



Resident Memory-Like Tumor-Infiltrating Lymphocytes (TIL_{BM}): Latest Players in the Immuno-Oncology Repertoire

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Resident memory T cells (T_{RM}) are a recently identified subset of long-lived memory T cells that are characterized in terms of their unique surface phenotype combined with a non-recirculating pattern of localization to non-lymphoid, peripheral tissues. T_{BM} have quickly become a key area of focus in understanding immune responses to microbial infection in so-called "barrier" tissues, and appear to be particularly critical for protection against repeat exposure at the same site. More recently, tumor-infiltrating T cells with canonical T_{BM} features are being identified in human cancers, in particular cancers of epithelial origin, and their presence is broadly found to be associated with favorable longterm prognosis. Moreover, recent studies have shown that these "resident memory-like" tumor-infiltrating lymphocytes (referred to herein as TIL_{RM}) are uniquely activated in melanoma patients undergoing PD-1 directed checkpoint blockade therapy. Accordingly, there is much interest at present regarding the biology of these cells and their precise role in anti-cancer immunity. Herein, we review the current state of the literature regarding TIL_{RM} with a specific emphasis on their specificity, origins, and relationship to conventional pathogen-specific T_{BM} and speculate upon the way(s) in which they might contribute to improved prognosis for cancer patients. We discuss the growing body of evidence that suggests TIL_{RM} may represent a population of bona-fide tumor-reactive T cells and the attractive possibility of leveraging this cell population for future immunotherapy.

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BRIEF INTRODUCTION TO RESIDENT MEMORY T CELLS (T_{BM}) AND THE T_{RM}-DEFINING SURFACE MARKER CD103

In recent years, there has been growing recognition of the importance of a peripheral, nonrecirculating component of the immune system known as T_{RM} [for review see Ref. (1–3)]. T_{RM} have historically been defined by their peripheral tissue localization and lack of circulatory activity. More recently, there is increasing understanding of the unique surface phenotype(s) of T_{RM} and how the specific molecules that comprise this phenotype contribute to their (non-)circulatory nature. Although this phenotype can vary somewhat between tissues, disease states, and CD4 versus CD8 subsets, most T_{RM} in skin, lung, and GI tract typically express CD69, a molecule widely considered to be an indicator of recent activation, but which is also involved in downregulation of the receptor

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for sphingosine 1 phosphate (S1P1), thereby inhibiting the ability of T_{RM} to traffic out from peripheral tissue in response to S1P1 gradients (4). Likewise, T_{RM} frequently lack surface expression of CCR7, preventing them from trafficking in response to gradients of CCL19 and CCL21 (5). In addition, surface expression of CD103 (the $\alpha_{\rm E}$ component of the $\alpha_{\rm E}/\beta_7$ integrin molecule) (6, 7) is now widely considered to be a canonical marker of T_{RM}, and although T_{RM} populations can be comprised of variable proportions of both CD4 and CD8 cells, CD103 appears to be uniquely overexpressed by CD8 T_{RM} (8). CD103 expression is also biologically relevant to the non-recirculating phenotype of T_{RM} , as the ligand for $\alpha_E(CD103)/\beta_7$ integrin is E-cadherin expressed on epithelial cells (9). Although chemokines are thought to be the initial mediator of T cell recruitment into peripheral sites of inflammation, adhesive interactions between $\alpha_{\rm E}(\rm CD103)/\beta_7$ and E-cadherin is thought to be responsible for the long-term "retention" of antigen-specific T_{RM} at relevant sites (10, 11). This phenomenon is particularly well studied in the context of mucosal tissue infection, where the long-term retention of T_{RM} at the site of an initial infection is thought to provide durable and rapid protection against repeat attack by the same organism. Indeed, once T_{RM} populations are established, they can be retained at the original site of infection for months or even years, even in the complete absence of relevant antigen (12-14). This T_{RM} phenomenon can also be exploited by vaccination strategies that involve delivery of vaccine to the relevant mucosal tissue (15). Indeed, the historical field of "mucosal" immunity and the newer field of "T_{RM}-mediated" immunity are rapidly merging in terms of the memory T cell components.

In addition to mediating adhesion and T_{RM} formation, both $\alpha_E(CD103)/\beta_7$ and E-cadherin are also capable of intracellular signaling. For example, the intracellular domain of E-cadherin interacts with β -catenin which in turn interacts with the actin cytoskeleton, affecting cell shape and motility (16). Likewise, cross-linking of surface-expressed $\alpha_E(CD103)/\beta_7$ impacts the shape and motility of lymphocytes (17), enhances T cell proliferation and induces lysis of target cells (18). Thus through the combination of "inside-out" and "outside-in" signals, $\alpha_E(CD103)/\beta_7$ has the potential to profoundly impact T_{RM} effector function, in addition to augmenting peripheral memory formation.

MECHANISM OF CD103 UPREGULATION ON $T_{\mbox{\tiny RM}}$

TGF- β has long been known to play a key role in the regulation of $\alpha_{\rm E}({\rm CD103})/\beta_7$ surface expression on T lymphocytes (19, 20). Although TGF- β is often considered solely as an immunosuppressive factor, it is, in reality, a highly pleiotropic cytokine that is expressed in a multitude of (primarily peripheral) tissue types and has biological activities that are context specific (21). Interestingly, although TGF- β is required for upregulation of $\alpha_{\rm E}({\rm CD103})/\beta_7$ surface expression, TGF- β exposure alone is not sufficient (18, 22). Rather, it is the combination of TGF- β plus concurrent signaling through the TCR that results in dramatic and rapid $\alpha_{\rm E}({\rm CD103})/\beta_7$ expression. Indeed, the combination of these two signals makes perfect sense biologically as it would allow for large numbers of lymphocytes (with diverse specificities) to transiently traffic through TGF- β -rich sites of peripheral infection, but result in the $\alpha_{E}(CD103)/\beta_{7}$ -mediated retention of only those T cells with relevant specificity. This model of T_{RM} formation is supported by the finding that in CD103 knockout mice, numbers of T_{RM} are substantially reduced (10). Likewise, dysregulation of the SMAD signaling pathway downstream of the TGF- β -receptor results in reduced numbers of T_{RM} (23).

Although TGF- β -mediated upregulation of CD103 clearly plays an important role in the establishment of T_{RM}, it is certainly not the only mechanism. For example, it has also been reported that the formation of T_{RM} populations can be enhanced through signaling *via* the homeostatic cytokine, interleukin-15 (IL-15) (24, 25). However, dependency upon IL-15 for T_{RM} formation varies from tissue to tissue (26), implying that the requirement for IL-15 is not absolute and may be more complex than that of TGF- β . Moreover, as described above, CD4⁺ T_{RM} populations, in general, express much lower levels of CD103 than do CD8⁺ T_{RM}, thus they must maintain residency in a CD103-independent manner (27, 28).

TRM IN THE CANCER SETTING

In recent years, there has been growing appreciation that T_{RM} biology/immunology is not unique to the infectious disease setting. Indeed, it has long been speculated that T_{RM} play a key role in both allograft rejection and autoimmunity. For example, $\alpha_{\rm E}({\rm CD103})/\beta_7$ is expressed on the majority of tissue-infiltrating CD8⁺ T cells during transplant rejection (20, 29, 30) and graft versus host disease (22). In CD103-deficient mice, T cells are not able to infiltrate allogeneic islet cell transplants and allografts persist for long periods in vivo (30, 31) often surrounded by a characteristic "halo" of CD103-deficient CD8 T cells. In the autoimmune disease setting, islet infiltrating cells in both human diabetic patients (32, 33) and mouse models of autoimmune diabetes (34) are enriched for $\alpha_E(CD103)/\beta_7$ -expressing T_{RM}. Presumably, in each of these settings T_{RM} are derived *via* the same TGF-ß plus concurrent TCR signaling mechanism described above for infectious diseases.

 $\alpha_{\rm E}({\rm CD103})/\beta_7$ -expressing tumor-infiltrating T cells (TIL) are also now turning up, with increasing regularity, in various cancer settings, particularly in cancers of epithelial origin. This should really not be surprising considering the relationship between TGF- β and α_{E} (CD103)/ β_{7} and the frequent expression of TGF- β in cancers of various types. TGF-ß overexpression in cancer has been broadly considered as an immunosuppressive mechanism of tumor escape from immunological pressure (21, 35). However, an alternate hypothesis could be that TGF- β production by tumors is not so much an acquired trait as it is an amplification of the TGF-β that is expressed as part of the "normal" biology of epithelial tissues. Regardless of the mechanism, when tumor-reactive T cells enter these TGF-\beta-rich environments and then become activated through the TCR, there is full reason to assume they would upregulate $\alpha_{\rm E}(\rm CD103)/\beta_7$ on the cell surface, in the same manner that conventional T_{RM} do.

However, as described above, CD103 expression is only one part the larger phenotypic profile that defines T_{RM} . Whether CD103-expressing TIL are phenotypically identical to conventional pathogen-specific T_{RM} , or whether they are simply closely

related cousins is an issue that remains to be determined. For example, the phenotypic features that are known to be shared among conventional T_{RM} populations, regardless of their specificity and/or tissue location, are reported to be driven by the T_{RM} master transcriptional regulators Blimp-1 and Hobit (36). However, the expression of Blimp-1 and Hobit in tumorinfiltrating T_{RM} is yet to be reported. By contrast, the transcription factor Runx3, which influences the downregulation of mRNA transcripts associated with cellular migration (S1pr1, Klf2, and Ccr7) appears to be expressed in both conventional and tumor-infiltrating T_{RM} (37). Moreover, conventional pathogenspecific T_{RM} are thought to be retained in peripheral tissue after resolution of infection, acting as a vanguard against future reexposure. In this context, a large proportion of conventional T_{RM} are likely persisting in peripheral tissue in an antigen-free manner, until such time as they become re-challenged through re-exposure. By contrast, tumor-infiltrating T_{RM} (assuming they are tumor-specific) are resident within active tumor tissue and would thus be continuously exposed to antigen, which would likely result in a phenotype distinct from conventional "resting"

 T_{RM} . For these reasons and because the precise relationship between conventional T_{RM} and tumor-infiltrating T_{RM} is yet to be well-defined in the literature, in our laboratory we have adopted the term "TIL_{\text{RM}}" (resident memory-like TIL) to delineate these CD103-expressing tumor resident cells from conventional pathogen-specific T_{RM} .

Until recently, broader investigation into the global nature of TIL_{RM} infiltration in human tumors was severely hampered by the lack of an anti-human CD103 antibody that was suitable for IHC of formalin-fixed tissues. This situation changed in 2013 when a new antibody was, ironically, developed for diagnosis of hairy cell leukemia (38), a setting where CD103 is ectopically overexpressed. Since the introduction of this reagent, TIL_{RM} have now been reported to be present in at least eight different tumor settings including lung, breast, ovarian, endometrial, cervical, melanoma, colorectal, pancreatic, and bladder cancer (39–53) (see **Table 1**). In the majority of these reports, CD103 is used as a marker to delineate "intraepithelial" TIL, and more importantly, the presence of CD103⁺ TIL is associated with favorable prognosis.

TABLE 1 Summary of studies examining CD103+ TIL _{RM} as a prognostic indicator in solid cancers.		
Tumor histology	Summary	Reference
Bladder	A large proportion of TIL in the urothelium co-express CD8 ⁺ CD103 ⁺ . Carcinoma stromal tissue was highly enriched for CD8 ⁺ CD103 ⁺ TIL but not associated with increased E-cadherin expression	Cresswell et al. (50)
Colorectal	Microsatellite instable tumors show increased infiltration of CD8+ CD103+ TIL compared to microsatellite stable tumors	Quinn et al. (47)
Colon	CD103 expression is enhanced by antigen recognition and TGF- β signaling. T cell activation in the presence of TGF- β induces CD103 expression	Ling et al. (49)
Ovarian	CD103 ⁺ TIL were found to be abundant across all major ovarian cancer subtypes but highly enriched in high-grade serous cancer (HGSC), and their presence correlates with improved survival	Webb et al. (55)
Lung	CD103 ⁺ TIL correlate with improved early stage patient survival in non-small cell lung cancer (NSCLC) and intraepithelial TIL density. CD103 ⁺ TIL show enhanced effector function against autologous tumor	Djenidi et al. (39)
Ovarian	CD103 demarcates intraepithelial CD8 ⁺ TIL which co-express PD-1 and appear quiescent in the tumor microenvironment	Webb et al. (41)
Breast	High abundance of CD103 ⁺ TIL in ER negative (basal-like subtype) tumors within intraepithelial regions correlates with good prognosis	Wang et al. (40)
Melanoma	Interlesional TIL populations show an enriched gene signature indicative of a resident memory phenotype which is responsive to immune checkpoint blockade	Boddupalli et al. (48)
Endometrial	Abundance of CD8 ⁺ CD103 ⁺ TIL in endometrial tumor epithelium is a strong prognostic indicator in endometrial adenocarcnoma	Workel et al. (42)
Ovarian	CD103 ⁺ TIL collected from HGSC co-express PD-1 and CD27. TIL activated in the presence of HGSC upregulate CD103	Komdeur et al. (43)
NSCLC and head and neck squamous cell cancer	Cytotoxic T lymphocytes have an enriched resident memory gene signature. CD8+ CD103+ TIL co-express checkpoint receptors such as PD-1 and CTLA-4. Higher density of resident memory T cells (T _{FM})-like TIL are associated with improved patient outcome	Ganesan et al. (46)
Cervical	CD103 gene expression is associated with effector T cell function. Abundance of intraepithelial CD8 ⁺ CD103 ⁺ TIL correlates with improved patient survival	Komdeur et al. (44)
Pancreatic	Increased ratio of CD8 ⁺ CD103 ⁺ TIL to CD8 ⁺ CD103 ⁻ TIL correlates with improved patient survival	Lohneis et al. (51)
Melanoma	Presence of CD8 ⁺ CD69 ⁺ CD103 ⁺ TIL correlates with improved patient survival in melanoma. CD103 ⁺ TIL show high levels of expression of the inhibitory markers PD-1 and LAG-3	Edwards et al. (45)
Lung	Single-cell RNA sequencing of lung TIL showed distinct pre-exhausted and exhausted TIL phenotypes. Tumor resident T cells expressed high levels of CD69 and CD103 overall	Guo et al. (52)
Breast	Single-cell RNA sequencing of breast TIL revealed high TIL abundance was characterized by a T_{RM} -like phenotype and associated with improved patient survival in triple negative breast cancer	Savas et al. (53)

TIL $_{\mbox{\scriptsize RM}}$ CELLS IN THE GYNECOLOGIC CANCER SETTING

Our group first noted the presence of TIL_{RM} cells in the ovarian cancer (OvCa) setting during a flow cytometry-based survey of immune cells present in OvCa patient ascites (54). Interestingly, some but not all, ascites specimens contained CD103-expressing T cells, specifically within the CD8 subset and sometimes comprising as much as 80% of the cells in that compartment. The presence of these cells in a fluid-based tissue (ascites) initially seemed inconsistent with them being a T_{RM} population as T_{RM} are normally restricted to solid tissues. However, the ascites compartment in ovarian patients can contain large numbers of free-floating tumor cells plus abundant amounts of TGF-β. Thus it should not be surprising that tumor-specific T cells present in this fluid compartment could adopt a T_{RM} phenotype more typical of solid tissues. We have also found that these cells have a unique phenotype that includes upregulation of HLA-DR, Ki67, and PD-1, but a lack of CD69, CD137, or intracellular cytokines suggesting that they have been recently activated, but are not actively "engaging" with targets at the time of analysis. Although the cells were PD-1 positive (41), they lacked other markers of exhaustion and were capable of robust cytokine production after stimulation with PMA/ionomycin, ex vivo, suggesting that they were not terminally exhausted. These initial findings regarding CD103-expressing TIL_{RM} in OvCa were limited to flow cytometric analysis of small numbers of ascites specimens. However, once an IHC-suitable antibody was available, we followed up by analyzing larger cohorts of patients using tissue microarray technology and showed that CD103-expressing TIL_{RM} cells were also present in the solid tumors of some, but not all, OvCa patients (55). Moreover, we also demonstrated that infiltration of tumors by TIL_{RM} correlated strongly with a favorable 5-year disease-specific survival advantage in high-grade serous cancer (HGSC), the most lethal of OvCas (55, 56). This finding has now been replicated in three additional cohorts of OvCa patients (43, 57, 58) as well as in endometrial (42) and cervical cancers (44). Clearly, TIL_{RM} cells are playing an important role in the gynecologic tumor setting, as they are in other epithelial tumor settings.

EVIDENCE IN SUPPORT OF TIL_{RM} CELLS BEING "TUMOR-SPECIFIC"

Based upon their significant prognostic benefit and unique surface phenotype, we and others speculate that TIL_{RM} in OvCa as well as other cancers are highly likely to be tumor-specific (56). Unfortunately, at present there is a paucity of well-characterized tumor antigens in the HGSC setting to directly test this hypothesis. Nonetheless, our group has previously characterized the cellular immune response to the cancer/testis tumor antigen (NY-ESO-1) in a small cohort of HGSC patients (59) by IFN- γ ELISPOT. The specificity of one such patient was mapped to a well-known HLA-A2-restricted epitope (NY-ESO-1₁₅₇₋₁₆₅) for which MHC tetramer reagents are available. Combining tetramer staining with CD103 staining revealed that NY-ESO-1-specific CD8⁺ cells in this tumor sample were indeed CD103⁺ (54), confirming that tumor-specific cells fell within the TIL_{RM} compartment in this patient. However, the NY-ESO-1-specific cells in this sample comprised only a tiny proportion of the entire TIL_{RM} population, which had otherwise unknown specificity.

Similar results regarding tumor specificity of TIL_{RM} have been obtained in other cancer settings. One of the first studies to demonstrate tumor specificity of TIL_{RM} was in the non-small cell lung cancer setting wherein the authors found that CD8+CD103+ TIL selectively upregulated CD107a and granzyme B in the presence of autologous tumor cells and also specifically lysed autologous tumor cells when co-cultured in the presence of an anti-PD-1 blocking antibody (39). More recently, TIL_{RM} populations have been identified in the melanoma setting and have been shown to contain cells that stain with melan A-specific tetramers (45), again confirming the presence of tumor-specific T cells in the TIL_{RM} subset. Likewise, TIL_{RM} have been demonstrated to play a role in anti-tumor immunity in various murine tumor models. For example, using a murine model of melanoma it was reported that CD103 was required for establishment of gp100-specific TIL_{RM} populations at the tumor site (60). Interestingly, in this model gp100-specific TIL_{RM} cells even remained at the site after tumor resolution and provided long-term immunity against rechallenge, but also caused permanent vitiligo in the dermis. On a somewhat related note CD103⁺ TRM have also recently been reported to be abundant in human vitiligo specimens (61).

Despite the abundance of evidence supporting the likely tumor specificity of TIL_{RM}, one should also consider the alternate hypothesis, that because many of these epithelial tumor types originate from a tissue that could be directly or indirectly considered a mucosal barrier tissue, the TIL_{RM} populations could actually be conventional pathogen-specific T_{RM} "bystander" populations that have been amplified during tumor outgrowth. Indeed, this possibility has been raised in a very recent study designed to characterize the phenotypes of authentic tumor-specific TIL versus bystander virus-specific TIL present in human colorectal and lung tumors (62). Interestingly, in this study both the tumor-specific and bystander T cells were found to express features of T_{RM}, including CD103, whereas CD39 was found to be a more reliable marker for distinguishing between the two. Although this study does not contradict earlier findings demonstrating CD103 expression by tumor-reactive TIL, if correct, it suggests that TIL_{RM} populations may actually be more heterogeneous than previously thought. Indeed this might be particularly relevant in the gynecologic cancer setting as HSV-2 reactive T cells with a typical T_{RM} phenotype have been reported to be present in the cervical tissue of women with known HSV-2 infection (63) and the numbers of typical T_{RM} in the fallopian tube are reported to increase with age (64). Perhaps these pathogen-specific TIL_{RM} populations in previously healthy gynecologic barrier tissues simply "come along for the ride" once the tissue becomes cancerous, and perhaps even co-exist with nascent tumor-specific TIL_{RM} populations. Clearly, it remains a challenge to the field to more precisely define the specificity of TIL_{RM}.

THE "PARADOX" OF THE PROGNOSTIC EFFECT OF TIL_{RM}

As described above, the significant prognostic benefit conferred by $\rm TIL_{RM}$ in HGSC and other cancers implies that they are likely

to be tumor-specific or at least encompass tumor-specific populations. However, at the same time, this interpretation is somewhat paradoxical as these cells are present in tumor specimens that have been obtained from patients who have required clinical intervention (in the form of surgical de-bulking in the case of HGSC). This scenario suggests that if TIL_{RM} are indeed tumor-specific, they have ultimately lost the ability to control growth of the primary tumor. In recent years, it has become readily apparent that this paradox can be explained, at least in part, by various mechanisms of immune suppression and/or immune exhaustion. Indeed, the tumor microenvironment in OvCa, much like other cancers, has long been considered to be highly immunosuppressive due to the presence of soluble immune-inhibitory factors including IL-10, TGF- β , IDO, and PGE-2 (65). Likewise, the master immuneinhibitory switch molecule CTLA-4 has also been shown to be upregulated in the OvCa setting (66). In addition, inhibitory cells such as CD4⁺ Foxp3⁺ regulatory T cells (67), immunosuppressive B7-H4⁺ tumor-associated macrophages (68), and myeloidderived suppressor cells (69) have all been reported to be present in OvCa. More recently, the PD-1 immune checkpoint pathway has also been found to play a potential role in OvCa (70), as it has in many other cancer settings.

As mentioned above, our group has recently made the observation that the CD103⁺ TIL_{RM} in HGSC tumors (and ascites) are almost universally positive for PD-1 surface expression (41). By contrast, PD-1 surface expression does not seem to be a universal



FIGURE 1 | Proposed model of TIL_{RM} formation. **(A)** Precursor resident memory T cells (T_{RM}) populations are composed of previously activated CXCR3⁺ T cells which are attracted to the chemokines CXCL9/10 in the inflamed tumor environment. Within the epithelial tumor tissue, cells encounter TGF- β which promotes CD103 expression. In response to TCR engagement cells may express increased CD69 which in turn disrupts S1PR1 expression leading to a breakdown in the chemoattractant signal from S1P concentrations in the blood. TIL_{RM} cells bind to their target tumor cells with increased strength due to CD103 binding to its ligand E-cadherin, thus promoting their residency in the epithelial tissue. Similarly, precursor T_{RM} may traffic to the inflamed ascites environment and interact with epithelial tumor cells leading to TIL_{RM} formation. Finally, bystander precursor T_{RM} populations may traffic to the inflamed tumor and/or ascites environment and develop T_{RM}-like characteristics but with irrelevant antigen specificity. **(B)** Throughout cancer progression, the tumor microenvironment becomes increasingly inhospitable with increased tumor burden. Tumor cells upregulate immunosuppressive checkpoint receptors to avoid immune eradication. Following T cell activation and prolonged antigen stimulation T cells upregulate a variety of immune checkpoints which act to suppress anti-tumor immunity. TIL_{RM} may be inhibited due to the high expression of such checkpoint receptors and thus are likely candidates to respond to immune checkpoint blockade therapy.

characteristic of conventional T_{RM} where expression of PD-1 is reported to be dynamic and perhaps even restricted to certain tissue types (71, 72). This finding would suggest that unlike conventional T_{RM}, intra-tumoral TIL_{RM} may have become partially (or permanently) exhausted likely due to chronic stimulation with tumor antigen over a period of weeks to months. Indeed, we speculate that although CD103 expression may initially be beneficial to TIL_{RM} function by promoting retention within the tumor, CD103 may actually be detrimental in the longer term by causing T cells to become "trapped" within the tumor, thereby exacerbating the phenomenon of chronic Ag stimulation (see Figure 1). This scenario is supported by a recent finding in melanoma, wherein CD103⁺ TIL_{RM} selectively and specifically became activated and started expanding in patients who were undergoing anti-PD-1 immunotherapy (45). This finding suggests that TIL_{RM} may be critical players in dictating responsiveness to checkpoint blockade therapy, a topic which is currently undergoing intense scrutiny. Thus, more fully understanding the biology of TIL_{RM} becomes paramount in that context.

CONCLUSION AND FUTURE PERSPECTIVES

Resident memory T cells have rapidly gained a reputation as sentinels of peripheral immunity, primed to prevent infection *via* re-exposure to a previously encountered pathogen. However, the biology of T_{RM} is now spilling over into the field of oncology where T_{RM} are being detected in an increasing number of

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tumor settings. Whether all the functions and characteristics of conventional T_{RM} directly translate into the unique, dynamic and often hostile microenvironment of tumors has yet to be fully elucidated. Furthermore, what role TIL_{RM} play in preventing disease recurrence after standard treatments such as radiation and chemotherapy is essentially unknown territory. Clearly much remains to be learned about these cells. What is certain is the prognostic benefit that comes along with the presence of TIL_{RM} , implying that at best, they play a direct role in anti-tumor immunity, or at minimum, they are a surrogate indicator of a separate phenomenon that leads to favorable outcomes for patients with TIL_{RM} positive tumors. Future studies should explore the potential utility of these cells in cancer immunotherapy strategies, including checkpoint blockade, cancer vaccination, and cellular therapies. Of particular interest would be understanding methodologies to convert immunologically "cold" tumors to "warm" ones by coaxing the formation and putative anti-tumor activity of TIL_{RM}. One can even imagine that the TIL_{RM} phenomenon could be applied to the rapid growing field of chimeric antigen receptor (CAR) T cell technology as it transitions into the solid tumor setting, by facilitating the retention of CAR T cells in solid tumor targets. Clearly, we are still in the early days of understanding TIL_{RM} biology, but the potential implications for immuno-oncology are significant.

AUTHOR CONTRIBUTIONS

All authors listed have made a substantial, direct, and intellectual contribution to the work and approved it for publication.

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