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Human skin is protected by four functionally and phenotypically discrete populations of resident and recirculating memory T cells

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Abstract

The skin of an adult human contains approximately 20 billion memory T cells. Epithelial barrier tissues are infiltrated by a combination of resident and recirculating T cells in mice but the relative proportions and functional activities of resident versus recirculating T cells have not been evaluated in human skin. We discriminated resident from recirculating T cells in human engrafted mice and lymphoma patients using alemtuzumab, a medication that depletes recirculating T cells from skin, and then analyzed these T cell populations in healthy human skin. All non-recirculating resident memory T cells (T_{RM}) expressed CD69, but the majority were $CD4^+$, $CD103^-$ and located in the dermis, in contrast to studies in mice. Both $CD4^+$ and $CD8^+ CD103^+$ T_{RM} were enriched in the epidermis, had potent effector functions and had a limited proliferative capacity compared to $CD103^- T_{RM}$. T_{RM} of both types had more potent effector functions than recirculating T cells. Induction of CD103 on human T cells was enhanced by keratinocyte contact, depended on $TGF\beta$ and was independent of T cell keratinocyte adhesive interactions. We observed two distinct populations of recirculating T cells, $CCR7^+/L\text{-selectin}^+$ central memory T cells (T_{CM}) and $CCR7^+/L\text{-selectin}^-$ T cells, which we term migratory memory T cells (T_{MM}). Circulating skin-tropic T_{MM} were intermediate in cytokine production between T_{CM} and effector memory T cells. In patients with cutaneous T cell lymphoma, malignant T_{CM} and T_{MM} induced distinct inflammatory skin lesions and T_{MM} were depleted more slowly from skin after alemtuzumab, suggesting T_{MM} may recirculate more slowly. In summary, human skin is protected by four

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functionally distinct populations of T cells, two resident and two recirculating, with differing territories of migration and distinct functional activities.

Introduction

Research in both mice and humans has confirmed that skin and other epithelial barrier tissues are populated by a combination of non-recirculating resident memory cells (T_{RM}) as well as T cells that recirculate in and out of tissues (1, 2). However, the relative proportions and functional activities of resident and recirculating T cells in skin have never been comprehensively studied, especially in humans. T_{RM} cause psoriasis and mycosis fungoides and are implicated in a variety of other human autoimmune and inflammatory conditions (3). A better understanding of resident and recirculating T cells could lead to novel therapies for T cell mediated inflammatory diseases.

Our prior studies of patients with cutaneous T-cell lymphoma (CTCL) found that treatment with alemtuzumab, a humanized anti-CD52 antibody, depleted all circulating T cells and purged the skin over time of recirculating T cells but spared a population of non-recirculating T_{RM} in the skin (2). Alemtuzumab depletes T cells in the blood stream but not in the skin because it depletes by antibody dependent cellular cytotoxicity (ADCC) requiring the presence of neutrophils and/or NK cells, cell types that are rare in human peripheral tissues such as the skin but frequent in the circulation (2, 4).

In these studies, we have studied resident and recirculating T cell subsets in a human engrafted mouse model and in human patients with CTCL, utilizing alemtuzumab as a tool to deplete recirculating T cells from skin. This has allowed us to discriminate between resident and recirculating T cell populations in human skin and to study their relative frequencies and effector functions. We find that human skin is protected by two distinct populations of recirculating T cells and two distinct populations of resident memory T cells, each with different functional capacities.

RESULTS

Skin T cells in a human engrafted mouse model recapitulate T cell populations in healthy human skin

In order to selectively study recirculating and resident T cells in human skin in an accessible model, we grafted NSG mice with human neonatal foreskin, infused them IV with allogeneic PBMC, allowed a dermatitis to develop and then treated these mice with alemtuzumab to deplete recirculating T cells from skin (Fig. 1A). Neonatal foreskin was used in these studies because, based on mouse studies showing that T_{RM} are generated in skin following cutaneous infections, we suspected neonatal human foreskin would have few resident memory T cells (1, 5, 6). This was in fact the case. Human neonatal foreskin contained resident antigen presenting cells (APC) including CD1a⁺ Langerhans cells and CD11c⁺ dendritic cells (DC, Fig. 1B,C) but had very few T cells (Fig. 1C,D). There were four-fold more CD11c⁺ DC than T cells in neonatal foreskin (Fig. 1C) and 45-fold fewer T cells in foreskin than in healthy adult human skin (Fig. 1D). Additionally, T cells isolated

from foreskin lacked expression of the T_{RM} markers CD69 and CD103, which were robustly expressed by a subset of T cells from adult human skin (Fig. 1E). Neonatal foreskin was therefore an excellent source of T cell depleted human skin in which the migration and differentiation of newly generated T_{RM} could be studied. Allogeneic T cells injected IV into grafted mice migrated specifically into the human skin graft and induced an inflammatory dermatitis characterized by migration of T cells into both the dermis and epidermis (Fig. 1F–H). In contrast, mice grafted with human foreskin but not injected i.v. with allogeneic T cells lacked inflammation (Fig. 1I). T cells were evident within the graft by week 1 after infusion and the T cell infiltrate was maximal by week 3 (Supp. Fig. 1). In mice, T_{RM} generated by viral infections with HSV and vaccinia virus co-expressed CD69 and CD103 (5, 6). A subset of T cells infiltrating the human skin graft up-regulated CD69 and CD103 after migration into the graft (Fig. 1J). CD69 was expressed within the first week after T cell entry into the graft but up-regulation of CD103 was slower and peaked at week three (Fig. 1J). We compared the T cells in foreskin grafts with those isolated from healthy adult human skin, a tissue known to contain both skin-tropic $CCR7^+/L\text{-selectin}^+$ central memory cells (T_{CM}) and non-recirculating T_{RM} (2, 7). Grafts from mice contained $CCR7^+/L\text{-selectin}^+$ skin tropic T and T_{CM} cells that expressed CD69 and CD103 in proportions similar to those isolated from antigen-experienced adult human skin (Fig. 1K). A population of $CD69^+$ T cells that lacked CD103 expression, a subset not reported in mice, was also observed in skin grafts and in healthy human skin (Fig. 1K). The percentages of T cells expressing CD69 and CD103 in the grafted skin was not significantly different from the percentages expressed by T cells from healthy adult human skin (Fig. 1L).

Alemtuzumab depletes recirculating T_{CM} from skin grafts but spares a population of T cells in skin

Our previous studies in human CTCL patients treated with alemtuzumab demonstrated that this medication depleted T cells in blood and purged the skin of recirculating T cells over time but spared a population of non-recirculating resident memory T cells in skin (2). Similar to our results in CTCL patients, treatment of foreskin-grafted, PBMC-injected mice with alemtuzumab depleted circulating T cells from the blood but spared a subpopulation of T cells in skin (Fig. 2A,B). In contrast, A-dmDT(390)-bisFv(UCHT1), a recombinant immunotoxin composed of diphtheria toxin conjugated to anti-CD3, completely depleted T cells in both blood and skin (Fig. 2A,B). In CTCL patients, alemtuzumab depleted the skin of recirculating $CCR7^+/L\text{-selectin}^+$ central memory T cells (T_{CM}), a population that is also observed in healthy skin (2, 7). Likewise, alemtuzumab depleted both $CD4^+$ and $CD8^+$ T_{CM} from skin grafts on human engrafted mice (Fig. 2C,D).

Distinct populations of non-recirculating T_{RM} exist in human engrafted mice and human skin

Having established that the cell populations in human engrafted mice were comparable to those in healthy human skin and that alemtuzumab depleted recirculating T cells in this model as it did in human patients (Figs. 1,2), we set out to study the non-recirculating T_{RM} populations left within the human skin grafts after alemtuzumab treatment. The vast majority of T_{RM} remaining in skin after alemtuzumab depletion of recirculating T cells expressed $CD69^+$ and a subset coexpressed CD103 (Fig. 3A). In human engrafted mice,

CD4⁺ T cells were four-fold more frequent than CD8⁺ T cells both before and after depletion of recirculating T cells with alemtuzumab, suggesting that CD4 cells predominated in both the recirculating and resident T cell populations in this model (Fig. 3B). Approximately half of CD8⁺ T_{RM} expressed CD103 but the majority of CD4⁺ T_{RM} did not (Fig. 3C). When T_{RM} were evaluated as a whole, the most frequent population were CD4⁺ CD103⁻ T cells, which made up approximately 66% of the total T_{RM} population (Fig. 3D). CD8⁺ T cells made up 17% of the T_{RM} population and only 26% of the total T_{RM} population expressed CD103. CD69⁺ T cells were evident in both the epidermis and dermis of grafts from alemtuzumab treated human engrafted mice and only a subset of these T cells, in both the epidermis and the dermis, coexpressed CD103 (Fig. 3E). In order to confirm these results in human patients, we studied the T cells remaining in the skin of CTCL patients after alemtuzumab therapy. We found that, similar to results in human engrafted mice, T_{RM} remaining in skin after alemtuzumab virtually all expressed CD69 and a subset of both CD4 and CD8 T cells co-expressed CD103 (Fig. 3F). In addition, a small population of L-selectin⁺CCR7⁻ existed in human engrafted mice, healthy skin, and the skin of CTCL patients (Figs. 2C, S3A, E–F). These cells were not depleted by alemtuzumab and co-expressed CD69, suggesting they are T_{RM} (Figs. 2C, S3A, E–F).

Antigen experienced human skin contains two phenotypically and functionally distinct populations of T_{RM}

Studies in human engrafted mice and alemtuzumab treated CTCL patients demonstrated there were two distinct populations of CD69⁺ non-recirculating T_{RM} in skin, one that expressed CD103 and one that did not. However, the relative percentages and effector functions of cells in the mouse model and in patients with a Th2 biased T-cell lymphoma were unlikely to reflect those observed in healthy human skin (8–11). We therefore extracted and studied the T cells from healthy adult human skin, a tissue in which T_{RM} have presumably accumulated over many years of infections and antigen exposures.

In mouse skin, the most frequently observed and well described T_{RM} subset are intraepithelial CD8⁺ CD103⁺ T cells and T cells of this phenotype have also been observed after HSV infection in humans (5, 6, 12, 13). In order to study the location and proportion of T_{RM} in human skin, we enzymatically split skin into epidermis and dermis and isolated the T cell populations by collagenase digestion (Fig 4A–C). CD4⁺ and CD8⁺ T cells co-expressing CD69 and CD103⁺ were evident in both the epidermis and the dermis, although CD103⁺ T_{RM} of both subsets were more frequent in the epidermis. CD69⁺CD103⁻ T_{RM} were more prevalent in the dermis and both CD4⁺ and CD8⁺ T cells had this phenotype. CD69⁻ T cells, referred to as recirculating T cells, were the least frequent T cell population in both the epidermis and the dermis (Fig. 4B,C).

We next studied the functional capacities of T_{RM} and recirculating subsets from whole skin, epidermis and dermis. In general, cytokine production by the two distinct T_{RM} subsets was greater than observed in recirculating T cells, consistent with their role as resident immune sentinels (Fig. 4D). Effector cytokine production by CD103⁺ T_{RM} also tended to be more robust than in CD103⁻ T_{RM}, most notably with respect to production of IFN γ , TNF α and IL-22. Taken together, these results demonstrate that two phenotypically and functionally

distinct populations of T_{RM} exist in both the $CD4^+$ and $CD8^+$ T cell compartments in human skin. $CD103^+$ T cells were enriched in the epidermis and had a tendency for increased effector cytokine production, whereas $CD103^- T_{RM}$ were more frequent in the dermis and had slightly lower but still very potent effector functions when compared to recirculating T cells.

In order to study the relative proliferative capacity of the two T_{RM} subsets in healthy skin, we first isolated T cells from the epidermis and dermis as $CD103$ enriched and depleted populations respectively. T cells isolated from the epidermis retained $CD103$ expression but rapidly lost functional activity and underwent apoptosis after removal from the skin environment, similar to the behavior of $CD103^+ T_{RM}$ isolated from mouse brain after infection with vesicular stomatitis virus (14) (Fig. S2A–E). In order to study antigen responses and proliferative capacity without removing T cells from skin, we injected stimulatory anti- $CD3$ and anti- $CD28$ antibodies or antigen derived from heat killed *Candida albicans* or *Staphylococcus aureus* directly into explants of human skin. Injected skin explants were cultured for 2 weeks after which T cells were collected. Proliferation was assayed by staining for Ki-67 and cytokine production was assayed by intracellular flow cytometry following short-term stimulation with PMA and ionomycin (Fig. 5). In both $CD4^+$ and $CD8^+$ T cells, proliferation was greatest in the $CD103^- T_{RM}$ subset in response to both antibody and antigen stimulation, although differences only reached significance in $CD4^+$ T cells (Fig. 5A,C). Production of $TNF\alpha$ tended to be greater in the $CD103^+$ population (Fig. 5B,D). Taken together, these studies suggest that $CD103^+ T_{RM}$ had superior effector functions but a limited proliferative capacity.

T cell $CD103$ induction is enhanced by keratinocyte contact, depends on $TGF\beta$ and is independent of T cell keratinocyte adhesion

In mouse models of viral skin infection, $CD103$ expression by T cells required entry into the epidermis and depended on $TGF\beta$ (15, 16). In human engrafted mice, T cells that had migrated into the epidermis had a higher rate of $CD103$ positivity than dermal T cells (Fig. 6A,B). To determine if keratinocyte contact can induce human T cell $CD103$ upregulation, we cultured T cells stimulated with $\alpha CD3/\alpha CD2/\alpha CD28$ beads on keratinocyte or fibroblasts monolayers either in direct contact or separated by transwells. $CD103$ upregulation was enhanced by direct contact with keratinocytes (Fig. 6C). To determine if adhesive interactions between T cells and keratinocytes participated in the signaling required for induction of $CD103$, we included function blocking antibodies against E-cadherin, LFA-1 and $\beta 1$ integrins in T cell and keratinocyte monolayer cultures (Fig. 6D). Blocking adhesive interactions had no effect on $CD103$ upregulation but neutralizing antibodies against $TGF\beta$ decreased $CD103$ induction. Additional studies in transwells and in direct co-cultures demonstrated that neutralization of $TGF\beta$ decreased the $CD103$ upregulation induced by keratinocyte contact and that recombinant human $TGF\beta$ markedly increased $CD103$ induction in both the presence and absence of contact with keratinocytes (Fig. 6E,F). Although adhesive interactions between keratinocyte E-cadherin and $CD103$ did not participate in stimulating $CD103$ induction, these interactions did function to anchor T cells to keratinocyte monolayers.

Human skin also contains two distinct populations of recirculating memory T cells

We had previously reported that healthy human skin was patrolled by a skin tropic population of central memory T cells that co-expressed both skin homing receptors (CLA/CCR4) and central memory markers (CCR7/L-selectin) and were depleted by alemtuzumab therapy (2). However, when reviewing our results, we observed that both healthy skin and the skin of CTCL patients contained a distinct population of CCR7⁺/L-selectin⁻ T cells (Fig. 7A,B). CCR7⁺/L-selectin⁻ T cells were confined to the dermis and absent from the epidermis (Fig. S3A). Approximately a third of CCR7⁺/L-selectin⁻ T cells co-expressed CD69, suggesting they may be T cells in a transitional state, but the remaining CCR7⁺/L-selectin⁻/CD69⁻ T cells made up a significant proportion of T cells in both healthy skin and in CTCL patients (Fig. 7B). These T cells were depleted in CTCL patients by alemtuzumab therapy, demonstrating that they represent a recirculating T cell population (Fig. 7A,B). In mice, CCR7 expression is required for exit from peripheral tissues via lymphatics and a recirculating population of CD4⁺/CCR7⁺/L-selectin^{lo} T cells was found in Kaede mice to recirculate out of skin in a CCR7-dependent manner (17–19). CCR7⁺/L-selectin⁻ cells, which we will refer to as migratory memory T cells (T_{MM}), were present in similar proportions in both healthy skin and among the benign, nonmalignant T cells in patients with CTCL and these cells were completely depleted by alemtuzumab therapy (Fig. 7B). T_{MM} made up the majority of both CD4⁺ and CD8⁺ CLA⁺ skin tropic T cells in the blood of healthy human donors, confirming that T cells do recirculate between the blood and skin (Fig. 7C, representative histograms in Fig. S4). We next isolated skin tropic (CLA⁺) T_{CM}, T_{MM} and effector memory T cells (T_{EM}) from human blood and compared their relative functional capacities after stimulation with PMA and ionomycin followed by comprehensive cellular phenotype by CyTOF mass cytometry (Fig. 7D). In general, effector cytokine production by T_{MM} was intermediate between that of T_{EM} and T_{CM}. Interestingly, production of Th1, Th17 and Th22 proinflammatory effector cytokines was highest in T_{EM}, but Th2 cytokine production (IL-4 and IL-13) was highest in T_{CM}. TNF α was produced at high levels by all T cell subsets. Although not significantly different, IL-2 production, previously reported as a feature of T_{CM} that may render them resistance to apoptosis, tended to be higher in the recirculating T_{CM} and T_{MM} populations than in T_{EM} (20).

T_{MM} are associated with the development of expanding skin lesions in patients with CTCL and are depleted from skin more slowly than T_{CM}

CLA⁺ skin tropic CCR7⁺/L-selectin⁻ T_{MM} clearly recirculate between human blood and skin, as evidenced by their presence in both sites and their depletion by alemtuzumab in patients with CTCL. T_{MM} leaving skin via the lymphatics would be expected to enter skin draining lymph nodes and eventually re-enter the circulation via the thoracic duct. However, these cells lack L-selectin and likely cannot access lymph nodes from blood the way naïve T cells and T_{CM} are thought to, by migrating across high endothelial venules (21, 22). Instead, these T cells would be expected to home directly back to skin. In order to study the biologic behavior of these cells in human beings in vivo, we studied CTCL patients with malignant T cells that expressed markers of T_{MM}. We have previously observed that patients with classic Sézary syndrome, composed of the clinical triad of diffuse erythema of the skin, enlargement of the lymph nodes draining both skin and other organs and circulating malignant T cells in the blood, have malignant T cells with the phenotype of skin tropic

central memory T cells (2, 23). $CLA^+ T_{CM}$ would be expected to recirculate between the skin, blood and lymph nodes according to our current understanding of trafficking, and this is precisely where malignant T cells accumulate in Sézary patients. Moreover, we found that malignant T cells of a $CLA^+ T_{CM}$ phenotype induced in patients a diffuse erythema of the skin, and that this diffuse erythema was the best clinical predictor of complete remission in response to alemtuzumab therapy (Fig. 8A)(24). However, patients have a myriad of clinical presentations and we have found that some patients present with erythematous skin lesions that have ill-defined borders, in contrast to patients with Sézary syndrome, who present with diffuse erythema, and patients with mycosis fungoides (MF), who present with fixed, recurrent and sharply demarcated inflammatory skin lesions (Fig. 8). We found that patients with malignant T cells that had the phenotype of $CCR7^+/L\text{-selectin}^- T_{MM}$ tended to have only involvement of the lymph nodes draining the skin and to have expanding inflammatory skin lesions with ill-defined borders. (Fig. 8B–D). In one patient, the malignant T cells in blood and skin were a combination of both T_{CM} and T_{MM} . This patient had a background of diffuse erythema as well as more discrete skin lesions on presentation (Figure 8E). After 18 days on alemtuzumab, re-biopsy of the skin demonstrated that T_{CM} had been partially depleted but large numbers of T_{MM} remained in the skin. This was associated with resolution of the diffuse erythema but persistence of the erythematous skin lesions. Taken together, the lack of lymph node disease, the clinical appearance of these lesions and the fact that T_{CM} were depleted more rapidly from the skin than T_{MM} after alemtuzumab therapy suggest that T_{MM} do recirculate between the blood and skin, are excluded from the lymph nodes, and may recirculate more slowly than T_{CM} .

Discussion

The skin surface of a healthy adult human being contains nearly 20 billion diverse memory T cells with potent effector functions (7). Our previous studies in alemtuzumab treated CTCL patients demonstrated that human skin contains a combination of resident and recirculating memory T cells but the relative frequencies and functional activities of the subsets remain uncharacterized (2). In the current studies, we used alemtuzumab as a tool to deplete recirculating T cells from skin in order to discriminate between recirculating and resident memory T cell populations.

We developed a human engrafted mouse model to study human resident and recirculating T cell populations in skin. Immunodeficient mice were grafted with human neonatal foreskin, a tissue that contains resident Langerhans cells and DC but has few T cells. The paucity of resident T cells in the graft allowed us to be more confident that T cells observed in the skin after allogeneic i.v. PBMC infusion were derived from the i.v. injected T cells. T_{RM} are normally generated as a result of cutaneous infections. We took advantage of the fact that graft resident APC were likely to activate T cells migrating into the graft from the blood via direct allo-recognition, a direct recognition of MHC I and II by allogeneic $CD8^+$ and $CD4^+$ T cells, respectively. Direct allo-recognition induces a robust polyclonal T cell response, engaging up to 10% of T cells (25). In other words, we utilized the allogeneic T cell response to engage the T cell receptor and initiate the T_{RM} differentiation program within grafted skin. We observed the development of $CD69^+$ and $CD103^+$ T cells in grafted skin in proportions that were similar to those in adult human skin. However, antigen recognition of

allogeneic tissues is likely ongoing in human engrafted mice and some CD103⁻ T cells may be in the process of transitioning to CD103⁺ T_{RM}.

We next treated these mice with alemtuzumab in order to discriminate between recirculating and resident memory T cell subsets. Similar to our previous results in CTCL patients treated with alemtuzumab, alemtuzumab purged the skin of recirculating T_{CM} but spared a population of T cells in skin. In contrast, immunotoxin A-dmDT (390)-bisFv(UCHT1), an anti-CD3 recombinant diphtheria immunotoxin currently in trials for the treatment of CTCL, depleted T cells completely in both blood and skin of grafted mice (26).

We observed that virtually all T cells remaining in the skin of grafted mice following alemtuzumab expressed CD69 and similar results were observed in alemtuzumab treated CTCL patients. In mice, nonrecirculating T_{RM} generated from experimental infections expressed CD69 and populations of T cells in human peripheral tissues have been shown to express this marker as well (1, 7, 27). CD69 suppresses the activity of sphingosine 1 phosphate receptor 1 (S1P1) and S1P1 down-regulation in mice was found to be necessary for retention of CD8⁺ T_{RM} in skin (16, 28). However, it has never been conclusively demonstrated in humans that CD69 identifies the non-recirculating T cell pool in peripheral tissues. Our studies in human engrafted mice and in human patients treated with alemtuzumab demonstrate that CD69 is an excellent marker for nonrecirculating T_{RM} in human skin.

Most of the T_{RM} generated by viral infections in mice are sessile CD8⁺ T cells expressing CD69 and CD103 localized to the epithelium, although dermal CD4⁺ T_{RM} with more mobility have also been observed following HSV infection (1, 5, 6, 29). Our studies in engrafted mice, CTCL patients and healthy human skin demonstrate that the majority of human skin T_{RM} are dermal CD4⁺ T cells that lack CD103 expression. However, we do find that CD103⁺ T cells, both CD4⁺ and CD8⁺, are enriched in the epidermis. With respect to their functional capacities, CD103⁺ T_{RM} had more potent effector functions but a poorer proliferative capacity. T cells isolated from the epidermis, which are enriched for CD103⁺ T_{RM}, rapidly underwent apoptosis when removed from the skin microenvironment and had a poor proliferative capacity; these findings are very similar to the behavior of CD103⁺ T_{RM} isolated from the brains of mice following infection with vesicular stomatitis virus (14).

Upregulation of CD103 by skin T cells in mouse models has been shown to occur after entry into the epidermis and to depend on TGFβ (15). A second study suggested IL-33 and TNFα may also contribute (16). We observed in human engrafted mice that T cells that had migrated into the epidermis had higher levels of CD103 than dermal T cells. Hypothesizing that contact with keratinocytes might participate in CD103 upregulation, we found that direct culture of stimulated blood T cells with keratinocyte monolayers enhanced CD103 expression in a manner that was TGFβ dependent but did not require adhesive interactions between T cells and keratinocytes. Moreover, exogenous TGFβ markedly enhanced CD103 induction in the presence or absence of direct keratinocyte contact. Although signaling from integrin and CD103-E-cadherin mediated adhesive interactions did not enhance CD103 upregulation, CD103-E-cadherin interactions did serve to tether T cells to keratinocyte monolayers. Taken together, these studies suggest that CD103 in human T cells is

upregulated by TGF β produced within the epidermal compartment. Although adhesive interactions with keratinocytes do not induce CD103 upregulation, CD103 likely participates in tethering T cells within the epidermal compartment.

In summary, we find that human skin is populated by a combination of CD103⁺ and CD103⁻ T_{RM}, with CD4⁺ T_{RM} predominating. The preponderance of CD4⁺ T_{RM} in human skin may reflect the diversity of antigen exposures encountered by individuals throughout their lifetimes. Most mouse models of T_{RM} generation utilize viruses to induce T_{RM} formation, including HSV and vaccinia infection, and these mice are kept in pathogen free barrier conditions to prevent other infections (5, 6). In contrast, our healthy adult human skin donors will have been exposed to a combination of viral, bacterial, fungal and possibly parasitic infections. It is worth noting that the antigen specificities of human skin T_{RM} are a valuable immunologic record of what infectious pathogens and antigens an individual has been exposed to during his or her lifetime. Moreover, encoded within the cytokine and effector functions of these T cells is a history of how these individuals overcame each pathogen. There are currently no techniques that allow us to extrapolate antigen specificity from the sequences of the TCR receptor. However, once such techniques are developed, we will be able to investigate the immunologic history of an individual from the T cells remaining in their skin. Recent studies in mice showed that skin antigen exposures led individual naïve T cells to differentiate into roughly equal numbers of skin resident T_{RM} and recirculating T_{CM}, suggesting that immune reactions in skin generate TCR identical resident and recirculating T cell subsets (30).

We identified recirculating T cell subsets by studying the populations of T cells that were depleted from skin by alemtuzumab. Our prior work in patients with leukemic CTCL (L-CTCL) treated with alemtuzumab found that human skin and the skin of CTCL patients contained a population of T cells that coexpressed both skin homing addressins (CLA/CC4) and central memory markers (CCR7/L-selectin) (2). In fact, in many patients with L-CTCL, malignant T cells had this phenotype (23). Alemtuzumab depleted these T cells from blood and purged them from the skin over time as well, confirming that they recirculate. T_{CM}, as first described, were thought to be a subset of T cells that recirculated between the blood and lymph nodes and lacked tissue tropism (20). However, skin tropic T_{CM} exist in both the skin and blood in humans and would be expected to recirculate actively between the lymph nodes, blood and the skin. Indeed, a similar population of recirculating T cells was identified in the skin of Kaede mice (17). These T cells expressed co-expressed CCR7 and L-selectin at intermediate levels and recirculated between the blood, skin, and both skin draining and distant lymph nodes. These T cells had quantitatively lower levels of L-selectin expression than T cells from the circulation but expression was sufficient to allow these cells to recirculate through noncutaneous, distal lymph nodes (17). Because these T cells could access systemic lymph nodes, we believe they represent the analogous population in mice to cells we refer to as skin tropic T_{CM}.

We also found evidence for an additional population of CCR7⁺/L-selectin⁻ recirculating T cells in human skin, which we refer to as migratory memory T cells (T_{MM}). This population of recirculating T cells was poorly reproduced in our engrafted mouse model but was present in the blood and skin of healthy individuals and CTCL patients and was depleted by

alemtuzumab therapy. In humans, CLA⁺ T_{MM} were the most prevalent subset of CLA⁺ memory T cells in the blood. T_{MM} would presumably be excluded from lymph nodes draining noncutaneous tissues because they lacked L-selectin. However, they would access skin draining lymph nodes as they migrated out of the skin into the lymphatics and from there into skin draining lymph nodes. We found support for this concept in the clinical findings that patients with CTCL and T_{MM} malignant T cells tended to have enlargement of only skin draining lymph nodes (termed dermatopathic nodes) but lacked diffuse lymph node involvement, a feature often observed in L-CTCL patients with T_{MM} malignancies.

With respect to their functional capacities, T_{MM} were intermediate between T_{CM} and T_{RM} in their production of inflammatory cytokines. Interestingly, we found that Th2 cytokine production was enriched in recirculating T cells (T_{MM} and T_{CM}) but that the production of Th1, Th2, Th17 and Th22 associated inflammatory cytokines was enriched in effector memory T cells (T_{EM}). IL-2 was expressed at higher levels by recirculating T cells, consistent with a possible role for this cytokine in preventing apoptosis and maintaining cell viability during their turbulent and continual migration through the bloodstream and tissues.

Although the data we have are limited, our findings in human patients treated with alemtuzumab suggest T_{MM} may recirculate more slowly through skin than T_{CM}. CTCL Patients with malignant T cells of a T_{MM} phenotype developed ill-defined but discrete skin lesions, whereas L-CTCL patients with malignant T cells of a T_{CM} phenotype had diffuse, often complete erythema of the skin (erythroderma). We hypothesize that the diffuse erythema caused by malignant T_{CM} arises from the rapid migration of these cells in and out of skin and also within the skin itself. In contrast, T_{MM} gave rise to ill-defined but discrete skin lesions, consistent with a slower migration of these T cells in and out of skin and within the skin itself. Moreover, in the one patient we were capable of studying, T_{MM} were depleted more slowly from the skin than T_{CM}. However, an alternative explanation may be that T_{MM} may be the subset of T cells that give rise to T_{RM}. In this case, the skin lesions we see developing in these patients may represent T_{MM} differentiating in situ into T_{RM} at the sites. The continued development of new expanding skin lesions in these patients likely results from recirculating T_{MM} entering new areas of skin from the blood stream and initiating new lesions. Whether the skin lesions generated by T_{RM} represent newly generated T_{RM} or simply a slower migration of T_{MM} through skin remains to be determined. Lastly, it is not clear why T_{MM} are underrepresented in human engrafted mice. One possibility is that T_{MM} may be a major source of T_{RM} and are therefore depleted as T_{RM} develop in the skin. A second possibility is that the subset of human T cells may not survive well within the mouse microenvironment.

One caveat to our findings is that although we and other groups studying resident and recirculating T cell populations use designations such as T_{RM} and T_{CM} to describe subsets of T cells, markers such as CD69 and CCR7 that are used to define these putatively stable subsets can change with T cell activation. Additional work will be needed to identify exactly how stable these T cell subsets are and to identify additional, less variable surface markers to identify them.

In summary, we present a thorough characterization of resident and recirculating memory T cells in human skin. We find that two discrete populations of T_{RM} are stationed in human skin and two distinct populations of T cells recirculate through skin. Each of these subsets has distinct recirculation and functional capacities and each is likely to play a unique role in protection against known pathogens. Both T_{RM} and recirculating T cells are known to contribute to human diseases. For example, psoriasis is a disorder of dysregulated T_{RM} and L-CTCL represents a malignancy of recirculating T cells (23, 31–33). Recent findings in mice that T_{RM} and recirculating T cells have very different patterns of gene expression highlight the fact that these cell types have a very distinct biology (15). Excepting the current study, the relative proportions of resident and recirculating T cells in human tissues remain uncharacterized. Moreover, the contribution of T_{RM} vs. recirculating T cells to most human inflammatory and autoimmune diseases and the effects of anti-inflammatory therapies on these T cell subsets are unknown. A better understanding of these resident and recirculating T cell subsets is therefore critical to understand healthy immune responses and identify therapies that preferentially target pathogenic T cells in autoimmune and inflammatory diseases.

Materials and Methods

Study design / Experimental design

This is an experimental laboratory study performed on human tissue samples. All studies were performed in accordance with the Declaration of Helsinki. Blood from healthy individuals was obtained after leukapheresis, skin was obtained from healthy patients undergoing cosmetic surgery procedures. Blood and lesional skin from patients with CTCL were obtained from patients seen at the Dana-Farber/Brigham and Women's Cancer Center Cutaneous Lymphoma Program. L-CTCL and MF patients described in this manuscript met the WHO-EORTC criteria for L-CTCL/SS or MF (34). All tissues were collected with prior approval from the Partners and Dana Farber Institutional Review Board. Enumeration of T cell subsets in immunofluorescence studies was done in an investigator blinded fashion. Mechanistic studies on cells derived from blood and human tissues were performed using in vitro assays without blinding or randomization. Study components were not predefined.

Human engrafted mouse model

Human neonatal foreskins were grafted onto the backs of 6- to 8-week-old NOD/SCID/IL-2 receptor γ chain^{null} mice (Jackson Laboratories). One week later 5×10^6 allogeneic PBMC were injected i.v. after isolation from healthy donors by ficoll centrifugation. Antibody injections commenced 2–4 weeks after PBMC engraftment. Alemtuzumab (2.5 μ g) or isotype control (human IgG1, Biolegend) was injected i.p. 3 times/week for 3 weeks. Anti-CD3-diphtheria toxin (A-dmDT(390)-bisFv(UCHT1), 0.625 μ g) or isotype control (human IgG1) was injected i.v. once/day for 4 days. At the conclusion of the experiment, blood and skin grafts were harvested for analysis by flow cytometry, immunofluorescence and/or hematoxylin/eosin staining.

Skin and blood samples

The protocols of this study were performed in accordance with the Declaration of Helsinki, and were approved by the Institutional Review Board of the Partners Human Research Committee (Partners Research Management). Skin from healthy patients was obtained from patients undergoing cosmetic surgery procedures and blood from healthy individuals was obtained as discarded tissue following leukopheresis. Neonatal foreskins were obtained from infants undergoing circumcision at the Brigham and Women's Hospital. Blood and lesional skin from patients with CTCL were obtained from patients seen at the Dana-Farber/Brigham and Women's Cancer Center Cutaneous Lymphoma Program. L-CTCL and MF patients described in this manuscript met the WHO-EORTC criteria for CTCL. PBMC were isolated by ficoll centrifugation.

T cell isolation from skin

Full thickness human skin samples, fresh or after engraftment on mice, were extensively minced and then incubated for 2 hours at 37°C in RPMI 1640 containing 0.2% collagenase type I (Life Technologies) and deoxyribonuclease I (30 Kunitz units/ml, Sigma Aldrich). Thereafter, cells were collected by filtering the collagenase-treated tissue through a 40- μ m cell strainer. For some experiments, epidermis and dermis were separated by overnight incubation in dispase II (1.4 IU/ml, Roche Diagnostics) at 4°C followed by mechanical separation. Epidermis was then minced and incubated with dispase for an additional 30 min at 37°C with agitation followed by filtration through a 40- μ m cell strainer. Dermis was minced and treated with collagenase as per full thickness skin protocol. Cells from whole skin, dermis or epidermis were incubated at 37°C for 2 days in Iscove's DMEM supplemented with 20% fetal bovine serum, penicillin/streptomycin, glutamine and 2-mercaptoethanol prior to flow cytometry analysis. When indicated (Fig. S2), the pan-caspase inhibitor Z-VAD-FMK (R&D Systems) was included at low (10 μ M) or high (100 μ M) concentrations to inhibit apoptosis.

Flow cytometry

Analysis of T cells was performed using directly conjugated monoclonal antibodies obtained from BD Biosciences, eBioscience, Biolegend, Beckman Coulter or R&D Systems. Isotype-matched negative control antibodies were used to set the gates for positive staining. For analysis of cytokine production, T cells were stimulated with either control medium or 50 ng/ml PMA (Sigma Aldrich) and 750 ng/ml ionomycin (Life Technologies) plus 10 μ g/ml Brefeldin A (BD Biosciences) for four hours. Cells were surface stained, fixed, permeabilized, stained with anti-cytokine antibodies, and examined by flow cytometry. For analysis of proliferation, cells were surface stained, fixed, permeabilized, stained with anti-Ki-67 antibody, and examined by flow cytometry. Analysis of flow cytometry samples was performed on Becton Dickinson FACSCanto instruments and data were analyzed using FACSDiva software (V5.1).

Cryosection immunostaining

Human skin grafts were embedded in OCT, frozen, and stored at -80°C until use. 5- μ m cryosections were cut, air dried, fixed for 5 min in acetone, rehydrated in PBS, and blocked

with 20 µg/ml of human IgG (Jackson ImmunoResearch Laboratories) for 15 min at room temperature. Sections were incubated with primary antibody for 30 min, and then rinsed three times in PBS/1% BSA for 5 min. If necessary, secondary antibody was added for 30 min, followed by three rinses. Sections were mounted using Prolong Gold Antifade with DAPI (Life Technologies) and examined immediately by immunofluorescence microscopy. Antibodies were obtained from the following: Biolegend (CD3, CD69), LifeSpan BioSciences (CD103), Abcam (cytokeratin 15) and Jackson Immunoresearch (secondary antibodies). Sections were photographed using a microscope (Eclipse 6600; Nikon) equipped with a 40×/0.75 objective lens (Plan Fluor; Nikon). Images were captured with a camera (SPOT RT model 2.3.1; Diagnostic Instruments) and were acquired with SPOT 4.0.9 software (Diagnostic Instruments).

Proliferation and cytokine analysis of cells stimulated within explants

Ten mm punch biopsies of healthy adult skin were injected with 100 µl of PBS (control), α-CD3/α-CD28 antibody mixture (0.5 µg/ml each), or heat-killed *Staphylococcus aureus* (10⁶/ml, ATCC) or *Candida albicans* (5×10³/ml, ATCC). Biopsies were floated in 0.4 µm pore size Transwell inserts in X-VIVO 15 medium (Lonza). After 2 weeks cell that had migrated out of the biopsies were collected, rested overnight in X-VIVO 15 medium, and analyzed for cytokine production and proliferation by flow cytometry as described above. Only cells that had migrated out of the explants were analyzed.

In vitro studies of CD103 regulation

To obtain keratinocytes, healthy human skin was separated into epidermis and dermis by incubating with 1.4IU /ml of dispase II (Roche) in RPMI (Life Technologies) at 4°C for 12 hrs. Epidermis was peeled away from the dermis, minced and further digested with 1.4IU/ml dispase at 37°C for 30 min, then filtered in a cell strainer. Keratinocytes were cultured to confluence in 24-well flat-bottomed plates using keratinocyte-FCM media (Life Technologies). To obtain fibroblasts, the dermis was minced, digested with 0.2 % collagenase/DNase (Sigma Aldrich) at 37°C for 2 hours, then filtered in a cell strainer. Fibroblasts were cultured to confluence in RPMI supplemented with 15% FCS and 10 ng/ml EGF. CD3⁺CD4⁺ and CD3⁺CD8⁺ T cells were isolated using magnetic separation (negative isolation kits, Miltenyi Biotec) from healthy human PBMC. CD103 was expressed by a mean 3% of the isolated CD8 T cells but otherwise surface expression of CD69 and CD103 was <1% in these T cell fractions. 1×10⁵ T cells were stimulated with αCD3/CD2/CD28 T cell activation beads (Miltenyi Biotec) with 1:2 ratio for 24 hrs (day 0) and then added to keratinocytes or fibroblasts monolayers (day 1) in Iscove's Modified Dulbecco's Media (Corning) supplemented with 10% FCS and L-glutamine. For indirect cultures, transwells with 0.4 µm polyester membrane (Corning) were used to separate T cells from fibroblasts and keratinocyte monolayers. Neutralizing abs for TGFβ (R & D), E-Cadherin (Millipore), LFA-1 (Biolegend) and Integrinβ1 (R & D) and recombinant human TGFβ (R&D) were added to co-cultures system with final concentration of 10 µg/ml. T cells were collected by washing on day 6 after starting co-culture. Adherent T cells were counted by microscopy and collected by incubating the wells with trypsin/EDTA (Gemini Bio-Products) for 5 min at 37°C.

CyTOF mass cytometry analysis of peripheral blood T cells

T cells were isolated from healthy donor PBMC by depletion of non-T cells using the Pan T isolation kit and AutoMacs separator (Miltenyi Biotec). T cells were pre-treated with TAPI-2 (100 μ M, Sigma Aldrich) for 1 hr prior to stimulation and during the 4 hr of stimulation with PMA/ionomycin plus brefeldin A to prevent stimulation-induced cleavage of L-selectin (35). Following stimulation, T cells were stained with surface marker metal-conjugated antibodies (MCA, Fluidigm, Inc.) at concentrations found to be effective in prior antibody tests. After surface staining, cells were fixed, permeabilized and stained with MCA intracellular antibodies for 30 min at room temperature. The cells were then labeled overnight at 4°C with 125nM DNA intercalator (Fluidigm, Inc). Finally, the cells were diluted to the appropriate concentration in distilled water containing 10% EQ™ Four Element Calibration Beads (Fluidigm, Inc) to achieve an acquisition rate <500 events/s on the CyTOF2 instrument using the recommended default settings for the CyTOF2 (Fluidigm, Inc). CyTOF data were normalized to the beads through the CyTOF2 software after acquisition without bead removal and using default settings (Fluidigm, Inc). After built-in cell-identifying software created a FCS file, CyTOF data were analyzed in a manner similar to standard flow cytometry analysis by FCS express software 4.0 (De Novo Software) using DNA content and normalization beads to identify single-cell events.

Statistical analyses

Primary methods of data analysis included descriptive statistics (means, medians and s.d.). Differences between two sample groups were detected using the one tailed Wilcoxon–Mann–Whitney test, $\alpha=0.05$. For comparisons of multiple groups, a Kruskal–Wallis one-way analysis of variance with a Bonferroni–Dunn’s post test for multiple means test was used, $\alpha=0.05$.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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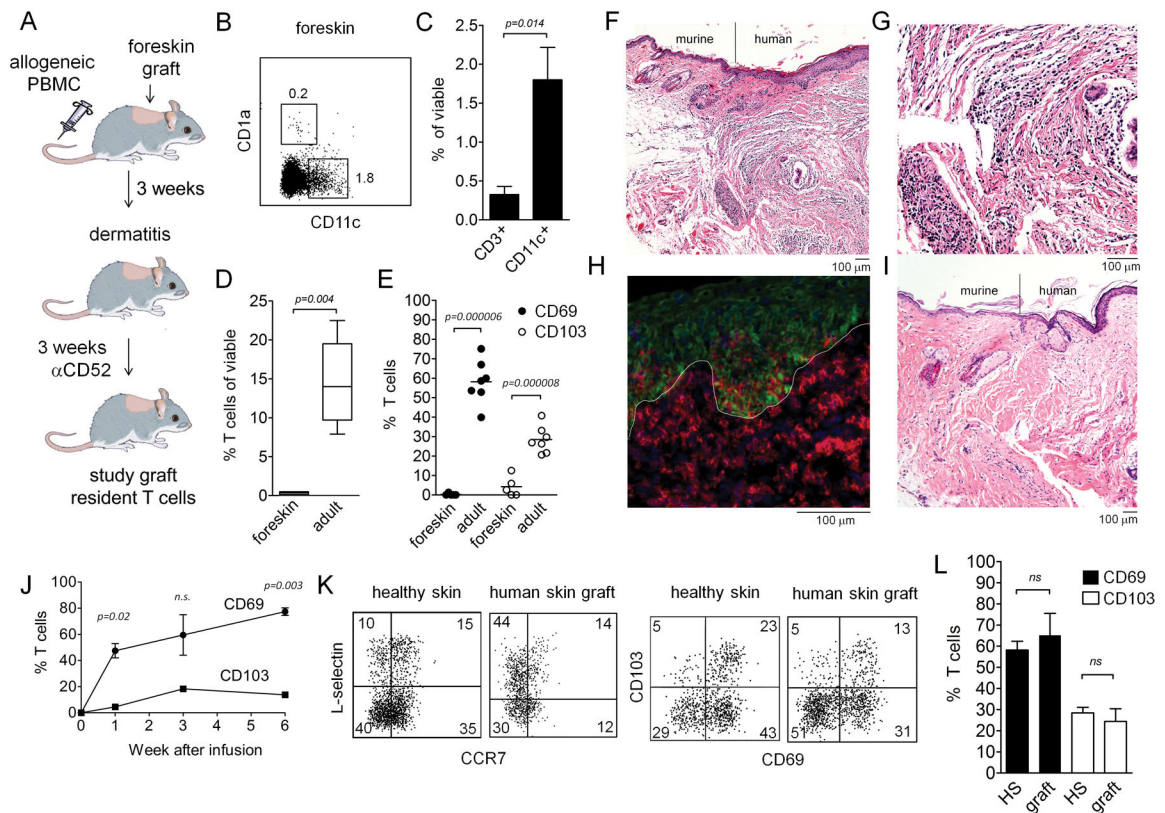


Figure 1. Skin T cells in a human engrafted mouse model recapitulate T cell populations in adult human skin

(A) A human engrafted mouse model was used to discriminate resident from recirculating T cells in skin. NSG mice were grafted with human foreskin and injected i.v. with allogeneic human PBMC, and alemtuzumab (α CD52) was used to deplete recirculating human T cells from grafted human skin. (B–E) human foreskins contained resident APC but few T cells. (B) Flow cytometry staining of collagenase digested foreskin demonstrated both CD11c⁺ DC and CD1a⁺ LC were present in foreskin. (C) Approximately four-fold more DC than T cells were present in foreskin, as assayed by the % viable cells of each cell type in collagenase digested foreskin. (D) T cells were 45-fold more frequent in healthy adult human skin than in foreskins; the percentage of CD3⁺ T cells in collagenase digested foreskin versus adult skin is shown. (E) The few T cells present in human foreskin lacked markers characteristic of T_{RM}. The percent of T cells expressing each marker in collagenase treated foreskin and adult skin are shown. The mean and SEM of four foreskins (C–E) and seven adult skin donors (D,E) are shown (test: T-test). (F–H) allogeneic PBMC injected i.v. into grafted mice migrated specifically into the human skin graft and induced an inflammatory dermatitis. Images are representative of eight PBMC injected and eight saline injected mice. (F) H&E stained section demonstrating the junction between mouse and human skin in grafted mice. (G) H&E stain, 20X view of inflammatory dermatitis. (H) A cryosections of human grafted skin immunostained for CD3 (red) and keratin (green) are shown. T cells were evident in both the dermis and the epidermis. The dermoepidermal junction is indicated by a white line. (I) in the absence of infused allogeneic PBMC, no

inflammatory infiltrate was observed and few or no T cells were visualized in skin. **(J)** CD69 up-regulation preceded CD103 up-regulation in T cells migrating into human skin grafts. Human skin grafts were harvested at the indicated time points, collagenase digested, and T cells were analyzed for CD69 and CD103 expression by flow cytometry. The mean and SEM of two grafted mice per time point are shown (test: T-test). **(K,L)** T cell populations in grafted skin recapitulate those observed in adult human skin. T cell expression of T_{CM} (CCR7/L-selectin) and T_{RM} (CD69/CD103) markers as assayed by flow cytometry are shown. T cells were isolated by collagenase digestion from healthy adult human skin or from skin grafts on PBMC infused mice. Representative histograms **(K)** and mean and SEM of expression of CD69 and CD103 are shown. The mean and SEM of three grafts and seven healthy skin donors are shown (test: T-test).

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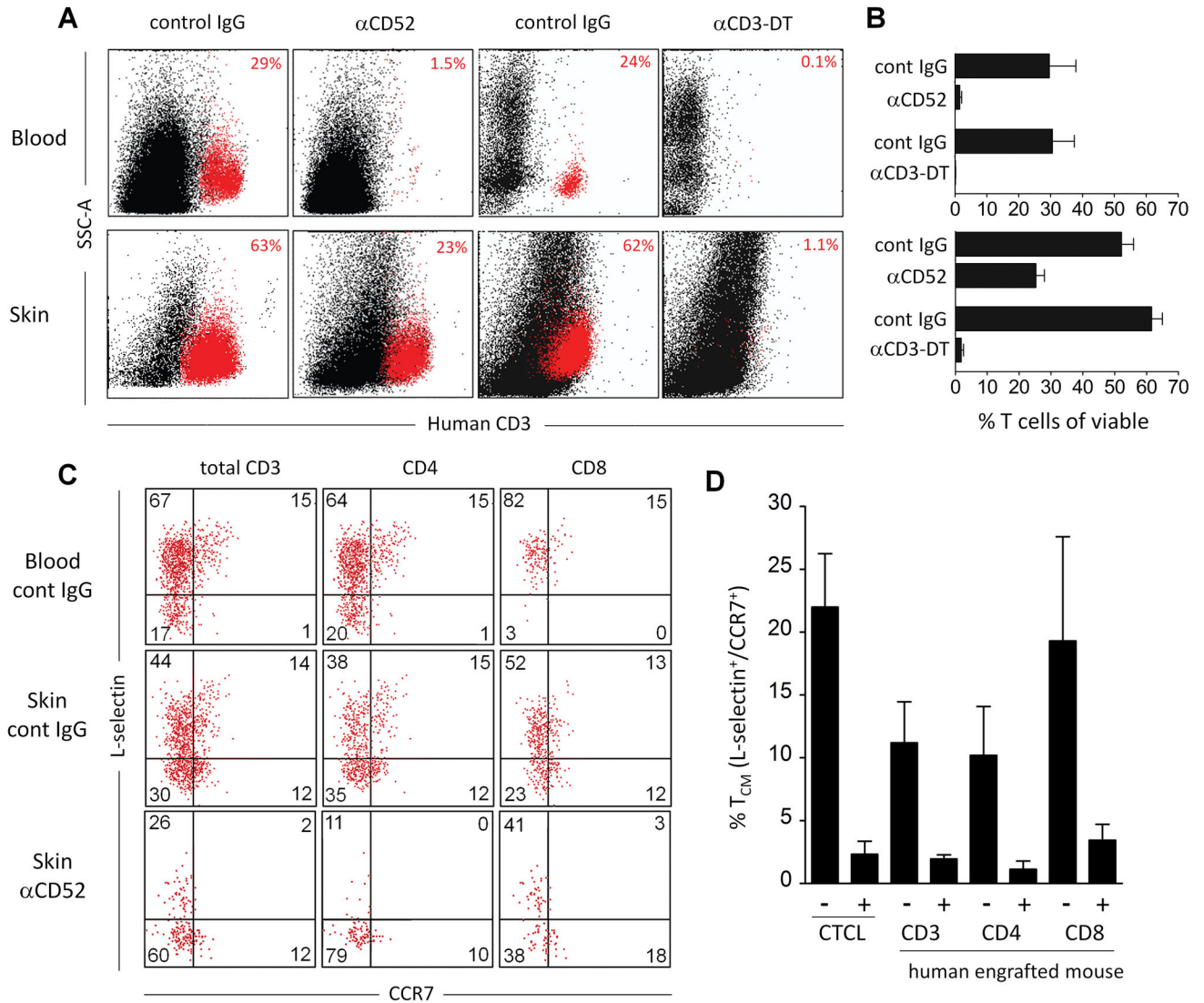


Figure 2. Alemtuzumab depletes recirculating T_{CM} from skin in human engrafted mice but spares a population of skin T cells

(A, B) Shown are human T cells isolated from peripheral blood (Blood) and collagenase digested foreskin grafts (Skin) stained for CD3 and analyzed by flow cytometry. Samples were obtained from human engrafted mice after treatment with either control IgG, alemtuzumab (α CD52) or with the anti-CD3-diphtheria immunotoxin A-dmDT(390)-bisFv(UCHT1) (α CD3-DT). Representative dot plots (A) and aggregate data (B) are shown from four control IgG (alemtuzumab control) injected mice, six alemtuzumab injected mice, two control IgG (α CD3-DT control) injected mice and two α CD3-DT treated mice. (C) $CCR7^+/L\text{-selectin}^+$ T_{CM} were present in the blood and skin of human engrafted mice and were depleted in the skin by alemtuzumab. T cells isolated from the blood and skin of control IgG treated mice and the skin of alemtuzumab treated (α CD52) human engrafted mice are shown. (D) Aggregate data showing depletion of $CCR7^+/L\text{-selectin}^+$ T_{CM} in skin grafts of human engrafted mice treated with control IgG (-) or alemtuzumab (+). Depletion of T_{CM} among the $CD3^+$, $CD4^+$ and $CD8^+$ T cell populations are shown. The mean and

SEM of three grafted mice per group are shown. For comparison, the numbers of CCR7⁺/L-selectin⁺ T_{CM} in the skin of five human CTCL patients (CTCL) before (-) and after (+) alemtuzumab therapy are shown.

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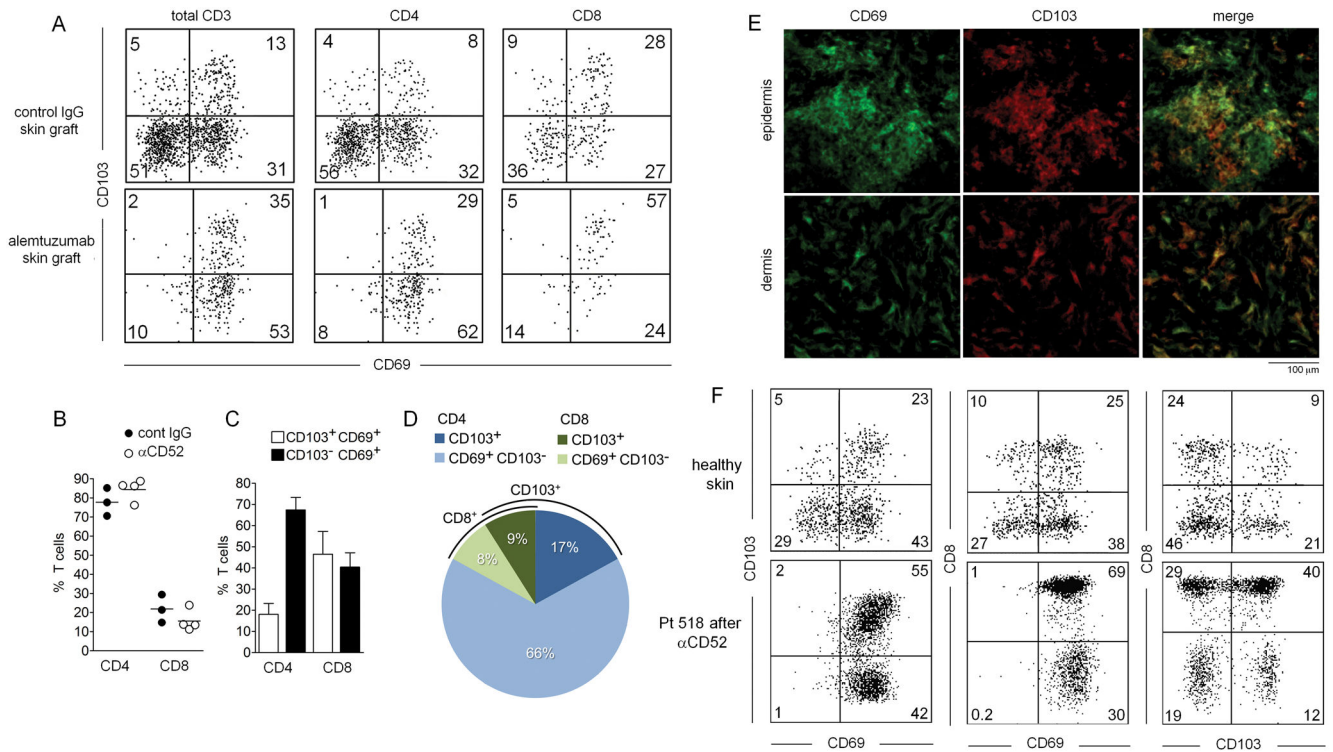


Figure 3. Distinct populations of non-recirculating CD103⁺ and CD103⁻ T_{RM} exist in human skin engrafted mice and in human skin

(A) T_{RM} cells remaining in the human skin grafts of mice after alemtuzumab treatment expressed CD69 and a subset coexpressed CD103. (B) The relative percentages of CD4⁺ and CD8⁺ T cells in human skin grafts did not change appreciably after alemtuzumab treatment. Shown are the percentage of total CD4⁺ and CD8⁺ T cells in three control IgG (cont IgG) and four alemtuzumab (αCD52) treated mice (test: T-test). (C) Most CD4⁺ T_{RM} cells were CD69⁺/CD103⁻ whereas the majority of CD8⁺ T cells expressed both CD69 and CD103. (D) CD4⁺/CD69⁺/CD103⁻ T cells were the most frequent T_{RM} population in alemtuzumab treated human engrafted mice. The relative percentages of the indicated T_{RM} subsets are shown. (E) CD103⁺ and CD103⁻ T_{RM} were found in both the epidermis and the dermis of human engrafted mice after alemtuzumab treatment. (F) Confirmatory studies in human patients demonstrated that CD103⁺ and CD103⁻ T_{RM} also exist in human skin following treatment with alemtuzumab. CD3⁺ T cells isolated from healthy skin (top row of histograms) and the skin of a CTCL patient after alemtuzumab treatment (bottom row of histograms) are shown. Similar to the findings in human engrafted mice, CD69 was expressed by virtually all T_{RM}, CD4⁺ and CD8⁺ T_{RM} of both subsets were observed, and CD4⁺ T cells were more frequently CD103⁻ whereas CD8⁺ T_{RM} were more frequently CD103⁺. Results are representative of data from three treated patients.

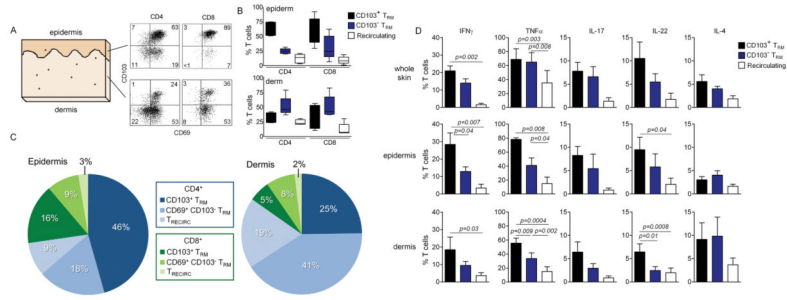


Figure 4. Two phenotypically and functionally distinct populations of TRM exist in healthy adult human skin

(A,B, C) Adult human skin was enzymatically separated into epidermis and dermis, T cells were then isolated and analyzed by flow cytometry for the expression of TRM markers CD69 and CD103. Representative histograms (A) and aggregate data from 5 donors (B, C) are shown. The percentages of CD69⁺/CD103⁺ (CD103⁺ TRM), CD69⁺/CD103⁻ (CD103⁻ TRM) and CD69⁻/CD103⁻ (Recirculating) T cells as a percentage of the separate CD4⁺ and CD8⁺ subsets are shown in (B) and proportions of each in CD3⁺ T cells as a whole (C) are shown. CD103⁺ TRM, both CD4⁺ and CD8⁺, were most frequent in the epidermis whereas CD103⁻ TRM, in both CD4⁺ and CD8⁺ T cells, were more frequent in the dermis. Recirculating T cells were the minority among both CD4⁺ and CD8⁺ T cell populations in skin. (D) CD103⁺ TRM were superior to other T cell subsets in the production of effector cytokines. Shown are the relative production of the indicated effector cytokines by CD69⁺/CD103⁺ (CD103⁺ TRM), CD69⁺/CD103⁻ (CD103⁻ TRM) and CD69⁻/CD103⁻ (Recirculating) T cells isolated from healthy human skin. T cell subsets were isolated from either non-fractionated skin (whole skin), the epidermis alone and the dermis alone. The mean and SEM of analyses from three donors (whole skin), and five donors (epidermis and dermis) are shown (test:one way ANOVA).

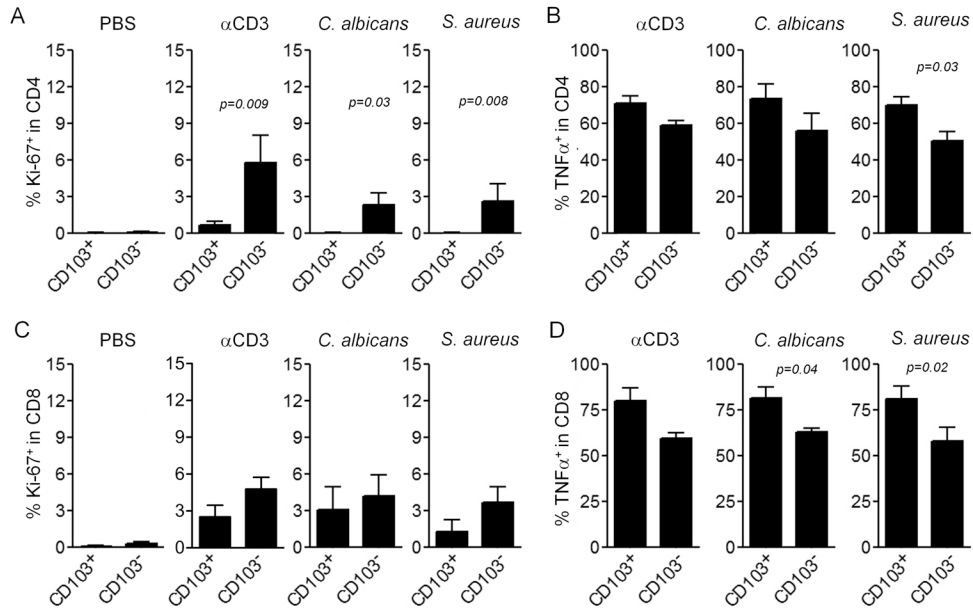


Figure 5. CD103⁺ T_{RM} have a lower proliferative capacity but increased effector function
 Skin explants were injected with PBS, stimulatory anti-CD3 and anti-CD28 antibodies (αCD3), or heat killed extracts of *C. albicans* and *S. aureus*. Two weeks later, T cells that had migrated out of the skin were collected and the proliferation of CD4⁺ T_{RM} (A) and CD8⁺ T_{RM} (C) were assayed by staining for Ki-67 and flow cytometry analysis. TNFα production was assayed by intracellular cytokine staining and flow cytometry analysis after stimulation with PMA and ionomycin. Results for CD4⁺ T_{RM} (B) and CD8⁺ T_{RM} (D) are shown. Results are shown for CD69⁺/CD103⁺ (CD103⁺ T_{RM}), and CD69⁺/CD103⁻ (CD103⁻ T_{RM}). The mean and SEM of three donors (PBS) and six donors (all other conditions) are shown (test: T-test).

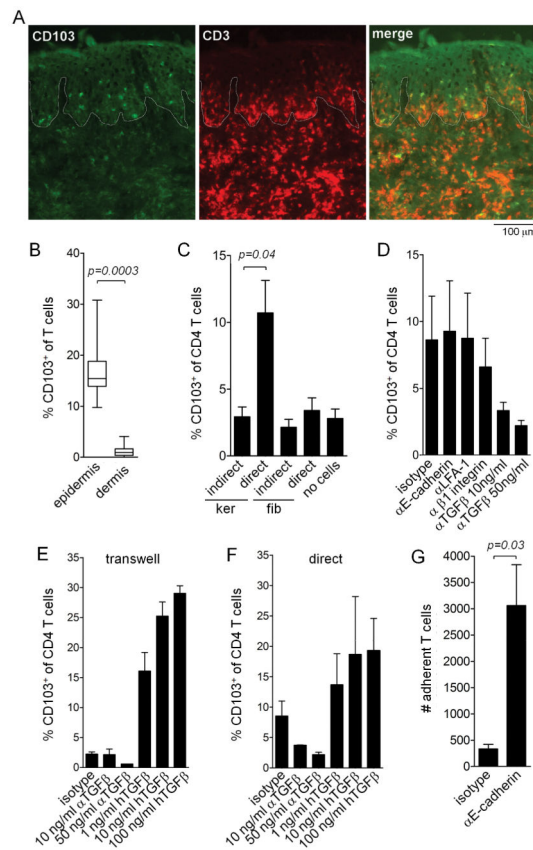


Figure 6. T cell CD103 induction is enhanced by keratinocyte contact, depends on TGF β and is independent of T cell keratinocyte adhesive interactions

(A,B) In human engrafted mice, T cells that migrating into the epidermis had a higher rate of CD103 up-regulation than dermal T cells. (A) Representative immunostains of human foreskin skin grafts 3 weeks after IV injection of CD103⁻ PBMC are shown. CD103⁺ T cells were primarily localized to the epidermal compartment. The dermal-epidermal junction is highlighted by a white line. (B) The mean and SEM of 8 separate experiments are shown (test: T-test). (C) Keratinocyte contact induced up regulation of CD103 in CD4⁺ T cells from human blood stimulated with α CD3/ α CD2/ α CD28 beads. CD4⁺ T cells were isolated from human peripheral blood by magnetic bead separation and then co-cultured in direct contact (direct) with confluent monolayers of human keratinocytes (ker) or fibroblasts (fib) or in transwells separated from keratinocyte or fibroblasts monolayers (indirect) for one week in the presence of α CD3/ α CD2/ α CD28 beads. CD103 expression was then assessed by immunostaining and flow cytometry analysis. The mean and SEM of four separate experiments are shown (test: ANOVA). (D). Blocking of adhesive interactions with keratinocytes had no effect on CD103 upregulation but neutralization of TGF β did decrease CD103 induction. CD4⁺ T cells were cultured on confluent human keratinocyte monolayers for one week in the presence of α CD3/ α CD2/ α CD28 beads and the indicated control and function blocking antibodies. The mean and SEM of four experiments are shown (test: ANOVA). (E,F) TGF β neutralizing antibodies decreased and exogenous TGF β increased CD103 induction. T cells were cultured in transwells across from keratinocyte monolayers (E) or indirect contact with keratinocyte monolayers (F) for one week in the presence of

α CD3/ α CD2/ α CD28 beads and the indicated neutralizing TGF β antibodies (α TGF β) or recombinant human TGF β (hTGF β). the mean and SEM of 2 separate experiments are shown (test: ANOVA). (G) Function blocking E-cadherin antibodies inhibited T cell adhesion to keratinocyte monolayers. CD4⁺ T cells were cultured on confluent human keratinocyte monolayers for one week in the presence of α CD3/ α CD2/ α CD28 beads in the presence of isotype control or E-cadherin function blocking antibodies. Nonadherent T cells were removed by rinsing and adherent T cells were then eluted from the wells by treatment with trypsin/EDTA. The mean and SEM of the number of adherent cells per well from three separate experiments are shown (test:T-test).

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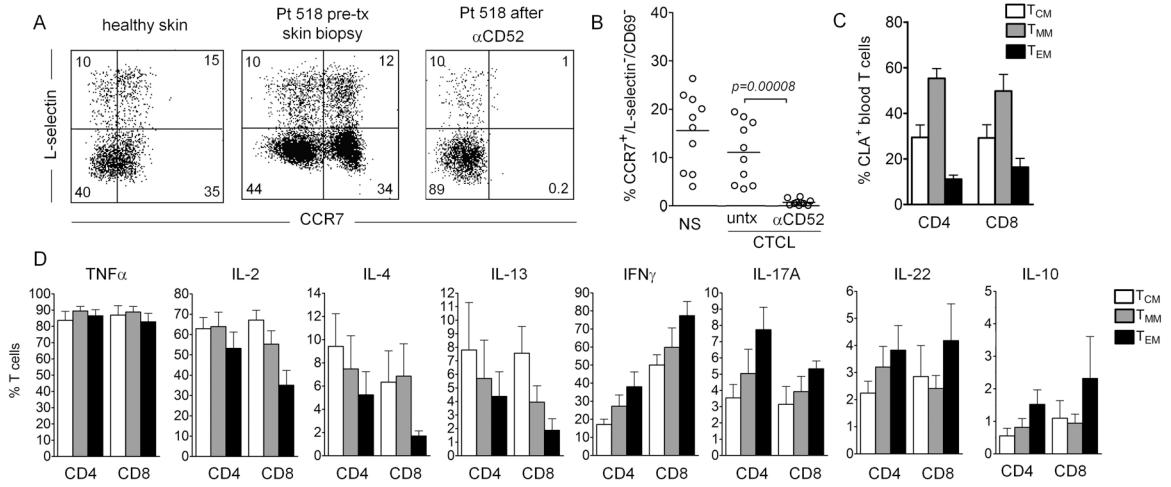


Figure 7. CCR7⁺/L-selectin⁻ T cells make up a second distinct population of recirculating T cells in human skin

(A) CCR7⁺/L-selectin⁻ T cells were present in healthy human skin and the skin of CTCL patients and were depleted by alemtuzumab therapy. (B) The proportion of T_{MM} (CCR7⁺/L-selectin⁻/CD69⁻ T cells) in 10 healthy skin donors (NS) and ten CTCL patients before (untx) and after (αCD52) alemtuzumab therapy are shown (test: T-test). (C) T_{MM} are the most frequent CLA⁺ memory T cell population in human blood. The relative percentages of CCR7⁺/L-selectin⁺ central memory (T_{CM}), CCR7⁺/L-selectin⁻ migratory memory (T_{MM}), and CCR7⁻/L-selectin⁻ effector memory (T_{EM}) T cells in the CD4⁺ and CD8⁺ T cells from human peripheral blood are shown. The mean and SEM of four donors are shown. Representative histograms are included in Fig. S4. (D) The effector functions of circulating T_{MM} are intermediate between those of T_{CM} and T_{EM}. For panels, C and D, T cells were isolated from peripheral blood, stimulated with PMA and ionomycin in the presence of an L-selectin shedding inhibitor, stained for surface markers and intracellularly for cytokine production and then analyzed by CyTOF mass spectrometry. The mean and SEM of four donors are shown (test: ANOVA).

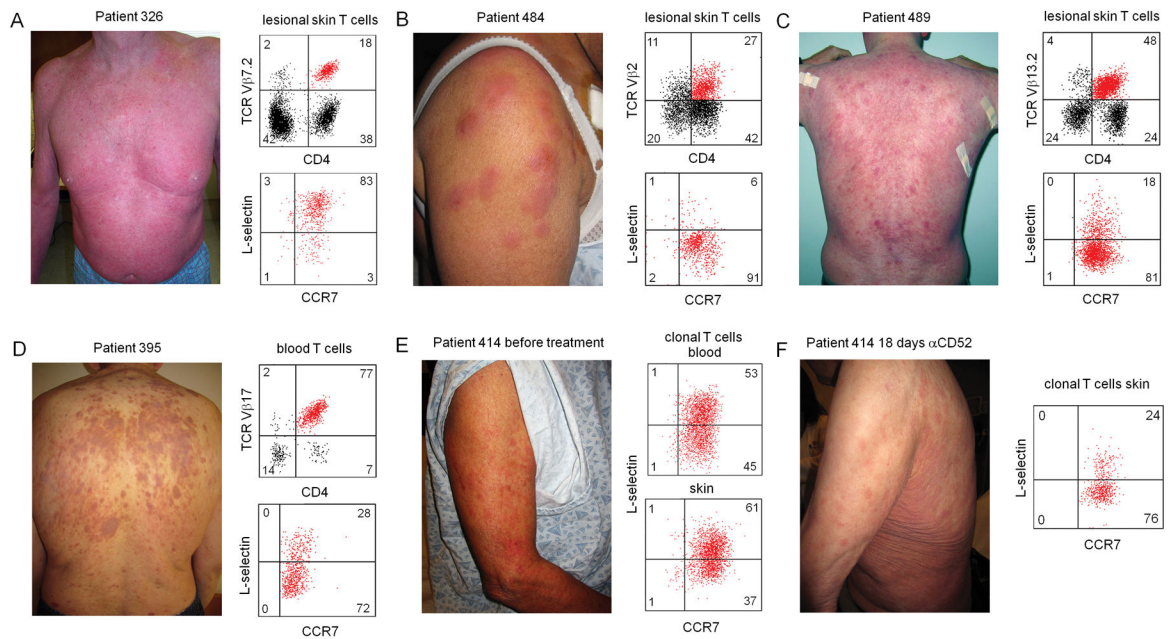


Figure 8. In human patients with CTCL, malignant T cells with a T_{MM} phenotype were associated with expanding skin lesions with ill-defined borders and were depleted more slowly from skin than T_{CM} after alemtuzumab therapy

(A) CTCL patients with malignant T cells of a T_{CM} ($CCR7^+/L\text{-selectin}^+$) phenotype had diffuse erythema of the skin. In contrast, CTCL patients in whom the malignant T cells have a T_{MM} phenotype ($CCR7^+/L\text{-selectin}^-$) in lesional skin (B,C) or blood (D), presented with expanding discrete skin lesions with ill-defined borders. (E) A patient who had malignant T cells of both T_{CM} and T_{MM} phenotype had both diffuse erythema on presentation as well as discrete skin lesions with ill-defined borders. The patient was placed on alemtuzumab therapy and the skin was re-biopsied 18 days later. (F) 18 days after alemtuzumab initiation, the discrete erythema had receded, leaving the ill-defined skin lesions. Analysis of T cells from the skin demonstrated a relative depletion of T_{CM} but persistence of T_{MM} in skin.