultured under neutral (nil) conditions, and similar levels were detected after 48 hours (supplemental Figure 1A-B, available on the *Blood* Web site; see the Supplemental Materials link at the top of the online article). Consistent with the flow cytometric analysis, expression of *IL21* in IL-12-activated STAT3_{MUT} naive CD4⁺ T cells was reduced compared with normal controls (supplemental Figure 1A-B). In contrast to *IL21*, *IFNG* was abundantly expressed by IL-12-treated STAT3_{MUT} naive CD4⁺ T cells; in fact, consistent with the intracellular staining data, *IFNG* expression by these cells exceeded that of normal naive CD4⁺ T cells at the 48-hour time point

(supplemental Figure 1C-D). These data suggest that IL-12 acts directly to rapidly induce IL-21 expression in naive CD4⁺ T cells.

IL-12 is capable of activating the JAK family protein tyrosine kinase TYK2,³⁸ which subsequently phosphorylates STATs, including STAT3.³⁹ Mutations in *TYK2* have been reported in 2 patients who developed susceptibility to various pathogens, including mycobacteria/Bacille Calmette-Guérin and herpes viruses.^{25,28} One of the contributing factors to disease pathogenesis is believed to be the unresponsiveness of their T cells to IL-12 with respect to induction of IFN γ expression.²⁵ To investigate this further, these 2 patients deficient for TYK2 were examined (supplemental Figure 2). CXCR5⁺CD45RA⁻ Tfh-like cells were detected in both patients at similar frequencies as healthy donors (supplemental Figure 2A-B,E; Table 3). When IL-21 expression in total

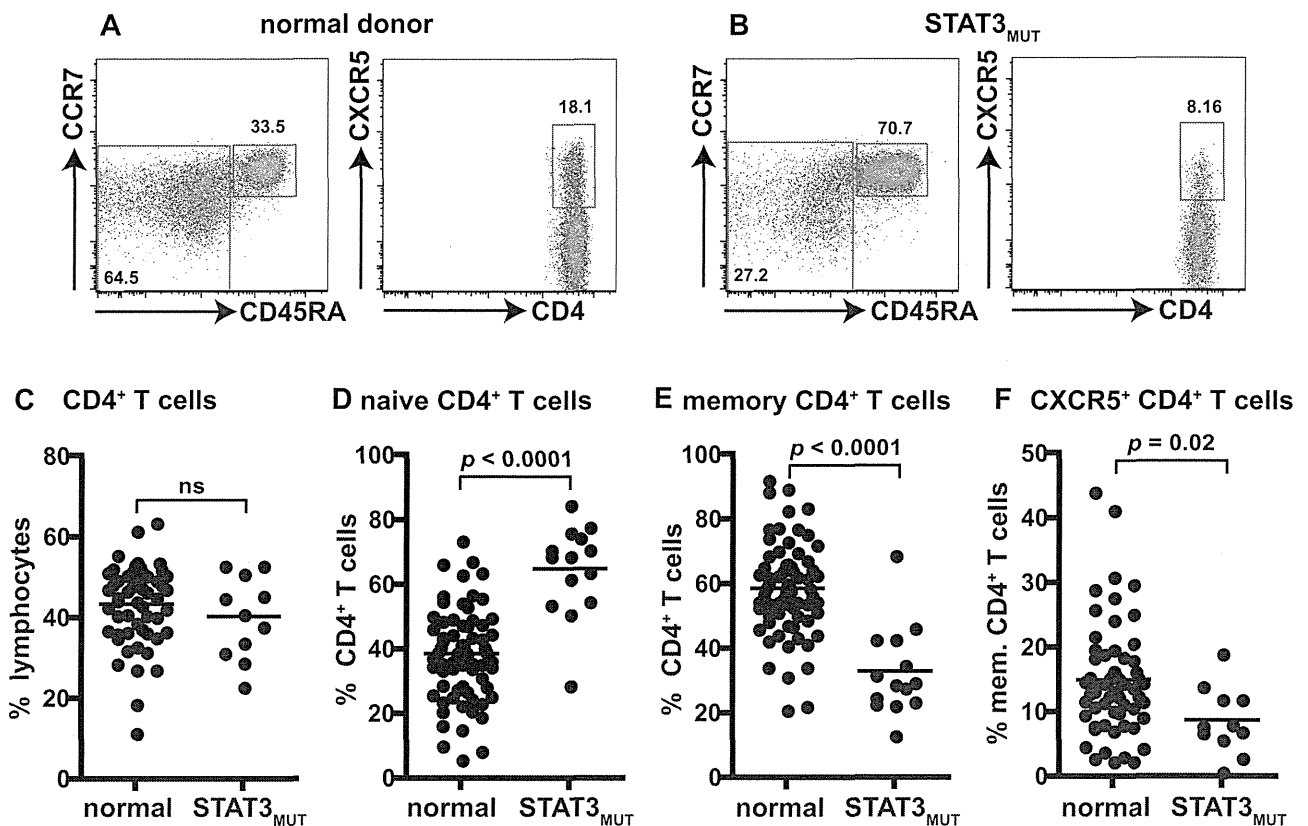


Figure 3. Mutations in *STAT3* compromise the generation of CD4⁺ memory and CXCR5⁺CD45RA⁻ Tfh cells. (A-B) The frequency of naive (CD45RA⁺CCR7⁺), memory (CD45RA⁺CCR7^{-/+}), and CXCR5⁺CD45RA⁻ CD4⁺ T cells in PBMCs was determined for healthy donors and STAT3_{MUT} patients. (A-B) Representative dot plots from 1 donor and 1 patient. (C-F) The frequency of (C) total CD4⁺ T cells, (D) naive (CD45RA⁺CCR7⁺), (E) memory (CD45RA⁺CCR7^{-/+}), and (F) CXCR5⁺CD45RA⁻ CD4⁺ T cells from all healthy donors (total CD4⁺ T cells, n = 54; naive CD4⁺ T cells, n = 70; memory CD4⁺ T cells, n = 70; CXCR5⁺CD45RA⁻ CD4⁺ T cells, n = 61) and STAT3_{MUT} patients (n = 14) was examined.

CD4⁺ T cells from 1 patient deficient for *TYK2* after culture with IL-12 was examined, the level of induction was ~2-fold less than that observed for normal CD4⁺ T cells (supplemental Figure 2F-G). We extended these studies by examining naive CD4⁺ T cells from the second patient deficient for *TYK2*. Compared with normal naive CD4⁺ T cells, induction of IL-21 expression by IL-12 in *TYK2*-deficient CD4⁺ T cells was severely reduced (supplemental Figure 2I). Induction of IFN γ was also dramatically compromised by *TYK2* mutations (supplemental Figure 2H,J). Taken together, these data suggest that mutations in *TYK2* and *STAT3*, which are both activated downstream of the IL-12R₂,^{21,32-35,38,39} cause a significant impairment in the ability of naive CD4⁺ T cells to differentiate into IL-21-expressing cells in response to IL-12.

STAT3 mutations impede division-linked differentiation of CD4⁺ T cells into IL-21-expressing effector cells

Because IL-12 can promote proliferation of human activated T cells,⁴⁰ and differentiation of CD4⁺ T cells into cytokine-expressing cells is linked to cell division,^{29,41} it was important to determine whether the reduction in expression of IL-21 in IL-12-stimulated STAT3_{MUT} CD4⁺ T cells resulted from reduced cell division or reflected an intrinsic defect in the differentiation program of these cells.

To do this, we examined proliferation by labeling naive CD4⁺ T cells with CFSE and tracked their division after in vitro stimulation by monitoring CFSE dilution. There was no difference in cell division between normal or STAT3_{MUT} naive CD4⁺ T cells

activated under neutral conditions (Figure 4E). Similarly, IL-12 increased proliferation regardless of whether the naive CD4⁺ T cells were derived from healthy donors or STAT3_{MUT} patients (Figure 4E). When IL-21 expression was examined on a per division basis, we found that it increased in normal CD4⁺ T cells after the first few divisions and then reached a maximum, being expressed in ~20%-30% of cells after 2-3 divisions (Figure 4F). Expression of IL-21 by STAT3_{MUT} CD4⁺ T cells also modestly increased with division; however, the frequency of IL-21⁺ cells in each division never exceeded 10% and thus was dramatically reduced compared with normal CD4⁺ T cells (Figure 4F). These data establish that the defect in IL-12-induced IL-21 induction in the absence of functional STAT3 was not because of a difference in proliferation, but rather because of an inability to efficiently acquire IL-21 during Tfh cell differentiation. Notably, when IFN γ expression was also analyzed on a per division basis, the frequency of cytokine-positive cells continued to increase with each cell division (Figure 4G). Furthermore, the heightened frequency of IFN γ ⁺ cells observed in STAT3_{MUT} compared with normal CD4⁺ T cells at the population level (Figure 4D and supplemental Figure 1) was also detected for cells that had undergone different divisions (Figure 4G).

IL-12 can induce additional characteristics of Tfh cells in STAT3 mutant CD4⁺ T cells

IL-12 not only promotes IL-21 expression in human naive CD4⁺ T cells but also induces additional features of Tfh cells such as

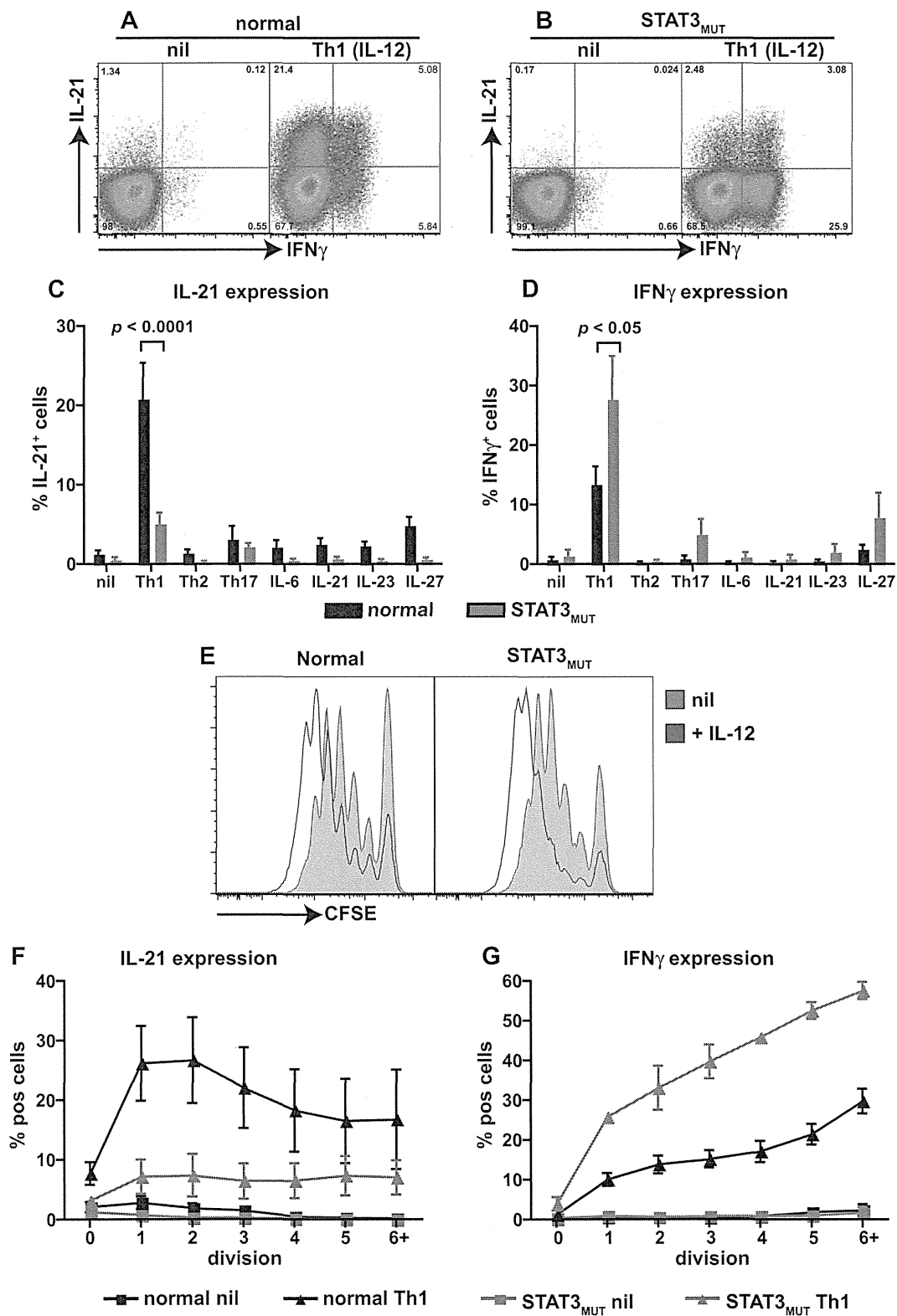
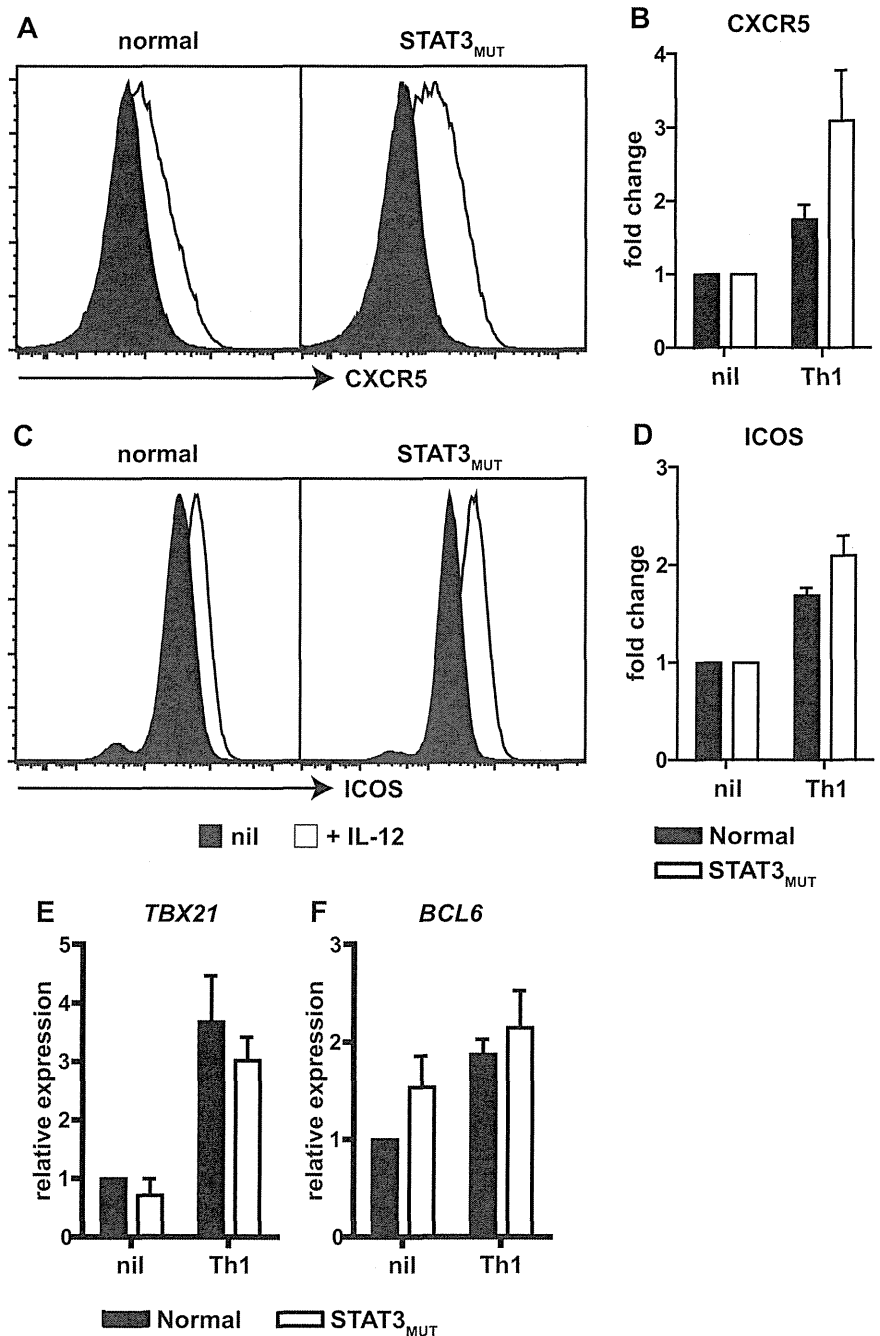


Figure 4. STAT3 mutations impair IL-12-induced expression of IL-21 in naive CD4⁺ T cells. (A-D) Naive CD4⁺ T cells isolated from healthy donors (n = 8) and STAT3_{MUT} patients (n = 9) were cultured for 5 days under neutral conditions (nil); polarizing Th1, Th2, or Th17 conditions; or in the presence of IL-6, IL-21, IL-23, or IL-27, after which time expression of intracellular IL-21 and IFN γ was determined. (A-B) Representative dot plots of cytokine expression by activated naive CD4⁺ T cells from 1 donor and 1 patient. (C-D) Percentage of activated normal and STAT3_{MUT} naive CD4⁺ T cells induced to express (C) IL-21 or (D) IFN γ in response to the indicated culture. The values represent the mean \pm SEM. (E-F) Naive CD4⁺ T cells isolated from healthy donors and STAT3_{MUT} patients were labeled with CFSE and cultured with anti-CD2/CD3/CD28 Abs in the absence (nil) or presence of Th1 polarizing conditions (+IL-12). After 5 days cells were harvested, and (E) proliferation and expression of (F) IL-21 and (G) IFN γ in cells that had undergone different divisions were then determined. (F-G) The values represent the mean \pm SEM (n = 3).

sustained expression of CXCR5 and ICOS, and a modest induction of the Tfh lineage restricted transcription factor *BCL6*.^{4,11} We

therefore wanted to explore whether the effects of IL-12 on these aspects of CD4⁺ T-cell activation were also compromised by

Figure 5. Induction of CXCR5, ICOS, and *BCL6* by IL-12 in CD4⁺ T cells is independent of STAT3. (A-D) Naive CD4⁺ T cells isolated from healthy donors and STAT3_{MUT} patients were cultured with anti-CD2/CD3/CD28 Abs in the absence (nil) or presence of Th1-polarizing conditions (+IL-12). After 4 days the cells were harvested, and surface expression of CXCR5 and ICOS was determined by flow cytometry and of *TBX21* and *BCL6* by qPCR. (A,C) representative histogram plots from 1 healthy donor and 1 STAT3_{MUT} patient. Expression of (B) CXCR5 (n = 3), (D) ICOS (n = 3), (E) *TBX21* (n = 4), and (F) *BCL6* (n = 5) after Th1 polarization is presented as fold-increase compared with the nil culture in each experiment. (B,D-F) The graphs represent the mean ± SEM of the indicated number of experiments.



mutations in *STAT3*. Compared with naive CD4⁺ T cells activated under neutral conditions, elevated expression of the T_H cell markers CXCR5 and ICOS was observed when either normal or STAT3_{MUT} cells were activated in the presence of IL-12 (Figure 5 A,C). In fact, compared with cells cultured under neutral conditions, IL-12 up-regulated CXCR5 and ICOS expressions by 2- to 3-fold on both normal and STAT3_{MUT} naive CD4⁺ T cells (Figure 5B,D). IL-12 also up-regulated *TBX21* (encoding T-bet) and *BCL6*, the transcription factors required to generate Th1 and T_H cells, respectively, to a comparable extent in normal and STAT3_{MUT} naive CD4⁺ T cells (Figure 5E,F). Thus, IL-12-mediated induction of CXCR5, ICOS, and *BCL6* in T_H-like cells either only require the residual levels of functional STAT3 that are available in STAT3_{MUT} CD4⁺ T cells or are STAT3-independent resulting from IL-12-induced activation of STAT4.^{20,21,34,38} This

latter scenario would underlie the normal induction of *TBX21* and IFN γ in STAT3_{MUT} Th1 cells.

Consistent with our previous findings,¹¹ most of the other culture conditions used in this study (ie, Th2- and Th17-polarizing conditions; exogenous IL-6, IL-21, IL-23) had no effect on expression of CXCR5 and ICOS on normal CD4⁺ T cells above that observed for the nonpolarizing culture (supplemental Figure 3A-B). However, IL-27 did modestly enhance ICOS expression (supplemental Figure 3B) and induce *TBX21* in naive CD4⁺ T cells (supplemental Figure 3C), and this was independent of STAT3. Interestingly, Th17 polarizing culture conditions induced the greatest levels of *BCL6* in naive CD4⁺ T cells, and this was substantially reduced in STAT3_{MUT} CD4⁺ T cells (supplemental Figure 3D), reflecting the contribution of STAT3 to the combined signaling of IL-6, IL-21, and IL-23 through their respective receptors.

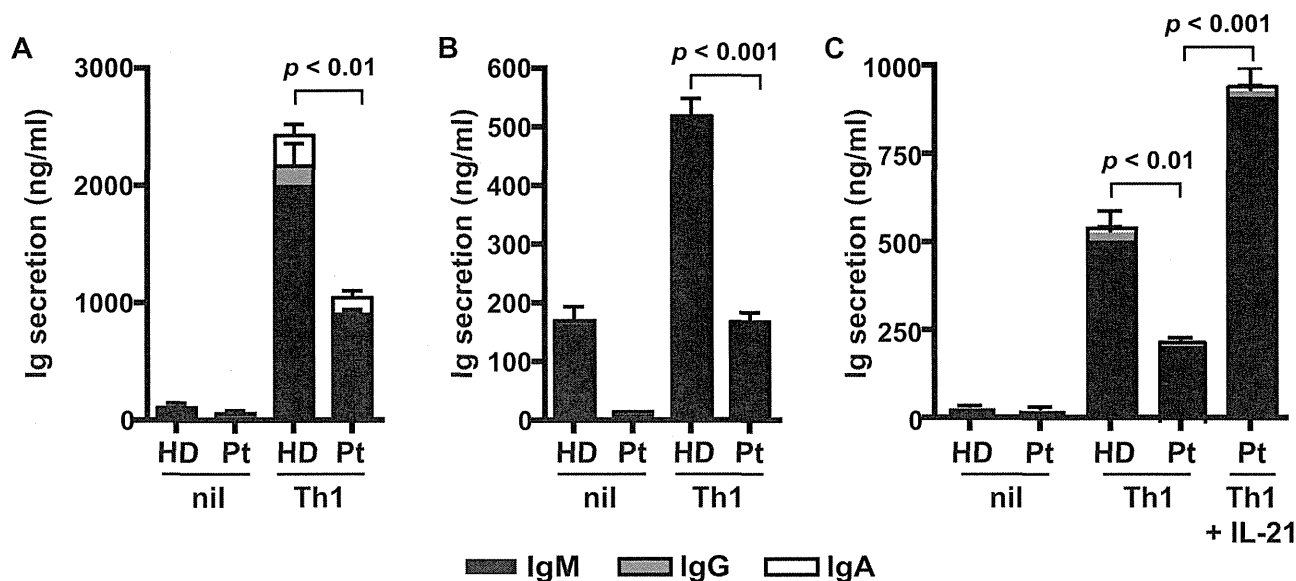


Figure 6. STAT3-deficient cells show impaired Tfh cell function in vitro. Naive CD4⁺ T cells isolated from healthy donors (HD) and STAT3^{MUT} patients (Pt) were cultured under neutral (nil) or Th1 polarizing conditions. After 5 days, the cells were harvested and treated with mitomycin C before being cocultured with allogeneic naive B cells, in the absence or presence of exogenous IL-21, for an additional 7 days. After this time secretion of IgM, IgG, and IgA was determined. (A-B) The data were derived from experiments that used naive CD4⁺ T cells isolated from different STAT3^{MUT} patients; (C) the data show the effect of exogenous IL-21 on the ability of STAT3^{MUT} CD4⁺ T cells to provide B-cell help. Each graph represents the mean \pm SEM of triplicate cultures; similar results were obtained in 5 (A-B) and 2 (C) experiments.

Defective IL-21 production by STAT3^{MUT} CD4⁺ T cells results in impaired Tfh cell function in vitro

We next questioned whether a reduction in IL-21 expression in STAT3^{MUT} CD4⁺ T cells would result in a detectable functional defect, such as a compromised ability of these cells to provide help to B cells. To investigate this, we established an in vitro B-cell helper assay in which naive CD4⁺ T cells were isolated from healthy donors and STAT3^{MUT} patients and activated under neutral or Th1-polarizing conditions. After 5 days, the CD4⁺ T cells were harvested and cocultured with naive B cells for an additional 7 days, after which time Ig secretion was quantified. Normal naive CD4⁺ T cells activated under Th1 conditions provided significantly more help to support Ig production by cocultured B cells than when these T cells were cultured under neutral conditions (Figure 6A-C). Although priming under Th1 conditions also increased the helper function of STAT3^{MUT} CD4⁺ T cells relative to those cultured under neutral conditions, it was significantly less than that of normal CD4⁺ T cells (Figure 6A-C). This reduction in B-cell help by IL-12-stimulated STAT3^{MUT} CD4⁺ T cells approximated their reduction in expression of IL-21 under these culture conditions (ie, ~50%-70%; compare Figures 4C and 6A-C), thereby suggesting that the impaired ability of STAT3^{MUT} CD4⁺ T cells to adequately promote B-cell differentiation resulted from insufficient production of IL-21. Indeed, the reduced ability of IL-12-primed STAT3^{MUT} CD4⁺ T cells to provide help to B cells could be overcome when IL-21 was added to the cocultures (Figure 6C). Thus, functional STAT3 deficiency compromises the ability of IL-12 to induce IL-21, which is subsequently detrimental to the capacity of these CD4⁺ T cells to mediate the differentiation of B cells into Ig-secreting cells.

STAT3 mutations abolish IL-21 expression in naive CD4⁺ T cells induced by IL-6, IL-21, IL-23, and IL-27

As previously shown, other cytokines (IL-6, IL-21, IL-23, IL-27) can also give rise to IL-21-expressing cells, albeit to a much lesser extent than IL-12.^{11,17,22} When STAT3^{MUT} naive CD4⁺ T cells were exposed to these cytokines, they were unable to up-regulate IL-21

expression (Figure 4C). Thus, although mutations in *STAT3* substantially reduced the ability of IL-12 to induce IL-21 expression in CD4⁺ T cells, the ability of IL-6, IL-21, IL-23, and IL-27 to do this was completely dependent on STAT3. Consistent with this, STAT3^{MUT} naive CD4⁺ T cells preactivated with IL-6, IL-21, or IL-23 were unable to support Ab production by cocultured B cells (supplemental Figure 3E-F).

Discussion

Lymphocyte differentiation is the outcome of the integration of signals from numerous external stimuli and the activation of specific transcription factors that regulate gene expression and ultimately cellular function. The differentiation of naive CD4⁺ T cells into Th1, Th2, Th17, and T-regulatory cells has been well-characterized for the roles of specific cytokines and transcription factors. The emergence of Tfh cells as the predominant subset of CD4⁺ T cells that mediate TD humoral immunity has been accompanied by the elucidation of the requirements for their generation and maintenance. Thus, engaging the TCR, ICOS, and the SLAM/SAP pathways by ligands present on APCs, together with signals mediated by STAT3 downstream of receptors for the cytokines IL-6, IL-21, and IL-27, coordinately induce expression of the transcription factors BCL6, IRF4, and c-MAF, which converge to yield Tfh cells.^{8,12} Despite these generalized findings, much controversy remains over the relative contribution of these individual components to Tfh cell formation; this is most apparent from subsequent studies that have challenged the role of IL-6 and IL-21 in this process.^{8,12,18,19} Furthermore, the molecular requirements for the generation of human Tfh cells remain incompletely defined.

We and others previously showed that IL-12 plays an important role in the differentiation of human Tfh cells, as evidenced by its ability to induce expression of IL-21 and to maintain expression of ICOS and CXCR5 on naive CD4⁺ T cells.^{11,22} Similar to studies in mice,^{7,14,17} IL-6, IL-21, and IL-27 also induce IL-21 expression in human naive CD4⁺ T cells, albeit to a much lesser extent than

IL-12.^{11,17,22} We have now substantially extended these findings by investigating the *in vivo* and *in vitro* development of T_{fh} cells in patients with loss-of-function mutations in *IL-12RB1*, *STAT1*, *TYK2*, and *STAT3*; that is genes that compromise cytokine-mediated intracellular signaling pathways probably involved in regulating human T_{fh} cell formation.

The specific ability of IL-12 to induce IL-21 expression in human naive CD4⁺ T cells was confirmed by showing that in the absence of a functional receptor (ie, in patients deficient for IL-12Rβ1) IL-12 was unable to give rise to IL-21-expressing cells (Figure 11). Interestingly, IL-12-mediated IL-21 expression partially depended on intact STAT3 signaling, because heterozygous mutations in *STAT3* (which render most STAT3 dimers nonfunctional) reduced IL-21 expression by 50%-75%. Induction of IL-21 expression in CD4⁺ T cells by IL-12 was also reduced in the absence of TYK2 but was unaffected by STAT1 deficiency. Further evidence that the generation of human T_{fh} cells is STAT1 independent was the finding that the frequency of circulating CXCR5⁺CD45RA⁻ T cells was unaffected by gain-of-function mutations in *STAT1* (22% ± 6.6%; n = 3). Although IL-12 is well-characterized for its ability to operate via STAT4-dependent pathways,²⁰ our finding of a requirement for STAT3 in IL-12 function is consistent with numerous studies that have documented STAT3 activation in human and murine T cells exposed to IL-12.^{21,33,34,38,42} TYK2 is similarly phosphorylated in IL-12-treated T cells.^{20,38,39} Despite the defect in IL-21 production, IL-12 could still induce STAT3_{MUT} naive CD4⁺ T cells to acquire other characteristics of T_{fh} cells such as increased expression of ICOS, CXCR5, and *BCL6* (Figure 5A-D). The intact induction of these phenotypic and molecular changes in T_{fh}-like cells, as well as residual expression of IL-21, in IL-12-treated CD4⁺ T cells derived from STAT3_{MUT} patients, are probably induced in a STAT4-dependent manner. Thus, IL-12 induces IL-21 expression predominantly through a TYK2/STAT3-dependent mechanism, with a minor contribution via STAT4 signaling. This is supported by the ability of both STAT3 and STAT4 to bind the promoters of the *IL21* and *BCL6* genes^{43,44} and also by the recent finding that IL-12 induces IL-21 in murine CD4⁺ T cells via STAT3- and STAT4-dependent pathways.⁴² Importantly, the reduction in IL-12-induced expression of IL-21 in STAT3_{MUT} CD4⁺ T cells translated to a functional defect in TD B-cell differentiation *in vitro* (Figure 6). Interestingly, although IL-12 could still induce normal levels of ICOS on STAT3_{MUT} CD4⁺ T cells, this could not compensate for the deficiency in IL-21 production by these T cells, which is the primary source of help for B-cell differentiation. These findings are consistent with a model in which ICOS has a dual role in T_{fh} cells, first, in their generation from naive precursors, and second, in enhancing IL-21 expression.^{12,14,45,46}

While the ability of IL-12 to induce IL-21 in naive CD4⁺ T cells was predominantly STAT3-dependent, induction of IL-21 by IL-6, IL-21, and IL-27; and the ability of CD4⁺ T cells primed with these cytokines to help B cells, were completely abrogated by mutations in *STAT3*. Thus, the ability of all cytokines currently identified to induce IL-21 in human CD4⁺ T cells, and subsequent B cell-helper function would be dramatically affected by *STAT3* mutations, that is, either strongly reduced or completely abolished. The importance of intact STAT3 signaling in generating human T_{fh} cells is reinforced by the significant reduction in the frequencies of circulating CD4⁺ CXCR5⁺ T cells in STAT3_{MUT} patients. By contrast, the frequency of these CXCR5⁺ T_{fh}-like cells was normal in patients deficient for IL-12Rβ1. This suggests that, although IL-12 induces the greatest frequency of IL-21-expressing CD4⁺ T cells, there is sufficient redundancy among cytokine signaling pathways involved in generating T_{fh} cells to overcome the inability of CD4⁺ T cells to give rise to T_{fh} cells in the absence of the IL-12 signaling. In other words, IL-6, IL-21, and IL-27 signaling

through STAT3 will still give rise to T_{fh} cells from *IL12RB1*-mutant CD4⁺ T cells. This is supported by the observation that humoral immune responses are intact in patients deficient for IL-12Rβ1.^{26,30,31} Given the critical role of T_{fh} cells in humoral immune responses, it makes teleologic sense that this level of redundancy evolved to protect against the detrimental effects of T_{fh} cell deficiency. In contrast, use of STAT3 by several cytokines in the generation of human T_{fh} cells provides an explanation for why STAT3_{MUT} patients exhibit defects in humoral immune responses (including reductions in circulating CD4⁺ CXCR5⁺CD45RA⁻ and CD45RA⁺ T_{fh}-like cells, memory B cells and an inability to mount protective Ab responses after vaccination or natural infection^{24,47,48}) that cannot be compensated entirely by IL-12-dependent STAT4 signaling. These clinical, cellular, and serologic features of STAT3 deficiency are reminiscent of patients with mutations in *ICOS* and *CD40LG*,^{49,50} which largely result from an absence of B-cell help by T_{fh} cells.^{49,50} Thus, although an intrinsic defect resulting from the inability of B cells to respond to cytokines such as IL-6, IL-10, and IL-21 would contribute to the functional Ab deficiency in STAT3_{MUT} patients,²⁴ this defect would be compounded further by the compromised generation and function of T_{fh} cells.

Overall, our findings have shed substantial light on the molecular requirements for generating human T_{fh} cells and have identified a signaling pathway that could be targeted to enhance T_{fh} cell generation in immunodeficient conditions. The corollary is that because dysregulated activation and/or generation of T_{fh} cells has been associated with autoimmunity in humans and mice, inhibiting this pathway may represent a novel approach to treating autoAb-mediated conditions.

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Authorship

Contribution: C.S.M. designed the research, performed experiments, analyzed and interpreted results, and wrote the manuscript; D.T.A., A.C., and E.K.D. performed experiments; J.B., S.B.-D., P.D.A., A.Y.K., D.A., D.E., K.M., S.S.K., Y.M., S.N., M.A.F., S.C., J.M.S., J.P., M.W., P.G., M.C.C., D.A.F., and J.-L.C. provided patient samples and clinical details; M.C.C. also genotyped STAT3-mutant patients; M.B. provided intellectual input; and S.G.T. designed the research, interpreted results, and wrote the manuscript. All authors commented on the manuscript.

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The kinase Btk negatively regulates the production of reactive oxygen species and stimulation-induced apoptosis in human neutrophils

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The function of the kinase Btk in neutrophil activation is largely unexplored. Here we found that Btk-deficient neutrophils had more production of reactive oxygen species (ROS) after engagement of Toll-like receptors (TLRs) or receptors for tumor-necrosis factor (TNF), which was associated with more apoptosis and was reversed by transduction of recombinant Btk. Btk-deficient neutrophils in the resting state showed hyperphosphorylation and activation of phosphatidylinositol-3-OH kinase (PI(3)K) and protein tyrosine kinases (PTKs) and were in a 'primed' state with plasma membrane-associated GTPase Rac2. In the absence of Btk, the adaptor Mal was associated with PI(3)K and PTKs at the plasma membrane, whereas in control resting neutrophils, Btk interacted with and confined Mal in the cytoplasm. Our data identify Btk as a critical gatekeeper of neutrophil responses.

Among 'professional' phagocytes with a sophisticated arsenal of microbicidal features, neutrophils are the dominant cells that mediate the earliest innate immune responses to microbes^{1–3}. Neutrophils migrate to the site of infection, sense and engulf microorganisms, produce reactive oxygen species (ROS) and kill the invading microbes via ROS by acting together with antimicrobial proteins and peptides^{1,2}. The enzyme responsible for the respiratory burst is NADPH oxidase, which catalyzes the production of superoxide from oxygen and NADPH. This enzyme is a multicomponent complex that consists of membrane-bound flavocytochrome *b*₅₅₈ (gp91^{phox} and p22^{phox}), cytosolic components (p47^{phox}, p67^{phox} and p40^{phox}) and a GTPase (Rac1 or Rac2)^{3–6}. Activation of NADPH oxidase is strictly regulated both temporally and spatially to ensure that the reaction takes place rapidly at the appropriate cellular localization. Activation of this system requires three signaling triggers, including protein kinases, lipid-metabolizing enzymes and nucleotide-exchange factors that activate the Rac GTPase^{3–6}.

Inadequate production of ROS is associated with various human pathological conditions. Deficiency of any component of the NADPH oxidase complex results in chronic granulomatous disease, in which bacterial and fungal infections are recurrent and life-threatening⁴. Abnormalities in the molecules involved in the signal-transduction pathway initiated by the recognition of pathogen-associated molecular patterns are accompanied by less production of ROS after exposure to specific stimuli and by susceptibility to bacterial infection. These abnormalities include deficiency in the kinase IRAK4, the adaptor MyD88 deficiency or the kinase NEMO (IKK γ)⁷. In contrast, many

other human disorders are believed to be associated with or induced by excessive production of ROS that causes DNA damage, tissue damage, cellular apoptosis and neutropenia^{8,9}.

Here we focus on determining the role of the kinase Btk in production of ROS and cellular apoptosis in human neutrophils, as 11–30% of patients with X-linked agammaglobulinemia (XLA), a human disease of Btk deficiency, have neutropenia^{10,11}, and Btk is a critical signaling component of phagocytic cells^{12–14}. The neutropenia of XLA is distinct from that of common variable immunodeficiency (CVID) in that the neutropenia is induced by infection, is usually ameliorated after supplementation with immunoglobulin and is not mediated by the autoimmune response^{10,11,14}. Although a few reports have suggested that myeloid differentiation is impaired in mice with X-linked immunodeficiency^{15,16}, the reason for the infection-triggered neutropenia is unknown. The role of Btk in human neutrophils remains largely unexplored.

Btk is a member of the Tec-family kinases (TFKs) that are expressed in hematopoietic cells such as B cells, monocytes, macrophages and neutrophils¹². It has a crucial role in cell survival, proliferation, differentiation and apoptosis, especially in cells of the B lineage. In humans with XLA, B cells fail to reach maturity and are presumably doomed to premature death by the *BTK* mutation that leads to the XLA phenotype¹⁷. Both mice with X-linked immunodeficiency that have natural mutations in *Btk* and mice in which *Btk* is targeted have B cell defects, but these are associated with much milder effects than those seen in XLA, which suggests species differences in the role of Btk^{18,19}.

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Btk is also an important signaling component of the innate immune system in phagocytic cells. Btk is involved in signaling via Toll-like receptors (TLRs) such as TLR2, TLR4, TLR7, TLR8 and TLR9, and is associated with the TLR adaptors MyD88, Mal (TIRAP) and IRAK1 (refs. 12–14,20–22). Defective innate immune responses have been observed in monocytes, dendritic cells, neutrophils and mast cells from Btk-deficient mice^{12,14}. Neutrophils from mice with X-linked immunodeficiency have poor production of ROS and nitric oxide¹⁵.

The contribution of Btk to the human innate immune system is less obvious. Stimulation via TLR2, TLR4, TLR7–TLR8 or TLR3 results in impaired production of tumor-necrosis factor (TNF) by dendritic cells from patients with XLA, whereas the TLR4-induced production of TNF and interleukin 6 (IL-6) by monocytes from patients with XLA remains intact^{23–25}. Neutrophils from control subjects and patients with XLA show no substantial differences in their phosphorylation of the mitogen-activated protein kinases p38, Jnk and Erk induced by engagement of TLR4 or TLR7–TLR8 or production of ROS induced by the same stimuli²⁶.

Here we evaluate the role of Btk in the production of ROS and cellular apoptosis in human neutrophils through the use of Btk-deficient neutrophils, a protein-delivery system based on a cell-permeable peptide, and specific kinase inhibitors. Unexpectedly, and in contrast to published observations of mice with X-linked immunodeficiency¹⁵, the production of ROS was substantially augmented in the absence of Btk in neutrophils stimulated via TLRs, the TNF receptor or phorbol 12-myristate 13-acetate (PMA) but not in monocytes or in lymphoblastoid B cell lines transformed by Epstein-Barr virus. Excessive production of ROS was associated with neutrophil apoptosis, which was reversed by the transduction of wild-type Btk protein. Btk-deficient neutrophils showed activation of key signaling molecules involved in the activation of NADPH oxidase, and this was accompanied by targeting of Rac2 to the plasma membrane. Mal was confined to the cytoplasm in association with Btk but was translocated to plasma membrane and interacted with protein tyrosine kinases (PTKs) and phosphatidylinositol-3-OH kinase (PI(3)K) in the absence of Btk. Here we present our findings on the mechanism by which Btk regulates the priming of neutrophils and the amplitude of the neutrophil response.

RESULTS

Excessive production of ROS in Btk-deficient neutrophils

To investigate the production of ROS in the absence of Btk, we monitored ROS in neutrophils, monocytes and Epstein-Barr virus-transformed lymphoblastoid B cell lines obtained from patients with XLA, healthy controls and patients with CVID (disease control) by staining with dihydrorhodamine 123 (DHR123) and a luminol chemiluminescence assay. PMA-driven production of ROS in Btk-deficient neutrophils was three to four times greater than that in neutrophils from healthy controls or patients with CVID, and we observed augmented production of ROS with a suboptimal dose of PMA, whereas the baseline production of ROS was similar (Fig. 1a–d). Similarly, and in contrast to published reports²⁶, engagement of TLR2 (with its ligand tripalmitoyl cysteinyl seryl tetralysine lipopeptide (Pam₃CSK₄)), TLR4 (with its ligand lipopolysaccharide) or the TNF receptor (with TNF) followed by stimulation with formyl-Met-Leu-Phe (fMLP), an agonist of G protein-coupled receptors, elicited augmented ROS responses in neutrophils from patients with XLA (Fig. 1e,f). The production of ROS was minimal after stimulation with the TLR9 agonist CpG-A in neutrophils from patients with XLA and was not significantly different from that of neutrophils from healthy controls. The observed phenomena were reproduced in Btk-deficient eosinophils but not in monocytes or Epstein-Barr virus-transformed lymphoblastoid B cell lines obtained from patients with XLA (Supplementary Fig. 1). These data indicated Btk-deficient neutrophils had excessive NADPH oxidase activity after various stimuli.

Augmented apoptosis in Btk-deficient neutrophils

Because high ROS concentrations are potentially harmful to cells, we investigated cell death induced by various stimuli in neutrophils from patients with XLA by staining with annexin V and the membrane-impermeable DNA-intercalating dye 7-AAD. Stimulation with PMA, TLR agonist plus fMLP, or TNF plus fMLP induced a significantly higher frequency of cells positive for annexin V among neutrophils from patients with XLA than among control neutrophils, whereas spontaneous cell death in the absence of stimulation was not significantly altered at 4 h in neutrophils from healthy controls versus those from patients with XLA (Fig. 2a,b). We observed cleavage of caspase-3, lower mitochondrial membrane potentials and degradation of proliferating

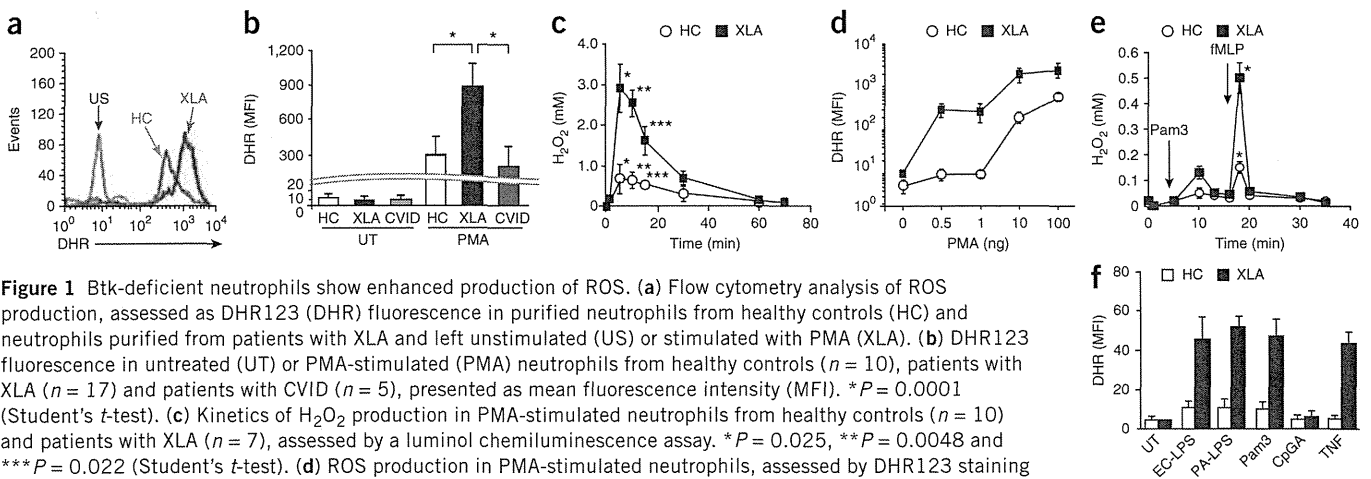


Figure 1 Btk-deficient neutrophils show enhanced production of ROS. (a) Flow cytometry analysis of ROS production, assessed as DHR123 (DHR) fluorescence in purified neutrophils from healthy controls (HC) and neutrophils purified from patients with XLA and left unstimulated (US) or stimulated with PMA (XLA). (b) DHR123 fluorescence in untreated (UT) or PMA-stimulated (PMA) neutrophils from healthy controls ($n = 10$), patients with XLA ($n = 17$) and patients with CVID ($n = 5$), presented as mean fluorescence intensity (MFI). $*P = 0.0001$ (Student's t -test). (c) Kinetics of H_2O_2 production in PMA-stimulated neutrophils from healthy controls ($n = 10$) and patients with XLA ($n = 7$), assessed by a luminol chemiluminescence assay. $*P = 0.025$, $**P = 0.0048$ and $***P = 0.022$ (Student's t -test). (d) ROS production in PMA-stimulated neutrophils, assessed by DHR123 staining and presented as a dose-response curve ($n = 5$ donors per group). (e) Kinetics of H_2O_2 production in neutrophils stimulated with Pam₃CSK₄ (Pam3) and fMLP, assessed by a luminol chemiluminescence assay ($n = 7$ donors per group). $*P = 0.005$ (Student's t -test). (f) DHR123 fluorescence in neutrophils incubated with lipopolysaccharide from *Escherichia coli* (EC-LPS) or *Pseudomonas aeruginosa* (PA-LPS), Pam₃CSK₄, CpG-A or TNF, followed by stimulation with fMLP ($n = 7$ donors per group). Data are representative of seventeen experiments (a) or are pooled from at least five (b,c,e,f) or four (d) independent experiments (mean and s.d. in b–f).

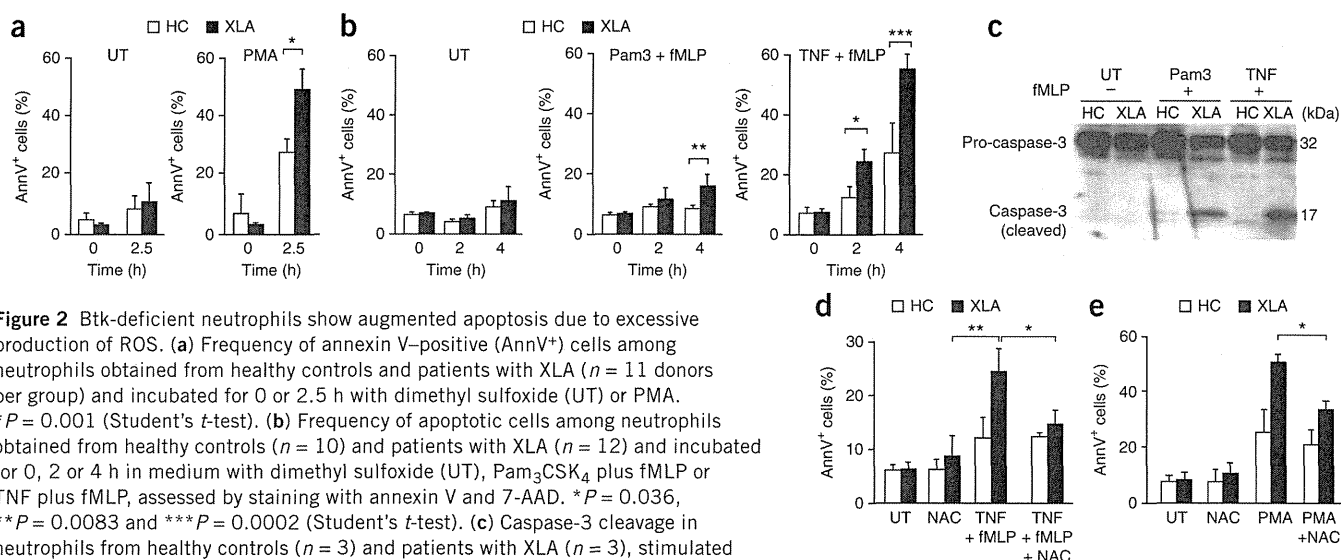


Figure 2 Btk-deficient neutrophils show augmented apoptosis due to excessive production of ROS. (a) Frequency of annexin V-positive (AnnV⁺) cells among neutrophils obtained from healthy controls and patients with XLA ($n = 11$ donors per group) and incubated for 0 or 2.5 h with dimethyl sulfoxide (UT) or PMA. $*P = 0.001$ (Student's t -test). (b) Frequency of apoptotic cells among neutrophils obtained from healthy controls ($n = 10$) and patients with XLA ($n = 12$) and incubated for 0, 2 or 4 h in medium with dimethyl sulfoxide (UT), Pam₃CSK₄ plus fMLP or TNF plus fMLP, assessed by staining with annexin V and 7-AAD. $*P = 0.036$, $**P = 0.0083$ and $***P = 0.0002$ (Student's t -test). (c) Caspase-3 cleavage in neutrophils from healthy controls ($n = 3$) and patients with XLA ($n = 3$), stimulated for 4 h as in b (above lanes). Pro-caspase-3 is the uncleaved form. (d,e) Frequency of apoptotic Btk-deficient neutrophils ($n = 5$ donors) stimulated for 2 h (d) or 2.5 h (e) with the antioxidant N-acetyl-cysteine (NAC). $*P = 0.043$ and $**P = 0.036$ (d) or $*P = 0.026$ (e); Student's t -test. Data are representative of three experiments (b) or at least five independent experiments (a,c–e; mean and s.d. in a,b,d,e).

cell nuclear antigen; hence, cell death was caused by apoptosis (Fig. 2c and Supplementary Fig. 2). Apoptosis assessed by these methods was augmented considerably for neutrophils from patients with XLA. The observed apoptosis was most probably triggered by ROS, as coincubation of neutrophils with N-acetyl cysteine, an antioxidant, rescued the cells from apoptosis induced by TNF plus fMLP or by PMA (Fig. 2d,e). We detected much more ROS release and stimulation-induced apoptosis of neutrophils from all patients with XLA regardless of the site or mode of their mutation (Supplementary Fig. 3). In addition, we found no correlation between genotype and the extent of neutrophil production of ROS. These data suggested that neutrophils from patients with XLA are susceptible to apoptosis triggered by pathogens.

Normalization of the ROS response by transduction of Btk

We next determined whether the enhanced apoptosis noted above was due to a defect in Btk itself or abnormal myeloid differentiation in the absence of Btk. For this, we prepared three recombinant Btk proteins (full-length Btk; Btk with deletion of the pleckstrin homology (PH) domain; and Btk with deletion of the kinase domain) fused to the cell-permeable peptide Hph-1 (Fig. 3a,b). We purified the products and transduced the proteins into neutrophils lacking Btk. The efficacy of transduction was more than 95%; and Hph-1–Btk expression was stable for at least 12–24 h (ref. 27). We adjusted the expression of Btk to that in neutrophils from healthy controls by incubating 1×10^6 cells for 1 h with $1 \mu\text{M}$ recombinant fusion protein. Transduction of full-length Btk into neutrophils from patients with XLA restored the production of ROS and the frequency of apoptotic cells after PMA stimulation to that observed for neutrophils from healthy controls (Fig. 3c,d). Transduction of the recombinant fusion of Btk with deletion of the PH domain only modestly reversed neutrophil overactivation (Fig. 3c), which indicated that appropriate cellular localization and interactions with other molecules were required for Btk function. Transduction of the recombinant fusion of Btk with deletion of the kinase domain minimally corrected excessive production of ROS (Fig. 3c), which suggested that the kinase activity of Btk or molecules that interacted via the kinase domain were critical for the regulation of ROS. We also confirmed the importance of the kinase domain

by an experiment that showed excessive production of ROS in normal neutrophils treated with $50 \mu\text{M}$ LFM-A13, an inhibitor of the kinase activity of Btk, but not in those treated with LFM-A11, a control compound (Fig. 3e). We also documented augmented apoptosis in control neutrophils treated with LFM-A13 (Fig. 3f). These data demonstrated that the enhanced production of ROS and apoptosis was directly related to a defect in Btk.

NADPH oxidase components in Btk-deficient neutrophils

The NADPH oxidase complex consists of the transmembrane component (gp91^{phox} and p22^{phox}), a cytosolic component (p47^{phox}, p67^{phox} and p40^{phox}) and Rac2 (refs. 3–6). The activity of NADPH oxidase is controlled by targeting of the cytosolic components to the plasma membrane or phosphorylation of the cytosolic components or both. To assess the mechanism of the excessive production of ROS in Btk-deficient neutrophils, we investigated the abundance, phosphorylation and subcellular localization of each component by immunoblot analysis.

The expression of each component of the NADPH oxidase complex was similar in neutrophils from patients with XLA and those from healthy controls (Fig. 4a). The amount of p47^{phox}, p67^{phox} and p40^{phox} in the cytoplasm and the membrane was not substantially different in neutrophils from patients with XLA and those from healthy controls (Fig. 4b). Similarly, the amount in the membrane-targeted fraction after stimulation with PMA was not very different in neutrophils from patients with XLA and those from healthy controls (Fig. 4c). Phosphorylation of Ser345 in p47^{phox} and of Thr154 in p40^{phox} are important for translocation of the cytosolic components to the membrane^{4,5,28}. Those modifications were not altered in Btk-deficient neutrophils (Fig. 4c). In contrast, we detected Rac2 in the plasma membrane of Btk-deficient neutrophils before stimulation with PMA. We observed four- to fivefold higher membrane expression of Rac2 in neutrophils from patients with XLA than in those from healthy controls in the resting state (Fig. 4b).

Typically, 10–15% of gp91^{phox} is located in the plasma membrane of unstimulated neutrophils, whereas the majority of the molecule resides in specific granules. Membrane expression increases after