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# Tumor-infiltrating lymphocytes in NSCLC: from immune surveillance to immunotherapy

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Lung cancer, predominantly non-small cell lung cancer (NSCLC), remains a principal driver of cancer-related morbidity and mortality worldwide. Despite advancements in surgery, radiotherapy, chemotherapy, and targeted treatments, outcomes remain poor in advanced NSCLC. The tumor microenvironment (TME) exerts a critical influence on therapy responses. Within the TME, immune cells such as T and B lymphocytes, dendritic cells, myeloid-derived suppressor cells, tumor-associated macrophages, neutrophils, and natural killer cells can drive both pro- and anti-tumor processes. This review integrates their classification, phenotypic plasticity, and roles in NSCLC, highlighting key preclinical and clinical evidence while discussing pathogenesis, prognostic significance, and therapeutic potential. We also summarize the current immunotherapeutic strategies for advanced NSCLC, including first- or second-line regimens with immune checkpoint inhibitors alone or combined with chemotherapy, anti-angiogenic agents, or additional checkpoint inhibitors, and future directions. By elucidating the interplay between the NSCLC immune microenvironment and emerging immunotherapies, this review emphasizes the need for novel combination regimens and robust predictive biomarkers to improve clinical outcomes and extend survival in advanced NSCLC.

## KEYWORDS

lung cancer, tumor microenvironment, immune cells, immunotherapy, CD8 + T cell, B cell

## 1 Introduction

Lung cancer remains the leading cause of cancer-related deaths worldwide, with non-small cell lung cancer (NSCLC) accounting for 80-85% of all cases (1, 2). Despite advances in surgery, radiotherapy, chemotherapy, and targeted therapies, the prognosis for advanced-stage NSCLC remains poor, with a five-year survival rate below 5% (3). Despite advancements in surgical resection, radiotherapy, chemotherapy, and targeted therapies, a majority of patients are diagnosed at locally advanced or metastatic stages,

resulting in a poor prognosis (4). Immunotherapy has emerged as a novel treatment strategy for solid tumors, including lung cancer; however, most patients derive significant benefit from these interventions (4, 5).

The tumor microenvironment (TME) is composed of tumor cells, immune cells, cancer-associated fibroblasts (CAFs), signaling molecules, and the extracellular matrix (ECM) (6–9). It can be broadly categorized into the immune microenvironment, dominated by immune cells, and the non-immune microenvironment, primarily comprising CAFs (10). In the NSCLC TME, tumor cells exert profound effects on infiltrating immune cells, shaping an immunosuppressive milieu that promotes tumor progression (11). This review provides a comprehensive overview of the classification and recent advances in the study of immune cells within the NSCLC immune microenvironment.

## 2 Immune cells in the NSCLC tumor immune microenvironment

### 2.1 Antigen-presenting cells

#### 2.1.1 Dendritic cells

DCs, originating from the myeloid lineage, serve as central regulators of antitumor immunity and are the most potent local antigen-presenting cells (APCs) (12). The immune checkpoint molecule B7-H3 has been identified as an independent predictor of poor prognosis in NSCLC patients, with significantly upregulated expression observed in tumor-resident DCs of NSCLC patients (13). Cytokine-induced killer (CIK) cells in combination with dendritic cells (DC-CIK) have been shown to induce apoptosis in Lewis lung carcinoma cells, potentially through the downregulation of 14-3-3 $\zeta$  and p-Bad proteins, thereby supporting the potential adjuvant role of DC-CIK in NSCLC combinatorial therapy (14, 15). Moreover, intratumoral injection of autologous dendritic cells engineered to express the CCL21 gene via an adenoviral vector elicited systemic antigen-specific immune responses, leading to increased CD8<sup>+</sup> T cell infiltration and upregulation of intratumoral PD-L1 expression (16). In addition, non-small cell lung cancer cells downregulate the expression of costimulatory molecules such as CD80 and CD86 and pro-inflammatory cytokines including IL-12 and IL-23, while promoting the secretion of the anti-inflammatory cytokine IL-10 by CD1c<sup>+</sup> dendritic cells, thereby impairing endogenous antitumor immunity (17). These findings provide a theoretical foundation for combining immunotherapy with *in situ* vaccination strategies, although further studies are needed to evaluate the efficacy of PD-1/PD-L1 checkpoint blockade in conjunction with DC-CCL21-based therapy.

#### 2.1.2 B cells

In NSCLC, tumor-infiltrating B cells (TIBs) are more abundant in tumor tissues than in adjacent non-tumor tissues, particularly in NSCLC (18). TIBs and CD4<sup>+</sup> TILs are primarily located within tertiary lymphoid structures (TLSs), which correlate with improved

prognosis in early-stage NSCLC, indicating TIBs' antitumor role (19). Germain et al. (20) revealed that somatic hypermutation and class-switch recombination occur in TLS germinal centers in NSCLC, underscoring the prognostic significance of B cell density. Low CD20<sup>+</sup> B cell and DCLAMP<sup>+</sup> dendritic cell densities predict high mortality risk, suggesting their utility in therapeutic stratification. TIBs can present tumor antigens to CD4<sup>+</sup> TILs, modulating their phenotypes into activated, antigen-related, and non-responsive subtypes, with activated TIBs linked to effector T cell responses (21). Beyond anti-CD20 therapies, B cell-targeted strategies remain preclinical, necessitating future research to enhance antitumor immunity. Conversely, TIBs also promote tumor progression by suppressing antitumor immunity, influenced by the TME. Regulatory B cells (Bregs) secrete IL-35 and TGF- $\beta$ , inhibiting immune responses (22, 23). In autoimmune and transplantation contexts, B cell depletion impairs T cell function, while Bregs promote Treg phenotypes (24, 25). These findings highlight the dual role of TIBs in NSCLC and suggest TIB-CD4<sup>+</sup> TIL interactions as potential immunotherapeutic targets.

### 2.2 Effector immune cells

#### 2.2.1 CD8<sup>+</sup> and CD4<sup>+</sup> T cells

Tumor-infiltrating lymphocytes (TILs), particularly CD4<sup>+</sup> and CD8<sup>+</sup> T cells and their immunomodulatory cytokines, are critical components of adaptive immunity within the TME (26, 27). The density of CD4<sup>+</sup>/CD8<sup>+</sup> T cell infiltration in the tumor stroma is strongly associated with NSCLC patient prognosis, with higher TIL levels correlating with improved overall survival (OS) and disease-free survival (DFS) (28). CD8<sup>+</sup> T cells, in particular, play a pivotal role in suppressing lung cancer progression, though their antitumor activity can be inhibited by Foxp3<sup>+</sup> regulatory T cells (Tregs) (29). The transcriptional profiles of these cells offer insights for immunotherapy optimization and biomarker identification for checkpoint blockade therapies (30). Inhibitory receptors such as PD-1, TIM-3, CTLA-4, and LAG-3 on CD8<sup>+</sup> T cells are linked to disease progression, as their overexpression impairs T cell activation and effector function (31). Ren et al. (32) demonstrated that PD-L1 expression in lung squamous cell carcinoma correlates with CD8<sup>+</sup> TIL infiltration and MET gene upregulation, highlighting potential therapeutic targets.

CD4<sup>+</sup> T cell subsets, including Tregs and Th17 cells, also significantly influence cancer pathogenesis (33–35). Tregs modulate cytokine and chemokine production, affecting immune cell recruitment and antitumor responses, thereby contributing to lung cancer progression (36). NSCLC patients exhibit higher CD4<sup>+</sup> T cell frequencies in tumors compared to normal tissues, with CD4<sup>+</sup>Foxp3<sup>+</sup> Tregs constituting a substantial TIL proportion (37). Cho et al. (11) identified Treg heterogeneity based on TNFRSF9 expression, linking activated Tregs to poor prognosis in lung adenocarcinoma. Th17 cells, characterized by IL-17 secretion, promote inflammation and lung cancer development, with IL-17 pathway alterations and genetic/epigenetic modifications increasing cancer susceptibility (12). Further research is needed to clarify the

Treg-Th17 balance in lung cancer pathogenesis and its prognostic implications (38).

### 2.2.2 Natural killer cells

NK cells are critical early effectors in anti-tumor immunity and constitute the first line of defense against malignancies (39). The activation and cytotoxic function of NK cells are tightly regulated by a balance between inhibitory and activating receptors expressed on their surface (40). Studies have indicated that the presence of NK cells does not significantly impact the clinical outcomes of NSCLC patients, possibly due to the TME's capacity to remodel intratumoral NK cells by reducing the expression of activating receptors while upregulating inhibitory receptors such as CTLA-4 (41). Additionally, NK cells express immune checkpoint molecules, including PD-1, LAG-3, and TIM-3 (42). Emerging evidence suggests that the lung TME may induce NK cell suppression, with CD8<sup>+</sup> T cell infiltration positively correlating with CTLA-4 expression in NK cells, implying the presence of an inhibitory NK cell population even in immunologically active tumors (43). Despite significant advancements in cancer immunotherapy, a substantial proportion of patients fail to derive clinical benefit, partly due to the absence of a tumor-specific NK cell response. Therefore, strategies aimed at reactivating NK cells in combination with other

therapeutic modalities may offer promising prospects for improving lung cancer treatment and prognosis (Figure 1).

## 2.3 Immunosuppressive and regulatory immune cells

### 2.3.1 Myeloid-derived suppressor cells

MDSCs are a heterogeneous population of bone marrow-derived cells with potent immunosuppressive properties, widely present in the peripheral blood and tumor tissues of cancer patients (44–46). Their abundance and proportion correlate with tumor size and malignancy. MDSCs suppress both antigen-specific antitumor immunity mediated by T cells and non-specific antitumor immunity mediated by NK cells and macrophages through the expression of high levels of arginase, nitric oxide synthase, and reactive oxygen species (47). MDSCs are classified into polymorphonuclear MDSCs (PMN-MDSCs) and monocytic MDSCs (M-MDSCs). PMN-MDSCs exhibit morphological and phenotypic similarities to neutrophils, whereas M-MDSCs resemble immature monocytes. In both murine and human tumors, PMN-MDSCs predominate and closely resemble neutrophils in phenotype (48). Clinical analyses indicate that the

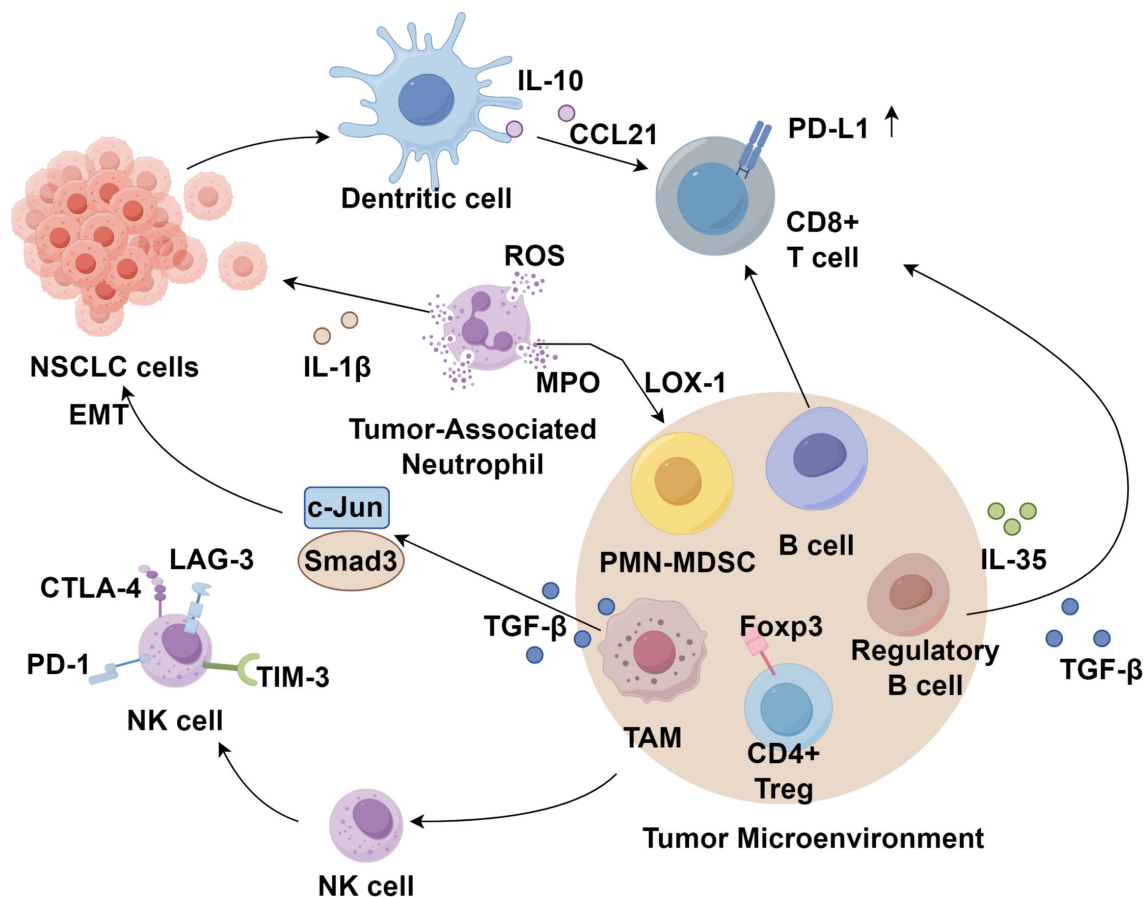


FIGURE 1  
Immune cells in the NSCLC tumor immune microenvironment.

expression of C-C chemokine receptor 5 (CCR5) is significantly elevated in peripheral blood M-MDSCs of NSCLC patients compared to healthy individuals. Additionally, the proportion of PMN-MDSCs and CCR5<sup>+</sup> M-MDSCs in circulation negatively correlates with recurrence-free survival but is not associated with their proportion in tumor tissues (49). Tian et al. (50) demonstrated that endoplasmic reticulum stress in neutrophils induces significant upregulation of lectin-type oxidized LDL receptor-1 (LOX-1), facilitating their transformation into PMN-MDSCs. Current evidence suggests that the abundance of LOX-1<sup>+</sup> PMN-MDSCs in NSCLC patients correlates with tumor size. However, further studies are warranted to evaluate their potential as prognostic biomarkers and therapeutic targets in NSCLC.

### 2.3.2 Tumor-associated macrophages

Macrophages infiltrating the tumor tissue undergo differentiation and maturation into tumor-associated macrophages (TAMs) under the influence of immunosuppressive cytokines derived from both the tumor and its microenvironment (51). TAMs play a crucial role in promoting tumor angiogenesis and lymphangiogenesis (52). TAMs exhibit distinct phenotypic polarization, primarily classified into M1 and M2. IL-4 can induce M2 polarization of macrophages within the TME, contributing to extracellular matrix (ECM) degradation and TME remodeling (53). Lung cancer cells also promote M2 polarization of TAMs, and the secretion of TGF- $\beta$  by TAMs activates the c-Jun and Smad3 pathways in lung cancer cells, leading to increased expression of SOX9, thereby facilitating epithelial-mesenchymal transition (EMT), tumor proliferation, migration, and invasion (54). A study by Alessandra et al. (55) demonstrated that the density of M2-polarized TAMs in the stroma of lung squamous cell carcinoma is significantly higher than that in lung adenocarcinoma. Furthermore, the density of M2 TAMs in the stroma of NSCLC correlates with lymph node metastasis, pathological staging, reduced disease-free survival (DFS), and OS (56, 57). These findings suggest that the stromal density of M2 TAMs serves as a potential indicator of tumor malignancy and clinical prognosis.

### 2.3.3 Tumor-associated neutrophils

At the cellular level, neutrophils constitute the most abundant immune cell population in NSCLC, accounting for approximately 20% of total immune cells (58). Studies have indicated that early-stage lung cancer can induce the formation of a distinct subset of tumor-associated neutrophils (TANs) with APC characteristics, capable of cross-presenting antigens and enhancing anti-tumor T cell responses (59). TANs also contribute to tumor proliferation, invasion, and metastasis through the synthesis and secretion of specific proteases. McLoed et al. (60) demonstrated that neutrophils facilitate lung cancer progression by expressing IL-1 $\beta$ , thereby mediating resistance to nuclear factor-kappa B (NF- $\kappa$ B) inhibitors. This mechanism represents a key factor in the limited therapeutic efficacy of NF- $\kappa$ B inhibitors in lung cancer. Contrary to these pro-tumorigenic effects, neutrophils also exert anti-tumor and anti-metastatic functions (61). The primary mechanism by which TANs inhibit tumor cell proliferation involves antibody-dependent

cellular cytotoxicity (ADCC), wherein antibodies recognize tumor cells via Fc receptors on TANs, leading to the release of cytotoxic mediators and subsequent tumor cell lysis. Moreover, TANs directly produce cytotoxic mediators such as reactive oxygen species (ROS) and myeloperoxidase (MPO), thereby inhibiting tumor cell proliferation (62). Studies have confirmed that interferon-beta (IFN- $\beta$ ) suppresses the expression of angiogenic factors such as vascular endothelial growth factor (VEGF) and matrix metalloproteinase-9 (MMP-9) in tumor-infiltrating neutrophils, leading to accelerated tumor vascularization and growth in IFN- $\beta$ -deficient models (63). However, as most research on the role of neutrophils in lung cancer is based on animal models, limited data are available regarding the phenotype and function of TANs in lung cancer patients, necessitating further investigation.

## 2.4 Functional plasticity of tumor-infiltrating immune cells

The tumor microenvironment (TME) is characterized by the remarkable plasticity of infiltrating immune cells, which can adopt either anti-tumorigenic or pro-tumorigenic phenotypes depending on local signals, cytokine milieu, and tumor-derived factors (64). Effector CD8<sup>+</sup> T cells mediate cytotoxicity against tumor cells but may become functionally exhausted, marked by sustained expression of inhibitory receptors, such as PD-1, TIM-3, LAG-3 and impaired cytokine production (65, 66). Similarly, macrophages exhibit phenotypic polarization along an M1–M2 spectrum: M1 macrophages promote antigen presentation and tumor clearance, whereas M2 macrophages support immunosuppression, angiogenesis, and metastasis (67, 68). B cells, dendritic cells, neutrophils, and NK cells also show phenotypic duality in NSCLC (69–71). This functional versatility enables tumors to co-opt immune responses and contributes to immunotherapy resistance (Table 1).

## 2.5 Immune–stromal crosstalk and extracellular matrix remodeling

In NSCLC, immune cell-mediated modulation of the extracellular matrix (ECM) and stromal compartment plays a pivotal role in tumor invasion, angiogenesis, and immune evasion (72, 73). Tumor-associated macrophages (TAMs), particularly of the M2 phenotype, secrete matrix metalloproteinases (MMPs) such as MMP-9 and MMP-2, which degrade collagen and other ECM components, thereby facilitating tumor cell migration and metastasis (74–76). Additionally, TAM-derived transforming growth factor-beta (TGF- $\beta$ ) stimulates cancer-associated fibroblasts (CAFs), promoting desmoplastic reactions and the deposition of fibrotic ECM, which impairs immune cell infiltration and fosters an immunosuppressive niche (77–80). Neutrophils contribute similarly by releasing neutrophil elastase, enzymes that remodel ECM and release growth factors stored in the matrix (81, 82). CD8<sup>+</sup> T cells and Th17 cells can also indirectly influence stromal



TABLE 1 Functional plasticity of key immune cells in the NSCLC tumor microenvironment.

Immune Cell Type	Anti-tumorigenic Phenotype	Pro-tumorigenic Phenotype
CD8 <sup>+</sup> T Cells	Effector T cells (high IFN- $\gamma$ , granzyme B, cytotoxic activity)	Exhausted T cells (PD-1 <sup>+</sup> , LAG-3 <sup>+</sup> , TIM-3 <sup>+</sup> , reduced cytotoxicity)
CD4 <sup>+</sup> T Cells	Th1 cells (support CD8 <sup>+</sup> T cell activation)	Tregs (Foxp3 <sup>+</sup> , suppress immune response via IL-10, TGF- $\beta$ )
B Cells	Antigen-presenting B cells, antibody-producing plasma cells	Regulatory B cells (IL-35, TGF- $\beta$ secretion, Treg promotion)
Macrophages	M1 macrophages (IL-12 <sup>+</sup> , TNF- $\alpha$ <sup>+</sup> , anti-tumor)	M2 macrophages (IL-10 <sup>+</sup> , TGF- $\beta$ <sup>+</sup> , promote EMT, angiogenesis)
Dendritic Cells	Mature DCs (high CD80/CD86, IL-12 secretion, antigen presentation)	Tolerogenic DCs (IL-10 <sup>+</sup> , low co-stimulatory molecules)
Neutrophils	N1 TANs (antigen-presenting, promote CD8 <sup>+</sup> responses, ADCC)	N2 TANs (secrete VEGF, MMPs, promote metastasis)
NK Cells	Activated NK (NKG2D <sup>+</sup> , perforin/granzyme production)	Dysfunctional NK (PD-1 <sup>+</sup> , CTLA-4 <sup>+</sup> , impaired cytotoxicity)

remodeling through cytokines such as IFN- $\gamma$  and IL-10, respectively, which modulate fibroblast activation and angiogenesis (66, 83). These immune–stromal interactions generate a dynamic feedback loop that alters the biophysical and biochemical properties of the TME, thereby enhancing immune resistance and diminishing immunotherapy efficacy (84). Understanding and targeting the immune-mediated reprogramming of the stroma represents a promising avenue for improving immunotherapeutic outcomes in NSCLC.

### 3 Immunotherapy for lung cancer

#### 3.1 Monotherapy with immune checkpoint inhibitors

Immune checkpoint inhibitors have emerged as a cornerstone of first-line treatment for advanced NSCLC with high PD-L1 expression and no actionable driver mutations. Pembrolizumab, atezolizumab, and cemiplimab as monotherapy options have been approved for clinical evidence (85). The landmark KEYNOTE-024 trial established pembrolizumab as a superior alternative to platinum-based chemotherapy, demonstrating significant improvements in objective response rate (ORR), progression-free survival (PFS), and overall survival (OS) in patients with PD-L1 TPS  $\geq 50\%$  (86, 87). Long-term follow-up confirmed sustained survival benefits, reinforcing its role as a first-line standard of care. The subsequent KEYNOTE-042 trial expanded eligibility to patients with PD-L1 TPS  $\geq 1\%$ , though the greatest survival advantage remained confined to those with PD-L1 TPS  $\geq 50\%$  (88, 89). Further validation came from IMpower110 and EMPOWER-Lung 1, which demonstrated that atezolizumab and cemiplimab, respectively, significantly improved survival outcomes compared to chemotherapy in PD-L1-high populations (90–93). These findings collectively underscore the efficacy of ICI monotherapy in patients with high PD-L1 expression. In patients with high PD-L1 expression, atezolizumab and cemiplimab monotherapy significantly improved survival outcomes compared to standard chemotherapy. These findings underscore the transformative impact of immune monotherapy in PD-L1-high expressors; however, its efficacy in patients with low or absent PD-L1 expression remains limited. Therefore, combination

immunotherapy strategies hold significant potential in expanding the patient population that may benefit from immune-based treatments.

#### 3.2 Combination immunotherapy

Immunotherapy combined with chemotherapy is the guideline-recommended first-line treatment for advanced NSCLC without driver mutations, regardless of PD-L1 expression. The KEYNOTE-189 trial (94, 95) demonstrated that pembrolizumab plus chemotherapy significantly improved PFS and overall survival (OS) in non-squamous NSCLC compared to chemotherapy alone. The KEYNOTE-407 trial (95, 96) extended these benefits to squamous NSCLC, establishing pembrolizumab-chemotherapy as a preferred regimen for both subtypes. Similarly, the EMPOWER-Lung 3 study (97, 98) showed cemiplimab plus chemotherapy improved OS, leading to FDA approval. The IMpower130 trial (99) also supported atezolizumab-chemotherapy for non-squamous NSCLC, with OS benefits. In China, the National Medical Products Administration (NMPA) approved camrelizumab, sintilimab, toripalimab, and sugemalimab combined with chemotherapy based on trials such as Camel (100, 101), Camel-sq (102), ORIENT-11 (103, 104), ORIENT-12 (105), RATIONALE-304 (106), RATIONALE-307 (107, 108), and GEMSTONE-302 (109). Toripalimab-chemotherapy was approved for non-squamous NSCLC based on the CHOICE-01 study (110), while penpulimab and serplulimab were approved for squamous NSCLC based on AK105-302 (111) and ASTRUM-004 (112) trials, respectively. Combination immunotherapy with anti-angiogenic agents has also shown promise. The IMpower150 trial (113, 114) demonstrated that atezolizumab plus bevacizumab, carboplatin, and paclitaxel (ABCP) improved PFS and OS compared to bevacizumab-chemotherapy (BCP), leading to FDA approval for metastatic non-squamous NSCLC. Dual immunotherapy regimens combining PD-1/PD-L1 inhibitors with CTLA-4 inhibitors have also demonstrated efficacy. These regimens have been FDA-approved for advanced NSCLC based on their clinical trial outcomes.

### 4 Conclusion

A deeper understanding of the NSCLC immune microenvironment has yielded critical insights into how distinct immune cell

populations can both foster and restrain tumor growth. T cells, in particular CD8<sup>+</sup> T cells, remain central to tumor elimination, but their function is frequently hampered by immunosuppressive cells and inhibitory checkpoint pathways. B cells, dendritic cells, and natural killer cells similarly exhibit dual, context-dependent roles, underscoring the complexity of anti-tumor immunity. Myeloid-derived suppressor cells, tumor-associated macrophages, and neutrophils further highlight the TME's capacity to promote immune evasion, metastasis, and therapy resistance. Recent clinical advances in immunotherapy have transformed the therapeutic paradigm for advanced NSCLC. Immune checkpoint inhibitors—alone or in combination with chemotherapy, anti-angiogenic agents, or other immunotherapeutic approaches—have significantly prolonged survival in select patient subgroups. Nevertheless, challenges remain, as many patients exhibit primary or acquired resistance, and the limited durability of responses underscores the need for more effective, personalized strategies.

We propose that a novel therapeutic paradigm should integrate immune contexture profiling, functional plasticity mapping of immune cells, and adaptive treatment strategies to convert immunosuppressive milieus into immunoreactive states. This review highlights that reprogramming key immune subsets—such as exhausted CD8<sup>+</sup> T cells, M2-like TAMs, and suppressive MDSCs—may represent a transformative approach in NSCLC treatment. Subsequent investigations must prioritize the discovery of robust biomarkers for prediction, elucidating pathways underlying immunosuppressive phenomena and tumor immune evasion, as well as optimizing combined therapeutic protocols that enhance immune checkpoint inhibition efficacy. Through a comprehensive strategy incorporating mechanistic understanding of immunocyte behavior, malignant transformation processes, and innovative treatment frameworks, the oncology community can advance toward attaining sustained clinical responses and enhanced patient outcomes in late-stage non-small cell lung carcinoma.

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