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REVIEW

The impact of lipids on the cancer–immunity cycle and strategies for modulating lipid metabolism to improve cancer immunotherapy



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Abstract Lipids have been found to modulate tumor biology, including proliferation, survival, and metastasis. With the new understanding of tumor immune escape that has developed in recent years, the influence of lipids on the cancer–immunity cycle has also been gradually discovered. First, regarding antigen presentation, cholesterol prevents tumor antigens from being identified by antigen presenting cells. Fatty acids reduce the expression of major histocompatibility complex class I and costimulatory factors in dendritic cells, impairing antigen presentation to T cells. Prostaglandin E2 (PGE2) reduce the accumulation of tumor-infiltrating dendritic cells. Regarding T-cell priming and activation, cholesterol destroys the structure of the T-cell receptor and reduces immunodetection. In contrast, cholesterol also promotes T-cell receptor clustering and relative signal transduction. PGE2 represses T-cell proliferation. Finally, regarding T-cell killing of cancer cells, PGE2 and cholesterol weaken granule-dependent cytotoxicity. Moreover, fatty acids, cholesterol, and PGE2 can improve the activity of immunosuppressive cells, increase the expression of immune checkpoints and promote the secretion of immunosuppressive cytokines. Given the regulatory role of lipids in the cancer–immunity cycle, drugs that modulate fatty acids, cholesterol and PGE2 have been envisioned as effective way in restoring antitumor immunity and synergizing with immunotherapy. These strategies have been studied in both preclinical and clinical studies.

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1. Introduction

Lipids comprise a complex group of biomolecules that differ in structure and function, and they have been classified into eight categories as follows: fatty acids (FAs), glycerolipids, glycerophospholipids, sphingolipids, sterols, prenols, saccharolipids, and polyketides^{1,2}. Lipids execute important physiological roles in energy storage and the modulation of membrane fluidity, as well as various in trafficking and signaling events^{3–7}. Among the various categories of lipids, FAs, cholesterol, and prostaglandins, which are arachidonic acid derivatives, have been shown to be critical in regulating tumor proliferation, survival, and metastasis^{8–11}. Increased fatty acid oxidation (FAO) and *de novo* lipid synthesis often occur in tumor and provide extra energy for tumor development and progression¹². Prostaglandin E2 (PGE2) enhances intestinal adenoma growth *via* activation of the Ras-mitogen-activated protein kinase cascade¹³. PGE2 can cause decreased expression of programmed cell death in human colonic cancer cells by increasing NF- κ B expression¹⁴. Cholesterol content within specific membrane regions can affect extrinsic (death receptor pathway) and intrinsic (mitochondrial) apoptotic pathways¹⁵. Moreover, cancer cells promote migration and invasion by balancing FA saturation levels¹⁶. With increased levels of cyclooxygenase-2 (COX-2)/PGE2, cancer cells become more migratory during epithelial–mesenchymal transition^{17,18}.

In addition to its role in mediating biological characteristics of tumor cells, a large amount of evidence indicates that lipids alter the function and status of immune cells in the tumor microenvironment (TME)^{19–22}. Lipids, especially cholesterol, FAs and prostaglandins, can affect any part of the cancer immune cycle, thereby disrupting normal antitumor immunity (Fig. 1). Regarding antigen presentation, cholesterol changes the formation of tumor-associated antigens (TAAs), which enables tumor cells to escape immune surveillance²³. FAs, especially saturated palmitic acid and short chain fatty acids (SCFAs), inhibit antigen presentation. They decrease the expression of major histocompatibility complex class I (MHC I) and costimulatory factor^{24,25}. During T-cell priming and activation, cholesterol keeps T-cell receptors (TCR) in a resting conformation by binding to the TCR β transmembrane region²⁶. In contrast, cholesterol facilitates the formation of a larger TCR signalosome and promote T-cell signal transduction as part of membrane rafts²⁷. In addition, PGE2 inhibits T-cell activation and cytotoxicity by downregulating interleukin 2 (IL-2), interferon- γ (IFN- γ) and granzyme B^{28–30}. Regarding T-cell killing of cancer cells, FAs enhance the immunosuppressive function of myeloid-derived suppressor cells (MDSCs) and PGE2 promotes the infiltration of regulatory T cells (Tregs)^{31,32}. Furthermore, FAs and PGE2 upregulate extrinsic immunosuppressive cytokines, such as interleukin 10 (IL-10) and transforming growth factor- β (TGF- β)^{33,34}. FAs and cholesterol can also promote the expression of immune checkpoints^{34,35}. Taken together, these data suggest that the abnormal accumulation of lipids hinders the establishment of a complete immune cycle and helps to establish an immunosuppressive environment. Strategies

that decrease the level of lipids can improve antigen presentation, promote T-cell priming and activation and disrupt the immunosuppressive state.

FAs, cholesterol, and PGE2 exhibit negative effects in the adaptive immune cycle. Therefore, recent preclinical experiments and clinical trials have tested the combination of lipid-based therapy and immunotherapy as a promising cancer treatment strategy^{36–40}. The aim is to block FA synthesis and absorption through methods including the inhibition of the activities of fatty-acid synthase (FASN), fatty acid transport protein 2 (FATP2), and peroxisome proliferator-activated receptors (PPARs). PPARs are molecular sensors of FA and play a crucial role in regulating FAO and FA storage⁴¹. Statins are the most common pharmacological intervention option for cholesterol regulation. They are inhibitors of hydroxyl methylglutaryl-coenzyme-A (HMG-CoA), the key catalytic enzymes in the rate-limiting step of cholesterol biosynthesis⁴². PGE2 synthesis is governed by COX-2, and PGE2 interacts with the E prostanoïd receptor (EPR) to activate downstream signals. Inhibitors of COX-2 and EPR can modulate the content and function of PGE2.

Given the importance of the complete immune cycle in the suppression of tumors and the profound impact of lipids on the immune system, here, we present a review describing how lipids affect the cancer adaptive immunity cycle and how lipid-based therapy regulates immune responses in cancer patients treated with immunotherapy. Moreover, we discuss potential innovative strategies for targeting immune cells with the modulation of lipids to develop potential combination immunotherapy targets.

2. Lipid-mediated cancer–immunity cycle: Cancer antigen presentation

Antigen presentation mainly includes two distinct events: tumor antigens are displayed on the cancer cell surface, and antigen presenting cells (APCs) take up antigens and cross-present to T cells. Lipids can act as TAAs directly to affect the immune response during initial step. Lipids can also modulate antigen presentation through APCs (Fig. 2).

Glycosphingolipids (GSLs) are widely present on tumor cells and cancer stem cells as a kind of TAAs that elicit an immune response in the host^{43,44}. In cancer cell plasma membranes, cholesterol can bind to GSLs to reorient GSL carbohydrate to a membrane parallel, rather than perpendicular conformation⁴⁴. Therefore, the GSL–cholesterol complex restricts tumor-associated GSL immunoreactivity and leads to ineffective immune surveillance and passive immunity in various tumors, such as breast carcinoma, prostate cancer, and colon carcinoma⁴⁴.

DCs are the most potent APCs and initiate the majority of adaptive immune responses, but DCs in tumors usually exert immunosuppressive phenotypes. Saturated palmitic acid treatment was found to significantly reduce MHC I expression and lower APC–T-cell conjugation, which in turn inhibited the ability of APCs to active T cells. However, continued monounsaturated oleic acid treatment can normalize antigen presentation by

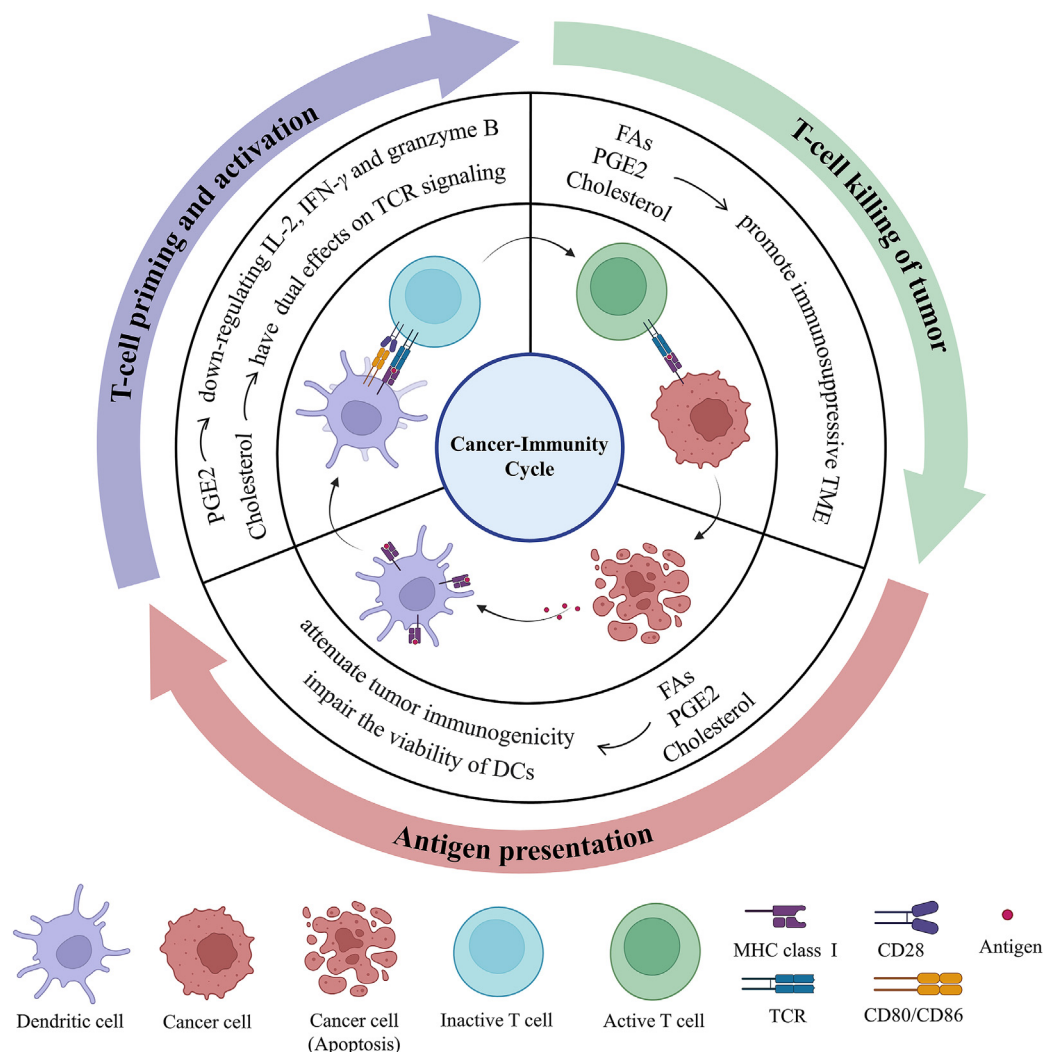


Figure 1 Lipids mediate the cancer–immunity cycle. The complete adaptive immune cycle is crucial for tumor immunotherapy, and mainly includes the following three components: (1) antigen presentation, (2) T-cell priming and activation, and (3) T-cell killing of tumors. Lipids, especially fatty acids, cholesterol, and prostaglandins, mainly have a negative effect on the cancer–immunity cycle. Combination drugs of modulating lipids with immunotherapy methods can potentially further improve the antitumor immune response in patients.

sequestering palmitic acid into triglyceride-rich lipid droplets²⁴. The study showed that saturated and unsaturated FAs exhibit differential regulation of antigen presentation and costimulatory molecule expression, so selective inhibition of FAs is needed. In addition, SCFAs, as butyrate and propionate, reduce the anti-cytotoxic T lymphocyte antigen 4 (CTLA4)-induced increase in the levels of the costimulatory molecules CD80 and CD86 on DCs and the inducible co-stimulator (ICOS) on T cells, which attenuates the therapeutic effect of anti-CTLA4 in metastatic melanoma²⁵. Due to the increased FA uptake mediated by macrophage scavenger receptor 1, triglyceride levels are elevated in peripheral blood DCs in non-small lung cancer and renal cell carcinoma patients. DCs with increased levels of triglycerides exhibit impaired capacity to process and present TAAs, and a significantly lower ability to stimulate T cells, but the mechanism is not clear⁴⁵. Other studies revealed that DCs containing high lipid levels not only exhibited lower MHC expression, but also lower costimulatory expression than DCs containing relatively low lipid level⁴⁶. In cancer stem cells, Arf1 ablation causes the inhibition of

lipolysis, which facilitates the engulfment of tumor-associated antigens on dendritic cells (DCs) and promotes DC antigen presentation, thereby enhancing T-cell infiltration and activation. Moreover, Arf1 ablation can efficiently synergize with programmed cell death 1 (PD-1) blockade to inhibit tumor growth⁴⁷. PGE2, which is derived from arachidonic acid, both reduced the accumulation of tumor-infiltrating dendritic cells (TIDCs), thus reducing antigen uptake, and inhibit TIDC maturation within TME. Such a TME induced CD8⁺ T-cell tolerance upon migration to the tumor draining lymph nodes in murine renal cell carcinoma²⁸.

Based on the above studies, it is speculated that modifying lipid metabolism can restore the vitality of DCs, which may be a promising anticancer immunotherapy. Blocking lipid synthesis with an acetyl-CoA carboxylase inhibitor restored the capacity of DCs and substantially enhanced the effects of cancer vaccines⁴⁵. Moreover, the activity of FASN, a key enzyme involved in *de novo* lipogenesis, can be blocked by cerulenin, which can restore TIDC capacity to extend control of ovarian cancer⁴⁶.

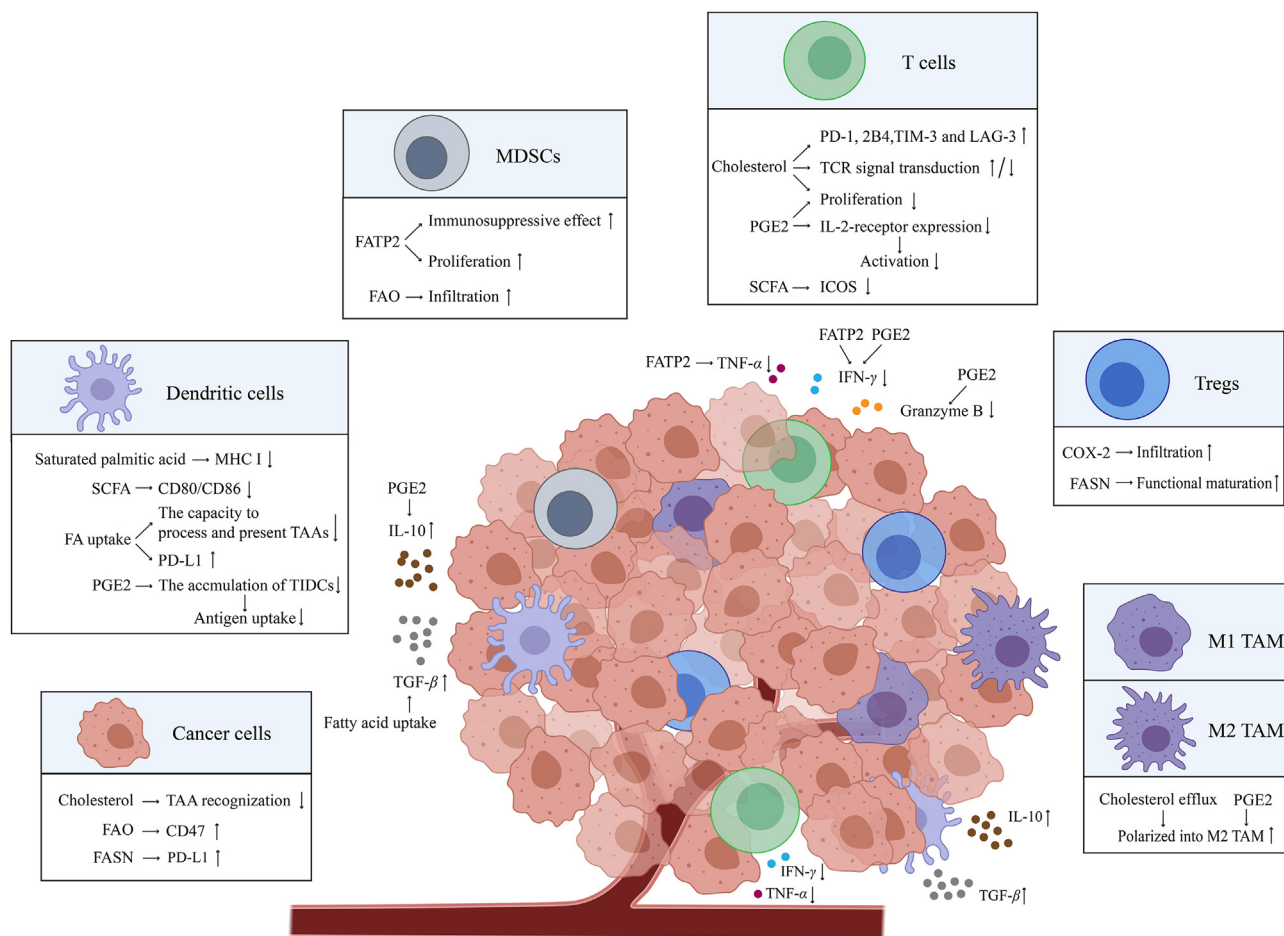


Figure 2 Lipids reshape tumor microenvironment. Lipids mainly regulate cancer cells, immune cells, and cytokines to form an immunosuppressive tumor microenvironment, which inhibits the immune response and promotes tumor immune evasion. In cancer cell plasma membranes, cholesterol binds to glycosphingolipids, a kind of TAA, to change their conformation so that they cannot be recognized. Moreover, FAO-derived acetyl-CoA upregulates CD47 transcription in glioblastoma multiforme to weaken macrophage phagocytosis. FASN stabilizes the PD-L1 protein by promoting PD-L1 palmitoylation. In DCs, saturated palmitic acid significantly reduces MHC I expression and lowers the conjugation of APCs and T cells and SCFA reduces the expression of costimulatory molecules CD80 and CD86 on DCs and ICOS on T cells during anti-CTLA4 treatment. Increased FA uptake reduces the ability of DCs to process and present antigens and increases the expression of PD-L1. PGE2 reduces the accumulation of TIDCs, which inhibits antigen uptake. In MDSCs, FATP2 promotes the production of PGE2, which causes the expansion of polymorphonuclear MDSCs and further suppression of the antigen-specific T-cell response. Moreover, FATP2-induced lipid accumulation generates ROS and promotes the immunosuppressive effects of MDSCs on T cells. FAO promotes the immunosuppressive function of tumor-infiltrating MDSCs. In T cells, cholesterol increases immune checkpoint expression (*e.g.*, PD-1, 2B4, TIM-3, and LAG-3) and regulates TCR signaling. PGE2 inhibits T-cell proliferation and decreases IL-2 receptor expression in T cells to inhibit the early step of T-cell activation. In regulatory T cells, *de novo* fatty-acid synthesis mediated by FASN contributes to the functional maturation of Tregs, and COX-2 expression levels are positively correlated with the levels of tumor-infiltrating Foxp3⁺ Tregs. In macrophages, cholesterol and PGE2 induce macrophages to polarize to M2 TAM. For cytokines secreted by immune cells, lipids not only promote the secretion of immunosuppressive cytokines, such as IL-10 and TGF- β , but also inhibit the secretion of cytotoxic cytokines, such as TNF- α , IFN- γ , and granzyme B. Abbreviations: TAA, tumor-associated antigen; FAO, fatty acid oxidation; PD-L1, programmed cell death ligand 1; DC, dendritic cell; TIDC, tumor-infiltrating dendritic cell; Treg, regulatory T cell; MHC I, major histocompatibility complex class I; APC, antigen presenting cell; SCFA, short-chain fatty acid; ICOS, inducible co-stimulator; CTLA4, cytotoxic T lymphocyte antigen 4; FA, fatty acid; PGE2, prostaglandin E2; MDSCs, myeloid-derived suppressor cells; FATP2, fatty acid transport protein 2; PD-1, programmed cell death 1; TIM-3, T-cell immunoglobulin and mucin-domain containing-3; LAG-3, lymphocyte activation gene-3; IL-2, interleukin-2; IL-10, interleukin-10; TNF- α , tumor necrosis factor α ; TGF- β , transforming growth factor- β ; FASN, fatty-acid synthase; TCR, T-cell receptor; M1 TAM, M1 tumor-associated macrophage; M2 TAM, M2 tumor-associated macrophage; COX-2, cyclooxygenase-2; IFN- γ , interferon- γ ; TME, tumor microenvironment; CoA, coenzyme-A.

3. Lipid-mediated cancer–immunity cycle: T-cell priming and activation

TCR signaling and specific cytokines are responsible for T-cell growth, proliferation, and differentiation. Prolonged proliferation

of T cells and their differentiation into effector T cells are critical for antitumor immunity^{48,49}. In the TME, T-cell proliferation is impaired, and effector T-cell differentiation and function are altered or prevented⁴⁸. Here, we discuss how lipids affect T-cell priming and activation in the TME (Fig. 2). By understanding

these regulatory mechanisms of lipids, we can reorientate T-cell priming and activation by modulating lipid metabolism.

Previous studies have shown that cholesterol has a contradictory effect on TCR signaling depending on different biological functions. The direct binding motif of cholesterol appears in a variety of membrane proteins so that it can directly bind to membrane proteins, such as metabotropic glutamate receptor, adenosine A1 receptor, and γ -aminobutyric acid receptor⁵⁰. On the surface of the T-cell membrane, cholesterol also directly modifies the TCR through allosteric signals to alter it into an inactive conformation that cannot be phosphorylated by active kinases⁵¹. Cholesterol sulfate, which has a structure similar to cholesterol, can replace the originally bound cholesterol, inhibit the phosphorylation of CD3, destroy the TCR polymer, and effectively inhibit signal transduction²⁶. In contrast, cholesterol plays a positive role in regulating TCR signaling as part of membrane rafts. Yang et al.²⁷ found that by inhibiting cholesterol esterification, plasma membrane cholesterol levels in CD8⁺ T cells can be increased, resulting in the formation of larger TCR microclusters. This feature is beneficial for improving the avidity of TCRs to tumor antigens and promoting the formation of a larger TCR signalosome. Therefore, acetyl-CoA acetyltransferase one is a key cholesterol esterification enzyme, and its pharmacological inhibitor (avasimibe) was used to treat melanoma in mice with significant antitumor effects. Combination therapy of avasimibe plus anti-PD-1 is superior to monotherapy in repressing melanoma and Lewis lung carcinoma (LLC) progression in mice²⁷. Moreover, blocking cholesterol synthesis with lovastatin, a reversible competitive inhibitor of HMG-CoA, can dose-dependently inhibit CD3-induced T-cell proliferation⁵².

Prostaglandins inhibit T-cell proliferation and attenuate T-cell cytotoxicity. In the presence of PGE2, when T cells interacted with renal cell carcinoma cells, T-cell proliferation was inhibited, and intracellular IFN- γ expression and cell surface CD28 expression were reduced. The COX-2 inhibitor NS-398 increases cytotoxic T lymphocyte responses to renal cell carcinoma cells²⁸. Moreover, PGE2 results in decreased IL-2 production, IL-2 receptor expression, and responsiveness to exogenous IL-2, which inhibit the early step of T-cell activation³⁰. In addition, T cells primed by PGE2-matured DCs exhibit less granzyme B expression and poor tumor cytotoxicity. This result was also demonstrated *in vivo*²⁹.

Cholesterol and PGE2 metabolic reprogramming are significantly associated with T-cell priming and activation. Therefore, this strategy is another effective means to increase the activation of T cells to improve the efficacy of existing immunotherapies.

4. Lipid-mediated cancer-immunity cycle: T-cell killing of tumor

The majority of T cells in the tumor microenvironment are depleted with an increase in inhibitory receptors, a decrease in effector cytokines, and impairment of cytotoxicity, potentially leading to tumor immune evasion. The regulatory mechanisms of T-cell exhaustion in the TME can be divided into the extrinsic pathway and intrinsic pathway⁵³. Regarding extrinsic regulatory mechanisms, cancer cells and stromal cells (tumor-associated DCs, Tregs, tumor-associated macrophages (TAMs), and MDSCs) are the main exogenous cells that regulate T-cell exhaustion⁵⁴, and both IL-10 and TGF- β are important extrinsic cytokines involved in the exhausted process of T cells^{55–58}. Regarding intrinsic regulatory mechanisms, the binding of immune checkpoint

receptors and cognate ligands can exhaust T cells and inhibit their ability to kill tumor cells normally, so tumor cells can escape the immune surveillance of the host. On T cells, the immune checkpoint receptors include PD-1, CTLA-4, T-cell immunoglobulin and mucin-domain containing-3 (TIM-3), B- and T-lymphocyte attenuator (BTLA), lymphocyte activation gene-3 (LAG-3) and T cell immunoreceptor with Ig and ITIM domains (TIGIT). Their corresponding ligands, such as programmed cell death ligand 1 (PD-L1), galectin-9 and B7 homolog 3 (B7-H3) protein are expressed on tumor cells or MDSCs^{59–61}. Much evidence suggests that lipids lead to the functional exhaustion of T cells and suppress adaptive immune responses through the extrinsic pathway and intrinsic pathway (Fig. 2).

4.1. Lipids regulate T-cell exhaustion by extrinsic mechanisms

Lipid metabolism is a key factor in regulating the function of MDSCs. Tumor-infiltrating MDSCs show increased FA absorption and thus increased levels of FAO as their energy sources. Such MDSCs represent a stronger immunosuppressive effect on CD8⁺ T cells than MDSCs with normal lipid content^{31,62}. Pharmacological inhibition of FAO both reduced the immunosuppressive functions of tumor-infiltrating MDSCs and blocked their production of inhibitory cytokines. Monotherapy with the FAO inhibitor etomoxir or ranolazine significantly repressed the tumor growth of LLC and MCA-38 colon adenocarcinoma in a T-cell-dependent manner and synergized the antitumor effect of adoptive T-cell therapy⁶². In mice bearing LLC tumor, CT26 colon carcinoma, and B16F10 tumor, polymorphonuclear-MDSCs showed increased FATP2 expression, which suppressed the antigen-specific T-cell response in the TME. This phenomenon occurred because FATP2 promoted arachidonic acid uptake, and overload of arachidonic acid resulted in higher production of PGE2. This process promoted the expansion of polymorphonuclear-MDSCs and further suppressed the antigen-specific T-cell response⁶³. In another study, FATP2-induced lipid accumulation in MDSCs generated ROS and promoted the immunosuppressive function of MDSCs on T cells by reducing T-cell proliferation and suppressing the ability of T cells to produce IFN- γ and tumor necrosis factor α (TNF- α)⁶⁴. Therefore, reducing lipid accumulation by lipofermata, an inhibitor of FATP2, can block immunosuppressive activity in MDSCs and relieve T-cell exhaustion. More importantly, lipofermata can also enhance the antitumor effect of immunotherapy with anti-PD-L1 in B16F10 and LLC tumor-bearing mice⁶⁴. PGE2, the derivative of FAs, was conducive to the formation of an immunosuppressive microenvironment in melanoma. Microsomal PGE synthase-1 was positively correlated with CD14 and integrin alpha M in MDSCs. Therefore, PGE2 depletion by PGE synthase knockout may reduce MDSC immunosuppressive activity on activated T cells⁶⁵. Modulating PGE2 production with a COX-2 inhibitor has been demonstrated to efficiently improve the antitumor effect of immune checkpoint inhibitors (ICIs). Persistent tumor regression was observed in mice carrying prostaglandin E synthase KO tumors or in patients with metastatic melanoma and non-small cell lung cancer treated with a combination strategy of COX inhibitors and ICIs^{65,66}. For pancreatic cancer, immunosuppression in TME is considered a major obstacle to the effectiveness of immunotherapeutic approaches to induce antitumor immune responses⁶⁷. Pharmacological inhibition of COX-2 reduces the accumulation of MDSCs, thereby transforming the immunosuppressive microenvironment of pancreatic cancer into a T-cell-permissive environment⁶⁸. This strategy increases cancer

responses to the combination of checkpoint blockers and CD40 agonists⁶⁸.

Tregs maintain immune tolerance but also lead to immunosuppression in the tumor microenvironment^{69,70}. High blood levels of butyrate and propionate, which are the SCFAs, were associated with a higher proportion of Tregs in tumor-draining lymph nodes in tumor-bearing mice (CT26 models)²⁵. *De novo* fatty-acid synthesis mediated by FASN contributes to the functional maturation of Tregs, and FASN deletion from Tregs inhibits colon tumor growth, but the addition of palmitate, a product of FASN, restores the suppressive function of Tregs⁷¹. A retrospective analysis of 100 patients with non-small cell lung cancer showed that COX-2 expression is positively correlated with tumor-infiltrating foxp3⁺ Tregs, and patients with elevated COX-2 expression have shorter recurrence-free survival than patients without COX-2 expression³².

Macrophages can be polarized into classically activated (M1) and alternatively activated (M2) macrophages; the former foster an inflammatory response against tumor cells, whereas the latter tend to exert an immunosuppressive phenotype⁷². The inhibition of cholesterol efflux mediated by the ATP-binding cassette transporter G1 induces M2 macrophages to transform into the M1 type and enhances their cytotoxicity to kill cancer cells⁷³. In glioblastoma multiforme, FAO-derived acetyl-CoA upregulated CD47 transcription to weaken macrophage phagocytosis⁷⁴. Inhibition of FAO by etomoxir, a carnitine palmitoyltransferase 1 (CPT1) inhibitor, synergized with an anti-CD47 antibody to control regrown tumors with elevated macrophage phagocytosis⁷⁴. In addition, PGE2 has been found to induce M2 macrophage polarization and recruit MDSCs to suppress host immunity, which in turn increases lung tumor stemness and epithelial–mesenchymal transition-like features⁷⁵. In an immune-competent mouse model of C57 with inflammatory lung, using lower doses of prostaglandin E synthase inhibitors, Cay10526 can significantly inhibit lung metastasis⁷⁵. Moreover, conditional depletion of COX-2 and PGE2 synthase in myeloid cells disrupt M2-like TAM function, which enhanced cytotoxic T-cell function in a mouse model of breast cancer through⁷⁶.

DCs engulf FA-carrying tumor-derived exosomes, which can increase the secretion of TGF- β ³⁴. Moreover, in bone marrow-derived DCs, increasing PGE2 levels can promote the secretion of IL-10 through stimulation of EP₂R and EP₄Rs, which inhibits T-cell proliferation³³. MF-766 blocks EP₄R, synergistically improving the efficacy of anti-PD-1 therapy in CT26 and EMT6 tumor-bearing mice⁷⁷.

These results indicate that therapeutic strategies blocking lipid synthesis can restore T-cell activity; enhance IFN- γ , IL-2, and TNF- α production; decrease the infiltration and function of immunosuppressive cells; and promote the infiltration of CD8⁺ T cells, natural killer cells and conventional DCs^{68,77}. Therefore, these results suggest that drugs that regulate lipid metabolism can not only restore lipid homeostasis in tumors but also exert anti-tumor effects by increasing tumor immunity. In addition, these drugs can be used as immune sensitizers in combination with other drugs.

4.2. Lipids regulate T-cell exhaustion by intrinsic mechanisms

It is worth noting that increased lipids levels can increase immune checkpoint expression, suggesting that drugs blocking lipid synthesis can be used as novel immune checkpoint inhibitors to coordinate immunotherapy. A study revealed that FA-

carrying tumor-derived exosomes significantly increase the expression of inhibitory checkpoint proteins on DCs, such as PD-L1 and signal regulatory protein α (SIRP α)³⁴. FASN activity stabilizes the PD-L1 protein *via* PD-L1 palmitoylation to cause cisplatin resistance in bladder cancer⁷⁸. In breast cancer, disruption of PD-L1 palmitoylation sensitizes breast cancer cells to T-cell killing and thus represses tumor growth⁷⁹. These studies suggest that inhibition of lipid metabolism by targeting PD-L1 palmitoylation with small molecules may provide new avenues for improving the therapeutic efficacy of anti-PD-L1/PD-1 treatment.

Tumor tissues are rich in cholesterol, which leads to high cholesterol content in tumor-infiltrating CD8⁺ T cells. Cholesterol increases endoplasmic reticulum stress and then activates the endoplasmic reticulum stress sensor X-box binding protein 1 (XBP1), which regulates PD-1 and 2B4 transcription in CD8⁺ T cells³⁵. Consequently, tumor-infiltrating CD8⁺ T cells can positively and progressively increase the expression of PD-1, 2B4, T-cell immunoglobulin and mucin-domain containing-3 (TIM-3), and lymphocyte activation gene-3 (LAG-3)³⁵. Reducing the level of cholesterol in CD8⁺ T cells can restore the less-exhausted phenotype of tumor-infiltrating CD8⁺ T cells and effectively enhance antitumor activity in a mouse melanoma model³⁵. Simvastatin targets cholesterol biosynthesis and inhibits long non-coding RNA snhg29-mediated YAP activation, which in turn decreases PD-L1 transcription. Moreover, simvastatin inhibits colon cancer proliferation and synergizes with anti-PD-L1 against tumors⁸⁰.

PGE2 in lung cancer tissue homogenate positively regulates the level of PD-1 in infiltrating CD8⁺ T cells by activating the EP2 and EP4 pathways⁸¹. In summary, exploring the mechanisms through which lipids regulate these extrinsic and intrinsic immunosuppression pathways may benefit our understanding of how immunosuppression is propagated in the tumor and which lipids are potential targets in lipid metabolism for immunotherapy.

5. Combination therapy

With the continuous development of research on the impact of lipids on the TME, the findings suggest the feasibility of combining drugs that modulating lipids and tumor immunotherapy in cancer treatment. Investigations of the combination of ICIs and drugs used to modulate FAs are currently mainly in preclinical research. PPAR- γ agonists (rosiglitazone and bezafibrate), that reduce FA storage, showed better inhibition of tumor growth and prolongation of survival time mainly in melanoma and lung cancer, when used in combination with immunotherapy, such as anti-PD-1 and cancer cell vaccines^{40,82,83}. Preclinical studies also suggested that statins could potentiate immunotherapy in the treatment of lung cancer, melanoma and breast cancer^{80,84–86}. Some retrospective studies have shown that the combined use of statins and PD-1 inhibitors was associated with longer tumor treatment fields and improved objective response rates (ORR) and progression-free survival (PFS). NSCLC patients treated with nivolumab and statins exhibited a statistically significant response ($P = 0.02$) and better ORR (40% *versus* 22%, $P = 0.04$) and PFS (median 7.8 *versus* 3.6 months, $P = 0.03$) than the non-statin group^{87,88}. In malignant pleural mesothelioma, patients who were treated with statins showed an increased ORR (22% *versus* 6%, $P = 0.05$) and PFS (median 6.7 *versus* 2.4 months, $P < 0.01$)⁸⁸. Moreover, the levels of cholesterol and SCFAs may be a biomarker of the efficacy of tumor immunotherapy^{89,90}.

Table 1 Clinical trials involving the combination of immunotherapy and lipid-mediated therapies.

Adaptive immune cycle	Clinical trial identifier	Status	Phase	Cancer type	Drug (target)	Immunotherapy
Antigen presentation	NCT04093323	Recruiting	2	HLA ⁻ A2 ⁺ refractory melanoma	Celecoxib (COX-2)	α -Type-1 polarized dendritic cells, recombinant interferon α -2b, PD-1 ligand inhibitor, PD-1 inhibitor
	NCT02151448	Completed	1/2	Peritoneal surface malignancies	Celecoxib (COX-2)	DC vaccine, interferon α -2b
	NCT01158534	Completed	2	Renal cell cancer	Celecoxib (COX-2)	Interferon α -2b
	NCT03710876	Active, not recruiting	3	Malignant pleural mesothelioma	Celecoxib (COX-2)	rAd-interferon
	NCT03403634	Completed	2	Metastatic carcinoma in the liver, colorectal carcinoma	Celecoxib (COX-2)	Interferon α -2b
	NCT00081848	Completed	1	Liver neoplasms	Celecoxib (COX-2)	Recombinant Fowlpox-GM-CSF
T-cell priming and activation	NCT02922764	Recruiting	1	Advanced solid malignancies and lymphoma	RGX-104 (LXR)	Ipilimumab (CTLA4)
	NCT03396952	Active, not recruiting	2	Cutaneous melanoma	Aspirin (COX)	Ipilimumab (CTLA-4)
T-cell killing of tumor	NCT00193648	Completed	1	Focal glomerulosclerosis	Rosiglitazone (PPAR γ)	Atezolizumab (PD-L1)
	NCT04114136	Recruiting	2	Melanoma renal cell, carcinoma, NSCLC, hepatocellular carcinoma, urothelial cancer, gastric adenocarcinoma, HNSCC, esophageal adenocarcinoma, microsatellite instability-high solid malignant tumor	Rosiglitazone (PPAR γ)	Nivolumab (PD-1)/pembrolizumab (PD-1)
	NCT02922764	Recruiting	1	Advanced solid malignancies, lymphoma	RGX-104 (LXR)	Nivolumab (PD-1)/pembrolizumab (PD-1)
	NCT03638297	Recruiting	2	Colorectal cancer	Aspirin (COX)	BAT1306 (PD-1)
	NCT03926338	Recruiting	1/2	Colorectal cancer	Celecoxib (COX-2)	Toripalimab (PD-1)
	NCT03396952	Active, not recruiting	2	Melanoma	Aspirin (COX)	Ipilimumab (PD-1)
	NCT03245489	Recruiting	1	Head and neck cancer	Acetylsalicylic acid (COX)	Pembrolizumab (PD-1)
	NCT02659384	Active, not recruiting	2	Ovarian neoplasms	Acetylsalicylic acid (COX)	Atezolizumab (PD-L1)

Combination strategies using immunotherapy and COX inhibitors were mainly carried out, and a small number of drugs modulating FAs and cholesterol were also implemented in clinical trials, which are summarized in [Table 1](#).

6. Conclusions

Dysregulated cellular metabolism is the hallmark of cancer, in which lipid metabolism is now an undisputedly thought to be factor in supporting cancer growth and progression. Combined with recent findings in the field of tumor immune escape and cancer immunotherapy, we review the role of aberrant lipids, especially, FAs, cholesterol, and prostaglandins, an arachidonic acid derivative, in regulating tumor immunity and immunotherapy by acting on the cancer–immunity cycle.

De novo lipid synthesis, increased FA uptake and FAO can impair the capacity of DCs to process TAAs, promote the infiltration of Tregs and secretion of TGF- β and increase the expression of PD-L1 and CD47 on tumor or immune cells. Long- or short-chain saturated FAs can reduce the expression of MHC I or costimulatory factors respectively, thereby inhibiting the ability of APCs to activate T cells. Lipid synthesis, FA uptake, and FAO can be blocked by inhibiting acetyl-CoA, PPAR γ , FATP2, and CPT1, respectively. These inhibitors synergize with immunotherapy in tumor-bearing mice (CT26, MC38, MCA-38 and LLC mice models). In clinical trials, only the PPAR γ inhibitor rosiglitazone in combination with PD-1/PD-L1 monoclonal antibodies to treat solid tumors has entered phase 1/2 ([Table 1](#)). Moreover, ASC40, a novel potent oral inhibitor of FASN, has entered phase three clinical trials in combination with

bevacizumab for the treatment of recurrent glioblastoma (NCT05118776) and exhibits promise as a drug candidate for synergistic immunotherapy. In addition, etomoxir, the pharmacologic inhibitor of CPT1, has been shown to synergize with the antitumor effect of adoptive T-cell therapy in preclinical research, but its clinical translation is less likely due to its severe hepatotoxicity⁹¹.

Cholesterol hindered DCs from recognizing GSL, a TAA, by changing its conformation, regulated TCR signaling, increased immune checkpoint expression in T cells and affected macrophage polarization to M2 macrophages. In preclinical research, simvastatin synergizes with anti-PD-L1 against colon cancer. However, the clinical trial investigating the combination of lovastatin and IFN α -2b was in a withdrawn state, but the reasons for the withdrawal were not specified (NCT00963664). RGX-104, a LXR agonist, protects cells from cholesterol overload⁹². It has been studied in combination therapy with ipilimumab, nivolumab, or pembrolizumab to treat advanced solid malignancies and lymphoma in phase 1, but no results have been published yet (Table 1).

PGE2 inhibited all three steps of the immune cycle. PGE2 reduces the accumulation of tumor infiltrating dendritic cell to inhibit antigen uptake. In addition, PGE2 inhibits T-cell proliferation and suppresses the secretion of IFN- γ , TNF- α , and granzyme B. Moreover, it increases the infiltration of immunosuppressive cells, such as Tregs and MDSCs, and the secretion of immunosuppressive cytokines, such as IL-10. It is worth noting that there are many clinical trials on the combination of COX inhibitors and immunotherapy against tumors (Table 1). This finding indicates the great potential of COX inhibitors in improving the efficacy of immunotherapy. COX-2-specific inhibitors may not affect the protective effect of COX-1 on the gastrointestinal tract, platelets, etc.⁹³. This finding indicates that the COX-2 inhibitor celecoxib can be safer when used in combination with immunotherapy.

In summary, understanding how lipids affect antigen presentation, T-cell priming and activation, and T-cell killing of tumors and how they alter the immune context within the TME may provide new therapeutic targets against tumors. Whether new drug candidates can achieve safety, efficacy and tolerability in pre-clinical and clinical trials when used alone or in combination with other cancer treatments remains to be verified.

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Author contributions

Mingming Zheng designed and wrote the paper. Wenxin Zhang, Xi Chen, Hongjie Guo, Honghai Wu, Yanjun Xu, and Qiaojun He revised the manuscript. Ling Ding and Bo Yang were responsible for the conception and design of the review.

Conflicts of interest

The authors declare no conflicts of interest.

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