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The therapeutic role of $\gamma\delta$ T cells in TNBC

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Triple-negative breast cancer (TNBC) is a subtype of breast cancer that presents significant therapeutic challenges due to the absence of estrogen receptor (ER), progesterone receptor (PR), and human epidermal growth factor receptor 2 (HER2) expression. As a result, conventional hormonal and targeted therapies are largely ineffective, underscoring the urgent need for novel treatment strategies. $\gamma\delta$ T cells, known for their robust anti-tumor properties, show considerable potential in TNBC treatment as they can identify and eliminate tumor cells without reliance on MHC restrictions. These cells demonstrate extensive proliferation both *in vitro* and *in vivo*, and can directly target tumors through cytotoxic effects or indirectly by promoting other immune responses. Studies suggest that expansion and adoptive transfer strategies targeting V δ 2 and V δ 1 $\gamma\delta$ T cell subtypes have shown promise in preclinical TNBC models. This review compiles and discusses the existing literature on the primary subgroups of $\gamma\delta$ T cells, their roles in cancer therapy, their contributions to tumor cell cytotoxicity and immune modulation, and proposes potential strategies for future $\gamma\delta$ T cell-based immunotherapies in TNBC.

KEYWORDS

$\gamma\delta$ T cell, gamma delta T cell, TNBC, immunotherapy, breast cancer

1 Introduction

Triple-negative breast cancer (TNBC) is a subtype of breast cancer distinguished by negative expression of estrogen receptor (ER), progesterone receptor (PR), and human epidermal growth factor receptor 2 (HER2) expression, comprising approximately 15–20% of all breast cancer cases (1). TNBC is notably aggressive, with a high rate of recurrence and metastasis, leading to poor prognoses and significant impacts on women's physical and mental health (2–4). Its characteristics, including a high mutation rate, extensive T cell infiltration, and elevated expression of programmed death-ligand 1 (PD-L1), make it a focal point in immunotherapy research (5, 6). While immune checkpoint inhibitors (ICIs) have shown some effectiveness in treating TNBC (7), most patients with advanced disease do not respond well, complicating the search for new therapeutic targets (8). Currently, tumor-infiltrating lymphocytes (TILs)—including subsets of helper CD4+ cells, B cells, NK cells,

$\gamma\delta$ T cells, and myeloid cells—are considered crucial biomarkers in tumor immunotherapy (9). Notably, $\gamma\delta$ T cell infiltration is regarded as the most favorable prognostic indicator (10). $\gamma\delta$ T cells originate from hematopoietic stem cells (HSCs) within the bone marrow and hold significant promise in the field of tumor immunotherapy. $\gamma\delta$ T cells are instrumental in tumor immunotherapy as they recognize and destroy tumor cells independently of specific antigen stimulation, unlike $\alpha\beta$ T cells, which are part of adaptive immune responses (11). $\gamma\delta$ T cells serve as a crucial link between innate and adaptive immunity, functioning as the frontline defense against tumors and playing pivotal roles in tumor progression (12, 13). These cells possess innate-like receptors enabling rapid responses to diverse pathogens, facilitating early immune defense even in the absence of prior antigen exposure (14). Furthermore, $\gamma\delta$ T cells actively participate in tissue surveillance at barrier sites, contributing significantly to the maintenance of tissue homeostasis (15, 16). Depending on the tumor microenvironment, different $\gamma\delta$ T cell subsets can display either anti-tumor or pro-tumor activities. Predominantly, $\gamma\delta$ T cells eliminate tumor cells by recognizing tumor-associated antigens via their T cell receptors (TCR) and can also augment the anti-tumor efficacy of other immune cells by secreting cytokines or expressing co-stimulatory molecules (17). These characteristics also facilitate their integration into combination therapies, including chemotherapy, radiotherapy, or other immunotherapies, to enhance treatment outcomes.

Although $\gamma\delta$ T cells have demonstrated therapeutic potential in various cancers, their role and effectiveness in treating TNBC remain in the exploratory stage. This article reviews the current research on $\gamma\delta$ T cells in TNBC treatment, discusses their possible therapeutic mechanisms, and examines the integration of this unique immune cell type into existing treatment paradigms, offering new hope for TNBC patients. By extensively analyzing the biological characteristics of $\gamma\delta$ T cells, their molecular interactions with TNBC, and the latest developments in preclinical and clinical research, we can enhance our understanding of this strategy's potential and challenges, thus paving the way for future research and the formulation of new treatment strategies.

2 Overview of $\gamma\delta$ T cells

2.1 Origin and distribution of $\gamma\delta$ T cells

T lymphocytes originate from pluripotent stem cells in the bone marrow. During embryonic and neonatal stages, some pluripotent or pre-T cells migrate to the thymus where, under the influence of thymic hormones, they differentiate and mature into immunologically active T cells. These mature T cells are then distributed to thymus-dependent areas of peripheral immune organs through the bloodstream and can recirculate through lymphatic vessels, peripheral blood, and tissue fluid, performing cellular immunity and immune regulation functions (18, 19). $\gamma\delta$ T cells, unique innate immune cells characterized by the expression of the $\gamma\delta$ heterodimer T cell receptor, are relatively rare, constituting

only 1% to 5% of peripheral blood T lymphocytes and are primarily found in mucosal tissues such as the skin, respiratory tract, digestive tract, and uterus (20). Human $\gamma\delta$ T cells originate in the medulla of the normal fetal thymus at 7–8 weeks, undergoing a developmental process similar to $\alpha\beta$ T cells, which includes functional TCR expression and negative selection to achieve self-tolerance. Unlike $\alpha\beta$ T cells, some $\gamma\delta$ T cells do not undergo positive selection, making them unrestricted by Major histocompatibility complex (MHC) in their antigen recognition and killing capabilities (21, 22). Various functional characteristics of $\gamma\delta$ T cells start to form in the thymus and gradually mature in the periphery. In the thymus, precursor cells differentiate into $\gamma\delta$ TCR+ thymocytes, which then exit to join the peripheral circulation as circulating $\gamma\delta$ T cells. These cells enter peripheral lymphoid organs and continue to develop under the influence of various hormones released by the thymus until they acquire the capabilities of mature immune cells (23, 24).

2.2 Genetic characteristics of $\gamma\delta$ T cells

T cells are classified into $\alpha\beta$ T cells and $\gamma\delta$ T cells based on the type of TCR expressed. $\gamma\delta$ T cells are a subpopulation of T cells characterized by their γ and δ chains in the T cell receptor, comprising 0.5–5% of all T cells. Unlike $\alpha\beta$ T cells, which rely on the recognition of target antigens presented on MHC molecules by the $\alpha\beta$ TCR to develop and function, $\gamma\delta$ T cells operate in an MHC-independent manner. $\alpha\beta$ T cells differentiate into effector cells upon recognizing peptide-MHC (pMHC) complexes, enabling cytotoxic activity or cytokine production to defend against pathogens and tumors (22). In contrast, $\gamma\delta$ T cells do not require antigen processing and presentation by antigen-presenting cells (APCs) for activation, allowing for rapid early immune responses (25). $\gamma\delta$ T cell effector functions are activated by TCRs and natural killer receptors (NKR) in response to stress-induced self-ligands (26). Moreover, similar to conventional $\alpha\beta$ T cells, $\gamma\delta$ T cells can differentiate into various effector profiles and produce different chemokines and cytokines, including IFN- γ , TNF- α , IL-17, IL-21, and IL-22 (27). Additionally, human $\gamma\delta$ T cells may possess antigen-presenting capabilities; for instance, blood V γ 9V δ 2 T cells can respond to microbial and tumor signals and initiate CD4+ and CD8+ T cells, akin to dendritic cells (DCs) (28).

2.3 Classification and biological characteristics of $\gamma\delta$ T cells

$\gamma\delta$ T cells are primarily classified into three main subgroups based on the TCR δ chain: V δ 1T cells, V δ 2T cells, and V δ 3T cells (29, 30). V δ 1T and V δ 3T cells are predominantly found in mucosal and tissue environments, such as the skin, intestines, liver, and spleen. V δ 1T cells demonstrate significant anti-tumor effects in conditions such as colorectal cancer, multiple myeloma, and chronic lymphocytic leukemia (31, 32), yet they can also exert potent immunosuppressive effects when infiltrating tumors (33). The role of V δ 3T cells in tumors remains less understood (34). V δ 2T cells, the most abundant subgroup, represent 50%–90% of all

$\gamma\delta$ T cells in peripheral blood and typically pair with $V\gamma 9$ TCR to form $V\gamma 9V\delta 2$ T cells, frequently utilized in clinical settings. $\gamma\delta$ T cells exhibit robust anti-tumor activities, making them valuable in adoptive immunotherapy for cancers. They are also capable of eliminating cancer stem cells in various tumors, including colon cancer, ovarian cancer, and neuroblastoma (35–38).

$\gamma\delta$ T cells are classified based on the expression of surface markers CD27 and CD45RA into four types: naive (CD27+CD45RA+, Tnaive), central memory (CD27+CD45RA–, TCM), effector memory (CD27–CD45RA–, TEM), and terminally differentiated (CD27–CD45RA+, TEMRA). Tnaive and TCM $\gamma\delta$ T cells, primarily located in peripheral lymphoid organs, lack immediate effector functions, whereas TEM and TEMRA $\gamma\delta$ T cells, commonly found at inflammation sites, exhibit immediate effector functions, including cytokine secretion and cytotoxic activities (39). Furthermore, $\gamma\delta$ T cells are also categorized based on function into $\gamma\delta$ T1, $\gamma\delta$ T17, follicular helper ($\gamma\delta$ Tfh), regulatory ($\gamma\delta$ Treg), and memory $\gamma\delta$ T cells (40). Notably, $\gamma\delta$ T1 cells produce IFN γ and are recognized as positive prognostic markers in cancer (41). $\gamma\delta$ T17 cells secrete IL-17, which may promote tumor progression (42), while $\gamma\delta$ Treg cells, prevalent in TILs across various cancers, exert immunosuppressive effects (43). An understanding of the classification and biological characteristics of $\gamma\delta$ T cells is crucial for developing effective immunotherapies that leverage their potential.

3 Recognition pathways and functions of $\gamma\delta$ T cells

3.1 T Cell receptor-mediated recognition pathway

$\gamma\delta$ T cells identify various antigens through a TCR-dependent mechanism to detect and activate against tumor cells (Figure 1). Specifically, the $V\gamma 9V\delta 2$ TCR recognizes non-peptide phosphoantigens (P-Ag) such as microbial-derived (E)-4-hydroxy-3-methyl-but-2-enyl pyrophosphate (HMBPPP), an intermediate in the non-mevalonate pathway of isoprenoid biosynthesis, which is a potent activator of $\gamma\delta$ T cells (44, 45). Similarly, host-derived isopentenyl pyrophosphate (IPP) acts as a P-Ag and stimulates $V\gamma 9V\delta 2$ T cell responses. Bisphosphonates inhibit the farnesyl pyrophosphate synthase (FPPS) in the isoprenoid biosynthesis pathway in target or APCs, causing IPP and its metabolites to accumulate and be targeted by $V\gamma 9V\delta 2$ T cells (46, 47). Mookerjee-Basu et al. and Scotet et al. have documented that tumors sensitive to $V\gamma 9V\delta 2$ display and interact with a complex similar to mitochondrial ATP synthase, specifically F1-ATPase (48, 49). The activation of $V\gamma 9V\delta 2$ T cells is further enhanced in the presence of apolipoprotein (apoA-I), as F1-ATPase can complex with apoA-I to present phosphoantigens

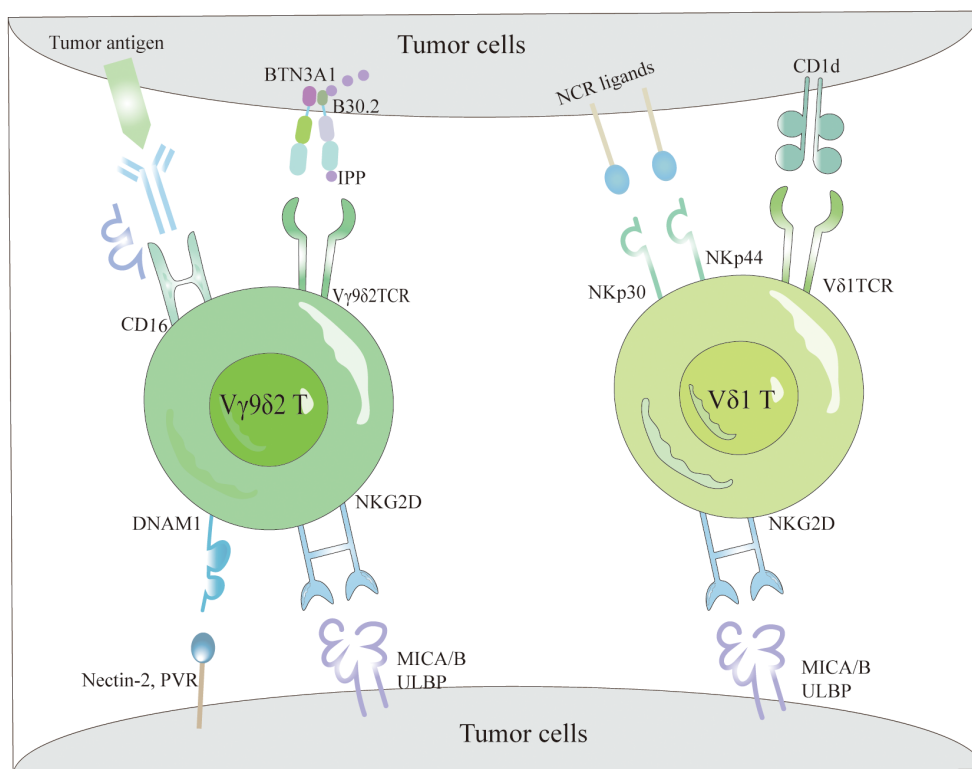


FIGURE 1
The tumor cell recognition of $\gamma\delta$ T cells. $\gamma\delta$ T cells recognize tumor cells based on the expression of $\gamma\delta$ TCRs and NKR. Specifically, $V\gamma 9V\delta 2$ TCR detects elevated intracellular levels of phosphorylated antigens, such as IPP, via BTN3A1, facilitating efficient tumor cell elimination. Additionally, they interact with tumor cells through the binding of NKG2D and DNAM-1 receptors to their respective ligands (MICA/B and ULBPs). Conversely, $V\delta 1$ T cells recognize lipid antigens presented by CD1d via their $V\delta 1$ TCR and utilize NKG2D and NCRs (NKp30 and NKp44) along with their corresponding ligands for tumor cell recognition. Moreover, $V\gamma 9V\delta 2$ T cells effectively target and eradicate tumors via the CD16-mediated ADCC mechanism.

recognizable by V γ 9V δ 2 TCR (50). Research indicates that detecting P-Ag in target cells necessitates a surface protein with intracellular and extracellular domains, specifically butyrophilin 3A1 (BTN3A1 or CD277), which binds HMBPP to its intracellular B30.2 domain. This binding induces an extracellular conformational change, facilitating the recognition of target cells by V γ 9V δ 2T cells through inside-out signaling (51, 52). Additionally, few ligands for V δ 1 TCR have been identified, with lipid antigens presented by the MHC-like molecule CD1d binding to V δ 1 TCR (Figure 1), a relationship elucidated by the crystal structure of the V δ 1 TCR with CD1d-sulfatide (53). This capability of $\gamma\delta$ T cells to recognize tumor cells via TCR underscores their integral role in both innate and adaptive immunity.

3.2 NK cell receptor pathway

The mechanisms by which $\gamma\delta$ T cells identify tumor cells are not limited to TCR interactions but also include their reliance on NKR (54) (Figure 1). These cells primarily identify tumor cells through NKR such as NKG2D, DNAM-1, NKp30, NKp44, and NKp46, which bind to specific ligands present on tumor cells (55). NKR not only regulate the activation and function of NK cells but also facilitate immune surveillance by $\gamma\delta$ T cells, enabling the distinction between transformed and infected cells. NKG2D, a C-type lectin receptor, binds ligands that are typically absent in most normal tissues but are overexpressed on tumor cells, thereby enabling $\gamma\delta$ T cells to recognize and eliminate tumor cells (56). Identified ligands for NKG2D in human cells include MHC class I chain-related proteins (MICA/MICB) and six UL16-binding proteins (ULBP1–6) (57). Contrary to previous beliefs that natural cytotoxicity-triggering receptor (NCRs) were exclusive to NK cells, recent data reveal their presence in T cells and NK-like cells (58, 59). Although V δ 1+ and V δ 2+ cells naturally lack NCR expression, it can be selectively enhanced in V δ 1+ cells through AKT-dependent signaling triggered by γ C cytokines (IL-2 or IL-15) and TCR stimulation (60). The NCRs expressed in V δ 1+ $\gamma\delta$ T cells (Figure 1), predominantly NKp44 and NKp30, endow these cells with heightened abilities for targeted cytotoxicity against tumor cells and for secreting IFN- γ (55, 61).

3.3 CD16 pathway

One pathway through which V γ 9V δ 2T cells exert their anti-tumor effects involves CD16, also known as Fc γ RIII (Figure 1), a low-affinity type III receptor that specifically binds to the Fc portion of Immunoglobulin G (IgG) (62). This receptor mediates antibody-dependent cellular cytotoxicity (ADCC) and cytokine production, including TNF- α (63). Research indicates that V γ 9V δ 2T cells treated with zoledronic acid and IL-2 can express CD16 (64). The interaction of CD16 with the Fc portion enables V γ 9V δ 2T cells to detect and destroy tumor cells expressing IgG through ADCC activation (65, 66). Depending on the presence or absence of CD16 expression, V γ 9V δ 2T cells are classified into two types: CD16- and CD16+. The CD16- subset produces higher cytokine

levels, expresses fewer killer inhibitory receptors (KIRs), and exhibits lower cytotoxicity, while the CD16+ subset has higher KIR levels and significant direct cytotoxic capabilities (67). The presence of CD16 enhances the recognition capabilities of V γ 9V δ 2T cells against IgG-expressing tumor cells, particularly in the CD16+ subgroup, which shows enhanced direct cell-killing ability.

4 Anti-tumor effects of $\gamma\delta$ T cells

$\gamma\delta$ T cells have the unique capability to recognize and destroy tumor cells without relying on traditional antigen presentation mechanisms, which is particularly advantageous in targeting tumors like TNBC that lack specific antigen presentation. $\gamma\delta$ T cells can directly lyse tumor cells via two independent pathways (Figure 2): firstly, by secreting perforin and granzymes (68, 69); secondly, by inducing cell death through the Fas/FasL pathway and tumor necrosis factor-related apoptosis-inducing ligand (TRAIL) (70, 71). On their surface, $\gamma\delta$ T cells express FasL, which can trigger programmed apoptosis in tumor cells by forming a Fas trimer upon binding to Fas (72). This interaction leads to the activation of the death effector domain (DED) and Fas-associated death domain-containing protein (FADD), subsequently activating downstream caspases that result in cellular destruction and death (73–75). Similarly, TRAIL induces apoptosis via caspase activation through FADD (71, 75). $\gamma\delta$ T cells also indirectly exert anti-tumor effects by activating other immune cells and possess antigen-presenting capabilities that stimulate the activation and proliferation of $\alpha\beta$ T cells (76, 77) (Figure 2). They promote the proliferation and differentiation of CD8+ T cells and regulate TNF- α and IFN- γ secretion, enhancing tumor clearance rates (78, 79). Additionally, research by Bansal et al. found that $\gamma\delta$ T cells expressing CXCL13 and CXCR5 demonstrate follicular helper T cell (Tfh) characteristics, particularly in the presence of IL-21, which aids B cell support (80, 81). Caccamo et al. discovered that V δ 2 $\gamma\delta$ T cells can activate B cells (Figure 2), leading them to produce significant amounts of immunoglobulins even without antigen stimulation, and facilitate the formation of germinal centers (82, 83). Moreover, Maniar et al. showed that zoledronic acid-activated $\gamma\delta$ T cells enhance NK cell-mediated cytotoxicity (Figure 2) against tumor cells, a process reliant on the interaction between CD137 ligand on $\gamma\delta$ T cells and CD137 on NK cells (84). Furthermore, $\gamma\delta$ T cells can modulate the production of cytokines such as interferon- γ by NK cells, influenced by DC-like cells (85).

The interaction between $\gamma\delta$ T cells and DCs is reciprocal (Figure 2). $\gamma\delta$ T cells facilitate the maturation of DCs, while mature DCs trigger the activation and proliferation of $\gamma\delta$ T cells, thereby enhancing their anti-tumor capabilities (86). Typically, Fas/FasL mediates cell apoptosis; however, notably, DCs express high levels of the Fas inhibitor cFLIP. Elevated cFLIP levels convert the pro-apoptotic signal of Fas into one that promotes dendritic cell maturation and function (87). Additionally, activated $\gamma\delta$ T cells secrete IFN- γ and TNF α , stimulating the expression of CD86 and MHC-I molecules on DCs surfaces, which supports further DCs maturation (88). Conversely, mature DCs boost $\gamma\delta$ T cell proliferation and augment their cytotoxic and immunoregulatory

abilities by secreting cytokines such as IL-1 β , IL-12, IL-18, IFN- γ , and TNF- α (89). DCs also facilitate contact-dependent activation of $\gamma\delta$ T cells through CD86-CD28 interactions and the expression of specific $\gamma\delta$ T cell activation ligands, like P-Ag (90). These intricate interactions underscore the versatile role of $\gamma\delta$ T cells in modulating anti-tumor immune responses.

5 The protumor function of $\gamma\delta$ T cells

While $\gamma\delta$ T cells serve an effector role in anti-tumor immune responses, their function is frequently inhibited by the tumor microenvironment (TME), which may even lead these cells to differentiate into subgroups that foster tumor progression. Research has demonstrated that tumor-infiltrating $\gamma\delta$ T17 cells (Figure 2) exacerbate tumor development by secreting IL-17 (91). IL-17, a pro-inflammatory cytokine, promotes chronic tissue inflammation, thus aiding tumor progression. In cancers such as pancreatic, liver, non-small cell lung, and breast cancer, elevated IL-

17 levels correlate with increased metastasis and poor prognosis (92). IL-17 enhances tumor progression through several mechanisms: it activates the PI3K/AKT signaling pathways to stimulate tumor cell proliferation (93), induces the production of matrix metalloproteinases (MMPs) which degrade the extracellular matrix, facilitating tumor cell invasion and distant metastasis (94), and promotes angiogenesis by stimulating angiogenic factors including VEGF, IL-6, and IL-8 (95, 96). Furthermore, IL-17 recruits myeloid-derived suppressor cells (MDSCs) into the tumor, sustaining an immunosuppressive TME (97, 98) (Figure 2). A study by Ma and colleagues revealed that MDSCs could also drive $\gamma\delta$ T cells to produce IL-17 via IL-1 β and IL-23, perpetuating a malignant cycle of tumor progression (99).

$\gamma\delta$ T cells potentially regulate tumor immune responses. In breast cancer, V δ 1+ $\gamma\delta$ T cells infiltrate and inhibit the proliferation of inactive T cells and the functionality of effector CD4+ and CD8+ T cells. They also suppress the maturation of DCs (Figure 2) and the proliferation of anti-tumor V δ 2+ cells, fostering an immunosuppressive environment (100). Furthermore, studies

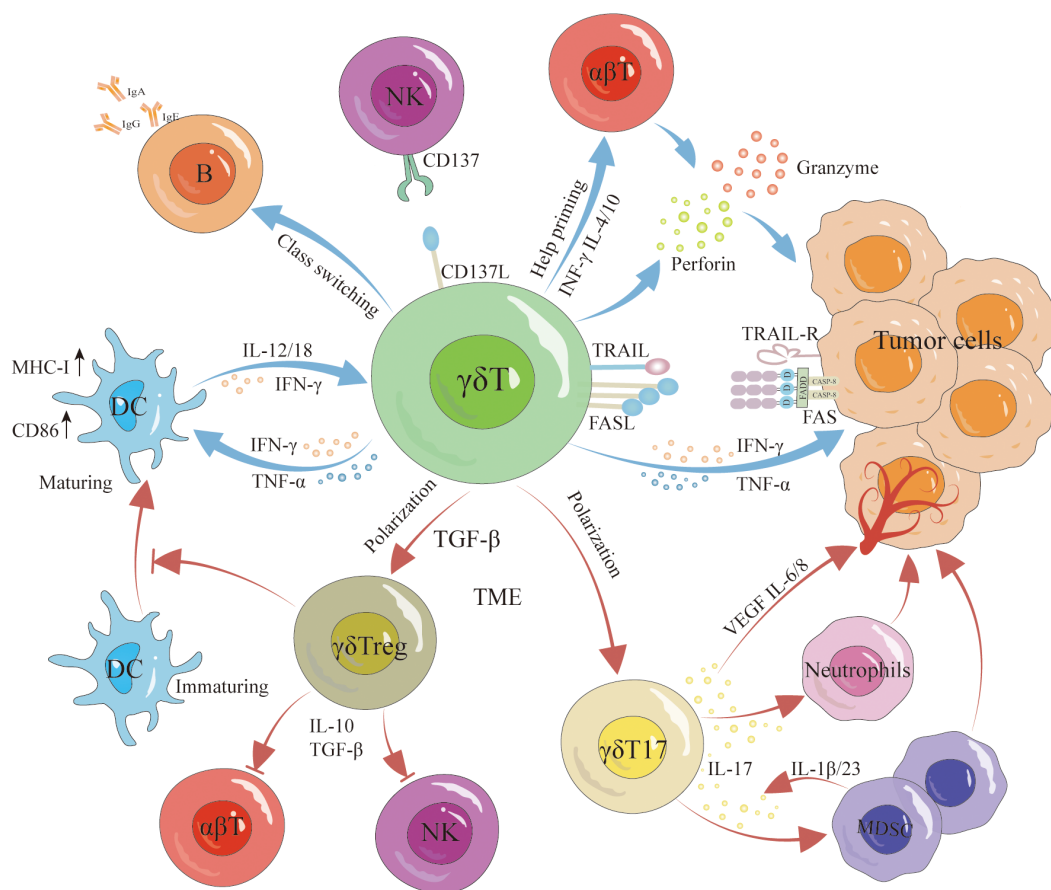


FIGURE 2

Antitumor and protumor functions of $\gamma\delta$ T cells. $\gamma\delta$ T cells exert antitumor effects through various pathways, involving both direct and indirect mechanisms. Direct antitumor effects include lysing the tumor via the perforin-granzyme pathway, Fas/FasL pathway, and TRAIL signaling. Furthermore, they indirectly hinder tumor growth by promoting DC maturation, inducing B cell activation and proliferation, and augmenting NK cell-mediated cytotoxicity against malignant cells. However, it is noteworthy that $\gamma\delta$ T cells can differentiate into $\gamma\delta$ Treg cells and $\gamma\delta$ T17 cells, promoting tumor growth. Specifically, $\gamma\delta$ Treg cells secrete IL-10 and TGF- β to suppress $\alpha\beta$ T cell and NK cell responses to tumors while concurrently inhibiting DC maturation. $\gamma\delta$ T17 cells promote tumorigenesis by secreting IL-17 to recruit MDSCs and stimulate the production of VEGF, IL-6, and IL-8, inducing angiogenesis and expediting tumor growth.

have shown that $\gamma\delta$ T cells can differentiate into FoxP3+ $\gamma\delta$ Treg cells under the influence of $\gamma\delta$ TCR monoclonal antibodies and TGF- β (101). These $\gamma\delta$ Treg (Figure 2) cells, similar to traditional Treg cells, secrete inhibitory cytokines such as IL-10 and TGF- β , which suppress the immune response of $\alpha\beta$ T cells and NK cells against tumors. Specifically, breast cancer-derived $\gamma\delta$ Treg cells can induce immunosenescence in naive T cells, effector T cells, and DCs. They also inhibit the proliferation of T cells from human peripheral blood mononuclear cells (PBMCs) (102), and by inducing CD86/CTLA-4 and PD-L1/PD-1 interactions, they alter the tumor environment's structure and reduce effector T cell activity (103). It is clear that the TME influences the phenotype and function of $\gamma\delta$ T cells, with $\gamma\delta$ T17 cells exacerbating inflammation, promoting angiogenesis, and recruiting MDSCs and other inhibitory cells, thereby enhancing tumor progression. Correcting the TME and reversing the negative regulatory effects of tumor-infiltrating $\gamma\delta$ T cells are critical for leveraging $\gamma\delta$ T cells in cancer immunotherapy.

6 The role of $\gamma\delta$ T cells in TNBC

$\gamma\delta$ T cells, as crucial components of TILs, play a significant role in regulating tumor immune responses. Notably in TNBC, research by Chabab et al. has revealed that $\gamma\delta$ T cell infiltration often exceeds that in other breast cancer (BC) types, a phenomenon closely linked to TNBC's higher mutation rate (104). Multiple studies suggest that the presence of TILs in TNBC often correlates with a more favorable prognosis (105, 106). Wu et al. have shown that the abundance of V δ 1+ $\gamma\delta$ T cells, rather than the total number of $\gamma\delta$ T cells, is critical in determining the treatment response in TNBC patients. These infiltrating V δ 1+ T cells, with their cytotoxic capabilities and ability to produce IFN- γ , operate through an intrinsic mechanism as they respond to MICA and cytokines IL-12 and IL-18. Further research indicates that the density of V δ 1+ T cells positively correlates with patients' progression-free survival (PFS) and overall survival (OS) (107). Craven et al.'s data analysis also supports this finding, linking $\gamma\delta$ TILs with prolonged OS (108). Conversely, Janssen et al. highlight that in TNBC, V δ 2+ T cells are the predominant $\gamma\delta$ TILs subgroup, actively contributing to anti-tumor effects by secreting IFN- γ and TNF- α . Their studies also reveal that $\gamma\delta$ TILs are not sources of IL-17 in TNBC, unlike in colorectal cancer, suggesting that TNBC's $\gamma\delta$ TILs may not depend on IL-17 for promoting tumor growth (109). Moreover, evidence suggests that $\gamma\delta$ T cells may facilitate breast tumor development through their immunoregulatory functions, correlating with poor prognosis in breast cancer. Within the TME of human breast cancer, a minority (<20%) of V δ 1 T cells that express CD73 and can produce IL-10, adenosine, and IL-8 exhibit immunosuppressive effects (110). These findings suggest that while some $\gamma\delta$ T cell subgroups are immunosuppressive, their impact is often masked by those with anti-tumor activity. Consequently, further investigation into the role of $\gamma\delta$ T cells in TNBC or its subtypes, and their influence on disease progression and treatment responses, is critically important.

7 The therapeutic potential of $\gamma\delta$ T cells in TNBC

TNBC presents significant diagnostic and treatment challenges due to its ambiguous biological characteristics. The IMpassion130 study marked a pivotal shift into the era of immunotherapy for breast cancer, making TNBC the most extensively studied cancer type in this domain (111). Recently, enhanced insights into the tumor microenvironment and immune evasion mechanisms have established immunotherapy as a viable approach for TNBC. $\gamma\delta$ T cells, which recognize tumor antigens without MHC restrictions and exhibit potent cytotoxic effects, can be substantially expanded *in vitro* and *in vivo*, demonstrating significant potential in tumor immunotherapy. The presence of TILs in TNBC has been linked to favorable prognoses, highlighting that adoptive cell therapy (ACT) provides new therapeutic avenues. ACT primarily encompasses therapies such as chimeric antigen receptor (CAR)-T, TCR, and TILs therapies, which all operate on similar principles (112).

The induction and adoptive transfer of $\gamma\delta$ T cells, particularly targeting V δ 2 and V δ 1 subtypes, represent a promising avenue in cancer immunotherapy research. Commonly, the combination of P-Ag such as BrHPP and HMBPP or nitrogen-containing bisphosphonates (N-BP) like zoledronic acid (ZOL) with IL-2 is utilized for both *in vivo* and *in vitro* expansion of V δ 2 $\gamma\delta$ T cells. This approach has been extensively implemented and clinically validated for therapeutic safety (21, 113–115). ZOL not only facilitates the transformation of V δ 2T cells into TEM phenotypes but also significantly boosts their cytotoxic capabilities, thus enhancing tumor suppression (116). *In vivo*, V γ 9V δ 2 T cells stimulated by P-Ag or N-BP also demonstrate the capacity to target and eliminate multiple tumor cells. However, the clinical response rate is typically lower compared to that observed in cases of overt metastasis (117, 118). In the context of neoadjuvant therapy for breast cancer, combining letrozole with ZOL to expand $\gamma\delta$ T cells *in vivo* has demonstrated substantial patient benefits (119). Additionally, *ex vivo* expansion of $\gamma\delta$ T cells for adoptive immunotherapy has shown notable anti-tumor effects in animal models (120). Vitamin C (VC) and its derivatives also positively influence the proliferation and activation of V δ 2T cells, particularly at high doses, where VC augments the cytotoxic effects of CD8+ T cells and synergistically boosts the efficacy of immunotherapy alongside immune checkpoint inhibitors (121–123). A novel protocol for V δ 2 T cell expansion developed by Xu et al., integrating ZOL, IL-2, IL-15, and VC, has proven effective in enhancing cell proliferation, differentiation, and cytotoxicity, significantly curtailing tumor growth and extending survival in mice (124).

Recent advancements in cancer immunotherapy have introduced an approach involving Delta One T (DOT) cells derived from V δ 1+ T cells. These cells are activated through TCR agonists and cytokines, resulting in substantial proliferation (12, 125). DOT cells have shown promising therapeutic effects against various tumor types, an outcome further enhanced by increased expression of NKp30, NKp44, NKG2D, and DNAM1 (126, 127). Additionally, Raute et al. have found that primarily expanded V δ 1+

and V δ 2+ T cells can target triple-negative breast cancer stem cells (BCSC) derived from patients. However, these BCSCs may differentiate *in vivo* into cells that lack stem cell-like properties and $\gamma\delta$ T cell activation ligands, thereby escaping effective $\gamma\delta$ T cell-mediated destruction. Nonetheless, $\gamma\delta$ T cells can still marginally kill these differentiated cells *in vivo* by recognizing P-Ag. Significantly, the cytotoxic effect is enhanced by ZOL, suggesting that a combination of $\gamma\delta$ T cells and ZOL might represent an effective strategy against both triple-negative BCSCs and non-stem tumor cells (128). In conclusion, the expansion of $\gamma\delta$ T cells offers new hope for treating patients with TNBC. Although this therapy remains under research, its potential for broad application in TNBC treatment holds significant promise as an effective therapeutic option.

CAR-T cell therapy, a form of adoptive cell therapy, merges the antigen specificity of antibodies with the effector functions of T cells, showing considerable potential to improve survival rates in TNBC patients (129). Upon reintroduction into the patient, CAR-T cells initiate cytotoxic immune responses by recognizing tumor-associated antigens. Initially employed in refractory hematologic malignancies (130), this technology has been extensively studied in TNBC, targeting antigens like ROR1, c-Met, EGFR, FR α , and MUC1. These targets provide specific foci for CAR-T cell therapy. However, the efficacy in solid tumors is often hampered by challenges such as tumor antigen heterogeneity, the immunosuppressive tumor microenvironment, and the limited infiltration and persistence of CAR-T cells within tumors (131, 132). Moreover, research has also ventured into expressing CARs on other effector cells like CAR- $\gamma\delta$ T cells, which specifically recognize and target tumor cell surface antigens, delivering a cytotoxic response (133). Capsomidis et al. have shown that CAR- $\gamma\delta$ T cells not only migrate efficiently to tumor sites but also exhibit strong cytotoxicity directed by specific tumor antigens (134). Demonstrated in xenograft mouse model studies, these cells exhibit significant anti-tumor activity both *in vitro* and *in vivo*, suggesting they could form a novel treatment approach for TNBC (135).

Abnormal signaling of immune checkpoint molecules has been observed to disrupt the normal function of the TCR and alter the phosphorylation levels of intracellular proteins via ITIM motifs and SHP-1/2. Consequently, this interference inhibits the proliferation and activation of $\gamma\delta$ T cells, leading to a reduction in their cytotoxicity (136). Therefore, the combined application of $\gamma\delta$ T cells and ICIs may emerge as an effective strategy to enhance the therapeutic efficacy of TNBC. Additionally, bispecific antibodies have demonstrated potential in augmenting the efficacy of $\gamma\delta$ T cell immunotherapy by significantly enhancing cytotoxicity through the fusion of tumor-binding and T-cell splicing structural domains (137). Oberg et al. reported that the administration of $\gamma\delta$ T cells expanded *in vitro* with specific bispecific antibodies effectively slowed the growth of pancreatic and colon cancers in preclinical models (138, 139). Additionally, bispecific molecules (GABs) linking the extracellular domain of the tumor-reactive V γ 9V δ 2

TCR to a CD3 binding structure have been shown to promote T-cell infiltration into the tumor microenvironment, thereby inhibiting tumor growth *in vivo* (140). Thus, the bispecific splicer holds great potential as a form of $\gamma\delta$ T cell-based immunotherapy. If successful in clinical trials, this treatment could offer a powerful and relatively inexpensive therapeutic option for TNBC patients. Another innovative approach involves transducing $\alpha\beta$ T cells with a high-affinity V γ 9V δ 2 TCR, termed T cells with defined $\gamma\delta$ TCR (TEG) (12). TEG demonstrates the ability to target a wide array of solid and hematological tumors and, in addition to exerting cytotoxic effects, exhibits paracrine activity that induces functional maturation of dendritic cells (14). Consequently, TEG can effectively target a broad spectrum of tumor cells owing to the wide reactivity of the V γ 9V δ 2 TCR, thereby addressing the limitations of low persistence and impaired activation of $\gamma\delta$ T cells within the tumor microenvironment (12, 125). These novel therapeutic strategies instill hope for patients with triple- TNBC by expanding treatment options and possibilities for the future.

8 Major challenges and clinical implications of utilizing $\gamma\delta$ T cells

One of the major challenges for $\gamma\delta$ T cells is their scarcity in the immune system, as they represent only a small fraction of the total T cell population (20). This limited presence hampers the ability to fully exploit their therapeutic potential, especially when compared to the more abundant $\alpha\beta$ T cells. Additionally, the specific tissue distribution of $\gamma\delta$ T cells, predominantly located in the peripheral regions of non-lymphoid tissues, complicates their accessibility and clinical application. Despite considerable efforts, expanding various clones of $\gamma\delta$ T cells to clinically relevant numbers remains a significant obstacle to their widespread use in cellular immunotherapy (141). However, the V δ 1 subpopulation has been shown to predict a favorable prognosis in triple-negative breast cancer, as supported by protein or gene level analyses (9, 107). Another significant challenge is the limited role of $\gamma\delta$ T cells in the TME. Although several studies have shown that $\gamma\delta$ T cells can modulate immunosuppressive cells within the TME, the scarcity of nutrients, the presence of suppressor molecules, and hypoxia may still constrain their therapeutic potential (142, 143). Suppressive molecules produced by tumor cells and other cells in the TME, such as TGF- β (144), prostaglandin E2 (PGE2) (145), adenosine (146), and soluble NKG2D ligands (147), can interfere with $\gamma\delta$ T cell proliferation and function.

Despite these challenges, ongoing clinical trials are assessing the safety and antitumor efficacy of $\gamma\delta$ T cells. However, the clinical utility of V γ 9V δ 2 T cells may be hindered by susceptibility to T cell exhaustion and activation-induced cell death (AICD) (148). Moreover, in rare cases, stimulation of $\gamma\delta$ T cells with phosphoantigens and ZOL may lead to adverse effects including fever, fatigue, eosinophilia, thrombosis, elevated liver transaminases, hyperglycemia, gastritis, musculoskeletal pain, and

nephrotoxicity (149, 150). Thus, the safety and tolerability of $\gamma\delta$ T-cell therapies require meticulous consideration and monitoring throughout design and implementation. In conclusion, despite numerous challenges, $\gamma\delta$ T cell therapy holds significant promise as a novel immunotherapeutic approach. By comprehensively investigating the tumor microenvironment, devising effective therapeutic strategies, and employing advanced immunological techniques to expand and activate $\gamma\delta$ T cells, the efficacy of this therapy can be enhanced to yield improved clinical outcomes for patients with tumors.

9 Summary and future perspectives

TNBC, characterized by the absence of ER, PR, and HER2 expression, limits patients' options for hormone or targeted therapies, thereby necessitating new treatment strategies. $\gamma\delta$ T cells, key components of immune defense, can target and destroy tumor cells independently of traditional MHC-mediated antigen presentation, thereby exerting significant anti-tumor effects. However, their potential pro-tumor activities, including suppressing anti-tumor responses, enhancing tumor angiogenesis, and secreting IL-17, are subjects of ongoing debate (68, 151). Although the tumor microenvironment is thought to recruit numerous $\gamma\delta$ T cells that may promote tumor progression, single-cell RNA sequencing from fresh breast cancer tissues and patients' peripheral blood reveals that $\gamma\delta$ T cells generally correlate with favorable clinical outcomes, with tissue-infiltrating $\gamma\delta$ T cells being more active and cytotoxic than their blood counterparts (152). Recent mouse studies have noted pro-tumor and pro-metastatic effects in $\gamma\delta$ T cells producing IL-17, although such cells are rare in humans. Evidence suggests that $\gamma\delta$ TILs contribute minimally to IL-17 secretion compared to Th17 and CD4+ T cells in the TME (109). This underscores the potential of $\gamma\delta$ T cell-based immunotherapy as a novel strategy for breast cancer treatment. Research indicates that $\gamma\delta$ T cells can inhibit TNBC progression through direct cytotoxic actions and by modulating immune responses. The presence of $\gamma\delta$ TILs in the TNBC microenvironment is strongly associated with favorable patient prognoses, underscoring their vital role in anti-tumor immunity. While various methods to expand $\gamma\delta$ T cells have shown promise in anti-tumor therapy, specific studies on their application in TNBC remain limited. Furthermore, enhancing the activity or specificity of $\gamma\delta$ T cells through CAR technology presents opportunities to improve their therapeutic potential in TNBC treatment.

Future research will persist in investigating the use of $\gamma\delta$ T cells in treating TNBC, with a particular emphasis on effectively expanding and activating these cells, overcoming the immunosuppressive tumor microenvironment, and enhancing their tumor-homing capabilities. Moreover, the use of engineered $\gamma\delta$ T cells, such as CAR- $\gamma\delta$ T cells and TEGs, either alone or in combination with checkpoint inhibitors, holds promise for

enhancing the response rate and anti-tumor effects of $\gamma\delta$ T cell therapy. Considering the complexity and therapeutic challenges of triple-negative breast cancer (TNBC), researchers must explore novel methods or techniques to achieve effective expansion of both V γ 9V δ 2 T cells and V δ 1 T cells. Specifically, the advancement of CAR- $\gamma\delta$ T cell therapy necessitates additional clinical data to support its use in TNBC treatment. As our understanding of the tumor microenvironment and immune evasion mechanisms expands, coupled with cutting-edge immunotherapies, TNBC treatment is poised to become increasingly personalized, significantly enhancing patient prognosis and quality of life.

Author contributions

WL: Conceptualization, Visualization, Writing – original draft, Writing – review & editing. XZ: Writing – original draft, Writing – review & editing. CR: Writing – original draft, Writing – review & editing. SG: Writing – review & editing. QH: Writing – original draft, Writing – review & editing. ML: Writing – original draft, Writing – review & editing. XL: Funding acquisition, Project administration, Resources, Supervision, Writing – review & editing.

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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