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Review

Turning cold tumors into hot tumors by improving T-cell infiltration

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Abstract

Immunotherapy, represented by immune checkpoint inhibitors (ICls), has greatly improved the clinical efficacy of malignant tumor therapy. ICl-mediated antitumor responses depend on the infiltration of T cells capable of recognizing and killing tumor cells. ICls are not effective in "cold tumors", which are characterized by the lack of T-cell infiltration. To realize the full potential of immunotherapy and solve this obstacle, it is essential to understand the drivers of T-cell infiltration into tumors. We present a critical review of our understanding of the mechanisms underlying "cold tumors", including impaired T-cell priming and deficient T-cell homing to tumor beds. "Hot tumors" with significant T-cell infiltration are associated with better ICl efficacy. In this review, we summarize multiple strategies that promote the transformation of "cold tumors" into "hot tumors" and discuss the mechanisms by which these strategies lead to increased T-cell infiltration. Finally, we discuss the application of nanomaterials to tumor immunotherapy and provide an outlook on the future of this emerging field. The combination of nanomedicines and immunotherapy enhances cross-presentation of tumor antigens and promotes T-cell priming and infiltration. A deeper understanding of these mechanisms opens new possibilities for the development of multiple T cell-based combination therapies to improve ICl effectiveness.

Key words: cold tumor; immune checkpoint inhibitors; T-cell infiltration; T-cell priming; nanomedicine

Introduction

Recently, immune checkpoint inhibitors (ICIs), such as nivolumab and pembrolizumab, have been applied to an increasing number of cancer types, forming a paradigm treatment in clinical trials [1, 2]. Although ICIs have shown clinical activity in a wide range of tumor types, a substantial percentage of patients still do not respond to ICI therapy [3]. ICI-mediated antitumor responses rely on the expression of PD-L1 in tumors and the infiltration of T cells capable of recognizing and killing tumor cells. Immune cells such as CD8+ T cells are associated with prolonged survival of cancer patients and increased efficacy of immunotherapy [4]. A lack of T cells in tumors can lead to resistance to immunotherapy [5]. The success of chimeric antigen receptor (CAR) T-cell infusions for patients' leukemia and lymphoma also

demonstrates the importance of T cells in antitumor immunity [6]. Considering the potential mechanisms of cancer immunotherapy, the infiltration of CD8+ T lymphocytes in tumors is important for the therapeutic response to ICIs.

According to the spatial distribution of cytotoxic immune cells in the tumor microenvironment (TME), a tumor is classified into one of three basic immunophenotypes: immune-inflamed, immune-excluded and immune-desert phenotypes (Figure 1) [3]. Immune-inflamed tumors, also named "hot tumors", are characterized by high T-cell infiltration, increased interferon- γ (IFN- γ) signaling, expression of PD-L1 and high tumor mutational burden (TMB) [7]. Tumors with an inflamed phenotype tend to be more responsive to ICIs [8, 9]. Immune-excluded tumors

and immune-desert tumors can be described as "cold tumors". In immune-excluded tumors, CD8+ T lymphocytes localize only at invasion margins and do not efficiently infiltrate the tumor [10]. In immunedesert tumors, CD8+ T lymphocytes are absent from the tumor and its periphery [10]. In addition to poor T-cell infiltration, "cold tumors" are characterized by low mutational load, low major histocompatibility complex (MHC) class I expression and low PD-L1 expression [7]. Immunosuppressive cell populations, including tumor-associated macrophages (TAMs) and T-regulatory cells (Tregs) and myeloid-derived suppressor cells (MDSCs), are also present in cold tumors [7]. These features suggest that cold tumors lack innate immunity or that the innate antitumor immune features present in "cold tumors" may be ineffective due to the exclusion of immune cells [3]. In contrast to the inflamed phenotype, cold tumors rarely respond to ICI monotherapy [9].

Driving T cells into the TME is a gradual process (Figure 2): tumor cell death and antigen release, antigen-presenting cell (APC) processing presentation of tumor antigens, and APC and T-cell interactions lead to T-cell priming and activation [11]. Ideally, once activated, these tumor-specific T cells exit lymph nodes and travel through the bloodstream to tumor site [11]. The production of T cells and their physical contact with tumor cells is crucial for the success of antitumor immunity [12]. Once infiltrating the tumor bed, cytotoxic T lymphocytes (CTLs) specifically recognize antigenic peptide-MHC complexes on the surface of tumor cells, form immune synapses, and release perforin and granzyme to

destroy the tumor cells [13]. In addition, CTLs contribute to the apoptosis of tumor cells through the Fas/FasL pathway and suppress tumors by inducing ferroptosis and pyroptosis [14]. Dead tumor cells release additional tumor antigens and thereby amplify the T-cell response [11].

With the development of nanotechnology, immunotherapy based on nanomedicines and biomaterials offers new opportunities for the future. Nanomedicines offer unique advantages in oncology treatment, such as improved drug precision and bioavailability and reduced immunotherapy-induced side effects [15]. In addition, nanomedicines promote selective accumulation in tumors through enhanced permeability and retention (EPR) effects or include high-affinity ligands to achieve active targeting of tumors [15]. Nanotechnology-based immunotherapy enhances tumor-specific immune responses, promotes the infiltration of CTLs, and inhibits tumor metastasis and recurrence.

Given the importance of T-cell infiltration, understanding the mechanisms of T-cell homing to the tumor is necessary. To improve the clinical benefit of immunotherapy, ICIs may be combined with strategies that convert "cold tumors" to "hot tumors", which may make these tumors more sensitive to ICI therapy. In this review, we summarize the various mechanisms of T-cell infiltration disorders and current approaches to directing T cells into tumors. Finally, we summarize recent advances, challenges and opportunities for nanomedicine-based local therapeutic strategies to enhance T-cell infiltration and discuss further prospects in this field.

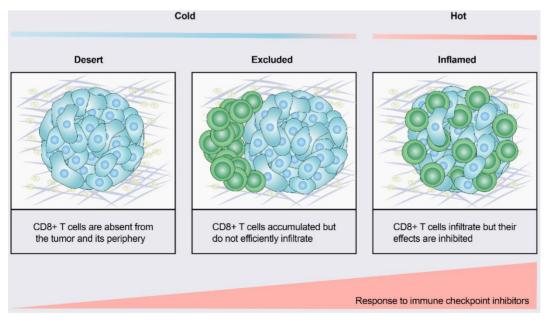


Figure 1. Tumor immune phenotypes. Based on the spatial distribution of CD8+ T lymphocytes in the tumor microenvironment (TME), a gradient of three immunophenotypes is observed: the immune-desert, immune-excluded and immune-inflamed phenotypes. In the immune-desert phenotype, immune cells are absent from the tumor and its periphery. In the immune-excluded phenotype, immune cells accumulate but do not efficiently infiltrate. In the immune-inflamed phenotype, immune cells infiltrate but their effects are inhibited. Notably, the three different phenotypes have different response rates to immune checkpoint inhibitors.

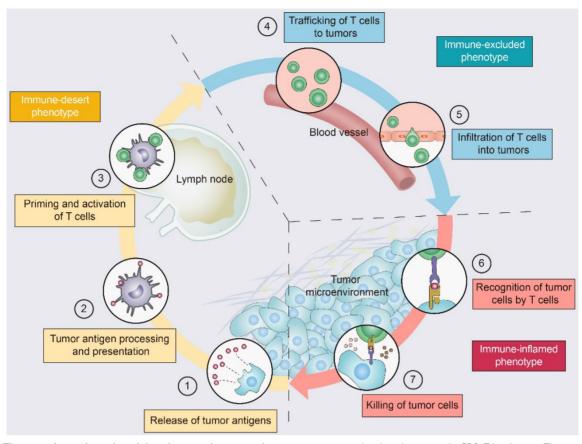


Figure 2. The tumor-immunity cycle and three immunophenotypes. Antitumor immunity is mediated to a large extent by CD8+ T lymphocytes. The tumor-immunity cycle consists of the following steps: (1) tumor antigen release, (2) tumor antigen processing and presentation, (3) T-cell priming and activation, (4) trafficking of T lymphocytes through the bloodstream to tumors, (5) infiltration of T lymphocytes into the tumor parenchyma from the vasculature or tumor periphery, (6) recognition of tumor cells, and (7) cytotoxic T lymphocyte (CTL) destruction of tumor cells by granule exocytosis or through the Fas/FasL pathway. Dead tumor cells release additional antigens, allowing the tumor-immunity cycle to continue. Notably, tumors with the immune-desert phenotype (yellow) cannot pass steps 1-3 due to the absence of T lymphocytes in both the tumor and its margins. Tumors with the immune-excluded phenotype (blue) cannot exceed steps 4-5 due to a lack of T lymphocytes in the tumor bed. Tumors with the immune-inflamed phenotype (red) cannot exceed steps 6-7 due to T-cell exhaustion and checkpoint activation. Adapted with permission from [11], copyright 2013 Elsevier.

Table 1. Mechanisms of defective T-cell priming

Mechanisms	Examples	References
Lack of tumor antigens	Lack of neoantigens	[16]
	Low mutational burden	[16, 17]
Insufficient antigen	Deletion of B2M	[24, 25]
processing or presentation	Deletion of TAP	[23]
	Loss of MHC class I (Lysosomal	[26]
	pathway)	
Dysfunction of DC-T cell	Loss of BATF3 DCs	[32]
interaction	Impaired DC activation (Loss of	[35]
	FLT3L or GM-CSF)	
	Overexpression of cosuppressive	[28]
	signals (CTLA-4)	
	Overexpression of STC1	[29]

Mechanisms of the "cold tumor" phenotype

ICI response rates are low in "cold tumors", as characterized by the absence of T-cell infiltration. In the process of driving T cells into tumors, there are many factors that can influence T-cell priming and T-cell homing to the tumor bed, leading to a noninflamed T-cell phenotype and failed antitumor immunity (Figure 3).

Defective T-cell priming

Lack of tumor antigens

The most direct cause of T-cell priming disorders is insufficient T-cell recognition due to a lack of tumor antigens. Table 1 summarizes the mechanisms of defects in T-cell priming. In general, targeted tumor antigens can be classified into two broad categories: nonmutated self-antigens and neoantigens generated nonsynonymous somatic mutations Self-antigens include nonmutated proteins that are aberrantly expressed or overexpressed in tumor cells, such as tumor-associated antigens (TAAs) and cancer/testis antigens (CTAs). Although self-antigens also elicit a tumor immune response, the primary target of the immune response is neoantigens, also termed tumor-specific antigens (TSAs). Neoantigens are specific to tumor cells and arise from somatic mutations in cancer genomes [16]. The recognition of tumor neoantigens may promote T-cell priming and infiltration and can lead to a long-term clinical response [11, 16].

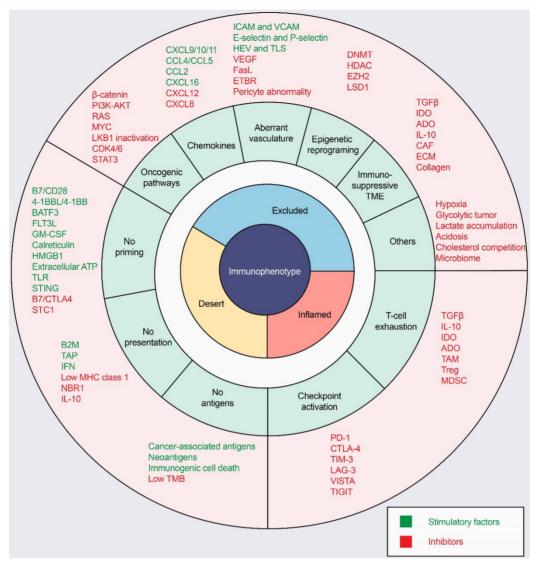


Figure 3. Mechanisms of three distinct tumor phenotypes. Three different phenotypes are associated with specific biological mechanisms. Tumors with the immune-desert phenotype (yellow) may lack T-cell priming due to the absence of tumor antigens, defective antigen processing and presentation machinery, or impaired DC-T-cell interactions. Tumors with the immune-excluded phenotype (blue) may exhibit activation of oncogenic pathways, aberrant chemokines, aberrant vasculature and hypoxia, or an immunosuppressive tumor microenvironment (e.g., stromal barriers). Tumors with the immune-inflamed phenotype (red) can be infiltrated by many immune cells, but these immune cells are suppressed due to checkpoint activation. ADO: adenosine; ATP, adenosine triphosphate; B2M: beta-2-microglobulin; BATF3: basic leucine zipper ATF-like transcription factor 3; CAFs: cancer-associated fibroblasts; CRT, calreticulin; CTLA4, cytotoxic T lymphocyte-associated antigen-4; CXCL: CXC-chemokine ligand; DNMT: DNA methyltransferase; ECM: extracellular matrix; ETBR: endothelin B receptor; EZH2: enhancer of zeste homolog 2; FLT3L: Fms-like tyrosine kinase 3 ligand; GM-CSF: granulocyte-macrophage colony-stimulating factor; HDAC: histone deacetylase; HEV: high endothelial venule; HMGBI: high mobility family protein B1; ICAM: intercellular adhesion molecule; IDO: Indoleamine 2,3-dioxygenase; IFN: interferon; IL: interleukin; MDSC: myeloid-derived suppressor cell; MHC: major histocompatibility complex; PD-I, programmed cell death protein I; PD-LI, PD-I ligand; STCI: stanniocalcin I; TAM: tumor-associated macrophage; TAP: transporter associated with antigen processing; TGFβ: transforming growth factor-β; TIM3, T cell immunoglobulin and mucin domain-containing 3; TLR: Toll-like receptor; TLS: tertiary lymphoid structure; TME: tumor microenvironment; Treg: T-regulatory cell; VCAM: vascular cell adhesion molecule; VEGF: vascular endothelial growth factor.

The TMB is broadly characterized as the number of total nonsynonymous single-nucleotide mutations in a tumor. In general, tumors with a higher TMB are believed to carry a higher neoantigen load that can be recognized by T cells, making them more likely to prime the immune system [17]. Significant associations between high TMB and improved response to ICIs have been reported in a variety of tumor types [18, 19]. The TMB has been used as a novel biomarker to predict the efficacy of programmed cell death protein 1 (PD-1) inhibitors [17, 19]. Consistent with the importance of the efficacy of

ICIs, a high TMB was associated with greater immune cell infiltration [20]. Furthermore, a mutliomics network analysis revealed that in tumors characterized by recurrent mutations, such as melanoma and colorectal cancer, mutation or neoantigen burden was positively correlated with CTL infiltration [21]. Considering the relationship between high TMB and tumor-specific T cells, low mutational load or neoantigen load contributes, in part, to the lack of immune infiltration and the acquisition of ICI resistance. However, in tumors characterized by recurrent copy number alterations,

such as breast cancer, a correlation between tumor-specific T-cell infiltration and neoantigen load is lacking [21]. An investigation of data on 266 melanomas in The Cancer Genome Atlas (TCGA) revealed no difference in antigen expression between cold and hot tumors [22]. This finding suggests that other mechanisms, in addition to those involving low TMB, contribute to the absence of T-cell infiltration.

Defects in tumor antigen processing and presentation machinery (APM)

After recognizing tumor antigens, APCs process the antigens and express the corresponding antigen peptide-MHC class I complex on its surface. However, alterations in the APM, such as downregulation of MHC-I molecule expression or the absence of beta-2-microglobulin (B2M), limit the presentation of antigen peptide-MHC class I complexes in the presence of tumor antigens. During antigen processing and presentation, transporters associated with antigen processing (TAP) transport cytosolic cleaved antigens to the endoplasmic reticulum for binding to the MHC. The deletion in TAP is related to defects in the antigen presentation process, which further affects the priming of T lymphocytes [23]. B2M, the invariant chain of the MHC, is critical for the successful folding and transport of MHC-I to the cell surface [5]. Knocking down the B2M gene in the M202 and M233 human melanoma cell lines resulted in the absence of MHC-I molecules expressed on their surface, and the absence of tumor-specific T-cell recognition and cytotoxicity [24]. Identical results were observed in a B2M-knockout mouse model of lung cancer that showed resistance to PD-1 blockade [25]. In addition, the lysosomal pathway has been linked to the reduced infiltration of CD8+ T lymphocytes. In pancreatic ductal adenocarcinoma (PDAC), the autophagyassociated receptor NBR1 induces the degradation of MHC-I on the cell surface of tumor cells, which in turn affects T-cell responses [26]. These findings suggest that defects in tumor antigen processing and presentation pathways inhibit T-cell priming and the effectiveness of cancer immunotherapies.

Dysfunctional DC-T cell interactions

Dendritic cells (DCs) are professional APCs with the unique ability to acquire antigens, migrate to secondary lymphoid organs (e.g., lymph nodes and spleen), and initiate the *in vivo* immune response. DC activation requires that pattern recognition receptors (PRRs) on their surface recognize "danger signals", including pathogen-associated molecular patterns (PAMPs) and damage-associated molecular patterns (DAMPs) [27]. This recognition enables DCs to

present a tumor antigen peptide-MHC class I complex to T cells upon contact with them. DCs also express costimulatory signals such as B7 (including CD80 and CD86), providing the secondary signaling necessary for T-cell activation [28]. Tumor cells can mediate diminished phagocytosis of DCs by trapping "danger signals". For example, stanniocalcin 1 (STC1), an intracellular checkpoint, can trap DAMPs (e.g., calreticulin (CRT)) and inhibit DC phagocytosis and T-cell activation, contributing to tumor immune escape. Furthermore, STC1 is associated with low T-cell activation and poor survival in melanoma patients [29].

DCs are generally classified into two broad categories: plasmacytoid DCs (pDCs) characterized by the production of IFN-α, and conventional DCs (cDCs), which effectively stimulate T-cell proliferation [30]. CDCs are further categorized into two distinct subsets: BATF3-dependent DCs and IRF4-dependent DCs [30]. BATF3 DCs have the ability to cross-present tumor-derived antigens through the MHC-I pathway and thus initiate T cells [31]. Furthermore, BATF3 DCs are the primary source of CXC-chemokine ligand 9 (CXCL9) and CXCL10, two key chemokines required to recruit CD8+ T cells expressing CXCR3 to tumors. There is a significant correlation between BATF3 DC markers (e.g., BATF3 and IRF8), the expression of CXCL9, CXCL10, and CXCL11 and the CD8+ effector T-cell phenotype in melanoma [22, 32]. In the absence of BATF3 DCs, CD8+ effector T cells fail to migrate to tumors and antitumor immunity is thus defective [32]. This finding validates the notion that BATF3 DCs may be essential for the priming and recruitment of the endogenous T cells necessary to counteract tumors.

The regulation of Fms-like tyrosine kinase 3 ligand (FLT3L) and granulocyte-macrophage colonystimulating factor (GM-CSF) is important for the differentiation and recruitment of DCs [33]. FLT3L is a growth factor that promotes the differentiation of hematopoietic progenitor cells from the bone marrow to the DC lineage [33]. Tumor-derived FLT3L increased the infiltration of BATF3 DCs and CD8+ T lymphocytes in mouse tumors and enhanced migratory and resident DC subsets in draining lymph nodes (DLNs), suggesting a mobilizing effect of FLT3L on DC cells [34]. Deficiency of FLT3L or GM-CSF resulted in a reduced number of DCs in secondary lymphoid organs and attenuated T-cell immune responses [35]. Given the important role of DC-T cell crosstalk in naïve T-cell priming, impaired DC activation, a lack of DCs, and the overexpression of cosuppressive signals can lead to impaired T-cell activation.

Table 2. Mechanisms of deficient T-cell homing to the tumor bed

Mechanisms	Examples	References
Oncogenic Pathway	WNT/β-catenin activation	[32, 36]
Activation	Loss of PTEN	[38]
	RAS activation	[39, 138]
	MYC activation	[40]
	LKB1 inactivation	[42]
Aberrant chemokine	Absence of CXCL9 and CXCL10	[36]
	Epigenetic regulation	[48-50]
	CXCL12 overexpression	[52]
Aberrant	Downregulation of adhesion molecules	[55-57]
Vasculature	(ICAM, VCAM, P-and E-selectin)	
	VEGF overexpression	[59]
	FasL upregulation	[60]
	Pericyte abnormality	[56, 61]
Hypoxia	HIF-1	[64-66]
Immunosuppressive	TGFβ overexpression	[72, 73]
TME	ADO, IDO	[68, 78]
	CAF	[82-84]
Metabolic	High glycolytic activity	[90-93]
competition	Lactate accumulation and acidification	[95, 96]
	Cholesterol competition	[98, 99]

Deficient T-cell homing to the tumor bed

Oncogenic pathway activation

For the purposes of this review, we summarize the mechanisms by which T cells are prevented from homing to the tumor bed (Table 2). There is growing evidence showing that the activation of tumor cell oncogenic pathways is related to the "cold tumor" phenotype and the potential for immunotherapy resistance. In WNT/ β -catenin-positive melanoma tumors, reduced production of CCL4 results in decreased recruitment of BATF3 DCs to the TME [36]. Ultimately, in the absence of the CXCL9 and CXCL10 produced by BATF3 DCs, CTLs are not recruited to the tumor [32]. An analysis of human metastatic melanoma samples showed a negative correlation between CD8A expression and activation of the β-catenin signaling pathway [36]. Direct injection of BATF3 DCs helped restore T-cell infiltration in β-catenin-positive tumors and resulted in modest tumor suppression [36]. This outcome suggests that WNT/β-catenin signaling activation and defective BATF3 DC recruitment mediate T-cell exclusion and tumor cell escape from the immune system. Notably, activation of this oncogenic pathway excludes CTLs only when β -catenin is located into the nucleus [37]. This finding indicates that the exclusion mechanism of CTLs is related to a transcriptional program specifically induced by β -catenin. In addition, only 48% of "cold" melanomas show active β-catenin signaling, suggesting that other oncogenic pathways may mediate immune exclusion [36].

Loss of PTEN activates the PI3K/AKT pathway, which is related to a noninflamed T-cell phenotype and immune resistance of melanoma. Loss of PTEN expression has been found to reduce the lipidation of

the autophagosome protein LC3, resulting in decreased autophagic activity, which inhibits T-cell priming and the T-cell-mediated antitumor response [38]. CD8+ T-cell infiltration in PTEN-deficient melanoma was significantly reduced compared to that in PTEN-expressing tumors. The results from a TCGA dataset analysis indicated that the expression of T-cell effector molecules (e.g., IFN-γ and granzyme B) was significantly reduced in melanomas with low PTEN expression [38].

As the gene with the most common mutations associated with cancer progression, RAS can lead to the activation of multiple signaling pathways, such as MAPK and PI3K, driving tumorigenesis [39]. In addition, oncogenic K-RAS mutations mediate inflammation and crosstalk with the TME. For example, oncogenic K-RAS mutations induce tumor-promoting inflammation through the production of inhibitory cytokines (e.g., IL-6 and IL-8), the activation of NLRP3 inflammasome, and the release of chemokines (e.g., CCL5 and CCL9) [39].

Furthermore, oncogenic signaling through MYC enhances the expression of CD47 and PD-L1 on tumor CD47 binds to inhibitory receptor signal-regulated protein-a (SIRPa) on the surface of APCs such as macrophages and DCs, which can prevent phagocytosis of tumor cells and interfere with antigen uptake [31, 40]. Oncogenic KRAS and MYC synergistically induce immune regulation. For example, co-activation of KRAS and MYC in a mouse lung cancer model leads to the production of CCL9 and IL-23. This mediates stromal reprogramming, promotes angiogenesis, and excludes T and B cells and NK cells from tumors [41]. It has also been found that inactivating mutations of LKB1 in non-small cell lung cancer (NSCLC) are related to increased neutrophil and decreased T-cell infiltration in a preclinical mouse model [42]. In addition, CDK4/6 and STAT3 activation is associated with a noninflamed T-cell phenotype [43-45]. Taken together, these results reveal that the activation of oncogenic pathways can affect not only tumor cells but also T cell-mediated antitumor immunity.

Chemokines and their epigenetic regulation

The interaction between some chemokine receptors on effector T lymphocytes and corresponding chemokines may affect the trafficking of effector T lymphocytes to tumor sites. The lack of several chemokines, including CXCL9, CXCL10, CCL4, CCL5, CXCL16 or CX3CL1, has been reported to lead to T-cell exclusion [46, 47]. Considering the importance of the TH1-type chemokines CXCL9 and CXCL10 to T-cell recruitment, certain tumors show low levels of CXCL9 and CXCL10 expression, which

may explain the reduced infiltration of effector T lymphocytes into these tumor beds [32, 36]. For example, BATF3 DCs are the major sources of CXCL9 and CXCL10, and a lack of BATF3 DCs leads to low expression of CXCL9 and CXCL10. In addition, epigenetic regulation in tumors is also important for maintaining low expression levels of these cytokines. DNA methyltransferase (DNMT) and enhancer of zeste homolog 2 (EZH2) can mediate DNA methylation and histone lysine methylation, respectively, to suppress the expression of CXCL9 and CXCL10 in ovarian cancer [48]. Similar results have been confirmed in colon cancer [49]. In preclinical models, treatment with epigenetic modulators promoted tumor infiltration of effector T cells and enhanced the effect of anti-PD-L1 [48]. In addition to CXCL9 and CXCL10, CCL5 expression is positively related to CD8+ T-cell infiltration [21, 47]. The binding of CCL5 to CCR5 promotes the recruitment of CD8+ T cells. However, DNA methylation leads to deletion of CCL5 expression, which in turn contributes to the absence of CD8⁺ T-cell infiltration [50]. In mouse models of NSCLC, the combined use of DNMT inhibitors and histone deacetylase (HDAC) inhibitors increased the expression of endogenous retrovirus (ERV), which in turn induced type I IFN responses. This combination treatment reversed the immune resistance of NSCLC models by downregulating oncogenic MYC signaling, leading to an increase in CCL5 and promoting T-cell infiltration into tumors

However, some chemokines are detrimental to the trafficking of T cells to tumor beds. Stromal cells, especially cancer-associated fibroblasts (CAFs), are the main producers of CXCL12. CXCL12 produced by CAFs misdirects CTLs to the extratumoral stroma and prevents CTLs from entering the tumor [52]. Furthermore, elevated CXCL8 expression has been reported to be associated with a reduction in the number of T cells in tumors, increased neutrophil and monocyte infiltration, and limited responses to ICIs [53, 54]. These results reveal the regulatory effect of chemokine receptor and ligand interactions on CTL homing to tumors and their integration into the TME.

Aberrant vasculature and hypoxia

Adequate T-cell infiltration in the tumor bed is not only dependent on the recruitment of the appropriate chemokines but is also controlled by the tumor vasculature. During the trafficking of CD8+ T lymphocytes to a tumor, they must enter the tumor circulatory system, adhere to vascular endothelial cells and migrate across the vessel wall [11]. The recruitment of CD8+ T cells to tumors requires the action of vascular endothelial adhesion molecules,

including P-and E-selectin, intercellular adhesion molecules (ICAMs), and vascular cell adhesion molecules (VCAMs) [55, 56]. However, downregulation or ineffective aggregation of adhesion molecules on tumor endothelial cells leads to endothelial cell anergy and reduced effector T-cell trafficking to tumor sites [55, 57]. Endothelin binds to a corresponding receptor on endothelial cells, endothelin B receptor (ETBR), and reduces ICAM-1 production, thereby inhibiting CD8+ T cell adhesion to endothelial cells [58]. Additionally, endothelial growth factor (VEGF), produced by tumor and stromal cells, stimulates the proliferation of endothelial cells, leading to new vessel formation, often accompanied by impaired tissue perfusion and increased vascular permeability [59]. VEGF also decreases the expression of important molecules, such as VCAM-1, on the cell surface of the endothelium, ultimately preventing T cells from migrating to the TME [59]. Another mechanism through which tumor endothelial cells can inhibit T-cell migration is modulation of immune cell activity or viability. IL-10, prostaglandin E2 (PGE2), and VEGF induced FasL upregulation in tumor endothelial cells to kill tumor-associated T cells, and anti-FasL attenuated this killing effect. Acetylsalicylic acid (ASA), which inhibits COX and PGE2 activity, and anti-VEGF antibodies promoted CD8+ T lymphocyte infiltration in the TME and improved prognoses Furthermore, pericyte abnormalities and inadequate coverage prevent the maintenance of endothelial cell integrity, resulting in the dysfunctional leakage and flow characteristic of tumor vasculature [56, 61]. However, other structures can promote the translocation of CD8+ T lymphocytes from blood vessels to tumor sites. The formation of high endothelial venules (HEVs) and related tertiary lymphoid structures (TLSs) facilitates T-cell migration to the TME and is often associated with better prognoses [46, 62].

In addition, impaired vascular tight junctions and increased permeability result in the promotion of hypoxia, acidosis and necrosis, which inhibit immune effector T-cell functions and antitumor immunity [56]. As a hallmark of cancer, hypoxia is caused by increased oxygen demand due to tumor cell proliferation and inadequate blood supply due to angiogenesis [63]. Hypoxia-inducible factor 1 (HIF1) is a key transcription factor activated by hypoxia [64]. Hypoxia inhibits T-cell infiltration in several ways. hypoxia promotes the recruitment of immunosuppressive cells to the TME [65]. Second, hypoxia-induced CCL28 and VEGF promote angiogenesis and affect T-cell trafficking [56, 66]. Finally, the expression of two ectonucleotidases,

CD39 and CD73, can be upregulated in tumors in response to hypoxia and transforming growth factor-β (TGFβ) [67]. CD39 and CD73 catalyze the sequential conversion of ATP to extracellular adenosine (ADO) [68]. ADO binds to the adenosine A2A receptor (A2AR) and inhibits the production of cytokines such as IL-2 and the development and proliferation of T cells [69]. The inhibition of A2AR increased T-lymphocyte infiltration and led to improved tumor control in mouse melanoma models, suggesting a potential effect of the ADO signaling pathway in promoting T-cell exclusion [70]. In addition, ADO can weaken antitumor immunity by inhibiting the effector functions of NK cells and DCs and by promoting the recruitment and polarization of MDSCs and Tregs [71].

The TME: immunosuppressive cells and factors

The immunosuppressive microenvironment at including dense tumor sites, stroma immunosuppressive cells and factors, can prevent T-cell priming and infiltration in "cold tumors". TGFβ is a potent immunosuppressive cytokine that promotes immune escape and blocks the acquisition of the TH1-effector phenotype [72]. CAFs, which are abundant in the TME, are the main producers of TGFβ. Increased TGFβ production by CAFs is associated with T-cell exclusion from the tumor and a poor response to atezolizumab [73]. TGFβ limits the proliferation of CD4+ T lymphocytes by inhibiting the production of IL-2 and induces the conversion of naïve CD4+ T lymphocytes into Tregs [74, 75]. TGFβ also negatively affects DC differentiation and antigenpresenting functions, which interfere with T-cell priming [76]. In summary, TGFβ hinders antitumor immunity by affecting T-cell differentiation and function and preventing T-cell infiltration into tumors.

Tryptophan metabolism is often dysregulated in a broad range of cancers and is associated with immune resistance. Indoleamine 2,3-dioxygenase (IDO) in tumor cells converts the essential amino acid tryptophan into kynurenine, which blocks the priming of T lymphocytes and facilitates the development of Tregs [77]. IDO also recruits and activates MDSCs and inhibits the accumulation of tumor-specific T lymphocytes in tumors [78]. IDO inhibitors such as epacadostat and navoximod have been used in combination with ICIs with promising results in clinical trials [79]. However, the failure of epacadostat in combination with pembrolizumab in the phase III clinical ECHO-301 study indicates that the effectiveness of drugs targeting IDO needs to be further considered [80].

CAFs are key cellular components in the tumor stroma and can promote tumor growth [81]. CAFs are predominantly located at the infiltrating edges of tumors, regulating tumor metastasis and influencing angiogenesis by synthesizing and remodeling the extracellular matrix (ECM) and producing cytokines, and transforming tumor margins into immune "cold" zones [52, 82]. CAFs led to immunosuppression and T-cell exclusion through several mechanisms. First, CAFs produce extracellular matrix that forms a physical barrier to prevent T-cell infiltration into the tumor area [83]. Second, CXCL12 produced by CAFs has been shown to inhibit T-lymphocyte infiltration within tumors in a pancreatic cancer model [84]. Third, CAFs can also reduce T-cell responses and exert immunosuppressive effects through production of TGFβ and IL-6 [82]. Reprogramming CAFs is an effective strategy to "normalize" the TME. This strategy reduces ECM levels, decompresses blood vessels, and increases the degree of T-cell penetration to improve cancer treatment [85].

In addition, TAMs exclude T cells from the tumor by regulating the ECM and mediating the nitration of CCL2 and CCL5 [86, 87]. TAMs affect T-cell recruitment by promoting abnormal angiogenesis through the production of VEGF and matrix metalloproteinase-9 (MMP9) [88]. Cytokine colony-stimulating factor-1 (CSF-1) and CSF-1R interactions are capable of promoting myeloid cell differentiation towards an immunosuppressive M2 macrophage phenotype. Targeting TAMs with CSF1R inhibitors reduces the number of TAMs and increases the infiltration of effector lymphocytes such as CD8+T cells [89].

Tumor cells are typically characterized by a high rate of glucose uptake and active glycolysis, even in the presence of oxygen. This phenomenon is known as the "Warburg effect". In this process, glucose is rapidly consumed and the abundance of lactate in the TME increases. The glucose-deficient, lactate-rich TME exerts metabolic stress on infiltrating T cells, leading to local immunosuppression and ICI resistance [77]. Glucose deprivation in the TME metabolically mediates T cell hyporeactivity, inhibits mTOR activation, and reduces glycolytic capacity and IFN-y production [90]. In addition, glycolytic activity and T-cell infiltration are negatively correlated in a variety of tumors [91-93]. Consistent with this observation, high glucose-transporter 1 (GLUT-1) expression in renal cell carcinoma are associated with low infiltration of CD8+ T cells [92]. These results suggest an association of glycolytic (Warburg) tumors with a noninflamed T-cell phenotype. Interestingly, in addition to tumor cells, stromal cells, such as CAF and TAM, can also promote lactate accumulation in the TME through the so-called "Reverse Warburg effect" [91]. Targeting glucose metabolism and lactate production in tumor and stromal cells, such as inhibition of LDH-A, may be an effective strategy to promote T-cell infiltration [94]. Lactate accumulation and acidification of the TME suppress antitumor immunity. Lactate-induced acidosis impairs the differentiation of monocytes to DCs and inhibits the antigen-presenting function of DCs, which in turn inhibits T-cell activation [95]. High concentrations of lactate and acidification in the TME inhibit monocarboxylate transporter 1 (MCT1)-mediated lactate release from T cells and suppress the proliferation of T cells that utilize aerobic glycolysis [96]. In addition, lactate inhibits the chemotaxis and antitumor activity of CTLs and promotes tumor immune escape [96]. Inhibition of lactate production or restoration of physiological pH of the TME can reverse the inhibitory effect of lactate on antitumor immunity. For example, neutralizing tumor acidity with sodium bicarbonate in combination with ICIs or adoptive cellular therapy (ACT) can effectively promote T-cell infiltration and improve antitumor responses in a variety of mouse tumor models [97].

In addition to glucose, metabolic competition between tumors and immune cells includes amino acids and fatty acids. For example, the high rate of cholesterol esterification in tumors inhibits T cell receptor (TCR) aggregation and immune synapse formation [77]. The cholesterol esterification key enzyme ACAT1 inhibitor avasimibe can promote the proliferation of CD8+ T cells and exhibit good antitumor effects [98]. New studies have also confirmed that inhibition of PCSK9, a key protein regulating cholesterol metabolism, upregulates MHC-I levels on the surface of tumor cells, increases intratumoral infiltration of CTLs, and synergistically inhibits tumor growth with anti-PD1 antibodies [99]. Considering the interaction between metabolism and immune cell metabolism, navigating metabolic pathways to reduce metabolic stress on T cells is a promising strategy to improve the efficacy of immunotherapy.

Therapeutic approaches to drive T cells into tumors

ICIs have revolutionized cancer treatment by activating T-cell-based antitumor immunity. However, a significant number of patients show a poor response to ICIs due to the multiple mechanisms mediating T-cell exclusion. Several approaches have been shown to drive T cells into tumors. These approaches "fire up" "cold tumors" to improve the efficacy of ICIs (Figure 4 and Table 3).

Table 3. Examples of therapeutic approaches to drive T cells into tumors

Main mechanismsTherapeutic approachesReferencesPromotes T-cell primingImmune adjuvants (TLR agonists, STING agonists)[102, 105]Oncolytic viruses[109, 111]Chemotherapy and radiotherapy[114, 119]Epigenetic modification inhibitors (DNMT inhibitor, HDAC inhibitor, EZH2 inhibitor)[140-142]Metabolic intervention[94, 97, 99]Local thermal ablation therapy (Radiofrequency ablation)[123]Photothermal therapy and photodynamic therapy (Ragher approaches)[165, 166]High-intensity focused ultrasound[167, 168]Promotes T-cell expansionAdoptive cellular therapy (TILs, CAR-T cells) vaccines[34, 125, 126]Promotes T-cell trafficking and infiltrationOncogenic pathway inhibitors[38, 130, 138]Epigenetic modification inhibitors[48, 49, 51]Antiangiogenic therapy (anti-VEGF)[146]TGFβ inhibitors[72, 73, 148]			
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Therapeutic approaches to promote T-cell priming

Immune adjuvants

Innate immune sensing pathways play critical roles in the development of antitumor immunity. The PRR family includes Toll-like receptors (TLRs), NOD-like receptors (NLRs), RIG-I-like receptors (RLRs), and C-type lectin receptors (CLRs) [27]. When TLRs are stimulated, DCs can produce a variety of proinflammatory cytokines, including tumor necrosis factor (TNF), IL-1, and type I IFNs [27]. DCs are the main sources of type I IFNs, which facilitate the expression of MHC-I on the surface of tumor cells and the maturation of DCs, thereby promoting T-cell priming [100].

In contrast to therapies such as vaccines and CAR-T cells, immune adjuvants harness the endogenous antigen repertoire in the tumor and have been used to enhance the immune response for the treatment of malignant tumors. Local administration of the TLR7/8 agonist imidazoquinoline with coupled nanoparticles significantly activated the DCs of secondary lymphoid organs, upregulated the expression of MHC-II, CD40, and CD86 on their surface, and expanded the number of tumor-specific CD8+ T lymphocytes, which inhibited tumor growth [101]. In clinical trials of patients with advanced malignant melanoma, combined treatment with the TLR9 agonists SD-101 and pembrolizumab resulted in increased type I IFN production and CD8+ T-cell infiltration and potentially improved clinical efficacy [102].

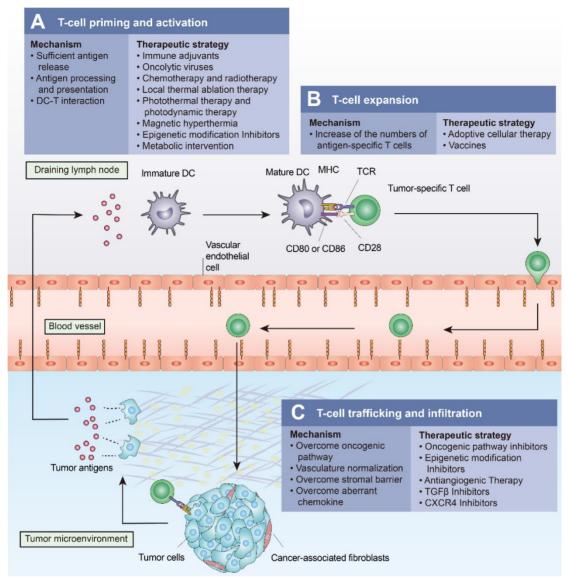


Figure 4. Approaches to turn a "cold tumor" into a "hot tumor". Some representative approaches that lead to increased T-cell infiltration and improved efficacy of immune checkpoint inhibitors are highlighted here. (A) Oncolytic viruses, local thermal ablation therapy (e.g., radiofrequency ablation), chemotherapy, and radiotherapy are all capable of inducing immunogenic cell death (ICD) to promote T-cell priming and activation. Local administration of immune adjuvants such as TLR agonists promotes the activation of dendritic cells (DCs). Epigenetic modification inhibitors can promote T-cell priming by increasing the expression of tumor antigens and by restoring antigen processing and presentation mechanisms. (B) Cancer vaccines and adoptive cellular therapies, such as CAR-T cells, can promote the expansion of tumor-specific T lymphocytes. (C) Intrinsic oncogenic pathway inhibitors, epigenetic modification inhibitors, antiangiogenic therapies, TGFβ inhibitors, and CXCR4 inhibitors promote T-cell trafficking and enable T cells to infiltrate the tumor more effectively.

As DNA receptors in the cytoplasm, cyclic GMP-AMP synthase (cGAS) in DCs, macrophages, and other immune cells recognizes aberrant DNA in the cytoplasm and catalyzes the formation of cGAMP, which subsequently activates the STING signaling pathway [103]. Activation of the STING signaling pathway mediates the expression of proinflammatory cytokines (e.g., type I IFNs) and chemokines (e.g., CXCL10) in a TBK1-IRF3-dependent manner, thereby initiating the antitumor immune response [104]. The systemic cGAMP mimetic SR-717 activated the STING signaling pathway, promoted CD8⁺ T-cell, NK cell, and CD8α⁺ DC activation, and significantly inhibited tumor growth [105]. SR-717 also induced PD-L1

expression in a STING-dependent manner, revealing the significance of the combination of STING agonists and ICIs for tumor treatment. In mouse tumor models with a low response to PD-1 blockade, the combination of PD-1 blockade and the STING agonist MSA-2 increased the infiltration of tumor CD8⁺ T lymphocytes and better inhibited tumor growth [106].

Oncolytic viruses (OVs)

OVs are now being recognized as emerging therapeutics with potent anticancer activity. In addition to selective tumor lysis, they can activate both innate and adaptive immune responses, resulting in alterations in the TME. First, lysis of

tumor cells by OVs induces immunogenic cell death (ICD), leading to a massive release of intracellular TAAs, PAMPs, and DAMPs [107]. Three DAMPs are released during ICD: passively released high mobility family protein B1 (HMGB1), actively secreted extracellular ATP, and cell surface-expressed CRT [108]. These DAMPs act as adjuvants to promote DC uptake and cross-present tumor antigens to T lymphocytes in DLNs. OVs also improve the function of DCs by stimulating their production of type I IFNs. Immune adjuvants interact with tumor antigens in tumor residues and act as individualized in situ vaccines to promote T-cell priming [109]. Second, OVs stimulate the production of CXCL9 and CXCL10 and upregulate the expression of selectins and integrins, providing key signals for T-cell trafficking. In addition, the degradation of the ECM by OVs disrupts the physical barrier to T-cell infiltration [109]. OVs can also deplete the immunosuppressive effects of CAFs, TAMs and MDSCs, significantly altering the TME [109].

Talimogene laherparepvec (T-VEC) was first shown to be effective as an oncolytic virotherapy for melanoma [110]. Combination therapy with T-VEC and pembrolizumab increased CD8+ T-lymphocyte infiltration, IFN-γ expression and the therapeutic effect of PD-1 blockade in patients with advanced malignant melanoma [111]. Similar effects have been observed with combination therapy of coxsackievirus and pembrolizumab [112]. These studies suggest that combination therapy with OVs and ICIs can improve CD8+ T-cell infiltration and activation and help to overcome the resistance of cancer to ICIs in patients. Combined with T-cell therapy, promoting T-cell proliferation and infiltration into the local TME is a potential direction for the development of OVs.

Chemotherapy and radiotherapy

It was previously thought that chemotherapy and radiotherapy exert their antitumor effects by directly killing tumor cells. However, accumulated evidence suggests that tumor suppression by chemotherapy and radiotherapy also relies on stimulating the immune system. When radiotherapy is administered to a local tumor, distant tumors outside the irradiated field also shrink. This phenomenon is termed the "abscopal effect", which indicates the significance of the immune system in radiotherapy-mediated antitumor responses [113]. After radiation therapy causes damage to tumor cells, ROS and endoplasmic reticulum (ER) stress mediate cellular stress and lead to ICDs [114]. This cascade promotes DC activation, increases the production of TNFa and IL-1 and produces endogenous cancer vaccines in vivo [114]. After radiation therapy,

endothelial cells express ICAM1, VCAM1, and E-selectin, which facilitate the attraction of immune cells [114]. Radiation also promotes the trafficking of effector T lymphocytes to the tumor site by inducing tumor cells to express and release chemokines (e.g., CXCL10 and CXCL16) [115]. However, considering the adverse effects of radiation therapy, it is necessary to optimize the radiation dose and fractionation levels during radiation therapy. Fractionated radiotherapy at individual doses of less than 8-10 Gy helps to induce sufficient ICD without increasing hypoxia or immunosuppression, inducing a de novo antitumor response [115]. In preclinical studies, stereotactic body radiotherapy (SBRT) increased effector infiltration in tumors and DLNs and was associated with higher survival rates [116].

addition to radiotherapy, many chemotherapeutic agents can exert their immunostimulatory effects by enhancing immunogenicity and increasing T-cell infiltration [117]. ICD inducing chemotherapy has been shown in various mouse models to contribute to the transformation of "cold tumors" to "hot tumors" in response to ICIs [118-120]. New immunotherapies that reduce the toxic side effects of systemic chemotherapy and enhance the immunogenicity induced by chemotherapeutic agents deserve further exploration. A new cocktail therapy involves local chemoimmunotherapy by mixing chemotherapeutic agents with immune adjuvants and alginate (ALG). The ICD-inducing chemotherapeutic agents produced tumor vaccines in situ in response to adjuvant stimulation. In situ gelation of the drug adjuvant ALG enables slow release of the drug, thereby reducing systemic toxicity. The combination of ICIs further amplifies the immune response and inhibits tumor metastasis and recurrence [121].

Local thermal ablation therapy

Image-guided thermal ablation has been developed as a promising method for the treatment of solid tumors. Currently, the commonly used thermal ablation methods include radiofrequency ablation (RFA), laser ablation (LA), microwave ablation (MWA), and high-intensity focused ultrasound (HIFU) ablation. RFA is widely used in the treatment of solid tumors, especially hepatocellular carcinoma (HCC). RFA utilizes the conversion of radiofrequency alternating current into heat to ablate the tissue surrounding the needle electrode and stimulate tumor-specific T-cell responses [122]. However, problems with incomplete ablation and tumor recurrence are drawbacks of RFA. The combination of the tyrosine kinase inhibitor sunitinib and RFA improves HCC treatment. RFA caused the release of TSA in situ in tumors but caused the upregulation of

PD-1 expression in T lymphocytes, which was related to the exhaustion of CD8+ T lymphocytes. Sunitinib inhibition of hepatocyte growth factor (HGF) inhibited the upregulation of PD-1 expression in tumor T lymphocytes. Inhibition of VEGF by sunitinib also promoted DC activation and inhibited tumor angiogenesis. Combination therapy ultimately led to a remarkable increase in CD8+ T lymphocytes and DCs as well as a decrease in Tregs, thus overcoming the drawbacks of monotherapy [123]. In addition, HIFU, as a new minimally invasive ablation therapy, has become a hotspot of cancer treatment. HIFU delivers acoustic energy to the target tissues in a noninvasive and precise manner, generating high temperatures that cause coagulative necrosis of the tumor tissue. HIFU promotes ICD and the activation of T cells [124]. HIFU also promotes antigen transfer to lymph nodes and T-cell migration to tumors through mechanical destruction of the mesenchyme [124].

Therapeutic approaches to increase the number of antigen-specific T cells

Adoptive cellular therapy (ACT)

ACT enhances the immune response of effector T including tumor-infiltrating cancer, lymphocytes (TILs) and CAR-T cells [6]. In TIL therapy, tumor-infiltrating lymphocytes are isolated from cancer patients, expanded in vitro, and then reinfused into the patient. However, due to the small infiltrating lymphocytes of downregulation of MHC molecules, TILs are only used for the treatment of a few tumor types, such as malignant melanoma [125]. CAR-T cells involve genetic modification of T lymphocytes to express the CAR to target tumor cells expressing a specific antigen [6]. For example, CD19-specific CAR-T cells have become the gold standard for the treatment of B-cell malignancies. In addition, CAR-T cell therapy for leukemia and lymphoma was approved by the FDA in 2017 [6]. In contrast to TILs, CAR-T cells are not limited by the MHC and can further enhance the immune response to tumors through the addition of costimulatory domains (e.g., CD28, OX40, and 4-1BB) [6]. This strategy leverages the direct recognition of tumor antigens by CAR-T cells and has the potential to treat "cold tumors" lacking pre-existing T-cell infiltration. CAR-T cells expressing IL-7 and CCL19 increased DC and T-cell infiltration in mouse solid tumor tissues and showed potent antitumor effects [126]. In addition, the use of FLT3L-secreting CAR-T cells and immune adjuvants led to similar results and induced host T-cell antigen epitope spread [34]. However, CAR-T cells exhibit limited clinical efficacy in solid tumors due to tumor antigen

heterogeneity and insufficient infiltration into tumors [6]. Additionally, the immunosuppressive microenvironment limits the tumor-killing effect of CAR-T cells, which makes it necessary to combine CAR-T cells with ICI therapy.

Cancer vaccines

Therapeutic vaccines such as peptide and tumor cell vaccines, nucleotide vaccines encoding new epitopes, and dendritic cell vaccines have been encouraging clinical advances. Therapeutic vaccines expand the pool of tumor-specific T cells, increase the transport of T lymphocytes to tumor areas, and have become emerging modalities for immunotherapy [127]. Sipuleucel-T, the first therapeutic cancer vaccine licensed by the FDA, has been used in the treatment of castration-resistant prostate cancer. Sipuleucel-T consists of the fusion protein PA2024 constructed from both prostatic acid phosphatase (PAP) and GM-CSF and autologous DCs, which enhance the antitumor effect [128]. neoantigens are highly tumor-specific immunogenic. The combination of the personalized neoantigen vaccine NEO-PV-01 with nivolumab significantly prolonged progression-free survival in an Ib clinical trial for patients with advanced malignant melanoma, NSCLC and bladder cancer. A neoantigen-specific T-cell response, T-cell trafficking to tumors and induction of epitope spread were also observed [129]. The combination of the mRNA personalized cancer vaccine RO7198457 atezolizumab showed clinical benefit in a phase Ib clinical trial for patients with advanced solid malignancies (NCT03289962). The vaccine induced a neoantigen-specific T-cell response in 77% of patients. This outcome demonstrates that the use of a personalized cancer vaccine combined with an immune checkpoint blockade can generate a specific immune response in patients. However, the complex operation, cumbersome process and high price remain limiting factors for the widespread use of personalized cancer vaccines in cancer treatment.

Therapeutic approaches to promote T-cell trafficking and infiltration

Oncogenic pathway inhibitors

The use of oncogenic pathway inhibitors helps reverse the inherent T-cell exclusion from tumors. PAK4 is abundantly expressed in "cold tumors" and plays a key role in the WNT/ β -catenin pathway. Knocking down PAK4 or applying the PAK4 inhibitor KPT-9274 in a mouse tumor model enhanced CTL infiltration in tumors and improved the therapeutic efficacy of a PD-1 blockade [130]. However, the

efficacy of treatments targeting WNT remains controversial. For example, endogenous inhibitors of the WNT pathway, such as some proteins of the dickkopf (DKK) family, have a role in promoting tumor immune escape and are associated with a poorer prognosis in some cancers [131, 132]. DKK2 inhibits WNT-β-catenin signaling by binding to the cell surface receptors LRP5 and LRP6 of the WNT pathway [133]. DKK2 expression is upregulated in human colorectal cancers (CRCs) and promotes tumor progression by inhibiting the activation of NK cells and CD8+ T cells [131]. This challenges the notion that inhibition of the WNT pathway will improve immunotherapy. Furthermore, recent studies have shown that activation of the WNT pathway in endothelial cells promotes T-cell infiltration into tumors and enhances the effectiveness immunotherapies such as the ACT, suggesting that there is still a need to further investigate the feasibility of using WNT inhibitors as immune adjuvants [134]. Indeed, clinical data using inhibitors of the WNT/β-catenin pathway do not support completely its putative function to boost immunotherapy in the

Activation of the PI3K-AKT pathway is associated with the inhibition of CTL infiltration and function. PI3K β inhibitors inhibit the activation of the AKT pathway in PTEN-deficient melanoma cell lines and enhance T-cell-mediated killing [38]. The combination of PI3K β inhibitors and ICIs significantly increased the number of infiltrating T lymphocytes in murine tumor models [38].

RAS was considered as an "undruggable" target in the past [135]. Monotherapies targeting the RAS oncogene have faced limited efficacy due to multiple mechanisms, such as feedback reactivation of the RAS downstream pathway [39]. However, recent studies have found that ARS-1620, a small-molecule inhibitor specifically targets KRAS-G12C mutants, significantly inhibited the growth of KRAS-G12C tumors [136]. This evidence suggests a new therapeutic avenue to inhibiting mutant RAS. In addition, MEK inhibitors in combination with ICIs led to an increase in tumor-infiltrating T cells and a decrease in the percentage of MDSCs, which in turn significantly inhibited tumor growth in TP53/KRASdriven lung cancer mouse models [137]. Oncogenic mutations of BRAF activate the RAF-MEK-ERK (MAPK) pathway. Inhibition of BRAF or MEK inhibits the production of inhibitory cytokines (e.g., IL-6, IL-10, and VEGF) or enhances the expression of differentiation antigen, melanocyte thereby promoting melanoma recognition by T cells [138, 139]. Three variant kinase inhibitors targeting MEK, cobimetinib. and binimetinib, trametinib,

clinically approved for therapeutic use in BRAF V600 mutant melanoma [135].

CDk4/6 can bind to cyclin D, which enables cells to enter S-phase through the RB-EF2 pathway and promotes tumor cell proliferation. Treatment of CT26 syngeneic mouse tumors with the CDK4/6 inhibitor abemaciclib led to an increase in tumor-infiltrating T lymphocytes, and a significant upregulation of T-cell activity, as evidenced by the increased expression of T-lymphocyte activation markers (e.g., IFNG, GZMB, CCL4 and CCL5). Abemaciclib also led to enhanced antigen presentation and had a synergistic effect when applied with anti-PD-1 therapy [44].

Epigenetic modification inhibitors

Epigenetic therapies can transform tumors from the immune "cold" state to the immune "hot" state through a variety of mechanisms. Epigenetic drugs can enhance the expression of multiple chemokines, such as CXCL9, CXCXL10, and CCL5, and promote T-cell trafficking to tumors [48, 49, 51]. Epigenetic therapy can also induce ERVs and suppress MYC signaling, thereby enhancing the expression of type I IFNs and related chemokines [51]. In addition, epigenetic therapies can enhance tumor immunogenicity by increasing the expression of tumor antigens such as CTA and by restoring MHC-I antigen processing and presentation mechanisms [140, 141]. The DNMT inhibitor guadecitabine upregulated MHC-I expression in breast tumor cells, enhanced IFN-y responses, and promoted T-cell recruitment to tumors. In addition, guadecitabine had a synergistic effect with an anti-PD-L1 antibody [142]. This outcome suggests the feasibility of combining epigenetic inhibitors with ICI strategies for future clinical application. A variety of epigenetic drugs have been approved by the FDA, such as azacitidine and decitabine (DNMT inhibitors), tazemetostat (an EZH2 inhibitor), and entinostat and vorinostat (HDAC inhibitors) [143].

Antiangiogenic therapy

Persistent angiogenesis caused dysregulated balance between pro-and angiogenic signals is one of the hallmarks of tumors [144]. Antiangiogenic therapy (AT) structurally and functionally overcomes tumor vascular abnormalities, improves tissue perfusion, and increases the infiltration of immune effector cells. AT-mediated immune reprogramming in turn improves vascular normalization, thereby creating an enhanced positive feedback loop [56, 145]. Bevacizumab is the first FDA-approved angiogenesis inhibitor. Increased infiltration of tumor-specific T lymphocytes was observed after combination therapy with bevacizumab and atezolizumab in patients with metastatic renal cancer. In addition, combination therapy resulted in downregulation of the expression of vascular signature genes (e.g., ANGPT2 and CD31) and upregulation of CD8+ T effector genes (e.g., CD8A, GZMB, and IFNG) and MHC-I, as well as chemokines (e.g., fractalkine) [146]. These results imply that the increase in tumor-specific T-cell infiltration may be due to enhanced lymphocyte trafficking mediated by the combination therapy. In addition, in the phase III IMbrave150 trial for patients with unresectable hepatocellular treatment with atezolizumab in combination with Bevacizumab significantly improved overall survival and progression-free survival outcomes compared with the results of sorafenib treatment [147]. Given the relationship between vascular normalization and immune reprogramming, combination therapy is expected to further reverse the immunosuppressive microenvironment and improve the efficacy of immunotherapy.

TGF_β inhibitors

Considering the immunosuppressive function of TGF β , therapy based on the inhibition of TGF β has been validated as an effective approach to promote T-lymphocyte infiltration. TGF β is related to a lack of immune response in the noninflamed T-cell phenotype. In a mammary cancer mouse model with immune-excluded phenotype, combined treatment with anti-PD-L1 and anti-TGF\$\beta\$ antibodies significantly reduced tumor burden and increased tumor-infiltrating T cells, especially CD8+ T effector cells [73]. Galunisertib, a small molecule that inhibits the activity of TGFBR1 kinase, has been the most widely tested compound. Galunisertib treatment increased T-cell infiltration and improved susceptibility to checkpoint therapy in a mouse colorectal model [72]. TGFβ impedes the generation of in situ tumor vaccines after radiotherapy. Treatment with the 1D11 antibody, which blocks systemic TGF β activity, enhanced the initiation of T-cell responses to endogenous tumor antigens after subcutaneous tumor irradiation [148].

CXCR4 inhibitors

CXCR4 is a receptor for CXCL12, which is overexpressed in a wide range of tumors. The CXCL12-CXCR4 axis plays an indirect role in the sequestration of CTLs from the tumor area to reduce CTL infiltration and mediates the infiltration of immunosuppressive cells into tumors [52]. In the PDAC model, inhibition of CAF-mediated CXCL12/CXCR4 axis with the CXCR4 inhibitor AMD3100 promoted T-cell accumulation and cancer regression

[84]. Previous studies have shown that immunotherapies such as pembrolizumab were not effective against "cold tumors" such as pancreatic cancer. However, in the COMBAT trial, the synergistic treatment of metastatic PDAC with the CXCR4 antagonist BL8040 and pembrolizumab increased tumor-infiltrating CD8+ effector T lymphocytes, reduced the density of MDSCs in tumors, and reduced the number of circulating Tregs [149]. These results reveal that regulating certain chemokines the homing of tumor-specific lymphocytes to the tumor and reverses immune resistance.

New cancer therapies based on nanotechnology

Nanomedicine

A variety of nanomedicines combined with immunotherapy contribute to the transition from "cold tumors" to "hot tumors" (Figure 5). Nanomedicines have three different targeting pathways: tumor cells, the TME, and the peripheral immune system [150]. Nanomedicine-based targeted tumor therapy includes passive and active targeting. Passive targeting can promote selective accumulation of nanomedicines in tumors through EPR effects [151]. However, recent studies have suggested that the passive targeting ability of nanomaterials may be associated with transcytosis [152]. Active targeting involves the use of targeted ligands (e.g., peptides, antibodies and transferrin) that specifically recognize specific receptors expressed by tumor cells [15].

Nanomedicines targeting tumor cells can induce ICD and enhance the tumor-immunity cycle [150, 153]. For example, doxil, a PEGylated liposome of the chemotherapeutic drug doxorubicin, promoted DC and CD8+ T cell proliferation via ICD and inhibited Treg infiltration [154]. Doxil was also synergistic with ICIs (anti-PD-1 and anti-CTLA-4) and showed higher efficacy than free doxorubicin [154]. When targeting the TME, nanomedicines inhibit immunosuppressive (e.g., M2TAMs and MDSCs) immunosuppressive molecules (e.g., TGFβ and ADO), and they can also augment the activity and function of effector immune cells (e.g., macrophages and CTLs) [150, 155]. The TGF β receptor inhibitor SB525334 was loaded onto liposomes targeting ACT T cells. This nanomedicine inhibited TGFB expression and promoted T-cell activation as well as tumor regression in melanoma mice [156]. Loading liposomes with IL-2 agonistic anti-4-1BB enhanced the tumor infiltration of CTLs as well as cytokine production and granzyme expression [157]. When targeting the peripheral immune system, nanomedicines are

designed to promote antigen presentation and CTL production in DLNs. Nanomedicine-based vaccines improved antigen delivery to lymph nodes, promoted antigen cross-presentation, and increased CTL activation levels [150, 158, 159]. Furthermore, nanomedicines have been engineered to directly promote T-cell priming by replacing APCs in secondary lymph nodes. The use of biomimetic magnetosomes as artificial APCs was characterized by magnetic nanoclusters encapsulated by leukocyte membranes and modified to stimulate signals on the membranes. Artificial APCs not only expanded and stimulated CTLs, but also guided reinfused CTLs efficiently into tumor tissues, thus inhibiting tumor growth [160].

Tumor phototherapy (PT)

PT, including photothermal therapy (PTT) and photodynamic therapy (PDT), has been developed as a potential treatment for solid tumors, especially superficial tumors. PTT uses photothermal agents (e.g., organic nanoparticles, gold nanoparticles and graphene oxide) with photothermal conversion capabilities to absorb near-infrared (NIR) lasers and

convert them into heat energy to kill tumor cells [15]. PDT involves the use of a laser to irradiate the tumor into which photosensitizers had been delivered to activate the photosensitizers and produce cytotoxic ROS. It can cause DNA damage in the nucleus and thus induce tumor cell death [15]. Compared to other ablation modalities (e.g., radiofrequency ablation and microwave ablation), PT is more selective and less toxic to surrounding tissues due to the accumulation of photosensitizers in the tumor and the controllability of the light.

PTT and PDT induce ICD through thermal and chemical damage, respectively, and enhance the infiltration of CTLs and the immunotherapeutic response [15]. PDT based on a layer-by-layer Apt/PDGs^s@pMOF nanoplatform enhanced the immunogenicity of triple-negative breast cancer in mice, selectively suppressed MDSCs, and promoted the transition to "hot tumors" [161]. However, a single PT treatment shows limited efficacy because of limited light penetration, a heat-shock response due to the PTT, and the dependence of PDT on oxygen. The combination of PDT and PTT can achieve promising synergistic antitumor effects. A hybrid

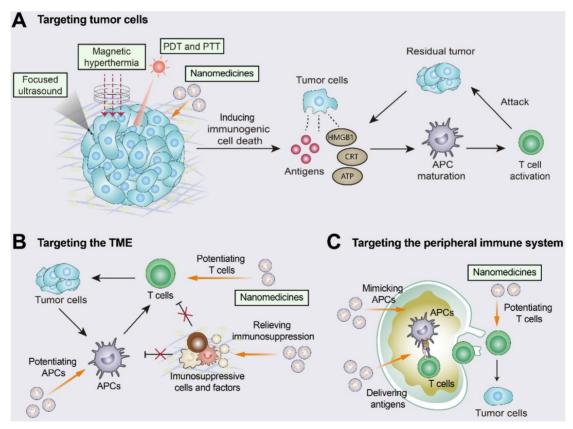


Figure 5. Improving T-cell infiltration with nanomedicines. Nanomedicines have three different targeting pathways: tumor cells, the TME, and the peripheral immune system. (A) Multiple approaches including photothermal therapy (PTT), photodynamic therapy (PDT), magnetic hyperthermia (MH), and high-intensity focused ultrasound (HIFU) can induce ICD by promoting the release of tumor antigens and damage-associated molecular patterns (DAMPs). The released DAMPs act as adjuvants to enhance the immunogenicity of the tumor and, together with the released tumor antigens, promote dendritic cell (DC) activation and T-cell priming. (B) When targeting the TME, nanomedicines inhibit immunosuppressive cells and immunosuppressive molecules (e.g., TGFβ) and enhance the activity of T cells. (C) When the peripheral immune system is targeted, nanomedicines are engineered to augment tumor antigen presentation and T-cell priming in lymph nodes. Adapted with permission from [150], copyright 2019 American Chemical Society.

nanoporphyrin (Pp18-Lips)-mediated synergistic PTT/PDT caused an increase in tumor-infiltrating CTLs and inflammatory cytokines (e.g., TNF-a and IFN-y) and a decrease in Tregs. Synergistic PDT/PTT produced a stronger antitumor immune response and stronger tumor suppression than PDT or PTT alone [162]. PT is also synergistic with immunotherapy, as PT enhances the immunogenicity of the tumor by mediating ICD, while immunotherapy enhances the "abscopal effect" of the treatment. PTT-mediated GOP@aPD-1 nanoparticles efficiently delivered anti-PD-1 to melanoma cells and combined ICI treatment with tumor-targeted PTT. This combination resulted in increased CD8+ T-cell infiltration, elevated efficacy of the PD-1 blockade in mouse melanoma, and inhibition of tumor growth [163].

Magnetic hyperthermia (MH)

MH refers to the selective heating of tumors by converting magnetic energy into thermal energy through the hysteresis and relaxation effects of magnetic nanoparticles (MNPs) in the presence of an alternating magnetic field (AMF) [164]. In addition to killing tumor cells with heat, MH also induces ICD, showing great potential in cancer therapy. Compared to PT and thermal ablation therapy, MH has no penetration depth limit and enables more effective targeting and more precise control of the heating temperature. The iron oxide nanomedicine ferumoxytol promoted the polarization macrophages into the proinflammatory phenotype. This switch induced the apoptosis of tumor cells, as indicated by increased caspase-3 Another typical example cleavage [165]. ferrimagnetic vortex-domain iron oxide nanorings (FVIOs), which have excellent nanomagnetic properties. FVIO-mediated mild MH induced the expression of CRT on the surface of 4T1 breast tumor cells and promoted T-cell activation. FVIOs caused a significant increase in CTL infiltration along with a decrease in MDSCs and were synergistic with anti-PD-L1 antibodies [166].

High-intensity focused ultrasound (HIFU)

The combination of HIFU and immunotherapy achieved significant therapeutic efficacy. Compared to immunotherapy alone or HIFU alone, the combination of HIFU and the TLR agonist CpG in mouse melanoma enhanced antigen cross-presentation in DLNs, release of type I IFNs, and expression of genes relevant to T-cell priming and stimulation (e.g., Eomes, Prf1 and Icos) [167]. However, HIFU remains under-researched and there is a lack of studies demonstrating tumor control with HIFU monotherapy [124]. The current state of the

field suggests that HIFU can promote T-cell priming and tumor regression, but induction of additional immune adjuvants may be necessary. The killing of tumor cells by HIFU has also been linked to cavitation effects, which can efficiently aggregate energy. A recent study showed that direct injection of microbubbles and plasmid DNA encoding IFN- β into tumors in a mouse model followed by application of low-frequency ultrasound (250 kHz) to break up the tumors was shown to remove a substantial number of tumor cells and simultaneously achieve massive infiltration of CTLs. The remaining tumor cells also formed membrane pores, allowing gene transfection of the cells and triggering antitumor immune responses [168].

Conclusions and future perspectives

Considering the relevance of T-lymphocyte infiltration in tumor sites to the prognosis of ICI therapy, the cause of the absence of a T-cell response is crucial to determine. In this review, we summarize various mechanisms that inhibit T-cell infiltration, such as defects in tumor antigen processing and presentation endogenous oncogenic processes, pathway activation, aberrant vasculature chemokines, and TME suppression. Other known and unknown factors affecting T-cell infiltration, such as TLSs and the microbiome, remain to be evaluated. TLSs have a lymph node-like function and are relevant to T-cell infiltration into tumors and a good response to immunotherapy [169, 170]. Exploring therapeutic strategies to enhance TLS formation and function may promote the activation of naïve T lymphocytes by DCs in close proximity to the tumor and improve the response to cancer immunotherapy. Compared to "cold tumors", "hot tumors" are more responsive to ICI monotherapy. Thus, promoting the conversion of "cold tumors" to "hot tumors" through interventions can help reduce resistance to ICIs. In addition, we then discuss a variety of therapeutic measures to improve T-cell infiltration, such as oncogenic pathway inhibitors, anti-vascular therapy, ACT, vaccines, oncolytic viruses, and cytotoxic therapies. The combination of ICIs with these therapies reverses T-cell exhaustion, enhances the "abscopal effects" of therapy, and demonstrates incremental clinical efficacy. However, the optimal dose and sequence of administration of combination therapy needs to be further evaluated to optimize T-cell function, promote T-cell memory, and avoid overactivation [171]. In addition, some issues still need to be addressed, such as the nonspecific distribution of drugs, and the treatment-induced systemic adverse effects.

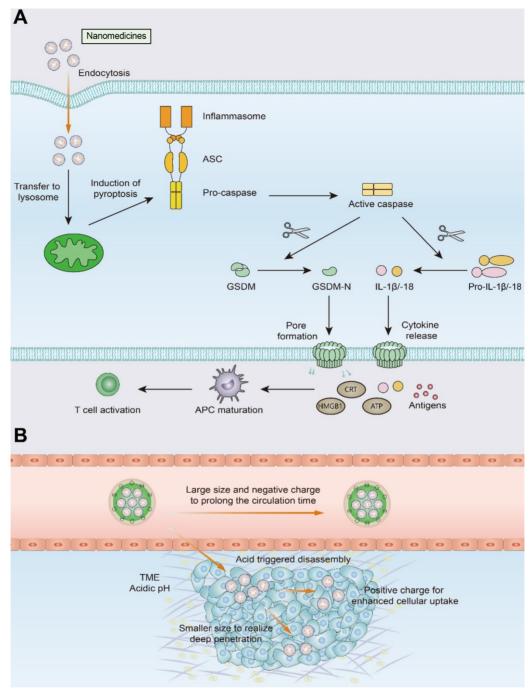


Figure 6. Schematic illustration of pyroptosis and size-transformable nanoparticles. (A) Multiple nanomedicines regulate the expression of caspase proteins that mediate the pyroptosis process. Activated caspases cut gasdermin (GSDM) into two fragments: the C-terminal domain and the N-terminal domain. Following cleavage, the gasdermin-N domains result in cell swelling with big bubbles. Gasdermin-induced pyroptosis results in the release of a massive quantity of proinflammatory molecules and activation of T cells. (B) Use of size-transformable nanoparticles to prolong the circulation time and realize deep penetration.

With the development of nanotechnology, the nanomedicine and biomaterial-assisted local therapies offers new opportunities for the future. The use of nanomedicines improves drug precision and bioavailability, reduces immunotherapy-induced side effects, and enables selective accumulation of drugs in tumors through EPR effects and active targeting. As mentioned above, PTT, PDT, MH, and HIFU are all capable of inducing a *de novo* antitumor response, which is achieved through the induction of ICD.

These approaches do not require prior knowledge of tumor antigens and induce the generation of endogenous personalized *in situ* vaccines. However, killing solid tumors by noninflammatory apoptosis or ablation does not make tumor cells sufficiently immunogenic. In contrast to apoptosis, pyroptosis is a proinflammatory form of cell death that leads to the release of a massive quantity of inflammatory molecules (e.g., IL-1 β and IL-18), mobilizing a robust antitumor T-cell response (Figure 6) [172-174]. The use

of nanotechnology to induce pyroptosis increases the immunogenicity of tumor cells and may effectively improve T-cell infiltration in tumors [172]. For designed Zhao et al biomimetic nanoparticles (BNPs) loaded with indocvanine green (ICG) and decitabine for photoinduced pyroptosis. Due to promoter methylation of the GSDME gene, GSDME expression is much lower in most tumor cells than in normal cells. As an inhibitor of DNA methylation, decitabine promotes caspase-3 cleavage to GSDME by upregulating GSDME expression, thus leading to tumor cell pyroptosis. Pyroptosis mediated by BNP results in the release of a large number of inflammatory molecules from tumor cells and induces DC maturation and T-cell activation in DLNs, demonstrating a robust immune response against primary and distant tumors [175].

However, certain challenges to nanomedicine use must be overcome, such as short blood circulation time, and insufficient penetration and accumulation in tumor tissues. The use of size-transformable nanoparticles in phototherapy or chemotherapy may achieve deep penetration of nanomedicines and improve T-cell infiltration into tumors [176, 177]. Nanocarriers with relatively large particle sizes utilize the EPR effect to prolong the circulation time of the nanomedicine and improve its accumulation in tumor Upon reaching the tumor site, nanomedicine undergoes size transformation in response to pH or enzymes and releases transformed small nanoparticles that exhibit effective tumor tissue penetration and can be thus efficiently internalized by tumor cells.

In addition, the use of small molecular weight nanobodies in diagnostic imaging allows for a more convenient and complete assessment of the extent of T-cell infiltration in the TME, providing new ideas for achieving the integration of diagnosis and treatment of tumors. A better understanding of these aspects will be beneficial for guiding personalized cancer immunotherapy and extending the benefits of ICI therapy to a broader group of patients.

Abbreviations

A2AR: A2A receptor; ADO: adenosine; ALG: alginate; AMF: alternating magnetic field; APC: antigen presenting cell; APM: antigen processing and presentation machinery; ASA: Acetylsalicylic acid; AT: Antiangiogenic therapy; ATP: adenosine triphosphate; B2M: beta-2-microglobulin; BNPs: biomimetic nanoparticles; CAFs: cancer-associated fibroblasts; CAR: chimeric antigen receptor; CDCs: conventional DCs; cGAS: cyclic GMP-AMP synthase; CLRs: C-type lectin receptors; CRCs: colorectal cancers; CRT: calreticulin; CSF-1: colony-stimulating

factor 1; CTAs: cancer/testis antigens; CTLA4: cytotoxic T lymphocyte-associated antigen-4; CTLs: cytotoxic T lymphocytes; CXCL: CXC-chemokine ligand; DAMPs: damage-associated molecular patterns; DCs: dendritic cells; DKK: dickkopf; DLNs: draining lymph nodes; DNMT: DNA methyltransferase; ECM: extracellular matrix; EPR: enhanced permeability and retention; endoplasmic reticulum; ERV: endogenous retrovirus; ETBR: endothelin B receptor; EZH2: enhancer of zeste homolog 2; FLT3L: Fms-like tyrosine kinase 3 ligand; FVIOs: ferrimagnetic vortex-domain iron oxide nanorings; GLUT-1: glucose-transporter 1; GM-CSF: granulocyte-macrophage colony-stimulating factor; GSDM: gasdermin; HCC: hepatocellular carcinoma; HDAC: histone deacetylase; HEVs: high endothelial venules; HGF: hepatocyte growth factor; HIF1: Hypoxia-inducible factor 1; HIFU: high-intensity focused ultrasound; HMGB1: high mobility family protein B1; ICAMs: intercellular adhesion molecules; ICD: immunogenic cell death; ICG: indocvanine green; ICIs: immune checkpoint inhibitors; IDO: Indoleamine 2,3-dioxygenase; IFN-y: interferon-y; IL: interleukin; LA: laser ablation; MCT1: monocarboxylate transporter 1: MDSCs: myeloid-derived suppressor cells; MH: magnetic hyperthermia; MHC: major histocompatibility MMP9: metalloproteinase-9; magnetic nanoparticles; MWA: microwave ablation; NIR: near-infrared; NLRs: NOD-like receptors; NSCLC: non-small cell lung cancer; OVs: oncolytic viruses; PAMPs: pathogen-associated molecular patterns; PAP: prostatic acid phosphatase; PD-1: programmed cell death protein 1; PD-L1: PD-1 ligand; PDAC: pancreatic ductal adenocarcinoma; PDCs: plasmacytoid DCs; PDT: photodynamic therapy; PGE2: prostaglandin E2; PRRs: pattern recognition receptors; PTT: photothermal therapy; radiofrequency ablation; RLRs: RIG-I-like receptors; SBRT: stereotactic body radiotherapy; SIRPa: signal-regulated protein-α; STC1: stanniocalcin 1; TAAs: tumor-associated antigens; TAMs: tumor-associated macrophages; TAP: transporter associated with antigen processing; TCGA: the Cancer Genome Atlas; TCR: T cell receptor; TGFβ: transforming growth factor-β; TILs: tumor-infiltrating lymphocytes; TLRs: Toll-like receptors; TLSs: tertiary lymphoid structures; TMB: tumor mutational burden; TME: tumor microenvironment; TNF: tumor necrosis factor; Tregs: T-regulatory cells; TSAs: tumor-specific antigens; T-VEC: talimogene laherparepvec; VCAMs: vascular cell adhesion molecules; VEGF: vascular endothelial growth factor.

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Competing Interests

The authors have declared that no competing interest exists.

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