Chemokine Receptor Expression Identifies Pre–T Helper (Th)1, Pre–Th2, and Nonpolarized Cells among Human CD4\(^+\) Central Memory T Cells

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Abstract

We previously reported that central–memory T cells (T\(_{CM}\) cells), which express lymph node homing receptors CCR7 and CD62L, are largely devoid of effector functions but acquire characteristics of effector–memory T cells (T\(_{EM}\) cells) (i.e., CCR7\(^-\) T helper [Th]1 or Th2 cells) after stimulation with T cell receptor agonists or homeostatic cytokines. Here we show that three chemokine receptors identify functional subsets within the human CD4\(^+\) T\(_{CM}\) cell pool. T\(_{CM}\) cells expressing CXCR3 secreted low amounts of interferon \(\gamma\), whereas CCR4\(^+\) T\(_{CM}\) cells produced some interleukin (IL)-4, but not IL-5. In response to IL-7 and IL-15, CXCR3\(^+\) T\(_{CM}\) and CCR4\(^+\) T\(_{CM}\) cells invariably generated fully differentiated CCR7\(^-\) Th1 and Th2 cells, respectively, suggesting that they represent pre-Th1 and pre-Th2 cells. Conversely, CXCR5\(^+\) T\(_{CM}\) cells lacking CXCR3 and CCR4 remained nonpolarized and retained CCR7 and CD62L expression upon cytokine-driven expansion. Unlike naive cells, all memory subsets had a low T cell receptor rearrangement excision circle content, spontaneously incorporated bromodeoxyuridine ex vivo, and contained cells specific for tetanus toxoid. Conversely, recall responses to cytomegalovirus and vaccinia virus were largely restricted to CXCR3\(^+\) T\(_{CM}\) and T\(_{EM}\) cells. We conclude that antigen-specific memory T cells are distributed between T\(_{EM}\) cells and different subsets of T\(_{CM}\) cells. Our results also explain how the quality of primary T cell responses could be maintained by T\(_{CM}\) cells in the absence of antigen.

Key words: T cell subsets • memory maintenance • cytokines • differentiation • chemokine receptors

Introduction

Upon recognition of antigenic peptides on DCs, naive T lymphocytes proliferate and differentiate into a variety of effector cells depending on the stimulatory conditions and cytokine milieu (1, 2). Accumulating evidence indicates that during the T cell differentiation process, effector functions and homing potentials are coordinately regulated (3). For instance, developing Th1 cells acquire the capacity to produce IFN-\(\gamma\) and expression of chemokine receptors such as CCR5, CXCR3, and CXCR6 that drive them to sites of delayed-type hypersensitivity reactions. Conversely, developing Th2 cells acquire the capacity to produce IL-4 and express CCR3, CCR4, CCR8, and the prostaglandin D2 chemoattractant receptor CRTh2 (4–7), which are required to migrate at sites of allergic reactions (6, 8, 9).

Expression of the lymph node homing receptors CCR7 and CD62L (10, 11) has been used to define subsets of human memory T cells with distinct functional properties. T cells within the CCR7\(^+\) “central–memory” T cell (T\(_{CM}\) cell) subset show hypo-acetylated cytokine genes and have no or low effector functions, but efficiently differentiate to Th1 or Th2 effector cells after TCR stimulation in the presence of IL-12 or IL-4, respectively (12–14). It has been proposed that in secondary immune responses, T\(_{CM}\) cells generate L. Rivino, M. Messi, and D. Jarrossay contributed equally to this work.

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Abbreviations used in this paper: BrdU, bromodeoxyuridine; CFSE, carboxyfluorescein succinimidyl ester; PdBu, phorbol-12-13-dibutyrate; T\(_{CM}\) cell, central–memory T cell; T\(_{EM}\) cell, effector–memory T cell; TREC, T cell receptor rearrangement excision circle; TSST, toxic shock syndrome toxin; TT, tetanus toxoid; VV, vaccinia virus.

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Expression of the lymph node homing receptors CCR7 and CD62L (10, 11) has been used to define subsets of human memory T cells with distinct functional properties. T cells within the CCR7\(^+\) “central–memory” T cell (T\(_{CM}\) cell) subset show hypo-acetylated cytokine genes and have no or low effector functions, but efficiently differentiate to Th1 or Th2 effector cells after TCR stimulation in the presence of IL-12 or IL-4, respectively (12–14). In contrast, T cells of the CCR7\(^-\) “effector–memory” T cell (T\(_{EM}\) cell) subset show polarized cytokine gene acetylation patterns in vivo and rapidly produce high amounts of IFN-\(\gamma\) and IL-4 upon antigenic stimulation (12–14). It has been proposed that in secondary immune responses, T\(_{CM}\) cells generate...
new waves of effector cells in antigen-draining lymph nodes, whereas T_{CM} cells provide immediate protection against invading pathogens in peripheral tissues (3, 14).

The maintenance of T cell memory is controlled by cytokines that promote cell survival and slow homeostatic proliferation (15). In particular, IL-7 and IL-15 have been shown to regulate mouse CD8+ memory T cell survival and self-renewal in the absence of antigen (16, 17), whereas naive and CD4+ memory cells require IL-7 and TCR ligands (16, 18, 19), but do not respond to IL-15 (17). Conversely, human CD4+ memory T cells proliferate in response to IL-15 in a TCR-independent fashion and with slow kinetics (20, 21), suggesting different roles for IL-15 in mouse and human CD4+ memory T cell homeostasis. Notably, T_{CM} cells proliferating in response to IL-7 and IL-15 differentiate and generate Th1 and Th2 effector cells (21), but how uncommitted T_{CM} cells proliferating in the absence of antigen could maintain the quality of the primary response remained unclear.

The existence of T_{CM} and T_{EM} cell subsets has also been documented in mice (22, 23). In this experimental system, it has been possible to directly examine the kinetics of memory cell generation and the capacity of effector and memory subsets to reconstitute long-term memory (24), and there is growing evidence that T_{CM} cells have higher reconstitution potential (24, 25). In particular, effector Th1 cells, defined by their secretion of IFN-\gamma, were found to be short-lived and unable to reconstitute T cell memory. In contrast, a population of activated Th1 lineage cells, which did not secrete IFN-\gamma after primary antigenic stimulation, persisted for several months in vivo and developed the capacity to secrete IFN-\gamma upon subsequent stimulation (26).

Since the first description of T_{CM} and T_{EM} cells, it was evident that other chemokine receptors, as well as adhesion and costimulatory molecules, are expressed on different fractions of T_{CM} cells (12, 14). Heterogeneity of human CD4+ T_{CM} cells has further been documented using CXCR5, the receptor for CXCL13, a chemokine expressed in B follicles (27, 28). CXCR5+ T_{CM} cells lacked effector functions and cells specific for tetanus toxoid (TT), but contained residual T cell receptor rearrangement excision circles (TRECs), suggesting that they represent recently activated cells (29–31). Conversely, other recent studies claimed that both T_{CM} and T_{EM} cells possess high levels of effector functions, and that consequently neither CCR7 nor CXCR5 expression identify nonpolarized CD4+ memory T cells (9, 32–35). Understanding T_{CM} cell differentiation stage and potential is of importance for the homeostatic maintenance of memory T cells and for the conservation of T cell polarization in secondary responses.

Here we report that CXCR3 and CCR4 identify two novel subsets of pre-Th1 and pre-Th2 cells within T_{CM} cells. These cells possessed low IFN-\gamma- or IL-4-producing capacities when compared with CXCR3+ and CCR4+ T_{EM} cells and spontaneously differentiated to Th1 and Th2 effector cells in response to homeostatic cytokines IL-7 and IL-15 independently of conventional Th1 or Th2 cell-inducing stimuli. In contrast, T_{CM} cells lacking CXCR3 or CCR4 and expressing CXCR5 were nonpolarized cells whose differentiation to Th1 or Th2 cells is dependent on TCR triggering and signaling by polarizing cytokines.

**Materials and Methods**

**Cell Culture.** PBMCs were isolated from buffy-coated blood from healthy donors. Monocytes were depleted by adherence for 30 min and CD4+ T cells were isolated by negative selection with magnetic beads using Automacs (Miltenyi Biotec). Memory T cells were isolated by further depletion of naive T cells with anti-CD45RA beads (Miltenyi Biotec). Memory T cell subpopulations were purified to >95% by cell sorting after five-color staining as follows: anti-CXCR5 (R&D Systems) followed by anti-IgG2b PE (Biosystems), anti-CCR7 (R&D Systems) followed by anti-IgG2a FITC (Biosystems), and anti-CXCR3 Cy-Chrome, anti-CD45RA APC, and anti-CCR4 biotin followed by streptavidin-APC-Cy7 (BD Biosciences). Labeling of T cells with carboxyfluorescein succinimidyl ester (CFSE) was performed as described previously (21). Monocytes were purified by positive selection with anti-CD14 beads (Miltenyi Biotec). For DC differentiation, CD14+ cells were cultured for 4 d in complete medium (RPMI 1640 supplemented with 2 mM glutamine, 1% nonessential amino acids, 1% sodium pyruvate, 50 \mu g/ml kanamycin, 50 U/ml penicillin, and 50 \mu g/ml streptomycin; Gibco BRL) containing 10% FCS (Hyclone), 50 ng/ml granulocyte/macrophage colony-stimulating factor (Novartis), and 1,000 U/ml IL-4. The DCs were cultured for 24 h with 100 ng/ml lipopolysaccharide (Salmonella abortus equi; Sigma-Aldrich) and pulsed with 30 min with 100 ng/ml toxic shock syndrome toxin (TTST). CFSE-labeled 5 × 10^6 T cells were cultured with TSST-pulsed DCs in flat-bottom wells at a 5:1 ratio, and recombinant cytokines were used at either 25 ng/ml (IL-7 and IL-15; R&D Systems), 10 ng/ml (TNF, IL-6, IL-10, IL-4, and IL-12; BD Biosciences), or 1,000 U/ml (IL-2; Roche), whereas neutralizing antibodies to IL-4 and IL-12 (BD Biosciences) were used at 2 \mu g/ml.

**ELISA.** Intracellular Cytokine Staining, and IFN-\gamma Secretion Assay. Cytokine-producing capacity of FACS-purified subsets was assessed after stimulation of purified cell populations at 3 × 10^5/100 \mu l for 24 h with 50 nM phorbol-12-13-dibutyrate (PdBu) and 0.5 \mu g/ml ionomycin, or in wells coated with 2 \mu g/ml each of anti-CD3 (clone TR66) and anti-CD28 antibodies (BD Biosciences). Cytokine concentrations of supernatants were then assessed by ELISA according to a standard protocol and analyzed with the Softmax program. Intracellular IFN-\gamma was detected after stimulating cells in the presence of 10 \mu g/ml brefeldin A (Sigma-Aldrich) for the last 2 h and after fixation with paraformaldehyde and permeabilization with saponin. After saturation of nonspecific binding sites with 10% FCS, cells were incubated with APC-labeled antibody to IFN-\gamma and PE-labeled antibody to IL-2 or IL-4 (BD Biosciences), washed, and analyzed by flow cytometry on a FACS Calibur with CELLQuest software (Becton Dickinson). To sort live IFN-\gamma-producing cells, we stimulated cells for 60 h with 25 ng/ml cytokines (IL-7, IL-15, IL-12, TNF-\alpha, and IL-18), and IFN-\gamma-producing cells were identified with an IFN-\gamma secretion assay kit (Miltenyi Biotec) and purified by cell sorting.

**Recall Responses.** PBMCs from 50 ml of fresh blood from healthy volunteers were prepared, monocytes were isolated and either incubated for 16 h with a replication-deficient vaccinia virus (VV); provided by G. Sutter, Institute for Virology, Munich.
Quantification of signal joint TREC in sorted CD4
duction of proliferating T cells by intracellular staining.

In some experiments, the presence of pathogen-specific cells
could therefore be calculated as described previously
(36). In some experiments, the presence of pathogen-specific cells
was confirmed by restimulating cells with autologous monocytes
-treated as described above followed by assessment of cytokine pro-
duction of proliferating T cells by intracellular staining.

Ex Vivo Bromodeoxyuridine (BrdU) Labeling. The assay
was performed as described previously (25). In brief, fresh PBMCs
were cultured with 10 μg/ml BrdU (Sigma-Aldrich) for 16 h.
CD4+ cells were then positively selected with anti-CD4 mag-
etic beads and stained for CD45RA and chemokine receptor
expression. Cells were then fixed, permeabilized, treated with
DNase (Boehringer), stained with FITC-labeled anti-BrdU anti-
body (Becton Dickinson), and analyzed by flow cytometry.

Quantitative PCR of TREC. Memory CD4+ T cells were
isolated by MACS as described above, stained for CD4,
CD45RA, CCR7, and CXCR5, and sorted to >99.9% purity.
Quantification of signal joint TREC in sorted CD4+ T cell subsets
was performed by real-time quantitative PCR with the 5’
nuclease (TaqMan) assay using an ABI 7700 sequence detector
(Applied Biosystems). As described previously (37), 1–2 × 10^6
cells were lysed in 10 mM Tris, pH 8, containing 100 μg/ml of
proteinase K (GIBCO BRL) for 2 h at 56°C, and then for 15 min
at 95°C. PCR reaction of lysates was performed with 500 nM of
primers (CACATCCCTTTCAACCATGCT and GCCAGCT-
GAGGGTTTAGG) and 125 nM of probe FAM-ACAC-
CTCCTGTTTTTGTAAAGTGCCCACCT-TAMRA. PCR
conditions were as follows: 1 cycle of 2 min at 50°C, 1 cycle
of 10 min at 95°C, followed by 40 cycles of 30 s at 95°C, and 1 min
at 65°C. Levels of DNA were standardized by normalizing with
18S rRNA sequences.

Results

Subsets of Human CD4+ TCM Cells Identified by Expression
of CXCR5, CXCR3, and CCR4. Purified human CD4+
T cells were analyzed for chemokine receptor expression by
five-color staining. CD45RA+ cells expressed CCR7, but
were largely negative for the other chemokine receptors,
consistent with the view that they are predominantly anti-
gen-inexperienced “naive” T cells (not depicted). Con-
versely, the following three main subsets could be identified
in CD45RA+ cells according to CCR7 and CXCR5
expression: CXCR5+CCR7+ cells (CXCR5+ TCM),
CXCR5+CCR7− cells (CXCR5+ TCM), and CXCR5−
CCR7− cells (T EM; Fig. 1 A). Within these main subsets,
staining with antibodies to CCR3 and CCR4 revealed
further heterogeneity (Fig. 1 B). CXCR3 and CCR4 were
expressed on different populations of T EM cells, which
contain Th1 and Th2 effector cells (4, 9, 38). However,
CXCR3 and CCR4 were also expressed on some T CM cells,
especially within the CXCR5− subset (Fig. 1 B). Thus, the
following four major subsets of T CM cells were identified:
(a) CXCR3−CCR4−CXCR5+ T CM cells (“CXCR5+ T CM”),
(b) CXCR3−CCR4−CXCR5− T CM cells (“−CXCR5− T CM”),
(c) CXCR5−CXCR3+CCR4+ T CM cells (“CXCR3+ T CM”),
and (d) CXCR5−CCR4−CXCR3+ T CM cells (“CCR4+ T CM”).
Mean values ± standard deviations of the four sub-
sets in four healthy donors were 11 ± 5%, 18 ± 10%, 17 ±
8%, and 17 ± 12%, respectively.

Because CXCR3 and CCR4 have been associated with
differentiated Th1 and Th2 cells (7, 38), we analyzed expres-
sion of other surface markers that are acquired or lost
with T cell differentiation (12, 17, 39; Table I). As expected, naive T cells expressed uniformly CD27 and CD62L, but not the IL-2/15Rβ chain (CD122), whereas most TEM cells had lost CD27 and CD62L expression, but were CD122+.

CXCR5+ TCM cells had a phenotype similar to naive cells, whereas CXCR5− TCM cells expressed intermediate levels of CD27, CD62L, and CD122. In particular, CXCR3+ TCM cells were CD122+ and had partially lost CD62L expression, consistent with a more differentiated phenotype. The differences in CD122 expression were functionally relevant because they closely correlated with proliferation in response to IL-7 and IL-15 (Fig. 1 C, bottom). As reported for the CD8 compartment (25), the reduced accumulation of TEM cells was associated with a high rate of apoptosis (Table I). Similar results were obtained upon stimulation with anti-CD3 and anti-CD28 antibodies (not depicted).

We then compared the expansion potential of purified CFSE-labeled CD4+ naive and memory T cell subsets after TCR stimulation with TSST-loaded DCs because replicative capacity diminishes with T cell differentiation (25, 40). Proliferation and accumulation was high in naive cells, CXCR5+ TCM cells, and −/− TCM cells, intermediate in CCR4+ and CXCR3+ TCM cells, and low in TEM cells (Fig. 1 C, bottom). As reported for the CD8 compartment (25), the reduced accumulation of TEM cells was associated with a high rate of apoptosis (Table I). Similar results were obtained upon stimulation with anti-CD3 and anti-CD28 antibodies (not depicted).

Together, these results show that subsets of CD4+ TCM cells identified by CXCR5, CXCR3, and CCR4 expression differ in their proliferative response to cytokines and TCR ligands, and suggest that CXCR5+ TCM cells and −/− TCM cells are at an early stage of memory cell differentiation, whereas CCR4+ and CXCR3+ TCM cells have characteristics of more mature cell types.

CXCR3 and CCR4 Identify TCM Cells with Low IFN-γ and IL-4–Producing Capacities. Next, we analyzed effector cytokine–producing capacities of TCM and TEM cell subsets. Total TCM and TEM cell populations were sorted for CXCR3 and CCR4 expression, stimulated with PdBu and ionomycin, and secreted cytokines were quantified by ELISA (Fig. 2 A). TCM cells lacking CXCR3 and CCR4 failed to produce IFN-γ, IL-4, and IL-5, whereas double negative cells in the TEM subset produced all three cytokines. Consistent with the role of CXCR3 and CCR4 as Th1 and Th2 cell markers, CXCR3-expressing cells produced predominantly IFN-γ, whereas CCR4+ cells pro-

### Table I. Expression of CD27, CD62L, and IL-2/IL-15Rβ Chain (CD122) on Naive T Cells and Memory T Cell Subsets

| TCM | CXCR5+ | CXCR5− | CCR4+ |
|-----|--------|--------|-------|
| TCM | −/−    | CXCR3  | −/−   |
| CD27+ (%)b | 99 ± 1 | 98 ± 2 | 76 ± 9 |
| CD62Lb (%)b | 99 ± 1 | 89 ± 5 | 55 ± 15 |
| CD122 (MFI)c | 1 ± 1 | 3 ± 2 | 8 ± 3 |
| PI+ (%)d | 3 ± 2 | 7 ± 4 | 17 ± 9 |

aMean ± standard deviation of four healthy donors.
bPercent of positive cells.
cMean fluorescence intensity.
dPropidium iodide.

Figure 2. Ex vivo cytokine-producing capacities of CD4+ memory T cell subsets. (A) Purified CD4+ TCM and TEM cell subsets were stimulated with PdBu and ionomycin for 24 h and supernatants were analyzed for IFN-γ (diluted 1:4, white bars), IL-4 (black bars), and IL-5 (gray bars) by ELISA. Stimulation with anti-CD3 and anti-CD28 antibodies gave similar results (not depicted). Shown is the mean of four experiments with cells from different donors. (B) CXCR3+ CD4+ T cells were sorted for CCR7 and CD62L expression as indicated and IFN-γ production was assessed as described above. The mean of three independent experiments with three different donors is shown.
duced mainly type 2 cytokines. However, CXCR3+ TCM and CCR4+ TCM cells produced less effector cytokines than the corresponding TEM cell subset, and IL-5 production was entirely restricted to TEM cells. Similar results were obtained when CXCR5+ TCM and CXCR5+ TCM cell subsets were analyzed separately (not depicted). Because CXCR3+ TCM cells contained a considerable fraction of CD62L+ cells (Table I), we further analyzed IFN-γ-producing capacity of CXCR3+ TCM and TEM cells according to CD62L expression (Fig. 2 B). CXCR3+ TCM cells expressing CD62L produced only low amounts of IFN-γ, whereas CXCR3+ TCM cells lacking CD62L produced high levels of IFN-γ comparable to TEM cells. Thus, IFN-γ production among CCR7+ cells is largely restricted to a TEM cell-like subset of CD62L- CXCR3+ cells. Collectively, these results show that CXCR3 and CCR4 identify cells in the TCM cell pool that are nonpolarized or produce low levels of IFN-γ or IL-4.

Cytokine-stimulated CXCR3+ and CCR4+ TCM Cells Differentiate to Th1 and Th2 Cells. We previously showed that some cytokine-stimulated TCM cells spontaneously differentiate to Th1 or Th2 cells, whereas naive cells require TCR ligands or inflammatory cytokines for differentiation (21, 41). To understand whether the subsets defined by CXCR5, CXCR3, and CCR4 could discriminate cells with predetermined fates, we induced proliferation of purified CD4+ T cell subsets by either TSST-loaded DCs or IL-7 and IL-15 (Fig. 3 A). Because memory subsets showed different proliferative responses to IL-7 and IL-15 (Fig. 1 C), and acquisition of effector cytokine–producing capacities progressively increases with division number (42, 43), analysis was performed by gating on cells that had performed the same number of divisions (cytokines: 4; TSST: >7). Under both conditions of stimulation, CXCR5+ TCM cells that lacked CXCR3 and CCR4 expression remained nonpolarized, whereas CXCR5− TCM cells generated some Th1 and Th2 cells and acquired CXCR3 and CCR4 on a fraction of cells at the same time (Fig. 3 C). Cytokine-stimulated CXCR5+ TCM cells progressively lost CXCR5 expression, but homogeneously maintained high levels of CCR7 and CD62L, whereas a fraction of CXCR5− TCM cells progressively lost CCR7 and CD62L expression, thus acquiring the phenotype of TEM cells (Fig. 3 B). Notably, TEM cells remained CCR7− and maintained high effector functions under these conditions.

When CXCR3+ TCM and CCR4+ TCM cells were expanded with homeostatic cytokines, they maintained CXCR3 and CCR4 expression (Fig. 3 C) and spontaneously differentiated into Th1 and Th2 cells, respectively (Fig. 3 A). Thus, CCR4+ TCM cells produced high levels of IL-4 and also secreted IL-5 (not depicted), a type 2 cytokine produced exclusively by TEM cells (Fig. 2 A). Moreover, the amount of IFN-γ produced by CXCR3+ TCM cells stimulated with IL-7 and IL-15 was comparable to that produced by CXCR3+ TEM cells ex vivo, and a fraction of CXCR3+ TCM cells acquired expression of CCR5 (Fig. 3 C), a receptor for inflammatory chemokines expressed on Th1 effector cells (5). Together, these results demonstrate that TCM cells that lack CXCR3 and CCR4 expression are nonpolarized precursors, whereas CXCR3+ and CCR4+ TCM cells represent pre-Th1 and pre-Th2 cells that become fully differentiated Th1 and Th2 effector cells in response to homeostatic cytokines.
To exclude a selective outgrowth of preexisting IFN-γ–producing cells from CXCR3+ TCM cells, we wished to deplete cells with IFN-γ–producing capacity from cytokine-stimulated cultures. To this aim it was necessary to induce IFN-γ production of Th1 cell–polarized memory cells without activating TCR-dependent signaling. It is well established that TCR–independent IFN-γ production of activated Th1 cells can be induced by inflammatory cytokines IL-12 and IL-18 (44, 45). We found that production of IFN-γ by resting CD4+ memory cells in response to IL-12 and IL-18 required activation by IL-7 and IL-15, was boosted by TNF-α, and occurred with delayed kinetics (Fig. 4 A). In contrast, IL-2 production was restricted to TCR-stimulated cells (not depicted), confirming that IFN-γ production by cytokine-stimulated cells is TCR independent. Notably, TCR and cytokine stimulation induced a similar fraction of cells to secrete IFN-γ before cell division (Fig. 4 A, note empty/filled symbols), and IFN-γ production was in both cases restricted to CXCR3+ TCM and TEM cells (Fig. 2 and not depicted). Thus, mature Th1 cell–polarized memory cells can be identified by IFN-γ secretion after either TCR or optimal cytokine activation.

Next, we induced IFN-γ production by purified CFSE-labeled CXCR3+ TCM cells with cytokines, sorted undivided IFN-γ cells after 60 h, expanded them with IL-7 and IL-15, and analyzed effector cytokine–producing capacities of proliferating cells (Fig. 4 B). A large fraction of IFN-γ–CXCR3+ TCM cells differentiated under these conditions and acquired the capacity to produce high levels of IFN-γ. Sorting IFN-γ cells after 72 h gave similar results (not depicted). In contrast, CCR4+ TCM cells secreted IL-4 under the same conditions, whereas IFN-γ+ CXCR3+ TCM cells maintained high IFN-γ production. We conclude that CXCR3+ TCM cells lacking IFN-γ–producing capacity become Th1 cell effector cells after TCR–independent proliferation induced by cytokines.

![Figure 4](https://rupress.org/jem/article-pdf/200/6/725/993978/jem2006725.pdf)

**Figure 4.** (A) Kinetics and requirements of TCR- and cytokine-induced IFN-γ production. CFSE-labeled CD4+ memory T cells were stimulated for the indicated times with either anti-CD3 and anti-CD28 antibodies (squares), or with TNF-α, IL-12, and IL-18 in the absence (circles) or presence (triangles) of IL-7 and IL-15. IFN-γ production was analyzed by intracellular staining. Empty symbols indicate conditions with undivided cells, whereas filled symbols indicate conditions with dividing cells. The mean percentage of IFN-γ cells of three independent experiments is plotted. (B) Cytokine-stimulated CXCR3+ TCM cells lacking IFN-γ–producing capacity become Th1 cell effectors. Purified CFSE-labeled CXCR3+ TCM and CCR4+ TCM cells were stimulated with IL-7, IL-15, TNF-α, IL-12, and IL-18 for 60 h, and IFN-γ–secreting cells were purified by cell sorting. IFN-γ and IFN-γ–cells were then expanded for an additional 5 d with IL-7 and IL-15, briefly stimulated with PdBu and ionomycin, and analyzed for IL-4 and IFN-γ production by intracellular staining. One representative donor out of three is shown.

![Figure 5](https://rupress.org/jem/article-pdf/200/6/725/993978/jem2006725.pdf)

**Figure 5.** Effects of polarizing cytokines on TCR- and cytokine-induced differentiation. Purified CFSE-labeled CD4+ T cell subsets were stimulated with DC plus TSST or IL-7 plus IL-15 in the absence or presence of IL-12 and neutralizing anti-IL-4 antibody (Th1-condition) or IL-4 and neutralizing anti-IL-12 antibody (Th2-condition). After 7 d, cells were stimulated with PdBu and ionomycin and cells of the same division number were analyzed for IFN-γ and IL-4 production by intracellular staining. The percentages of IFN-γ cells (white bars), IL-4 cells (black bars), and of cells producing both cytokines (gray bars) are represented. One representative experiment out of five with different donors is shown.
We then analyzed the effects of polarizing cytokines on T cell differentiation induced by TCR agonists or homeostatic cytokines. IL-4 and IL-12 induced differentiation of TCR-stimulated CXCR5+ and especially −/− TCM cells into IL-4+ and IFN-γ-producing cells, respectively, whereas they failed to modulate T cell differentiation in IL-7 plus IL-15-activated T cells (Fig. 5, A and B). Under the same conditions of TCR stimulation, IL-4 induced CXCR5+ T cells to produce the opposite cytokine IL-4+ and promoted CCR4 expression, whereas IL-12 induced CCR4+ cells to produce IFN-γ (Fig. 5 A) and up-regulate CXCR3 (not depicted). Again, polarizing cytokines had little effect on the extent of T cell differentiation in IL-7 plus IL-15-stimulated cells (compare Figs. 3 A and 5 B). Similar results were obtained when IL-2 substituted for IL-7 and IL-15, and in the absence or presence of TNF, IL-6 and IL-10, DC-derived cytokines that strongly boost proliferation of TCM cells in response to IL-7 and IL-15 (not depicted; reference 21). Together, these findings suggest that flexibility of cytokine gene expression of human memory T cells requires TCR triggering and polarizing cytokines (13).

**Proliferation History, In Vivo Turnover, and Recall Responses of CD4+ Memory T Cell Subsets.** The proliferation history and in vivo turnover of the different memory T cell subsets was then assessed by measuring the amounts of TREC and the spontaneous BrdU incorporation of ex vivo–isolated cells. TREC carry a particular signal joint sequence (37) were quantified by TaqMan PCR in T cell subsets from five healthy donors (Fig. 6 A). As expected, CD4+ CD45RA+ naive T cells contained high levels of TREC, whereas B cells and T cell clones were negative (not depicted). Compared with naive T cells, CXCR5+ TCM and CXCR5− TCM cells contained much lower amounts of TREC, whereas T EM cells contained the lowest amount. Although there were considerable quantitative differences among individual donors, these data indicate that the different subsets of TCM cells have divided to a similar extent.

To measure the spontaneous BrdU incorporation, freshly isolated PBMCs were incubated with BrdU. CD4+ T cells were then purified and T cell subsets analyzed by intracellular staining with anti-BrdU antibodies. As shown in Fig. 6 B, CXCR5+ TCM, CXCR5+ TCM, and CCR4+ TCM cells spontaneously incorporated BrdU to a similar extent, whereas T EM and especially −/− TCM cells had a higher proliferation rate and naive cells were below the detection limit. These data indicate that memory T cells in different subsets slowly turn over under steady-state conditions in vivo.

To investigate whether the different memory subsets contained cells specific for recall antigens, T cell populations were isolated from smallpox- and/or tetanus-vaccinated donors and stimulated for 7 d with autologous monocytes that were incubated with TT or with an extract of CMV-derived proteins. To assess memory against smallpox, monocytes were infected with a replication-deficient VV. In these experiments, purified CFSE-labeled T cell subsets and CFSE dilution were used to read out proliferation of pathogen-specific T cells. From eight donors analyzed, seven responded strongly to TT, whereas one donor that had been boosted 20 yr ago had a low response (not depicted). Five donors responded strongly to CMV, and four of six donors that had also been vaccinated against smallpox had a detectable response to VV. Responses to autologous monocytes alone were undetectable or very low (not depicted).

We analyzed the distribution of TT−, CMV−, and VV−specific T cells among different memory subsets (one donor

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**Figure 6.** Proliferation history, in vivo turnover, and recall responses of CD4+ memory T cell subsets. (A) Total CXCR5+ TCM, CXCR5− TCM, and T EM cells were analyzed for their single joint TREC content and compared with naive cells from the same donor (naive cells: 100%). Bars indicate the mean TREC levels in memory subsets of five different donors. (B) Freshly isolated PBMCs were incubated with BrdU, CD4+ T cells were isolated, and BrdU incorporation was analyzed as a function of CD45RA and chemokine receptor expression by flow cytometry. The mean percentage of BrdU+ cells in a given subset of four donors is shown. (C) CD4+ naive and memory subsets were sorted, labeled with CFSE, and incubated with autologous monocytes that had either been infected with VV or incubated with TT or an extract of CMV-derived proteins. CFSE profiles of viable CD4+ CD14− cells on day 7 of one representative donor are shown. (D) Recall responses of memory subsets to TT of eight different donors were assessed as described above, and the frequency of TT-specific cells was calculated after 5 d. The frequency of TT-specific cells in the induced memory subsets of seven different TT-responsive donors was plotted against the time of the last boost.

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responding to all three pathogens is shown in Fig. 6 C). In all cases, pathogen-specific cells were detected in both the T_C and T_EM cell pools. TT-specific cells were undetectable among naive cells, but present in all memory subsets in six of seven responsive donors. Conversely, CMV-specific cells were largely restricted to CXCR3$^+$ T_C and T_EM cells in all five responsive donors, consistent with the notion that CMV infection promotes a Th1 cell response (46). Consistent with previous reports, VV-specific cells were less frequent (47, 48), but were detectable in CXCR3$^+$ T_C and T_EM cells and, interestingly, in three of four donors in CXCR5$^+$ T_C cells. Together, these results show that all memory subsets contain cells specific for recall antigens, and that the distribution of antigen-specific cells within T_C cell subsets varies for different pathogens.

Next, we compared precursor frequencies of TT-specific cells in different subsets in donors that had been boosted recently or several years ago (Fig. 6 D). TT-specific cells were relatively frequent among T_EM cells (>1:500) in all donors. Conversely, CXCR5$^+$ T_C cells were less frequent (<1:500), especially in donors that had not been boosted for several years, possibly explaining the failure of previous studies to detect TT-specific cells in this subset using thymidine incorporation (29, 31). Interestingly, the relative distribution among CXCR3$^+$ T_C and CCR4$^+$ T_C cells was highly variable, with some donors having higher numbers of CCR4$^+$ T_C cells and others containing predominantly CXCR3$^+$ T_C cells. These results indicate that TT-specific T cells are present in high frequency in T_EM cells, even several years after vaccination, and are distributed in different subsets of T_C cells.

**Discussion**

We have shown that the human CD4$^+$ T_C cell pool can be subdivided into subsets of nonpolarized cells and pre-Th1 and pre-Th2 cells based on chemokine receptor expression. These subsets have extensively divided in vivo and contain cells specific for recall antigens and with self-renewal capacity. Upon TCR-independent proliferation induced by homeostatic cytokines, T_C cell subsets are committed for different fates and become Th1, Th2, or remain nonpolarized cells, explaining how the quality of the primary immune response could be maintained by T_C cells in the absence of antigen.

Th cells expressing CXCR5 comprise CCR7$^-$ CD57$^+$ follicular Th cells in tonsils (29, 30, 49), and nonpolarized circulating CXCR5$^+$ T_C cells of unknown function and specificity that might have a recent activation history (31). Our results show that nonpolarized cells are present in both CXCR3$^+$ and CXCR5$^+$ T_C cell subsets and lack CXCR3 and CCR4 expression. Using CFSE dilution we were further able to show that CXCR5$^+$ T_C cells contained low numbers of TT-specific cells even several years after vaccination. Moreover, three of four smallpox-vaccinated, responsive donors contained VV-specific cells at low frequency in the CXCR5$^+$ T_C cell subset. Using quantitative PCR, we found that CXCR5$^+$ T_C and CXCR5$^-$ T_C cells contained comparable amounts of residual TREC3, whereas T_EM cells had slightly lower levels, suggesting that T_C cell subsets had divided to a comparable extent (approximately seven times). However, because naive and memory cells were identified by CD45 isoform expression that is not a stable marker (25, 50), the number of divisions performed by memory cells might be underestimated by our analysis. Spontaneous BrdU uptake indicated that CXCR5$^+$ T_C cells have an in vivo proliferation rate that is comparable to that of other memory subsets. The relative small cell size and the absence of CD69 on BrdU$^+$ cells suggest that this proliferation is driven by homeostatic mechanisms rather than by antigen. Why $^{-/-}$ T_C cells have a higher turnover than other memory subsets is currently unclear. They might be particularly fit because they combine a relatively high cytokine responsiveness with a low susceptibility to apoptosis (51). Alternatively, they might be preferentially located in cytokine-rich microenvironments or in proximity to DCs that boost proliferation in response to IL-7 and IL-15 (21). In any case, these results show that all CD45RA$^+$ subsets, including CXCR5$^+$ T_C cells, are memory cells that have extensively divided, slowly turnover in vivo, and contain cells specific for recall antigens.

Different viruses are known to induce CD8$^+$ memory cells belonging preferentially to different subsets (52, 53). Here we showed that although pathogen-specific CD4$^+$ T cells are present in both the T_C and T_EM cell pools, they have characteristic distributions in T_C cell subsets, reflecting the Th1/Th2 cell polarization induced by the pathogens or vaccinations. Thus, TT-specific cells were detected in all subsets, consistent with the notion that vaccination against tetanus induces a mixed Th1/Th2 cell response (54). Conversely, CMV and VV promote Th1 cell polarization (46, 47), and virus-specific cells were consequently detected in CXCR3$^+$ T_C cells but not in CCR4$^+$ T_C cells. In one donor, we were able to show that VV-specific T_EM cells were also CXCR3$^+$ (not depicted). Collectively, these results suggest that immune responses generate heterogeneous populations of memory cells that belong to different subsets and comprise a broad spectrum of differentiation stages. The distribution between CXCR3$^+$ and CCR4$^+$ subsets in the T_C and T_EM cell pools might be useful to monitor the quality of the memory response to different pathogens.

Human memory T cells can be subdivided into CCR7$^+$ T_C and CCR7$^-$ T_EM cells with different effector functions and homing potentials, suggesting a division of labor between these two subsets (12). However, several recent reports showed that antigen-experienced CCR7$^+$ cells possess immediate effector functions (9, 32–35). Although we identified here CCR7$^+$ memory cells with IFN-γ- and IL-4–producing capacities as CXCR3$^+$ T_C and CCR4$^+$ T_C cells, respectively, the following lines of evidence suggest that these cells are not fully differentiated effectors: (a) they had a higher expansion potential than T_EM...
cells and most cells had retained CD27 and CD62L expression; (b) IFN-γ production by CXCR3+ TCM cells was low and largely restricted to unconventional CCR7+ CD62L− cells; (c) although CCR4+ TCM cells produced some IL-4, production of IL-5, which acts on eosinophils at peripheral sites of allergic inflammation, was limited to T EM cells; (d) many CXCR3+ TCM and CCR4+ TCM cells had retained flexibility to differentiate to Th2 and Th1 cells, respectively, upon antigenic stimulation in the presence of appropriate polarizing cytokines; and (e) upon cytokine stimulation, they further differentiated, losing CCR7 and CD62L and acquiring nonlymphoid homing potential and high levels of effector functions. Together, these findings are consistent with the notion that nonlymphoid tissue homing potential and effector cytokine-producing capacities are progressively acquired upon T cell differentiation and reside predominantly in the T EM cell subset of the human CD4+ memory cell pool (3). However, because effector functions and nonlymphoid homing potentials are acquired in a stochastic manner (55), some cells have characteristics that are intermediate between T CM and T EM cells.

We previously proposed that cytokine-driven differentiation of T CM cells might be a mechanism to replenish short-lived T EM cells in the absence of antigen (21), but how nonpolarized T CM cells could faithfully maintain polarized Th1 or Th2 effector cell populations remained unclear. A recent report showed that CD4 T cell priming upon viral infection generated both short-lived effector cells and long-lived precursors that lacked effector functions, but spontaneously acquired IFN-γ-producing capacity when transferred into antigen-free hosts (26). We showed here that CXCR3+ T CM and CCR4+ T CM cells invariably differentiated to CCR7− Th1 or Th2 effector cells in an antigen-independent fashion, whereas CXCR5+ T CM cells remained nonpolarized and CCR7+. CXCR3 and CCR4 are preferentially induced under type 1 differentiation and reside predominantly in the TEM cell subset of the human CD4+ memory cell pool (3). How-ever, because effector functions and nonlymphoid homing potentials are acquired in a stochastic manner (55), some cells have characteristics that are intermediate between T CM and T EM cells.

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